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1 **A subset of highly responsive transcription factors upon tomato infection by pepino mosaic**
2 **virus**

3
4 **Dikran Tsitsekian^{a,1}, Gerasimos Daras^{a,1}, Dimitris Templalexis^a, Fengoula Avgeri^a, Leonidas**
5 **Lotos^b, Chrysoula G. Orfanidou^b, Vardis Ntoukakis^c, Varvara I. Maliogka^b, Stamatis**
6 **Rigas^{a,*}**

7
8 ^aDepartment of Biotechnology, Molecular Biology Laboratory, Agricultural University of Athens,
9 Athens 11855, Greece

10 ^b School of Agriculture, Plant Pathology Laboratory, Aristotle University of Thessaloniki,
11 Thessaloniki 54124, Greece

12 ^c School of Life Sciences and Warwick Integrative Synthetic Biology Centre, University of
13 Warwick, Coventry, UK

14 ¹ Contributed equally to this work.

15

16 * Corresponding author.

17 *E-mail address:* srigas@aua.gr (S.R.)

18 Tel: +30-210-529 4210

19

20 **e-mails:** DTs: dtsitsekian@aua.gr
21 GD: gdaras@aua.gr
22 DTem: dimitempl@aua.gr
23 FA: favgeri@aua.gr
24 LL: llotos@agro.auth.gr
25 CO: chorfani@agro.auth.gr
26 VN: V.Ntoukakis@warwick.ac.uk
27 VM: vmaliogk@agro.auth.gr
28 SR: srigas@aua.gr

29

30 **Short Title:** tomato responsive TFs against PepMV

31 **One-sentence summary:** A combinatorial research approach was applied primarily relying on
32 transcriptome analysis to identify highly responsive tomato TFs and understand the molecular
33 defensive mechanisms of the tomato-PepMV pathosystem.

34

35 **Keywords:** *Solanum lycopersicum*, RNA-seq, PepMV, transcription factors, C2H2 Zinc Finger,
36 *NAC*, *bHLH*

37

38 **Abbreviations**

39 DEGs: differentially expressed genes

40 TFs: transcription factors

41 PepMV: pepino mosaic virus

For Peer Review

42 **ABSTRACT**

- 43 • Plants have evolved well-tuned surveillance systems, including perplexed defense
44 mechanisms, to restrain pathogens. Transcription factors (TFs) are master regulators of
45 host molecular responses against plant pathogens. While pepino mosaic virus (PepMV)
46 constitutes a major threat to the global tomato production, there is still lack of
47 knowledge about the key TFs that regulate host responses against the virus.
- 48 • A combinatorial research approach was applied relying on tomato transcriptome
49 analysis, reverse-transcription quantitative-PCR validation, phylogenetic
50 classification, comparative analysis of structural features, *cis*-regulatory element
51 mining and *in silico* co-expression analysis to identify a set of 11 highly responsive
52 TFs involved in the regulation of host responses against PepMV.
- 53 • An endemic PepMV isolate, generating typical mosaic symptoms, modified the
54 expression of approximately 3.3% tomato genes resulting in 1,120 Differentially
55 Expressed Genes (DEGs). Functional classification of 502 upregulated DEGs revealed
56 that photosynthesis, carbon fixation and gene silencing were widely affected, whereas
57 618 downregulated genes had an impact on mainly plant defense and carotenoid
58 biosynthesis. Strikingly all 11 highly responsive TFs carried abiotic stress response *cis*-
59 regulatory elements, whereas 5 of them were better aligned with rice than *Arabidopsis*
60 gene homologs, suggesting that plant response against viruses may predate the
61 divergence to monocots and dicots. Interestingly, tomato *C2H2* family TFs, *ZAT1-like*
62 and *ZF2*, may have distinct roles in plant defense due to the opposite response pattern,
63 similarly to their *Arabidopsis* *ZAT10* and *ZAT12* homologs.
- 64 • These highly responsive TFs set the basis to study in deep the molecular responses of
65 tomato-PepMV pathosystem, opening a perspective to better comprehend viral
66 infections.

67 INTRODUCTION

68 Tomato (*Solanum lycopersicum*) is an important vegetable of high economic value due to its taste,
69 high-yield and elevated content of beneficial to human health compounds (Martinez-Valverde et
70 al., 2002). Tomato plants are susceptible to various pathogens that compromise both fruit yield
71 and quality. Viral diseases constitute a major threat due to practical restrictions regarding their
72 control (Hanssen et al., 2010). Viruses have a great potential to quickly evolve and adapt under
73 natural selection pressure due to their large population size, short generation time and high rates
74 of mutations or recombination events (Moya et al., 2004; Hanssen et al., 2010). The spread of
75 newly evolved strains constitutes a constant threat to modern agriculture.

76
77 Up to date, more than 312 viral species, including satellite viruses and viroids, have been reported
78 to infect tomato (Rivarez et al., 2021). Pepino mosaic virus (PepMV) together with the tomato
79 yellow leaf curl virus (TYLCV), tomato torrado virus (ToTV) and tomato brown rugose fruit virus
80 (ToBRFV) have recently emerged mostly in greenhouses with a devastating impact on tomato global
81 production (Hančinský et al., 2020; Hanssen et al., 2010). PepMV belongs to the *Potexvirus* genus
82 of the *Alphaflexiviridae* family (Virus Taxonomy: 2019 Release, ictvonline.org) and was first
83 reported in 1980 on pepino (*Solanum muricatum*) in Peru (Jones et al., 1980). Twenty years after
84 the initial report on tomato infections in the Netherlands, PepMV has become a major pathogenic
85 threat for greenhouse grown tomatoes worldwide (Hanssen et al., 2009; Alcaide and Arranda,
86 2021).

87
88 PepMV has a positive sense single stranded RNA genome of approximately 6400 nucleotides long
89 that contains five open reading frames (ORFs), including an RNA-dependent RNA polymerase
90 gene (*RdRp*), a triple gene block (*TGB*) and a coat protein gene (*CP*) (Hanssen and Thomma,
91 2010). Five PepMV strains have been reported with a sequence identity ranging from 78% to 95%:
92 the original Peruvian (LP), the European (EU), the American (US1), the Chilean (CH2) and the
93 new Peruvian (PES) (Hanssen and Thomma, 2010; Moreno-Pérez et al., 2014). Since its first
94 appearance, PepMV has been rapidly established in most tomato producing countries. Within a
95 short-time period, PepMV outbreaks were reported in the EU, UK, China, Canada and the USA
96 (Hanssen and Thomma, 2010). Initially, these reported outbreaks were related with the EU strain,
97 causing a rather mild series of symptoms. However, studies on the genetic structure of PepMV

98 populations in several European countries revealed that although the EU strain was presumed to
99 be dominant, the occurrence of other PepMV genotypes or mixed infections also became
100 frequently reported (Hanssen and Thomma, 2010). Even though the CH2 strain was evident in
101 Europe since 2006, mixed infections by both the EU and CH2 strains were also reported in tomato
102 crops (Hanssen et al., 2008; Hanssen and Thomma, 2010). Then a shift in PepMV populations was
103 gradually observed all over Europe, as the EU strain was likely overtaken by the CH2 one (Hasiów-
104 Jaroszevska et al., 2009; Hanssen and Thomma, 2010). This probably originated from a biological
105 advantage of CH2 strain over the EU, due to CH2 ability to spread more rapidly (Hanssen et al.,
106 2008; Hanssen and Thomma, 2010).

107
108 Typically, PepMV causes a wide range of symptoms, with fruit marbling being the most common
109 and economically important symptom (Spence et al., 2006). In addition, fruit discoloration, open
110 fruit, nettle-heads, leaf blistering or bubbling, leaf chlorosis and yellow angular leaf spots, leaf
111 mosaic and leaf or stem necrosis have been also associated with PepMV infections (Hanssen and
112 Thomma, 2010). The severity of PepMV symptoms likely depends on the environmental
113 conditions, as well as on the genetic features of the viral strains. Even minor nucleotide sequence
114 differences between isolates of the same genotype have been associated with enhanced
115 pathogenicity and severe symptoms (Hanssen and Thomma, 2010).

116
117 To cope with numerous pathogens, plants have developed a compendium of well-tuned
118 surveillance systems, including a sophisticated genetically controlled defense mechanism.
119 Successful defense relies on the well-timed and rigorous detection of the invading pathogen and
120 the subsequent induction of response pathways to avert pathogen attack. Transcriptional
121 reprogramming is a major part of plant defense response modulating a myriad of complex
122 molecular, biochemical, and physiological changes that occur in a highly synchronized manner
123 largely governed by transcription factors (TFs) (Moore et al., 2011). TFs are master regulators of
124 gene expression controlling a plethora of biological processes including defense and stress
125 responses (Tsuda and Somssich, 2015; Sukumari Nath et al., 2019). Previous studies have
126 identified several TFs implicated in regulating the expression of plant defense responsive genes
127 (Moore et al., 2011; Tsuda and Somssich, 2015; Sukumari Nath et al., 2019). While significant
128 progress has been made in terms of high-throughput analysis of tomato gene expression post

129 PepMV infection (Hanssen et al., 2011), the role of TFs that control the host's response remains
130 largely unknown.

131
132 Currently, tomato intensive cultivation, together with climate instability, promotes the outbreak
133 and dominance of biotic threatening agents. The most successful management strategy against
134 viral pathogens is through the application of preventive methods, given that the introduction of
135 genetic resistance in the plant host is time consuming, and cross-protection can be effective only
136 under well-defined and controlled conditions. Moreover, the exact defensive mechanisms of the
137 plant to overcome the disease are not completely understood. Thus, it is urgent to comprehend the
138 molecular networks controlling tomato responses against viral infections. The strain of PepMV,
139 alongside the environmental conditions, are the two most significant parameters affecting the
140 severity of disease symptoms. Herein, we report the transcriptome response of tomatoes against
141 PepMV infection by applying an RNA-seq approach to seek for differentially expressed
142 transcription factors (DETFs). This led to the identification of a subset of tomato highly
143 differentiated DETFs. Phylogenetic, structural and functional analysis of the DETFs showed that
144 they are highly conserved among vascular plants. Moreover, the highly differentiated DETFs
145 contained defense related elements in their regulatory regions. Our analysis reveals a subset of
146 tomato TFs with altered expression upon PepMV infection and broadens the practical implications
147 towards breeding of virus resistant crops.

148 MATERIALS AND METHODS

149 Plant material and growth conditions

150 Tomato (*Solanum lycopersicum*) Belladonna F1 hybrid seeds were planted in plastic pots
151 containing Potgrond P, a natural turf mixture (Klasmann-Deilmann GmbH, Geeste, Germany), and
152 grown at 22 °C in a Fitotron (Weiss Gallenkamp, Loughborough, UK) growth chamber under a
153 long day photoperiod with 16 h of light and 8 h of darkness per day and 100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ light
154 intensity.

155

156 Virus isolate and mechanical inoculation of tomatoes for transcriptome analysis

157 The TomCr3 isolate of PepMV was obtained from naturally infected tomato plants in 2019 grown
158 in a greenhouse in Ierapetra area of Crete island in Southern Greece (35.0119° N, 25.7407° E).
159 The virus was preserved in the laboratory through serial mechanical inoculations of tomato plants
160 (*Solanum lycopersicum L.*, hybrid *Belladonna*). The presence of PepMV was verified by enzyme-
161 linked immunosorbent assay (ELISA) with an antibody prepared by Agdia, Inc., and by reverse
162 transcription polymerase chain reaction (RT-PCR) as previously described (Ling et al., 2008).
163 Mechanical inoculations of tomato plants were performed by rub inoculation onto young leaves at
164 the fourth leaf stage. PepMV inoculum was prepared by grinding 0.5 g of TomCr3 infected leaves
165 in 5 mL of mechanical inoculation buffer (0.1M ($\text{Na}_2\text{HPO}_4/\text{KH}_2\text{PO}_4$) pH 7.4, 0.2% w/v Na_2SO_3).
166 The leaf extract was centrifuged at 1,000 x g for 5 s, 1.5 ml of the supernatant was transferred to
167 an Eppendorf tube which contained 0.2 g Silicon Carbide - 400 mesh (Merck KGaA) and was
168 vortexed thoroughly. A volume of 25 μL PepMV inoculum was placed onto the second leaf of
169 each plant (the inoculum was vortexed every 3 plants) and was rubbed with sterile hydrophobic
170 cotton. After 3 min the leaf was rinsed with tap water. Mock inoculations were performed with the
171 same procedure by using healthy tomato leaves for the inoculum and were used as a control. When
172 initial symptoms of PepMV infection became visible on the inoculated tomato plants, 10 days post
173 inoculation (dpi). Exactly 10 days post inoculation (dpi), approximately 0.1 g of tissue was cut
174 from the upper (non-inoculated) two or three leaves of each plant. The material was immediately
175 frozen in liquid nitrogen and stored at -80°C until further processing.

176

177 RNA Extraction, RNA Sequencing and Transcriptome Analysis

178 RNA preparation and RNA-seq analysis were performed as previously described (Daras et al.,
179 2019). Briefly, total RNA was extracted using the Direct-zol RNA Miniprep kit (Zymo research,
180 Irvine, CA, USA) with an on-column DNase treatment according to the manufacturer's
181 instructions from control and PepMV infected tomato tissues. The quantity and quality of RNA
182 were assessed using a NanoDrop ND-1000 spectrophotometer (Thermo Fischer Scientific,
183 Waltham, MA, USA) and agarose gel electrophoresis.

184

185 RNA-seq libraries were generated using the TruSeq Low Input kit according to the manufacturer's
186 instructions (Illumina). Sequencing was performed on BGISEQ-500 platform instrument at BGI
187 (Beijing Genomics Institute, Shenzhen, China). Raw reads were filtered into clean reads and
188 aligned to the tomato genome (GCF_000188115.3_SL2.50_ncbi_20180905). RNA-seq data were
189 analyzed using the SOAPnuke (version v.1.5.2) with parameters “-n 0.001 -l 20 -q 0.4 -A 0.25”
190 and the HISAT2 pipeline (version 2.0.4) with parameters “--phred33 --sensitive --no-discordant
191 --no-mixed -I 1 -X 1000”. Approximately 130 million reads were obtained for the control or
192 PepMV infected samples. On average, about 95% of the clean reads could be unambiguously
193 aligned to the SL2.50 reference genome sequence. For the detection of differentially expressed
194 genes (DEGs), clean reads were mapped to the reference genome with Bowtie2, and then the gene
195 expression level was calculated with RSEM with default parameters. Statistical analysis of
196 differential gene expression was conducted utilizing Poisson Distribution (Audic and Claverie,
197 1997) with fold change parameters greater than 2 and an FDR value ≤ 0.001 . Heatmaps of the
198 DEGs were constructed with Perseus software (version 1.6.8.0) using Euclidean distancing with
199 average linkage and without any constraints in the algorithm.

200

201 **Reverse-Transcription qPCR Analysis**

202 Reverse transcription (RT) was performed on 1 μg of total DNA-free RNA using Invitrogen
203 SuperScript IV VILO Master Mix (Thermo Fisher scientific, Waltham, MA, USA). Quantitative
204 gene expression analysis was performed using gene specific primers (Table S1) with a PikoReal
205 96 Real-Time PCR System (Thermo Scientific, Waltham, MA, USA) using the SYBR Select
206 Master Mix (Applied Biosystems, Waltham, MA, USA). For RT-qPCR analysis, *Glyceraldehyde-*
207 *3-phosphate dehydrogenase (GAPDH)* was chosen as a housekeeping gene and the control plants
208 were used as internal calibrators (Tsitsekian et al., 2021). Quantification of gene expression was

209 calculated as the expression of the gene of interest relative to *GAPDH* expression based on $2^{-\Delta Ct}$
210 method as previously described (Tsitsekian et al., 2021). The fold change was calculated by the $2^{-\Delta\Delta Ct}$
211 method as previously described (Ioannidi et al., 2016).

212

213 **Homologs identification, sequence alignment and phylogenetic analysis**

214 The analysis was performed as previously described (Tsitsekian et al., 2019), with slight
215 modifications. To identify and retrieve the tomato DETFs homolog genes of *Arabidopsis* and rice,
216 BLASTP searches were conducted using the tomato DETF protein sequences as the query search
217 in the publicly available Phytozome database v.13 (<https://phytozome-next.jgi.doe.gov/>,
218 Accessed: May 2021). Multiple sequence alignment of the identified amino acid sequences were
219 performed by the ClustalX 1.83 software. The results were exported as GCG/MSF format and
220 analyzed using the GeneDoc MFC Application version 2.6.0.2. The visualization of the
221 phylogenetic trees was made by the Interactive Tree of Life (<http://itol.embl.de>) online platform.

222

223 **Bioinformatics resources for *in silico* analysis**

224 Gene ontology (GO) and Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways
225 enrichment were performed using the DAVID tools database (Huang et al., 2009a; 2009b). For
226 protein conserved motifs analysis, the MEME-suite (Bailey et al., 2015) was used with default
227 settings. To identify the *cis*-elements within the promoter regions of the transcription factors the
228 PlantCARE database was applied (Lescot et al., 2002). The gene co-expression analysis of the
229 *Arabidopsis* homologs and the functional enrichment analysis of the resulting gene network were
230 performed by utilizing the STRING database (v.11.5; <https://string-db.org/>).

231

232 **Data availability**

233 The transcriptomic data of the tomato-PepMV pathosystem has been deposited in the Gene
234 Expression Omnibus database at the National Center for Biotechnology Information (NCBI) under
235 the accession number GSE203627.

236 RESULTS

237 **PepMV infection resulted in alterations of tomato gene expression**

238 An endemic PepMV isolate causing typical leaf mosaic symptoms, hereafter called TomCr3, was
239 obtained from greenhouse grown tomatoes (Fig. 1A). An experimental system was developed
240 inoculating tomato plants with the TomCr3 isolate and collecting leaf tissues for transcriptome
241 analysis 10 days post inoculation, when the symptoms became visible. Mechanical inoculations of
242 tomato hybrid line Belladonna F1 plants were performed by applying a TomCr3 extract onto the
243 first emerging leaf. To verify that the symptoms were caused by TomCr3 isolate, molecular
244 analysis of the *TGB2* and *TGB3* genes, known to be expressed by a single subgenomic PepMV
245 RNA segment, was performed (Fig. 1B). The PCR amplification of the reverse-transcribed viral
246 genomic RNA revealed the predominant expression of these two genes in the infected tissues.
247 Furthermore, multiple-sequence alignment of the *TGB2* and *TGB3* genes verified that TomCr3 is
248 a PepMV isolate (Figure S1).

249
250 To gain further insights into the molecular mechanisms controlling tomato defense response,
251 transcriptome analysis of PepMV infected and control plants was performed. RNA-sequencing
252 (RNA-seq) data analysis resulted in an average of 125M clean reads (Table S2A). Given that the
253 tomato genome encodes for 34,075 genes, the total genes identified by the analysis were
254 approximately 22,900 for the control and 22,500 for PepMV infected plants, which corresponded
255 to 31,300 and 30,200 transcripts, respectively (Table S2B). Overall, these quality metrics of the
256 RNA-seq approach demonstrated an efficient coverage of the tomato genome. The parameters of
257 false discovery rate (FDR) and \log_2 FC were set to $FDR < 0.001$ and $\log_2 FC \geq 1$ or ≤ -1 ,
258 respectively, to screen for differentially expressed genes (DEGs; Figure S2). These selection
259 criteria permit the identification of genes with a significant change of ≥ 2 fold in terms of expression
260 compared to the control. The analysis revealed 1,120 DEGs, which is equal to approximately 3.3%
261 of the tomato genes (Fig. 2A; Table S3A). This relatively small number of genes potentially
262 indicates a specific response of the tomato transcriptome against PepMV infection.

263
264 Based on the response pattern, the DEGs were categorized in two groups (Fig. 2 A,B). The first
265 group consisted of genes whose expression was upregulated by the virus infection and included
266 502 genes. The second group contained 618 genes that were downregulated upon PepMV

267 infection. Interestingly, these significant responsive genes highlighted a series of distinct
268 biological processes that were affected during the viral infection. The Gene Ontology (GO) and
269 Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways enrichment analyses that were
270 performed for the upregulated genes, revealed that photosynthesis, carbon fixation and post-
271 translational gene silencing (PTGS) were the most prominent enriched terms (Fig. 2C). On the
272 contrary, plant defense response, ABA signaling or response, carotenoid biosynthesis and the
273 activity of transcription factors were among the most highly enriched terms within the genes that
274 were downregulated (Fig. 2D). These results are consistent with the data of previous studies
275 relying on microarray analysis of PepMV infected tomato plants (Hanssen et al., 2011). Likewise,
276 photosynthesis and PTGS were reported to be particularly induced upon PepMV infection,
277 whereas the biosynthesis of carotenoids and the defense related responses were significantly
278 hampered. Taken together, our results are in agreement with previous reported data supporting the
279 accurate analysis of tomato transcriptome reprogramming upon PepMV infection.

280

281 **A subset of tomato TFs was substantially responsive against PepMV infection**

282 ~~As TFs are central regulatory hubs of the host transcriptome that respond against biotic stress~~
283 ~~factors, a deep analysis of the RNA-seq data was performed to comprehend the molecular networks~~
284 ~~that control tomato responses against PepMV infection. As TFs are central regulatory hubs of the~~
285 ~~host transcriptome responding against biotic stress factors, a deep analysis of the RNA-seq data~~
286 ~~was performed to identify the TFs that show differential expression and subsequently to~~
287 ~~comprehend the molecular networks that control tomato responses against PepMV infection.~~

288 Targeted analysis for TFs applying the initial cutoffs ($-1 \geq \log_2FC$ (PepMV/Control) ≥ 1 , FDR \leq
289 0.001) identified 84 differentially expressed TFs (DETFs), of which 38 were upregulated and 46
290 were downregulated upon PepMV infection (Fig. 3A; Figure S2; Table S3B). Given that the TFs
291 showing the highest alterations in terms of gene expression would be most likely involved in the
292 regulation of host's responses against the virus, a set of most strict selection criteria was applied.
293 These parameters excluded genes that showed a fold change of expression lower than 3 ($-1.59 \geq$
294 \log_2FC (PepMV/Control) ≥ 1.59) or a relatively low expression level (FPKM < 5), leading to the
295 identification of 11 highly responsive TFs (Fig. 3B; Figure S2). Notably, these TFs were
296 categorized in 7 distinct families that were further divided in almost two halves. The group of
297 upregulated TFs consisted of 5 members, whereas the rest of 6 TFs were downregulated, upon

298 PepMV infection (Fig. 3C, D). The *NAC* family was represented by three TFs, of which a pair was
299 upregulated, whereas *JA2L* was downregulated. The expression of two *bHLH* TFs, namely
300 *bHLH81* and *bHLH87*, was reduced. While *ZAT1-like*, a member of C2H2 TFs, was induced upon
301 PepMV infection, the expression level of another member, *ZF2*, was reduced. The remaining 4
302 DETFs were single members of the growth regulating factors (*GRF*), *B3*, ethylene responsive
303 factors (*ERF*) and *bZIP* families.

304
305 Given the great difference of these TFs expression upon PepMV infection compared to control
306 plants, a reverse-transcription quantitative polymerase chain reaction (RT-qPCR) method was
307 applied to validate their highly responsive pattern. Interestingly, RT-qPCR analysis supported the
308 classification of these TFs into a group of 5 upregulated genes that included two members,
309 *Solyc02g093420* and *GRF4*, that exhibited an almost 5-fold induction and into a second group of
310 6 downregulated genes including *JA2L* and *TAF-1-like* that showed the highest reduction of
311 expression (Fig. 4A). Moreover, the expression pattern of all 11 DETFs was almost identical
312 between both experimental approaches (Fig. 4B). As shown by the fold change of each gene
313 expression, the RT-qPCR analysis further confirmed the accuracy and repeatability of RNA-seq
314 data acquisition. Taken together, the application of stringent selection criteria led to the
315 identification of 11 highly responsive TFs in the tomato-PepMV pathosystem by distinct but
316 complementary gene expression analysis approaches, which most likely modulate critical host
317 responses against the viral pathogen infection.

318

319 **The evolutionary relationships of tomato DETFs in model plant species**

320 Rice (*Oryza sativa*) and *Arabidopsis thaliana* are representative model organisms of monocot and
321 dicot plant species, respectively. Their genome is often used as a reference to monitor the evolution
322 of homologous genes. Hence, a wide-genome BLAST query was performed to identify the highly
323 homolog counterparts of the responsive tomato TFs against PepMV infection (Table S4). To
324 understand the evolutionary history of the tomato TFs, a phylogenetic tree (Fig. 5 left panel) was
325 constructed and aligned with the exon-intron structure of the genes (Fig. 5 middle panel). In most
326 cases, the intron length of the tomato genes was relatively longer compared to rice and
327 *Arabidopsis*. Moreover, this complementary approach revealed that in terms of homology, the
328 tomato TFs were categorized into two subgroups. The first subgroup included 5 TFs that were best

329 aligned with the rice counterparts. These tomato TFs according to their tag numbers were [1], [2],
330 [3] and [4], which were induced by PepMV infection, and [6] whose expression was reduced. The
331 second subgroup consisted of 6 tomato TFs, which were highly homologous with the *Arabidopsis*
332 counterparts. Most of these were downregulated, namely [7], [8], [9], [10] and [11], whereas only
333 one [5] was stimulated upon the viral infection. Interestingly, the analysis revealed that the origin
334 of the first subgroup of tomato TFs is most likely defined before the evolutionary divergence
335 between the monocot and dicot species in angiosperms. These results are consistent with previous
336 phylogenetic analyses demonstrating that nearly half of the *Arabidopsis* and rice TFs can be
337 classified into clear orthologous groups supporting the notion that they originate from a common
338 ancestor of the monocot and dicot species (Xiong et al., 2005).

339
340 In addition to the phylogenetic classification, the TF homologs were further analyzed based on the
341 diversification of strongly conserved amino acid motifs within the members of each family (Table
342 S5). Strikingly, 4 out of the 7 families of TFs showed evident differences between the homolog
343 gene members within the family (Fig. 5 right panel). In the *ERF* family, the tomato *DREB3* TF,
344 like its *Arabidopsis* homolog, contained three conserved motifs. This is in contrast to the rice
345 counterpart, which had only two motifs. Surprisingly, some of the tomato TFs classified in the
346 remaining 3 families included members that showed higher structural homology to the rice
347 counterparts contrary to the other tomato members of the family that were more similar to the
348 dicotyledonous homologs of *Arabidopsis*. In particular, *GRF4*, belonging to the *GRF* family,
349 contained three distinct conserved motifs similar to its rice monocot homolog, but it deviated from
350 the dicot homolog of *Arabidopsis*, which bore just a pair of motifs. A somehow similar structural
351 representation was evident within the families of tomato *bHLH* and *NAC* TFs that included two
352 and three highly responsive TFs, respectively. The tomato *bHLH81*, which grouped best with its
353 rice homolog, had two motifs, whereas the tomato *bHLH87* contained one more rising to three
354 motifs like the *Arabidopsis* homolog. Likewise, two tomato TFs members of the *NAC* family, with
355 [2] and [4] as tag numbers, showed higher similarity to the rice monocot homologs and bore one
356 motif in the polypeptide sequence more than the tomato *JA2L* TF, which had two motifs similar to
357 the dicot *Arabidopsis* homologs. In line with the phylogenetic analysis, the divergence in terms of
358 the conserved structural motifs further supports that few of the tomato TFs derived from the
359 common ancestor that existed before the divergence between monocot and dicot plant species.

360

361 The tomato TFs have a potential enhanced response capacity to stress and defense

362 Regulatory elements are specific nucleotide motifs located at the promoter sequence that control
363 gene expression in a temporal, spatial, and/or cell type-specific manner. As the tomato TFs
364 responded to the viral infection, a comparison of their genomic regulatory regions, which cover
365 2000-bp upstream of the translation initiation site, was performed to understand the molecular
366 basis of these genes' expression. Remarkably, a number of *cis*-acting regulatory elements related
367 to stress or defense related hormone response were found (Fig. 6A; Table S6). The ABA and
368 abiotic stress responsive ABRE element were identified in the promoter regions of all highly
369 responsive tomato TFs. The SA-responsive (TCA-element) and JA-responsive elements (CGTCA-
370 motif and TGACG-motif) were also abundant in the promoter regions of 3 and 9 genes,
371 respectively. The ethylene responsive ERE motif or the wounding response WUN-motif were
372 present in the promoters of 8 genes. Several W-boxes were identified in the promoters of 7 genes,
373 indicating that these genes might be regulated by WRKY, whereas the TC-rich repeat defense and
374 stress response element was identified in 4 promoter regions. Taken together, these results
375 supported the notion that stress responsive *cis*-regulatory elements involved in defense or stress
376 responses are overrepresented in the promoters of the tomato highly responsive genes against
377 PepMV infection.

378

379 Since an adequate degree of gene homology was identified between tomato and *Arabidopsis* and
380 given that a plethora of experimental data available for *Arabidopsis*, a query for gene co-expression
381 networks was performed relying on the *Arabidopsis* homologs (Fig. 6B). This approach led to a
382 hub of transcriptional regulators *ZAT6*, *ZAT10* and *ZAT12*, comprised of the C2H2 zinc finger
383 gene family. The network also included two ribonuclease H-like superfamily proteins *CAF1a* and
384 *CAF1b*, the pathogen-induced transcription factors *WRKY33* and *WRKY40*, the Calcium-binding
385 EF-hand family protein *At4g27280* and *At4g29780* which encodes for a nuclease. Intriguingly,
386 based on the tomato transcriptome analysis, *ZAT1*-like the homolog of *Arabidopsis* *ZAT12* was
387 induced upon the infection of PepMV, whereas the expression of *ZF2* was downregulated. This
388 could possibly reflect different roles between *Arabidopsis* *ZAT12* and the *Arabidopsis* homologs
389 of tomato *ZF2*, namely *ZAT10* and *ZAT6*. Genetic and gene expression evidence in *Arabidopsis*,
390 support that the function of *ZAT10* may be linked to *ZAT12* during stress but possibly *ZAT10*

391 could be not only an activator, but also a transcriptional suppressor modulating the activation of
392 defense responses during different stresses, as well as stress combinations (Mittler, 2006).
393 Moreover, the *Arabidopsis* based approach highlighted the role of tomato TFs in defense and stress
394 response. Functional enrichment analysis of the gene co-expression network showed that the Gene
395 Ontology (GO) biological process enriched terms were associated with plant defense or with
396 response to stress factors (Fig. 6C). Concerning the Cellular Component (CC) terms analysis, two
397 processes tightly linked with gene expression regulation were overrepresented: the nucleus and the
398 P-bodies (processing bodies), which are cytoplasmic protein complexes involved in the
399 degradation and translational arrest of mRNA. In terms of Molecular Function (MF) term a sole
400 term was enriched related to the DNA-binding transcription factor activity. Overall, these results
401 support the notion that the responsive TFs most likely drive the reprogramming of tomato
402 transcriptome to cope with a viral attack.

403 DISCUSSION

404 ~~Tomato is a worldwide highly consumed vegetable and a major dietary source of mainly~~
405 ~~antioxidants, which have been linked to many health benefits. Tomato intensive cultivation,~~
406 ~~together with climate instability, promotes the outbreak and dominance of biotic threatening~~
407 ~~agents. Nowadays, viruses have become a critical threat to greenhouse grown tomato cultivars and~~
408 ~~thus it is considered extremely difficult to apply a successful crop protection protocol. PepMV has~~
409 ~~a severe impact on fresh market tomato production and is a major disease of greenhouse tomato~~
410 ~~crops globally. The strain of PepMV, alongside the environmental conditions, are the two most~~
411 ~~significant parameters affecting the severity of disease symptoms. Currently, the most successful~~
412 ~~management strategy against viral pathogens is through the application of preventive methods,~~
413 ~~given that the introduction of genetic resistance in the plant host is time consuming, and cross-~~
414 ~~protection can be effective only under well-defined and controlled conditions. Moreover, the exact~~
415 ~~defensive mechanisms of the plant to overcome the disease are not completely understood. Thus,~~
416 ~~it is urgent to comprehend the molecular networks controlling tomato responses against viral~~
417 ~~infections.~~

418

419 Tomato is a worldwide highly consumed vegetable and a major dietary source of mainly
420 antioxidants, which have been linked to many health benefits. Nowadays, viruses have become a
421 critical threat to greenhouse grown tomato cultivars and thus it is considered extremely difficult to
422 apply a successful crop protection protocol. PepMV has a severe impact on fresh-market tomato
423 production and is a major disease of greenhouse tomato crops globally. Upon a pathogen attack,
424 the plant host responds through reprogramming its transcriptome to cope with the infection. TFs
425 are master regulators of host molecular responses against various pathogens. While tomato
426 transcriptomic responses have been previously reported upon PepMV infection by applying
427 microarrays (Hanssen et al., 2011), there is still lack of knowledge about key TFs that could be
428 involved in host response against the virus. To fill this gap, NGS-based RNA-Seq was applied as
429 a hyper-analytical complementary approach to identify TFs, which generally are expressed in low
430 levels. The experimental design relied on the efficient implementation of an infection system of
431 tomato plants by an endemic isolate of PepMV, producing reproducible results and leading to a
432 coordinated sequence of symptom emergence.

433

434 Our transcriptomic analysis revealed that the expression of about 3.3% of genes composing the
435 tomato genome was modified upon virus infection, corresponding to 1,120 DEGs. Classification
436 of these genes in functional categories revealed that 502 genes, which showed stimulation of gene
437 expression, were mainly categorized in biological processes associated with photosynthesis,
438 carbon fixation and post-transcriptional gene silencing (PTGS). However, the most highly
439 enriched terms within the 618 downregulated genes were related to plant defense response, ABA
440 signaling or response, carotenoid biosynthesis and the activity of transcription factors.
441 Interestingly, these results are in conjunction with the data of a previous tomato GeneChip array
442 analysis (Hanssen et al., 2011). In particular, photosynthesis, defense response, RNA silencing and
443 carotenoid biosynthesis, likely responsible for the marbled leaf phenotype which is typical for a
444 mosaic disease, were also among the biological processes that were previously reported. This tight
445 correlation between both technical approaches demonstrates the core tomato responses against
446 PepMV infection and opens the way to elucidate the network of TFs being responsive upon the
447 viral attack.

448
449 In total 84 TFs were responsive against TomCr3 infection, showing either a two-fold induction
450 (38 TFs) or repression (46 TFs) in terms of gene expression. For the sake of a better overview, the
451 focus of the analysis was put to distinguish TFs that were highly responsive. By setting strict
452 selection criteria, the list of TFs with a fold change of transcriptional response greater than three
453 ended to 5 upregulated and 6 downregulated TFs classified in 7 families. The response pattern of
454 these TFs was also confirmed by RT-qPCR analysis. A phylogenetic classification distinguished
455 5 tomato TFs that were highly homologous with the rice counterparts, whereas the rest of 6 tomato
456 TFs were closer to the *Arabidopsis* homologs, showing the formation of distinct orthologous pairs.
457 Interestingly, phylogenetic analysis of all *Arabidopsis* and rice TFs revealed that approximately
458 half of these are classified into orthologous groups, supporting the idea that orthologous TFs may
459 have conserved fundamental functions between monocots and dicots (Xiong et al., 2005).
460 Likewise, the sorting of tomato TFs may suggest that these regulators of gene expression upon a
461 viral attack possibly originated early in the evolutionary history of higher plants, before the
462 divergence between monocot and dicot species in angiosperms that likely occurred 150 million of
463 years ago (Pires and Dolan, 2012).

464

465 Promoter analysis of the highly responsive TFs revealed that all contained defense and stress
466 response regulatory elements within their promoter sequences. Recent reports support the
467 functional involvement of these tomato TFs in defense or abiotic stress responses. In our analysis
468 the expression of *Jasmonic Acid 2-Like (JA2L)* was suppressed upon PepMV infection. *JA2L*
469 together with its homolog *JA2 (Jasmonic Acid 2)* belong to the NAC family of TFs. There are
470 reports demonstrating that both these NAC TFs are involved in the pathogen-triggered stomatal
471 movement upon tomato pathogen attack (Du et al., 2014). Upon perception of pathogen infection,
472 plants close stomata, a response, which is referred to as “stomatal defense” and is an integral part
473 of plant innate immunity (Melotto et al., 2006; Du et al., 2014). Likewise, viral infections are
474 strongly associated with a reduction in stomatal density and stomatal closure, thus appear to
475 negatively affect transpiration most likely to reduce viral movement around the host (Murray et
476 al., 2016; Manacorda et al., 2021).

477
478 Many pathogens use effectors to counteract stomatal defense, reopen plant stomata and facilitate
479 their entry into the plant leaf (Melotto et al., 2006; Du et al., 2014). Coronatine (COR) is a toxin
480 produced by a bacterial pathogen, *Pseudomonas syringae* pv tomato DC3000 (DC3000), and is
481 essential for the virulent strain for stomatal reopening and to overcome apoplastic defenses,
482 including the SA-mediated defense (Melotto et al., 2006). COR is a structural and functional mimic
483 of Jasmonic Acid (JA) and has been reported that COR hijacks the JA signaling pathway to
484 suppress plant defenses, including stomatal reopening and apoplastic defenses, through a hormonal
485 crosstalk (Du et al., 2014). Consequently, *JA2* acts in the ABA-mediated stomatal closure, whereas
486 *JA2L* functions in COR/JA-mediated stomatal reopening process (Du et al., 2014). On the basis of
487 these observations, it appears reasonable to assume that downregulation of *JA2L* expression in
488 virus infected tomato plants is most likely attributed to prevent stomatal reopening. Despite an
489 effect on stomatal density, there is evidence that plant infection by viruses may regulate stomatal
490 conductance most likely due to reduction in stomata aperture (Chaerle et al., 2006; Aguilar et al.,
491 2017; Manacorda et al., 2021). Consequently, tomato potentially responds by a *JA2L*-mediated
492 stomata closure mechanism, albeit this could be an indirect effect driven by disturbed hormonal
493 homeostasis. Nevertheless, further experiments are required to clarify whether stomatal closure is
494 a bona fide defense response by the plant against viruses or is a beneficial to the virus
495 reaction. Therefore, tomato potentially applies a defensive response against PepMV infection by a

496 ~~*JA2L*-mediated stomata closure mechanism to slow down transpiration, increase water use~~
497 ~~efficiency and reduce viral movement.~~

498
499 Despite *JA2L*, the expression of *bHLH81* and *DREB3A* TFs was also substantially reduced in
500 PepMV infected tomato plants. Intriguingly, the *Arabidopsis* homologs of both tomato *bHLH81*
501 and *DREB3A* genes responded differently upon a bacterial infection. The expression pattern of
502 *FBH3*, an *Arabidopsis* homolog of tomato *bHLH81*, was induced upon DC3000 infection.
503 Furthermore, *FBH3* subsequently regulated the expression of a group of downstream TFs that were
504 also induced upon bacterial infection (Lewis et al., 2015). *TINY2*, the functional homolog of
505 *DREB3A* in *Arabidopsis*, has been reported to be rapidly induced upon abiotic challenges such as
506 cold, drought, mechanical wounding and salinity (Wei et al., 2005). In addition, *TINY2* was also
507 induced upon ABA treatment. The opposite expression response pattern of *bHLH81* and *DREB3A*
508 potentially highlights distinct plant defensive strategies against viral and bacterial plant pathogens
509 or environmental cues.

510
511 An *Arabidopsis*-based approach revealed a co-expression interaction network among tomato TFs
512 of the *C2H2* zinc finger gene family. The tomato *ZAT1-like*, homolog of *Arabidopsis* *ZAT12*, was
513 induced upon PepMV infection, whereas the expression of tomato *ZF2*, which is homologous to
514 *Arabidopsis* *ZAT10* and *ZAT6* TFs, was downregulated. Remarkably, the *in silico* analysis revealed
515 that the *Arabidopsis* homologs of the *C2H2* zinc finger gene family, namely *ZAT10* (*At1g27730*),
516 *ZAT6* (*At5g04340*) and *ZAT12* (*At5g59820*), formed a single hub of gene expression regulators.
517 Nevertheless, these plant homologs have been previously documented to possess potential distinct
518 roles in plant defense or abiotic stresses. A similar expression pattern was observed, when *ZAT10*
519 and *ZAT12* were induced upon abiotic stresses in *Arabidopsis*, suggesting that both TFs could
520 share common functions in stress response (Mittler et al., 2006). However, characterization of
521 *ZAT10* and *ZAT12* transgenic lines of *Arabidopsis* subjected to abiotic stresses revealed an
522 interesting functional response of mainly *ZAT10*. Plants overexpressing *ZAT12* showed a tolerant
523 phenotype upon osmotic stress, contrary to *ZAT12* knockout lines, supporting the positive role of
524 *ZAT12* against stress conditions (Davletova et al., 2005). Likewise, overexpression of the *Brassica*
525 *carinata* *ZAT12* homolog (*BcZAT12*) in transgenic tomato plants conferred drought resistance and
526 suppression of the oxidative stress caused by heat-shock treatment (Shah et al., 2013; Krishna et

527 [al., 2021](#)). In striking contrast, *ZAT10* overexpression or knockout led to tolerant transgenic plants
528 against mainly abiotic stresses compared to the wild-type ([Mittler et al., 2006](#)). These observations
529 jointly suggest that while *Arabidopsis ZAT12* acts as a transcriptional activator, *Arabidopsis*
530 *ZAT10* can either act as a positive or a negative regulator of the defense related genes ([Mittler et](#)
531 [al., 2006](#)). Our results are consistent with a distinct response pattern among the *C2H2* TFs. As
532 *ZAT1-like* was induced, *ZF2* expression was downregulated against PepMV infection to coordinate
533 tomato molecular defensive responses.

534

535 Overall, we shed light on tomato transcriptome reprogramming upon viral infection of a PepMV
536 isolate through the application of an RNA-seq analysis approach. The analysis provided
537 complementary evidence about typical plant responses against a viral mosaic disease including
538 photosynthesis, carbon fixation, gene silencing, plant defense and carotenoid biosynthesis.
539 Moreover, a subset of highly responsive transcription factors was identified, which based on
540 previous research observations, constitute a group of gene transcriptional regulators with
541 functional features mainly related to abiotic challenges. The data particularly support a regulatory
542 network of gene expression composed of the *Arabidopsis ZAT10* and *ZAT12* homologs that show
543 distinct patterns of response upon viral infection. This work provides a platform for future studies
544 on plant viral infections opening the perspective to comprehend host responses related to the
545 mobilizing of molecular defense mechanisms.

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553 AUTHOR CONTRIBUTIONS

554 SR and GD conceived the research and designed the experiments. DTs. performed most of the
555 experiments and analyzed the data. DTem. helped in bioinformatics analysis together with FA.
556 LL, CO and VM performed PepMV isolation and plant inoculations. DTs prepared the initial draft.
557 SR and VM supervised the experiments and acquired funding. SR, VM, GD, LL and VN revised
558 the manuscript. All authors have read and approved the manuscript.

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717 **FIGURE LEGENDS**

718 **Figure 1.** A PepMV isolate caused highly virulent infections in tomato plants (A) Typical mosaic
719 symptoms of tomato leaves infected with TomCr3, the PepMV isolate (Scale bar = 1cm). (B)
720 Expression of PepMV *TGB2* and *TGB3* genes by Reverse-Transcription semiquantitative and
721 quantitative PCR (RT-qPCR) analysis. The expression of the viral genes was evident only in the
722 infected tomato tissues.

723
724 **Figure 2.** Overview of differentially expressed genes (DEGs) upon PepMV infection. (A) Scatter
725 plot showing differential gene expression between control and PepMV infected plants. The axes
726 represent \log_2 transformed RNA-seq gene FPKM values. Red and blue dots denote genes whose
727 expression was significantly up- or downregulated, respectively, upon PepMV infection. Grey dots
728 denote genes whose expression was unchanged. (B) Venn diagram showing the number of genes
729 that were significantly upregulated or downregulated in response to PepMV infection. (C-D)
730 Functional enrichment analysis of the upregulated (C) and downregulated (D) genes upon PepMV
731 infection. The enriched Gene Ontology (GO) terms of Biological Process (BP), Cellular
732 Component (CC) and Molecular Function (MF) are shown in blue, orange and green letters,
733 respectively, whereas the overrepresented KEGG pathways are shown in red. *p*-value is calculated
734 by Fisher's exact test, whereas the count value refers to the number of genes observed within each
735 term.

736
737 **Figure 3.** Differentially expressed transcription factors (DETFs) identified upon tomato infection
738 by PepMV. (A) Volcano plot showing the distribution of the 84 DETFs (2-fold change or greater
739 and $FDR \leq 0.001$). (B) Scatter plot of the highly responsive DETFs. The axes represent \log_2
740 transformed RNA-seq gene Fragments Per Kilobase of transcript per Million mapped reads
741 (FPKM) values. Red and blue dots denote genes whose expression was significantly up- or
742 downregulated, respectively (fold change greater than 3, $FDR \leq 0.001$ and $FPKM \geq 5$). Grey dots
743 denote genes whose expression was not modified. Genes within the dark grey box show low levels
744 of expression ($FPKM < 5$). (C and D) Identification data (C) and normalized gene expression
745 profiles (D) of the highly responsive tomato TFs upon PepMV infection.

746

747 **Figure 4.** Comparative expression analysis of the highly differentiated tomato TFs upon PepMV
748 infection. (A) Reverse-Transcription quantitative PCR (RT-qPCR) validation. Values are mean \pm
749 SD of four ($n = 4$) replicates derived by RT-qPCR analysis. Asterisks indicate significant
750 differences (t -test) between the PepMV infected and control plants ($P \leq 0.05$) (B) Comparison of
751 gene transcription results derived from RT-qPCR and RNA-seq transcriptome analysis. Magenta
752 bars represent the fold change of gene expression of the PepMV infected plants relative to the
753 control, calculated as the \log_2 transformed ratio of the FPKM values obtained by RNA-seq
754 analysis. Green bars represent the \log_2 transformed fold change expression values obtained by RT-
755 qPCR analysis. Positive values correlate with upregulated gene expression, whereas negative
756 values with downregulation of gene expression.

757
758 **Figure 5.** Phylogenetic classification and structural features comparative analysis of the highly
759 responsive tomato TFs compared with their homologs of rice and *Arabidopsis*. (Left panel)
760 Phylogenetic categorization of the tomato TFs relative to the highly similar monocot or dicot
761 counterpart. Each TF family has been marked with a color box. The tree distance is indicated on
762 top of every node. (Middle panel) Gene exon-intron structure of the tomato TFs and their monocot
763 or dicot homologs. The colored numbers in brackets indicate the tag number of each tomato DETF
764 as annotated in Fig. 3. The fonts in red and blue color present the tomato DETFs that are up- or
765 down-regulated, respectively, upon PepMV infection. The slash-separated numbers in parentheses
766 represent the exon/intron numbers of each TF homolog. (Right panel) Distribution of conserved
767 amino acid motifs within the tomato TFs and their homologs per family. Colored boxes indicate
768 distinct signature motifs shown in Table S5 that were identified within the members of each TF
769 family.

770
771 **Figure 6.** Evidence of the stress and defense response capacity of the tomato TFs. (A) Graphical
772 representation of stress or defense response *cis*-regulatory elements within the promoter regions
773 of the most differentiated tomato TFs upon PepMV infection. (B) An *Arabidopsis*-based approach
774 revealed a co-expression interaction network among tomato TF homologs. Purple circles indicate
775 the input *Arabidopsis* genes, whereas the grey circles depict the interactors identified by the
776 analysis. The gene IDs colored in red depict the tomato upregulated TFs and in blue the tomato
777 downregulated TFs. (C) Functional enrichment analysis of the *Arabidopsis* co-expression network

778 of TFs. The enriched Gene Ontology (GO) terms of Biological Process (BP), Cellular Component
779 (CC) and Molecular Function (MF) are shown in blue, orange and green letters respectively. The
780 count value refers to the number of genes observed within each term.

781

782

783 SUPPORTING INFORMATION

784 **Table S1.** Oligos used in this study for gene expression analysis

785

786 **Table S2.** RNA-sequencing data quality metrics

787

788 **Table S3.** Expression data of the differentially expressed genes (DEGs) identified by RNA-seq
789 analysis

790

791 **Table S4.** The rice (*Oryza sativa*) and *Arabidopsis thaliana* homologs of the highly responsive
792 tomato TFs against PepMV infection

793

794 **Table S5.** Identification of distinct signature motifs within the members of each TF family

795

796 **Table S6.** The *cis*-elements identified within the promoter regions of the 11 highly differentiated
797 DETFs

798

799 **Figure S1.** Multiple-sequence alignment of the *TGB2* and *TBG3* genes verified that TomCr3 is a
800 virulent isolate of PepMV. The TomCr3 sequence of 365 nucleotides was aligned with the
801 indicated five PepMV genotypes CH2, LP-2001, EU, US1 and PES with GenBank accession
802 numbers DQ000985, AJ606361, KJ018164, AY509926 and HG313806, respectively.

803

804 **Figure S2.** Outline of the experimental workflow leading to the identification of differentially
805 expressed transcription factors (DETFs) upon PepMV infection of tomato.

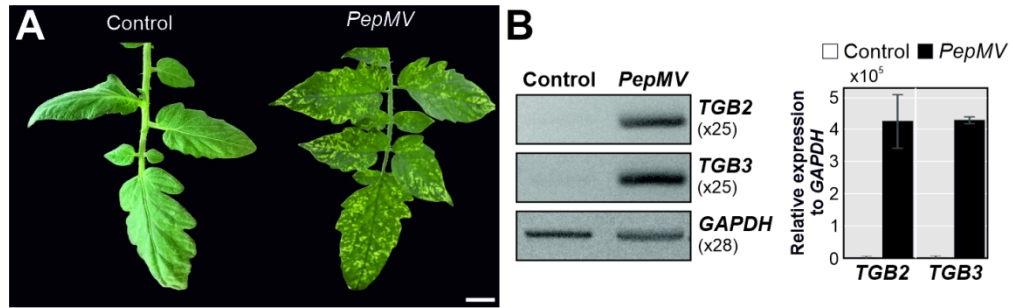
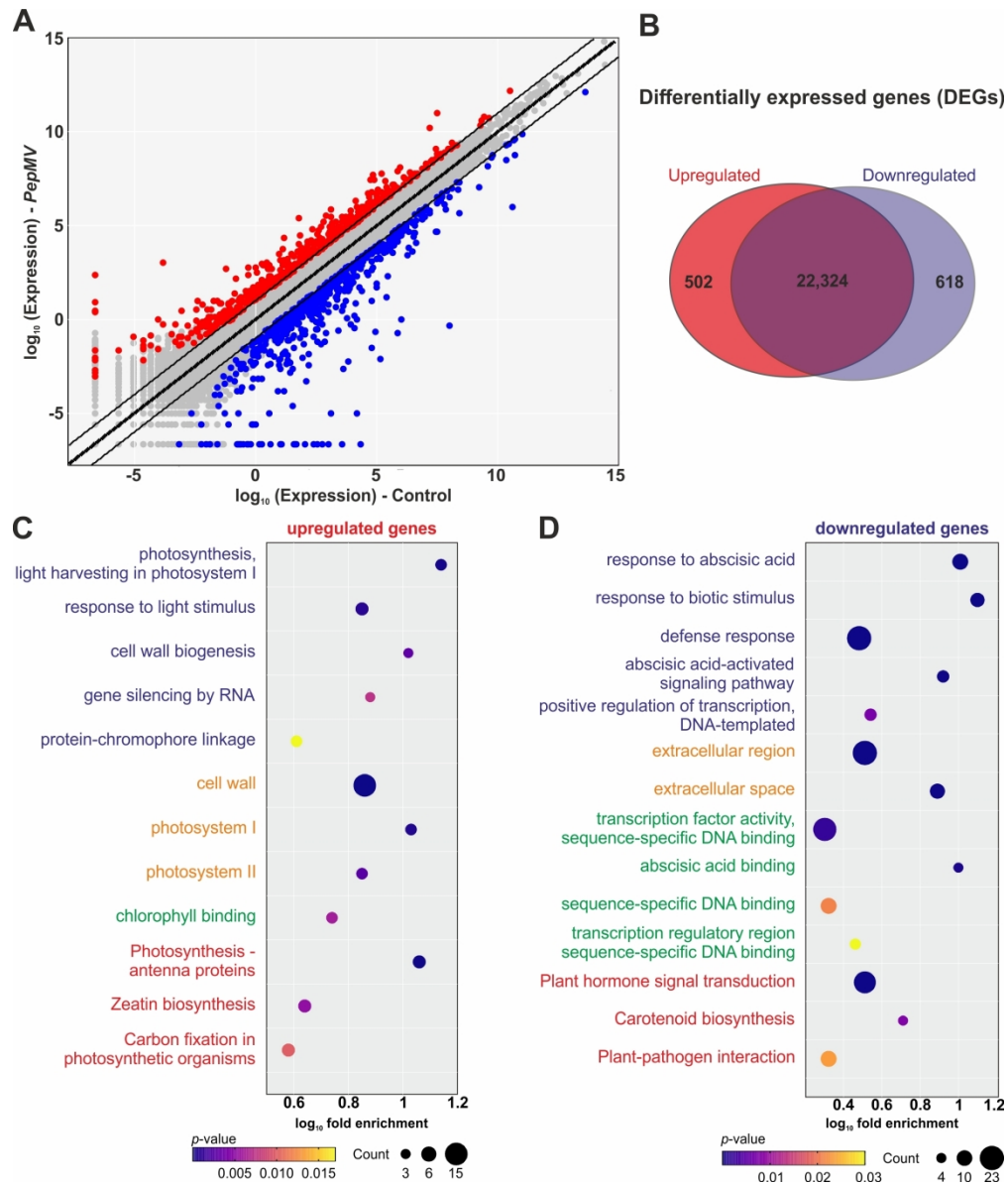


Figure 1

119x36mm (300 x 300 DPI)



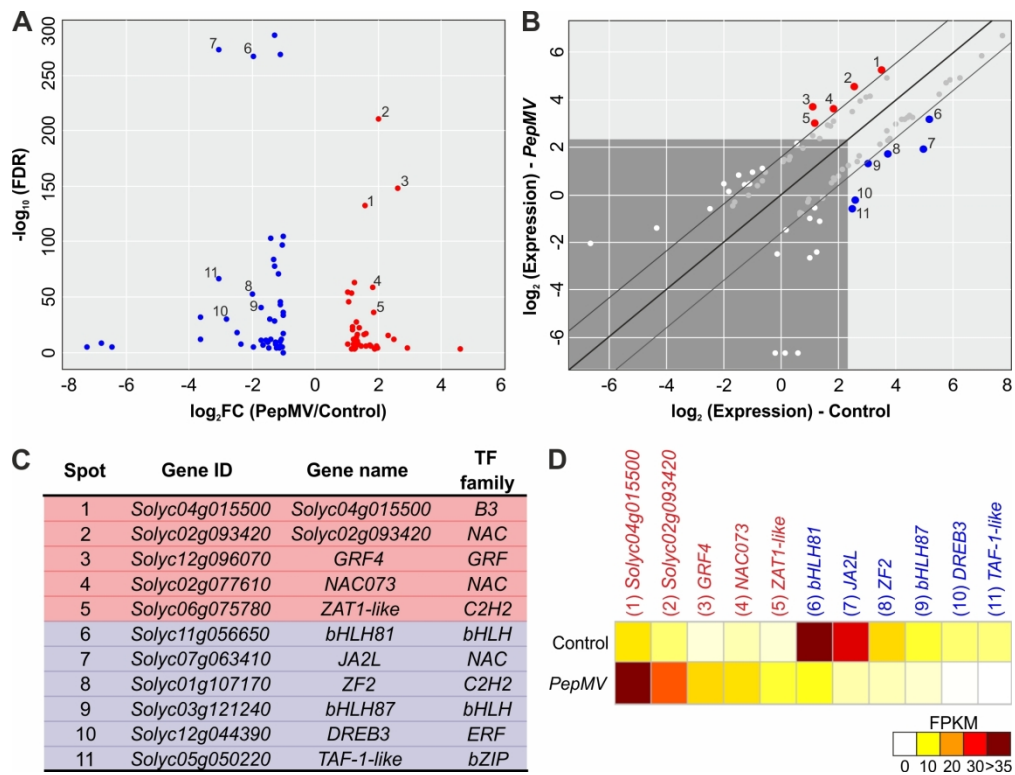


Figure 3

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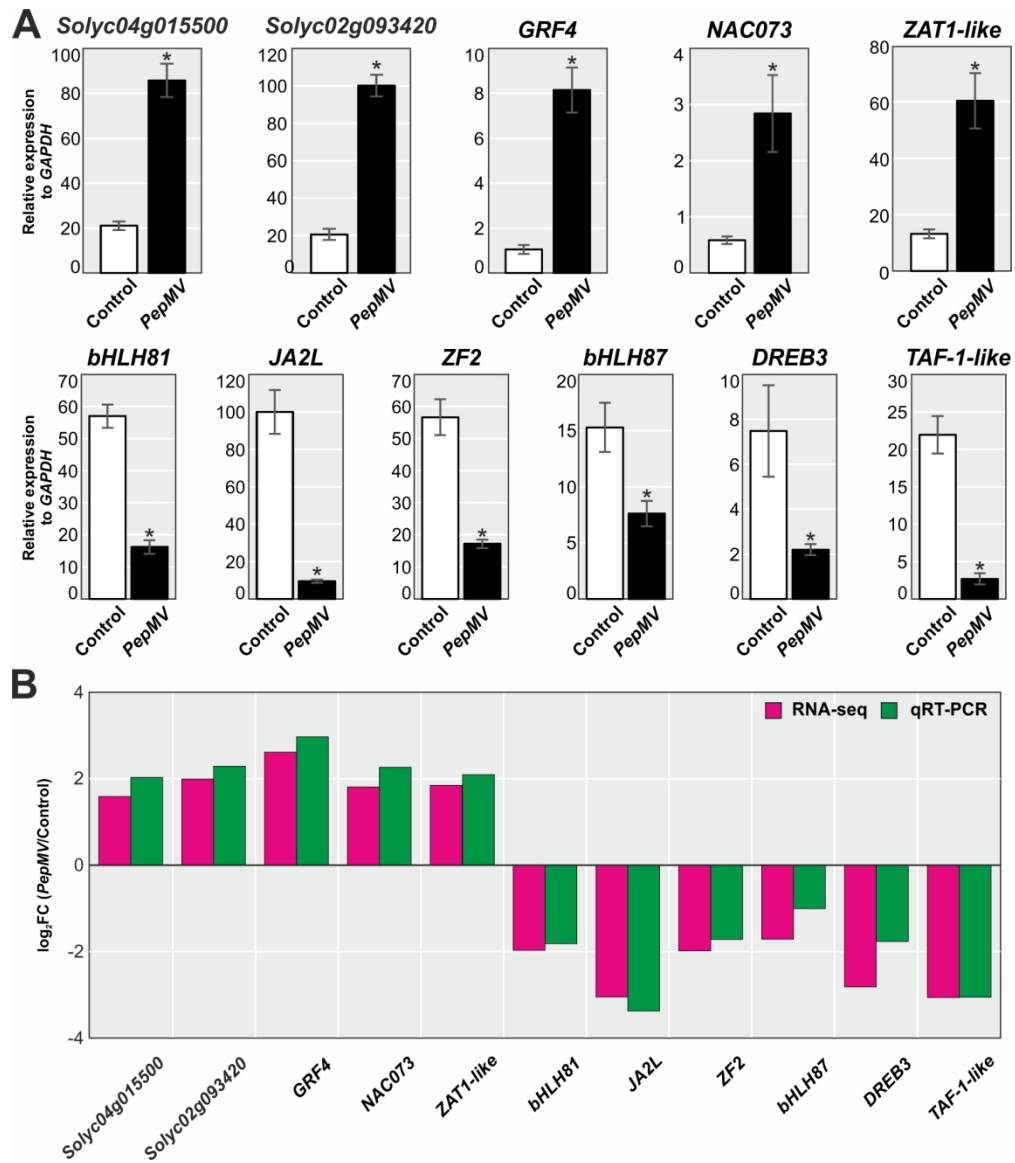


Figure 4

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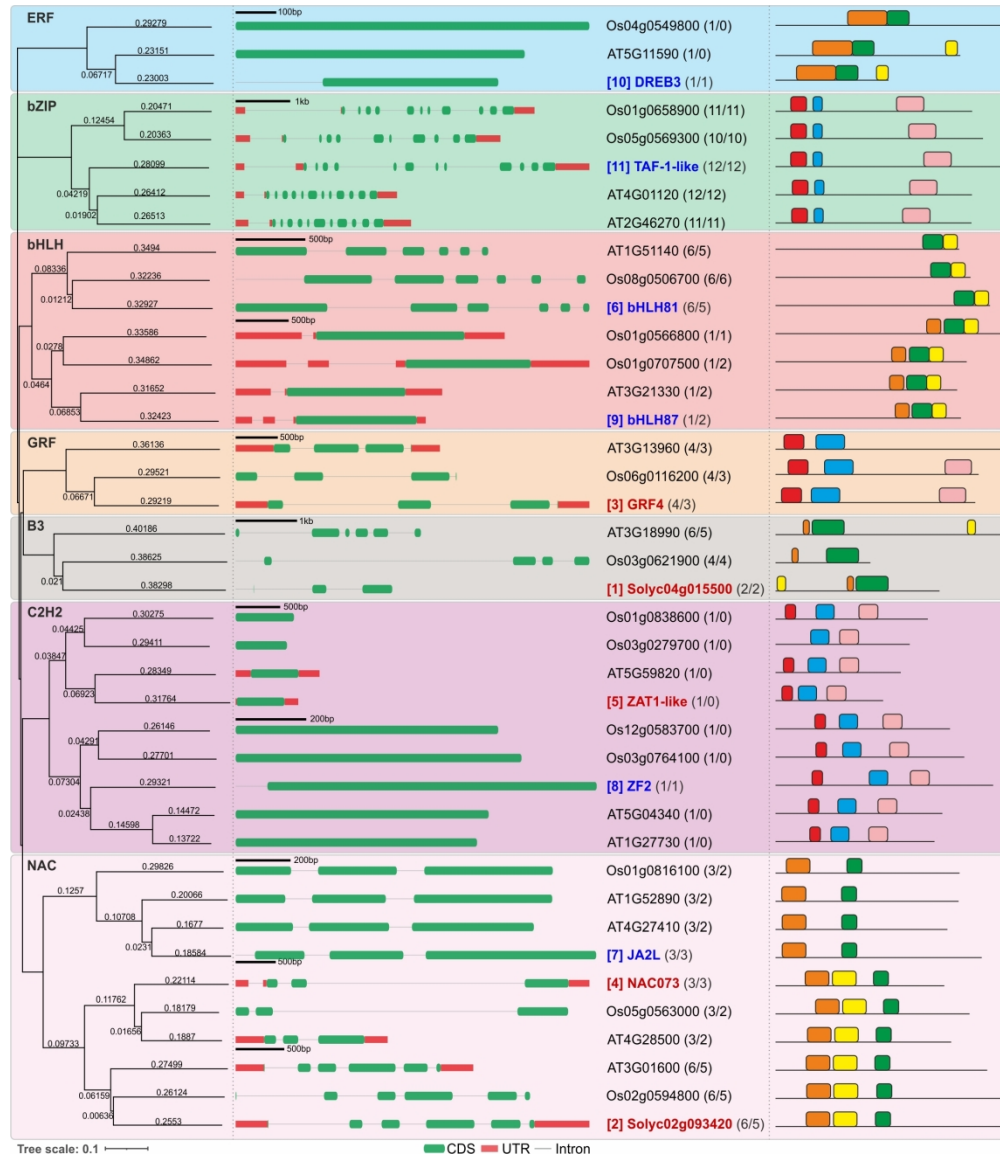


Figure 5

207x237mm (300 x 300 DPI)

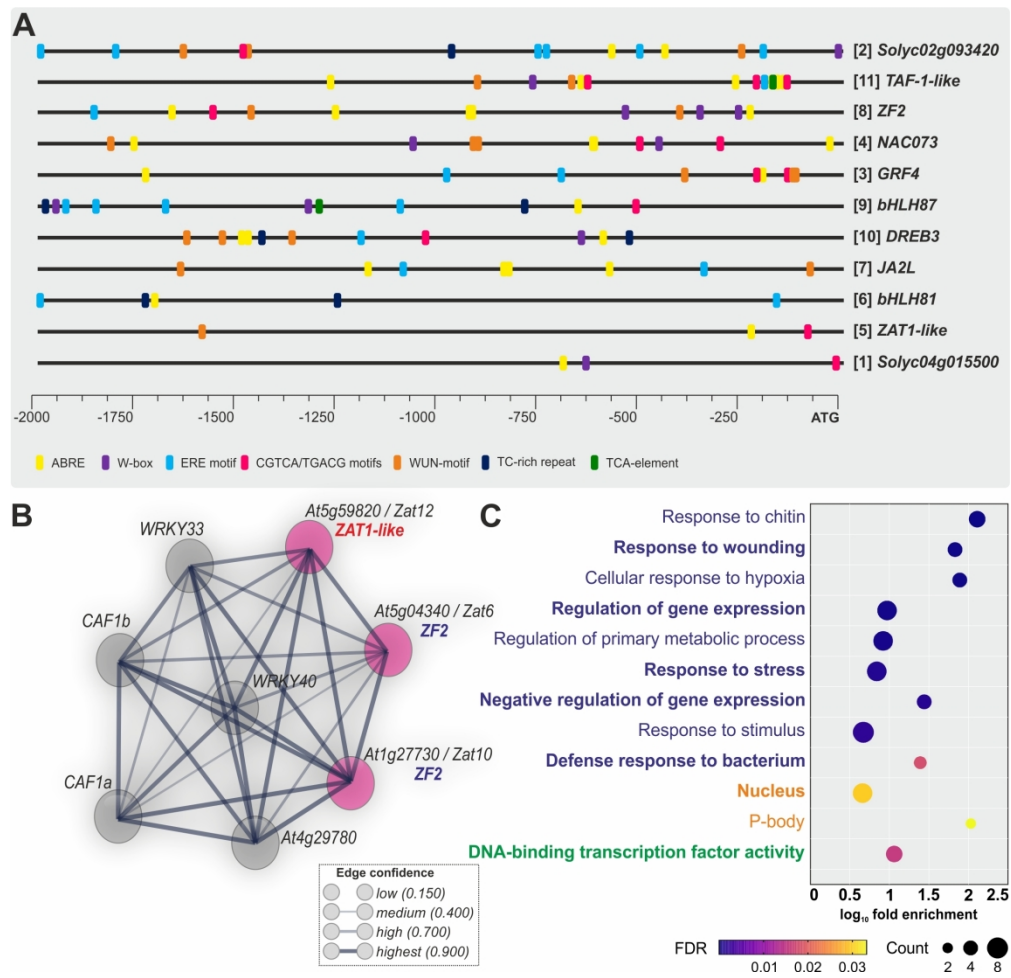


Figure S1

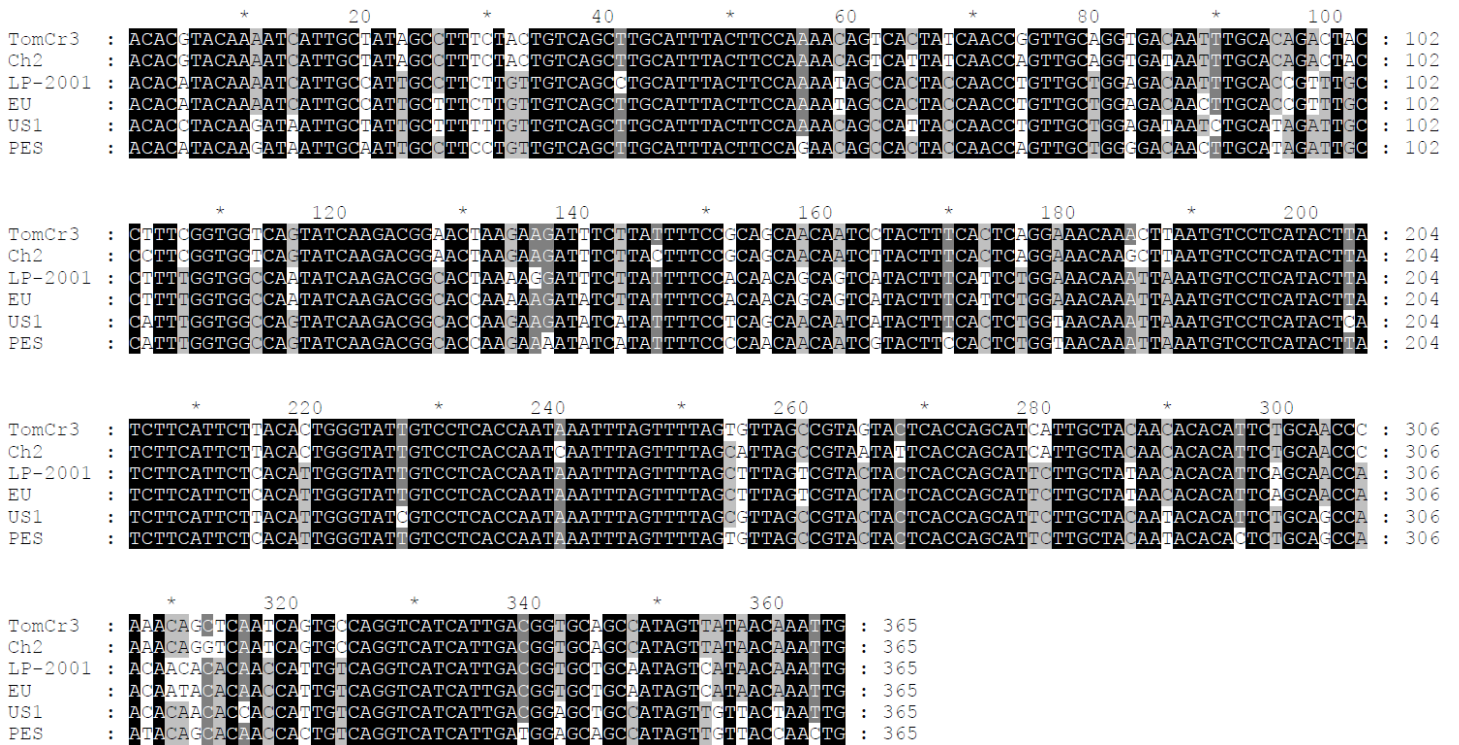


Figure S1. Multiple-sequence alignment of the *TGB2* and *TBG3* genes verified that TomCr3 is a virulent isolate of PepMV. The TomCr3 sequence of 365 nucleotides was aligned with the indicated five PepMV genotypes CH2, LP-2001, EU, US1 and PES with GenBank accession numbers DQ000985, AJ606361, KJ018164, AY509926 and HG313806, respectively.

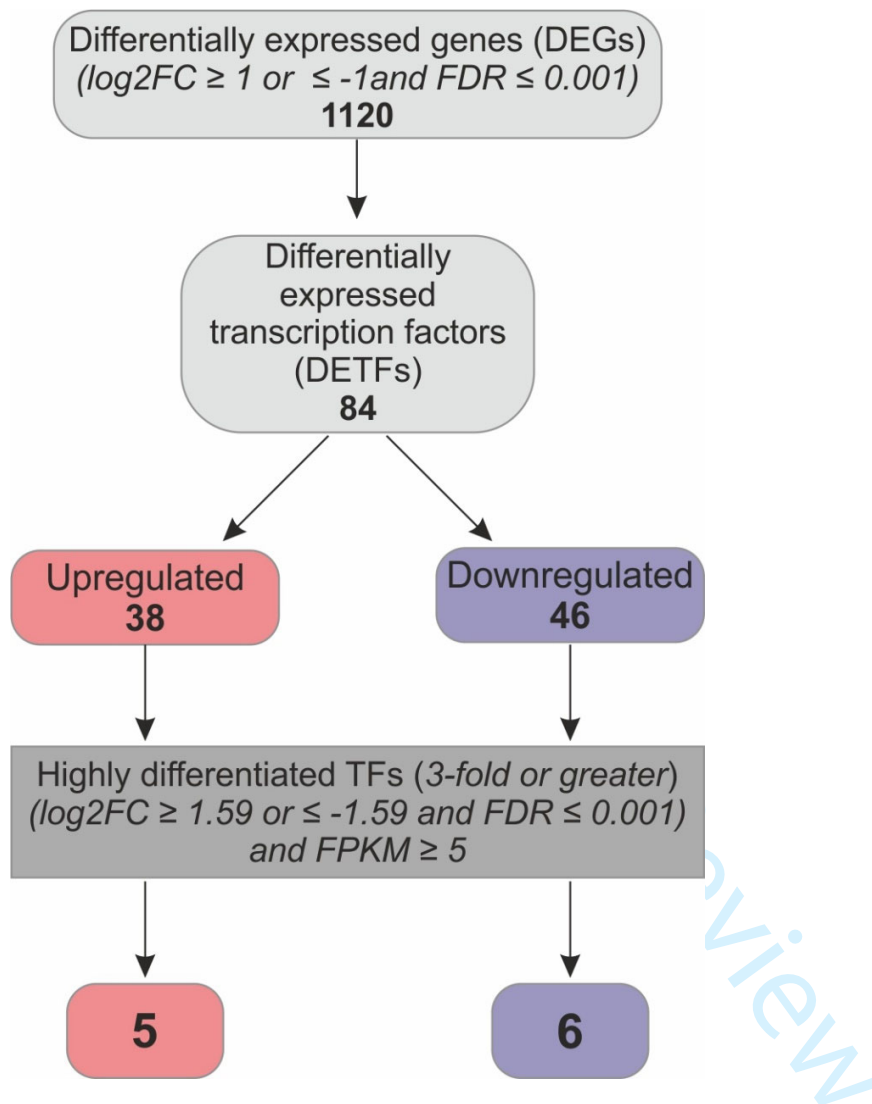
Figure S2

Figure S2. Outline of the experimental workflow leading to the identification of differentially expressed transcription factors (DETFs) upon PepMV infection of tomato.

Supplementary Table S1. Oligos used in this study for gene expression analysis

Entrez Gene ID	ITAG 4.0 Gene ID	Gene symbol		Sequence (5' → 3')	Tm	cDNA	gDNA
101248376	<i>Solyc04g015500</i>	<i>Solyc04g015500</i>	For Rev	GATGGCAAGCAGACAAGGGCAAG GTTTAGCAGTCTTACTTTCCATACCAC	64.77 62.32	202	791
101261180	<i>Solyc02g093420</i>	<i>Solyc02g093420</i>	For Rev	GTGACAATGAGGAAGCCAGTGTG CTCCATCTCACACTTAATTTAGCATCAG	62.96 62.80	206	379
101263989	<i>Solyc12g096070</i>	<i>GRF4</i>	For Rev	GACGAATGGCCTACAACCTAAGGA TGACTAGCATATCTGTTCCCTCAGC	61.12 61.45	216	309
101264084	<i>Solyc02g077610</i>	<i>NAC073</i>	For Rev	GAAGCAAAAGTACTTTCAGATAATCA CACCACCATCAATTTAGTGTGTA	57.24 59.73	213	2929
101254394	<i>Solyc06g075780</i>	<i>ZAT1-like</i>	For Rev	GTGGAATGGAGTTTTCTATGGGTC GATAAGAATGGAGGTACAGGTGGC	61.45 63.22	214	214
101263275	<i>Solyc11g056650</i>	<i>bHLH81</i>	For Rev	CACAGCAGATATGTTGGATTTTGC CATTATCAGTCCCTTTCATCCTAAC	59.73 62.07	153	248
101248026	<i>Solyc07g063410</i>	<i>JA2L</i>	For Rev	GGATTATGCACGAATACAGGCTCAGTGA GTAGAGATTCCAGCATATCATCAAACCTG	65.57 62.59	199	282
101055585	<i>Solyc01g107170</i>	<i>ZF2</i>	For Rev	GTTCTCACGTGTGCTCTATTTGTC TCTACCTCCGAGTCGATTGTCAG	64.98 61.45	284	284
101250001	<i>Solyc03g121240</i>	<i>bHLH87</i>	For Rev	GTACCAGGAGGAAGTAAAATGGATAC GAGAACATGTCAACTCTTTGGTTCG	60.10 61.45	221	221
101256853	<i>Solyc12g044390</i>	<i>DREB3</i>	For Rev	GAGAATTTGGTTGGGTACTTATCC CGTTCCATCGATTCACTGTGTCTC	59.73 63.46	228	228
101252588	<i>Solyc05g050220</i>	<i>TAF-1-like</i>	For Rev	GCAGAAGAATTGGCAGTGCAGGTCA CTCTTGCTAGTAGGTCTGCTGTGC	65.30 64.98	221	303
101258368	<i>Solyc05g014470</i>	<i>GAPDH</i>	For Rev	CAACATCATCCCTAGCAGCACT ACCCTTCAATTTACCCTCTGA	65.30 64.98	200	290

Supplementary Table S1. Oligos used in this study for gene expression analysis

Gene		Primer Sequence (5' --> 3')	Tm	PCR product
<i>TGB2</i>	For	GACTCCTAGAGCTGATCTCACTG	62.96	241
	Rev	GTGTAAGAATGAAGATAAGTATGAGGAC	61.08	
<i>TGB3</i>	For	CTTCATTCTTACACTGGGTATTGTCCTC	64.10	185
	Rev	GCTTTAAGAAGCTTCAGGTGTGTTTGGGA	62.32	

For Peer Review

Supplementary Table S2. RNA-sequencing data quality metrics. (A) RNA-seq clean reads quality metrics

Sample	Total Raw reads (Mb)	Total Clean reads (Mb)	Clean reads ratio (%)
Control	129.93	123.74	95.23%
<i>PepMV</i>	132.43	127.24	96.08%

For Peer Review

Supplementary Table S2. RNA-sequencing data quality metrics.
(B) Statistical summary of the identified genes and transcripts
between mock and PepMV infected plants

Sample	Total Gene Number	Total Transcript Number
Control	22931	31318
<i>PepMV</i>	22497	30192

For Peer Review

Supplementary Table S3. Expression data of the differentially expressed genes (DEGs) identified by RNA-seq analysis. (A) Expression data of the 1120 differentially expressed genes (DEGs) identified by RNA-seq analysis.

GeneID	Length	Control FPKM	PepMV FPKM	log2Fold Change (PepMV/Control)	FDR	Up/Down-Regulation (PepINF/Control)	Pvalue	Symbol	Description	Cellular Component	Molecular Function	Biological Process	Kegg Orthology
101256334	532	0.01	5.2	9.0223678	4.45E-34	Up	3.39E-35	LOC101256334	PREDICTED: Sol:GO:0005840,GO:000372;GO:000641				K02942//I
109118700	2174	0.01	1.83	7.5156998	1.77E-53	Up	9.35E-55	LOC109118700	PREDICTED: Sol:GO:0043231,GO:000372;GO:000945				K17964//I
109118699	2303	0.01	1.37	7.0980321	1.06E-42	Up	6.87E-44	LOC109118699	PREDICTED: Sol:NA	GO:000372	NA		K14398//c
101264023	1666	0.01	1.32	7.0443941	1.01E-27	Up	9.13E-29	LOC101264023	PREDICTED: Sol:NA	NA	NA		NA
101248021	2349	0.07	7.94	6.8256403	4.07E-234	Up	5.97E-236	LOC101248021	PREDICTED: Sol:NA	NA	NA		NA
109119317	977	0.01	0.31	4.9541963	0.00026	Up	9.01E-05	LOC109119317	PREDICTED: Sol:NA	NA	NA		NA
101248351	1087	0.01	0.24	4.5849625	0.000671	Up	0.000245	LOC101248351	PREDICTED: Sol:GO:0005634,GO:000098;GO:000635				K09422//t
101266758	1763	0.01	0.22	4.4594316	1.51E-05	Up	4.47E-06	LOC101266758	PREDICTED: Sol:NA	GO:004687	GO:000911		K00764//a
101261785	1996	0.01	0.22	4.4594316	2.12E-05	Up	6.34E-06	LOC101261785	PREDICTED: Sol:GO:0016021,GO:001522;GO:000682				K03327//I
104644719	730	0.03	0.51	4.0874628	0.0003	Up	0.000105	LOC104644719	PREDICTED: Sol:GO:0016021,GO:001678				NA
101249661	375	0.06	0.99	4.0443941	0.000719	Up	0.000263	LOC101249661	PREDICTED: Sol:GO:0005737,GO:004691;GO:003006				K19307//I
101261464	3007	0.02	0.31	3.9541963	1.28E-09	Up	2.67E-10	LOC101261464	PREDICTED: Sol:GO:0000502,GO:003022;GO:004217				K03028//I
101261911	2416	0.42	6.47	3.9453045	1.15E-147	Up	2.50E-149	LOC101261911	PREDICTED: Sol:GO:0005634,GO:000367;GO:000962				K10882//c
101268190	2061	0.01	0.15	3.9068906	0.00026	Up	9.01E-05	LOC101268190	PREDICTED: Sol:NA	NA	NA		K17604//z
109119186	2131	0.01	0.15	3.9068906	0.000101	Up	3.31E-05	LOC109119186	PREDICTED: Sol:NA	NA	NA		K19985//e
101260458	2133	0.01	0.14	3.8073549	0.00026	Up	9.01E-05	LOC101260458	PREDICTED: Sol:NA	NA	NA		K17604//z
101259485	583	3.41	41.3	3.5982981	3.83E-193	Up	6.62E-195	LOC101259485	PREDICTED: Sol:NA	NA	NA		NA
104647537	2901	0.01	0.12	3.5849625	0.0003	Up	0.000105	LOC104647537	PREDICTED: Sol:NA	NA	NA		K01528//I
101267923	885	0.07	0.83	3.5676845	5.54E-07	Up	1.43E-07	LOC101267923	PREDICTED: Sol:NA	GO:001672	NA		K00799//I
101247045	1400	0.04	0.46	3.523562	2.99E-06	Up	8.27E-07	LOC101247045	PREDICTED: Sol:GO:0016021,NA		NA		K02184//f
101251786	1089	0.12	1.33	3.4703199	1.91E-12	Up	3.27E-13	LOC101251786	PREDICTED: Sol:NA	NA	NA		K12486//s
100147717	742	180.41	1996.54	3.4681508	0	Up	0	LOC100147717	PREDICTED: Sol:NA	NA	NA		K03006//I
101261106	1901	0.05	0.54	3.4329594	2.36E-09	Up	5.03E-10	LOC101261106	PREDICTED: Sol:NA	NA	NA		K20929//I
104646534	737	0.05	0.54	3.4329594	0.000299	Up	0.000105	LOC104646534	PREDICTED: Sol:NA	NA	NA		K22184//I
104649450	729	0.19	1.95	3.3594028	1.02E-11	Up	1.82E-12	LOC104649450	PREDICTED: Sol:NA	NA	NA		NA
101265357	2515	0.04	0.38	3.2479275	4.77E-08	Up	1.12E-08	LOC101265357	PREDICTED: Sol:GO:0016021,NA		NA		NA
104647144	502	2.23	20.53	3.202618	3.73E-73	Up	1.51E-74	LOC104647144	PREDICTED: Sol:NA	NA	NA		NA
101244376	1229	0.04	0.35	3.129283	0.000247	Up	8.53E-05	LOC101244376	Solanum lycopersiGO:0009505,GO:004687;GO:000697				K00430//I
109118705	1015	0.04	0.33	3.0443941	0.000719	Up	0.000263	LOC109118705	PREDICTED: Sol:GO:0016021,GO:000467		NA		K13420//I
104648606	1296	0.04	0.33	3.0443941	0.000859	Up	0.000318	LOC104648606	PREDICTED: Sol:GO:0005634,GO:000367;GO:000635				NA

544271	1399	0.81	6.64	3.0351894	1.52E-67	Up	6.62E-69	ARG2	PREDICTED: Sol:NA	GO:004687	NA	K01476//a	
101255243	1134	0.1	0.81	3.0179219	2.33E-07	Up	5.82E-08	LOC101255243	PREDICTED: Sol:GO:0016021	GO:000552	NA	K01537//	
101245301	633	147.06	1164.57	2.9853206	0	Up	0	SN2	Solanum lycopersi	GO:0005618	NA	GO:000695	K03860//
101252520	604	0.57	4.51	2.9840936	4.89E-19	Up	6.03E-20	LOC101252520	PREDICTED: Sol:NA	NA	NA	NA	
101267575	1608	0.18	1.41	2.9696264	5.24E-17	Up	7.08E-18	LOC101267575	PREDICTED: Sol:GO:0016021	GO:001670	GO:001020	K20667//	
101253130	639	0.22	1.72	2.9668331	3.35E-08	Up	7.75E-09	LOC101253130	PREDICTED: Sol:NA	NA	NA	NA	
101252786	1384	0.05	0.38	2.9259994	0.000258	Up	8.93E-05	LOC101252786	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	K03094//	
101264863	1549	28.23	209.12	2.88903	0	Up	0	LOC101264863	PREDICTED: Sol:NA	GO:000848	GO:000905	K00815//t	
101256104	1933	0.05	0.37	2.8875253	7.90E-06	Up	2.27E-06	LOC101256104	PREDICTED: Sol:GO:0016021	GO:001670	NA	K07428//c	
101265400	1393	0.98	6.62	2.7559776	1.44E-60	Up	6.82E-62	LOC101265400	PREDICTED: Sol:NA	NA	NA	K04708//	
101250202	888	8.49	56.43	2.7326259	0	Up	0	LOC101250202	PREDICTED: Sol:GO:0000151	GO:006162	GO:004278	K11975//	
101264605	1130	26.87	176.02	2.7116711	0	Up	0	LOC101264605	Solanum lycopersi	GO:0016021	GO:000521	NA	K09872//a
104647308	424	0.3	1.96	2.7078192	1.12E-05	Up	3.25E-06	LOC104647308	PREDICTED: Sol:NA	NA	NA	NA	
101267701	746	1.17	7.57	2.6937848	3.76E-35	Up	2.78E-36	LOC101267701	PREDICTED: Sol:NA	NA	GO:000695	NA	
101260552	1027	0.09	0.58	2.688056	0.000164	Up	5.52E-05	LOC101260552	PREDICTED: Sol:NA	GO:001678	NA	NA	
101249807	1298	0.79	5.05	2.6763588	1.20E-41	Up	7.85E-43	LOC101249807	PREDICTED: Sol:GO:0005634	GO:000367	GO:001010	K09272//s	
104647943	495	3.58	21.93	2.6148743	5.44E-62	Up	2.53E-63	LOC104647943	PREDICTED: Sol:NA	NA	NA	NA	
101263989	1836	2.14	13.07	2.6105764	2.75E-148	Up	5.95E-150	LOC101263989	PREDICTED: Sol:NA	NA	NA	K20496//l	
101262976	1088	1.35	8.23	2.607933	1.25E-55	Up	6.44E-57	LOC101262976	PREDICTED: Sol:NA	NA	NA	K04733//i	
101255585	1481	7.41	44.89	2.5988486	0	Up	0	LOC101255585	PREDICTED: Sol:NA	NA	NA	K21596//c	
101268097	1680	1.23	7.33	2.5751549	5.23E-75	Up	2.05E-76	LOC101268097	PREDICTED: Sol:NA	GO:000365	GO:000635	K15032//	
544031	779	0.33	1.95	2.5629362	1.21E-09	Up	2.53E-10	ER5	Solanum lycopersi	NA	NA	GO:000920	NA
101256859	1826	36.94	217.37	2.5568971	0	Up	0	LOC101256859	PREDICTED: Sol:GO:0043231	GO:008002	GO:000815	K21371//c	
101252885	657	1.28	7.34	2.5196363	1.31E-27	Up	1.19E-28	LOC101252885	PREDICTED: Sol:NA	NA	GO:000695	NA	
101260563	1720	29.75	168.57	2.5023862	0	Up	0	LOC101260563	PREDICTED: Sol:GO:0009508	NA	GO:000635	NA	
101268894	789	1.94	10.95	2.4968023	5.63E-49	Up	3.20E-50	LOC101268894	Solanum lycopersi	GO:0005783	NA	GO:001010	NA
101245410	1412	0.25	1.4	2.4854268	3.92E-12	Up	6.84E-13	LOC101245410	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	K09286//	
101267770	2708	0.06	0.33	2.4594316	5.58E-06	Up	1.58E-06	LOC101267770	PREDICTED: Sol:NA	NA	NA	NA	
101243937	2971	0.14	0.77	2.4594316	2.50E-14	Up	3.88E-15	LOC101243937	PREDICTED: Sol:GO:0016021	GO:000552	NA	K04733//i	
101258262	4181	0.04	0.22	2.4594316	6.80E-06	Up	1.94E-06	LOC101258262	PREDICTED: Sol:GO:0016021	GO:000467	NA	K13420//	
104646731	1292	1.02	5.49	2.428237	3.69E-40	Up	2.48E-41	LOC104646731	PREDICTED: Sol:NA	NA	GO:000695	K10745//r	
101255344	1559	5.12	27.12	2.4051415	2.30E-236	Up	3.35E-238	LOC101255344	PREDICTED: Sol:GO:0005634	GO:000482	GO:000628	K03648//	
104645976	1005	6.95	36.52	2.3936019	1.08E-198	Up	1.82E-200	LOC104645976	PREDICTED: Sol:GO:0016021	GO:000815	GO:000665	K07252//	
101250114	964	0.18	0.93	2.3692338	1.13E-05	Up	3.30E-06	LOC101250114	PREDICTED: Sol:GO:0005634	GO:004698	NA	K08065//	
101260051	781	0.28	1.42	2.3423922	1.01E-06	Up	2.67E-07	LOC101260051	PREDICTED: Sol:NA	NA	NA	K04733//i	
101250357	2859	8.68	43.83	2.3361517	0	Up	0	LOC101250357	PREDICTED: Sol:NA	NA	NA	K01087//t	
104645999	393	1.48	7.37	2.3160674	1.82E-14	Up	2.79E-15	LOC104645999	PREDICTED: Sol:GO:0016021	NA	NA	NA	

101261864	1569	0.36	1.79	2.3138908	9.94E-16	Up	1.43E-16	LOC101261864	PREDICTED: Sol:GO:0005634.GO:000552GO:00325(K15601//
101254673	1667	0.54	2.61	2.2730185	2.56E-23	Up	2.67E-24	LOC101254673	PREDICTED: Sol:GO:0005794.GO:001641GO:00715(K16573//
101255673	1413	0.11	0.53	2.2684888	7.56E-05	Up	2.43E-05	LOC101255673	PREDICTED: Sol:GO:0005737.GO:00045(GO:00333(K01581//
101248826	1300	0.37	1.78	2.2662801	8.09E-13	Up	1.36E-13	LOC101248826	PREDICTED: Sol:NA NA NA NA
101266990	2251	0.11	0.52	2.2410081	3.66E-07	Up	9.26E-08	LOC101266990	PREDICTED: Sol:GO:0016021.NA NA K18171//
101252509	1590	3.38	15.92	2.2357452	4.19E-128	Up	1.05E-129	LOC101252509	PREDICTED: Sol:NA GO:00167(NA K20556//c
101256502	793	6.47	30.15	2.2203204	5.65E-116	Up	1.54E-117	LOC101256502	PREDICTED: Sol:GO:0046658.GO:00045(NA K03006//
543837	917	17.5	78.83	2.1713899	0	Up	0	PR-5	Solanum lycopersi NA NA NA K04733//i
101253069	2402	1.42	6.31	2.1517491	3.02E-74	Up	1.19E-75	LOC101253069	PREDICTED: Sol:GO:0016021.GO:00054(NA K13862//s
101246212	2240	2.01	8.91	2.1482299	1.77E-97	Up	5.55E-99	LOC101246212	PREDICTED: Sol:GO:0016021.GO:000552NA K04733//i
101247976	4798	6.03	26.68	2.1455288	0	Up	0	DCL2d	Solanum lycopersi NA GO:000372GO:00063(K11592//e
104646732	1753	0.1	0.44	2.1375035	8.84E-05	Up	2.87E-05	LOC104646732	PREDICTED: Sol:NA NA GO:00069(NA
101251725	479	1.48	6.45	2.123702	2.87E-14	Up	4.47E-15	LOC101251725	PREDICTED: Sol:NA NA GO:00097(K14488//
104646809	1314	0.12	0.52	2.1154772	0.000126	Up	4.17E-05	LOC104646809	PREDICTED: Sol:NA NA NA NA
101252867	609	63.03	269.88	2.0982075	0	Up	0	LOC101252867	PREDICTED: Sol:NA NA NA K03860//
101245904	632	24.36	103.14	2.0820179	9.89E-284	Up	1.20E-285	LOC101245904	PREDICTED: Sol:GO:0048046.GO:004712GO:00066(K03671//t
544212	1259	0.2	0.83	2.0531113	1.05E-05	Up	3.05E-06	LOC544212	PREDICTED: Sol:GO:0016020.NA GO:00096(K20628//e
101246976	471	58.04	240.23	2.0492969	0	Up	0	LOC101246976	PREDICTED: Sol:GO:0016021.NA NA NA
101263175	1195	40.04	165.22	2.0448745	0	Up	0	LOC101263175	PREDICTED: Sol:GO:0016021.GO:004435GO:007171K10666//
104646059	828	0.26	1.07	2.0410273	0.000112	Up	3.68E-05	LOC104646059	PREDICTED: Sol:NA NA NA K00799//
101262502	1131	0.22	0.9	2.0324215	4.10E-05	Up	1.27E-05	LOC101262502	PREDICTED: Sol:NA NA NA NA
101251987	1134	1.57	6.4	2.0273073	1.67E-32	Up	1.33E-33	LOC101251987	PREDICTED: Sol:NA NA NA NA
101246628	1396	27.34	111.14	2.023293	0	Up	0	LOC101246628	PREDICTED: Sol:GO:0009535.GO:001616GO:00097(K14172//l
101246149	699	2.79	11.29	2.0167085	5.96E-34	Up	4.56E-35	LOC101246149	PREDICTED: Sol:NA NA NA K09539//
101267371	763	0.22	0.89	2.0163018	0.000568	Up	0.000206	LOC101267371	PREDICTED: Sol:NA NA GO:00466(K14488//
101246713	1824	2.24	9.04	2.012824	1.40E-73	Up	5.60E-75	LOC101246713	PREDICTED: Sol:GO:0005576.GO:000462GO:00059(K01184//
101261180	2060	5.9	23.46	1.9914162	1.82E-211	Up	2.86E-213	LOC101261180	PREDICTED: Sol:GO:0005634.GO:000367GO:00063(NA
101262811	850	0.28	1.1	1.9740048	6.50E-05	Up	2.07E-05	LOC101262811	PREDICTED: Sol:GO:0005634.GO:000367GO:00063(K09260//
101254739	2441	0.5	1.96	1.9708537	1.90E-21	Up	2.12E-22	LOC101254739	PREDICTED: Sol:NA NA NA K02184//f
101268195	2375	0.24	0.94	1.9696264	1.75E-10	Up	3.42E-11	LOC101268195	PREDICTED: Sol:GO:0016020.NA NA K13412//c
101257220	735	0.46	1.8	1.9682911	2.28E-06	Up	6.22E-07	LOC101257220	PREDICTED: Sol:NA GO:00080(NA NA
101251607	1993	4.45	17.35	1.9630584	1.31E-148	Up	2.81E-150	LOC101251607	PREDICTED: Sol:NA GO:004687NA K05277//l
101246590	749	0.5	1.94	1.9560567	7.69E-07	Up	2.01E-07	LOC101246590	PREDICTED: Sol:GO:0005634.GO:000367GO:00063(K14516//e
101263653	1069	8.88	34.17	1.9440987	6.62E-151	Up	1.40E-152	LOC101263653	PREDICTED: Sol:NA NA NA K04733//i
101243792	951	0.49	1.88	1.939879	2.72E-08	Up	6.26E-09	LOC101243792	PREDICTED: Sol:NA NA NA NA
101262337	1127	0.2	0.76	1.9259994	0.000226	Up	7.75E-05	LOC101262337	PREDICTED: Sol:NA NA NA NA
101246961	581	4.82	18.31	1.9255267	5.28E-42	Up	3.45E-43	LOC101246961	PREDICTED: Sol:NA GO:00048(GO:000961NA

101244725	1949	0.27	1.02	1.9175378	3.09E-09	Up	6.63E-10	LOC101244725	PREDICTED: Sol:GO:0043231.GO:008002GO:000815K13495//c
778225	2192	3.69	13.9	1.9133922	4.28E-128	Up	1.08E-129	LOC778225	Solanum lycopersiGO:0009507.GO:004682GO:004665K02552//
101253556	1568	0.16	0.6	1.9068906	7.08E-05	Up	2.26E-05	LOC101253556	PREDICTED: Sol:NA GO:001672NA K13065//s
101250018	691	2.13	7.98	1.9055353	2.99E-22	Up	3.25E-23	LOC101250018	PREDICTED: Sol:NA NA NA K14306//
101267994	2074	0.76	2.84	1.9018196	7.53E-25	Up	7.40E-26	LOC101267994	PREDICTED: Sol:GO:0016021.GO:000552NA K05681//
101263993	2355	0.45	1.68	1.9004643	3.72E-17	Up	4.98E-18	LOC101263993	PREDICTED: Sol:GO:0005737.GO:000447GO:000605K01638//
543951	990	2.52	9.39	1.8977014	1.06E-37	Up	7.49E-39	TIP	Solanum lycopersiGO:0009705.GO:001525GO:003425K09873//a
101259870	1082	0.18	0.67	1.8961642	0.000681	Up	0.000249	LOC101259870	PREDICTED: Sol:GO:0005634.GO:000367NA K01870//i
101263261	1553	32.75	121.43	1.8905581	0	Up	0	Tom52	Solanum lycopersi NA GO:000552NA K10355//a
101255604	1892	11.97	44.34	1.8891856	0	Up	0	LOC101255604	PREDICTED: Sol:GO:0016021.GO:001670NA K07428//c
101244403	1537	8.42	30.8	1.8710382	2.16E-188	Up	3.80E-190	LOC101244403	PREDICTED: Sol:GO:0016021.GO:004687NA NA
101266172	1238	9.84	35.92	1.8680571	4.44E-175	Up	8.15E-177	LOC101266172	PREDICTED: Sol:GO:0009507.GO:000365GO:003255K15032//
101255488	3883	1.19	4.33	1.8634055	1.45E-68	Up	6.19E-70	LOC101255488	PREDICTED: Sol:GO:0043231.GO:000372GO:000945K17964//l
104646656	1363	0.21	0.76	1.8556101	4.38E-05	Up	1.37E-05	LOC104646656	PREDICTED: Sol:NA NA NA NA
104646221	533	3.35	12.1	1.852774	2.93E-24	Up	2.94E-25	LOC104646221	PREDICTED: Sol:NA NA NA NA
101253800	822	25.65	92.57	1.8515859	2.74E-288	Up	3.22E-290	LOC101253800	PREDICTED: Sol:GO:0009507.NA NA K07253//
101254394	1145	2.26	8.11	1.8433791	1.85E-36	Up	1.34E-37	LOC101254394	PREDICTED: Sol:NA GO:000367NA NA
101246807	1205	15.19	54.44	1.8415452	1.91E-252	Up	2.64E-254	LOC101246807	PREDICTED: Sol:NA NA NA K14496//a
101250317	1217	4.36	15.34	1.8148984	4.86E-71	Up	2.02E-72	LOC101250317	PREDICTED: Sol:GO:0016020.GO:000552GO:000726K04392//
101254591	1117	8.67	30.5	1.8147053	2.53E-128	Up	6.35E-130	LOC101254591	PREDICTED: Sol:NA GO:001645NA K09841//
101264084	1271	3.53	12.37	1.8091054	1.00E-59	Up	4.84E-61	LOC101264084	PREDICTED: Sol:GO:0005634.GO:000367GO:000635NA
101247679	4669	18.16	63.57	1.8075819	0	Up	0	DCL2c	Solanum lycopersi NA GO:000372GO:000635K11592//e
101255600	989	0.64	2.24	1.8073549	4.19E-09	Up	9.09E-10	LOC101255600	Solanum lycopersiGO:0009705.GO:001525GO:003425K09873//a
101255332	1580	0.14	0.49	1.8073549	0.000757	Up	0.000278	LOC101255332	PREDICTED: Sol:NA GO:000415GO:000650K00924//
101261675	1929	0.93	3.24	1.8006912	1.45E-24	Up	1.44E-25	LOC101261675	PREDICTED: Sol:GO:0043231.GO:008002GO:000815K13496//
101256037	989	0.31	1.07	1.7872707	8.37E-05	Up	2.71E-05	LOC101256037	PREDICTED: Sol:NA GO:004742GO:000912K16904//
101264232	3292	0.65	2.22	1.7720481	2.71E-28	Up	2.42E-29	LOC101264232	PREDICTED: Sol:GO:0016021.GO:001630NA K13466//
104647423	1047	2.98	10.17	1.7709354	3.04E-39	Up	2.08E-40	LOC104647423	PREDICTED: Sol:NA NA NA NA
101258746	1677	2.17	7.4	1.7698302	2.36E-46	Up	1.41E-47	LOC101258746	PREDICTED: Sol:GO:0005887.GO:001525GO:000335K03834//t
104647464	699	0.63	2.14	1.7641871	3.11E-06	Up	8.61E-07	LOC104647464	PREDICTED: Sol:GO:0005634.GO:000367GO:000635K09260//
101265385	1695	8.93	30.33	1.7640134	9.02E-189	Up	1.58E-190	LOC101265385	PREDICTED: Sol:GO:0016021.NA NA K07297//a
101268568	1750	1.46	4.95	1.7614602	1.54E-32	Up	1.22E-33	LOC101268568	PREDICTED: Sol:NA GO:001630NA K14963//
101250032	2636	0.11	0.37	1.7500217	8.00E-05	Up	2.58E-05	LOC101250032	PREDICTED: Sol:NA GO:004687NA K05278//f
101251556	1518	0.36	1.21	1.7489382	1.46E-05	Up	4.31E-06	LOC101251556	PREDICTED: Sol:NA NA NA NA
101248621	670	0.65	2.18	1.7458165	5.40E-06	Up	1.53E-06	LOC101248621	PREDICTED: Sol:NA NA GO:000975K14488//
101263764	2061	13.04	43.69	1.7443592	0	Up	0	LOC101263764	PREDICTED: Sol:NA NA NA K19041//
104647122	1093	0.26	0.87	1.7425038	0.000259	Up	8.96E-05	LOC104647122	PREDICTED: Sol:NA NA NA NA

101256761	1684	0.84	2.81	1.7421089	4.83E-18	Up	6.22E-19	LOC101256761	PREDICTED: Sol:NA	GO:001678	NA	NA	
101246259	1247	0.41	1.36	1.7299108	6.46E-07	Up	1.68E-07	LOC101246259	PREDICTED: Sol:GO:0005634	GO:000367	NA	NA	
101266674	1524	1.17	3.88	1.7295481	1.62E-21	Up	1.81E-22	LOC101266674	PREDICTED: Sol:NA	GO:000848	GO:000905	K00815//t	
101258730	1208	0.19	0.63	1.7293524	0.000724	Up	0.000265	LOC101258730	PREDICTED: Sol:NA	NA	NA	NA	
101258643	1095	0.39	1.29	1.725825	5.61E-06	Up	1.59E-06	LOC101258643	PREDICTED: Sol:GO:0016021	NA	NA	K04733//i	
104648572	2196	1.08	3.57	1.7248928	7.09E-29	Up	6.21E-30	LOC104648572	PREDICTED: Sol:GO:0016021	GO:000521	NA	K14638//s	
101260801	1641	4.09	13.51	1.7238549	4.32E-79	Up	1.62E-80	LOC101260801	PREDICTED: Sol:NA	GO:004687	NA	K05278//f	
101266787	1226	31.77	103.75	1.7073743	0	Up	0	LOC101266787	PREDICTED: Sol:GO:0009507	GO:000475	GO:000905	K01807//r	
101252998	2378	0.12	0.39	1.7004397	0.000467	Up	0.000167	LOC101252998	PREDICTED: Sol:NA	NA	NA	NA	
101249263	1223	4.04	13.1	1.6971396	1.64E-55	Up	8.44E-57	LOC101249263	PREDICTED: Sol:NA	GO:000395	NA	NA	
543803	1360	1428.5	4630.37	1.6966264	0	Up	0	ca2	Solanum lycopersi	NA	GO:000827	GO:001597	K01673//c
101244789	1261	1.52	4.92	1.694587	5.90E-22	Up	6.48E-23	LOC101244789	PREDICTED: Sol:NA	GO:001678	NA	K21026//a	
101256339	2359	1.44	4.66	1.6942611	1.91E-39	Up	1.30E-40	LOC101256339	PREDICTED: Sol:GO:0016021	NA	NA	NA	
104645965	908	0.6	1.94	1.6930222	1.93E-07	Up	4.77E-08	LOC104645965	PREDICTED: Sol:NA	NA	NA	NA	
101258619	1740	0.39	1.26	1.6918777	1.50E-08	Up	3.37E-09	LOC101258619	PREDICTED: Sol:GO:0016021	NA	NA	NA	
101245183	2288	6.29	20.19	1.682509	6.56E-160	Up	1.30E-161	LOC101245183	PREDICTED: Sol:NA	GO:000875	GO:000151	K21971//	
101246482	745	5.57	17.83	1.6785575	4.12E-44	Up	2.57E-45	LOC101246482	PREDICTED: Sol:NA	NA	NA	NA	
104644828	410	0.89	2.84	1.6740137	0.000154	Up	5.15E-05	LOC104644828	PREDICTED: Sol:NA	NA	NA	NA	
101264700	1780	41.45	132.26	1.6739328	0	Up	0	LOC101264700	PREDICTED: Sol:GO:0005737	GO:000374	NA	K03231//e	
101257381	1292	1.03	3.26	1.6622276	1.21E-15	Up	1.75E-16	LOC101257381	PREDICTED: Sol:NA	GO:004687	NA	K00475//	
109121128	738	0.48	1.51	1.6534422	0.000156	Up	5.25E-05	LOC109121128	PREDICTED: Sol:NA	NA	NA	NA	
101248089	1804	0.83	2.6	1.6473284	1.69E-16	Up	2.34E-17	LOC101248089	PREDICTED: Sol:GO:0016020	GO:001676	GO:004455	K07410//c	
101265704	1527	0.84	2.63	1.6466016	3.74E-14	Up	5.85E-15	LOC101265704	PREDICTED: Sol:NA	NA	NA	NA	
101244702	1624	8.27	25.86	1.644763	2.96E-139	Up	6.89E-141	LOC101244702	PREDICTED: Sol:NA	GO:000825	GO:000656	K01285//l	
104646463	1851	1.11	3.47	1.644376	3.99E-22	Up	4.36E-23	LOC104646463	PREDICTED: Sol:NA	GO:000827	NA	K07151//	
101251570	3311	0.24	0.75	1.6438562	5.51E-09	Up	1.20E-09	LOC101251570	PREDICTED: Sol:GO:0016021	GO:000467	NA	K13428//	
101260153	1622	0.44	1.37	1.6386005	2.73E-08	Up	6.28E-09	LOC101260153	PREDICTED: Sol:NA	GO:001687	NA	K14963//	
101256006	926	175.77	546.88	1.6375354	0	Up	0	CAB12	Solanum lycopersi	GO:0010287	GO:003146	GO:000976	K08910//l
101245470	885	2	6.22	1.6369146	2.29E-18	Up	2.91E-19	LOC101245470	PREDICTED: Sol:NA	NA	NA	NA	
101260301	1452	12.74	39.37	1.6277314	7.94E-186	Up	1.41E-187	LOC101260301	PREDICTED: Sol:NA	NA	NA	NA	
101260686	2924	9.22	28.41	1.6235602	2.75E-286	Up	3.28E-288	LOC101260686	PREDICTED: Sol:GO:0016021	GO:000552	NA	K20718//	
101250079	1338	3.55	10.92	1.6210819	1.01E-47	Up	5.85E-49	LOC101250079	PREDICTED: Sol:NA	GO:001674	NA	K13065//s	
101250825	605	5.47	16.82	1.620565	9.77E-32	Up	7.94E-33	LOC101250825	PREDICTED: Sol:NA	GO:001996	GO:001037	NA	
101246748	1304	0.44	1.35	1.617384	1.21E-06	Up	3.21E-07	LOC101246748	PREDICTED: Sol:GO:0005634	GO:000098	GO:000635	K09422//t	
101260060	1104	20.45	62.68	1.6159043	1.89E-219	Up	2.94E-221	LOC101260060	PREDICTED: Sol:GO:0031248	GO:000459	GO:000647	NA	
101243718	1117	0.76	2.32	1.6100535	3.18E-09	Up	6.83E-10	LOC101243718	PREDICTED: Sol:NA	GO:004687	NA	K05282//	
101267169	1192	1.47	4.48	1.6076826	1.21E-17	Up	1.59E-18	LOC101267169	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	K13434//	

101257326	1360	2.58	7.82	1.5997975	2.74E-34	Up	2.07E-35	LOC101257326	PREDICTED: Sol:GO:0016021.GO:000382GO:000911K07870//
101261071	1339	0.41	1.24	1.5966443	3.60E-06	Up	1.00E-06	LOC101261071	PREDICTED: Sol:GO:0009506.NA GO:001995K20628//e
543795	1258	36.83	111.38	1.5965369	0	Up	0	LOC543795	PREDICTED: Sol:GO:0016020.NA GO:000966K20628//e
101248376	1199	11.9	35.83	1.5902065	4.91E-134	Up	1.19E-135	LOC101248376	PREDICTED: Sol:GO:0005634.GO:000367GO:000635NA
101250851	3890	5.24	15.76	1.5886288	7.01E-197	Up	1.19E-198	LOC101250851	PREDICTED: Sol:GO:0016021.GO:000467GO:000646K00924//
101246318	1876	1.38	4.11	1.5744701	6.86E-25	Up	6.73E-26	LOC101246318	PREDICTED: Sol:NA GO:000842GO:000675K01082//
101253386	1807	0.33	0.98	1.5703157	2.10E-06	Up	5.71E-07	LOC101253386	PREDICTED: Sol:NA GO:000555(NA K13448//c
101262875	1482	31.39	93.15	1.5692507	0	Up	0	LOC101262875	Solanum lycopersiGO:0016021.NA GO:000681K15109//s
101254312	1737	14.13	41.88	1.5675	1.26E-224	Up	1.93E-226	LOC101254312	PREDICTED: Sol:GO:0043231.GO:000372GO:000945K06180//
101248387	650	15.2	45.04	1.5671355	5.56E-86	Up	1.94E-87	LOC101248387	PREDICTED: Sol:NA NA NA K01513//e
101257465	1288	24.58	72.79	1.5662553	1.33E-285	Up	1.59E-287	LOC101257465	PREDICTED: Sol:GO:0005622.GO:000521NA K14399//
101254471	1785	21.18	62.68	1.5653026	0	Up	0	LOC101254471	PREDICTED: Sol:NA GO:004687GO:004545K01728//
104648056	882	1.41	4.16	1.5608884	8.77E-12	Up	1.56E-12	LOC104648056	PREDICTED: Sol:NA NA NA K11273//c
101248663	3693	0.47	1.38	1.5539356	1.82E-16	Up	2.52E-17	LOC101248663	PREDICTED: Sol:GO:0005634.GO:000808GO:000625K10862//t
101251973	2333	6.39	18.72	1.5506926	3.13E-134	Up	7.59E-136	LOC101251973	PREDICTED: Sol:GO:0005634.GO:000482NA K08332//
101262525	788	8.06	23.6	1.5499351	5.97E-55	Up	3.10E-56	LOC101262525	PREDICTED: Sol:NA NA NA NA
101255793	1619	6.5	19	1.5474878	1.65E-93	Up	5.31E-95	LOC101255793	PREDICTED: Sol:GO:0005576.GO:000462GO:000597K01184//
100134889	1150	1	2.91	1.5410192	7.47E-11	Up	1.43E-11	GA2ox4	Solanum lycopersiNA GO:004687GO:000941K04125//
101268244	927	0.42	1.22	1.5384199	0.000272	Up	9.43E-05	LOC101268244	PREDICTED: Sol:NA NA GO:004662K14488//
101254986	847	50.65	145.68	1.5241687	0	Up	0	LOC101254986	PREDICTED: Sol:GO:0005623.GO:001503GO:004545K03676//
104647171	1374	0.43	1.23	1.5162498	6.15E-06	Up	1.75E-06	LOC104647171	PREDICTED: Sol:NA NA NA K03006//
101248064	934	19.21	54.92	1.5154741	1.44E-146	Up	3.15E-148	LOC101248064	PREDICTED: Sol:NA NA NA NA
101263304	1999	3.86	10.89	1.4963312	1.60E-63	Up	7.27E-65	LOC101263304	PREDICTED: Sol:NA GO:001675GO:000815K13495//c
544122	2382	6.77	19.02	1.4902895	3.23E-131	Up	7.96E-133	MFP1	Solanum lycopersiGO:0016363.GO:000367NA K20283//
101246986	1434	10.76	30.16	1.4869584	7.33E-124	Up	1.90E-125	LOC101246986	PREDICTED: Sol:NA NA NA NA
101263329	1226	0.39	1.09	1.4827821	9.03E-05	Up	2.93E-05	LOC101263329	PREDICTED: Sol:GO:0005739.NA NA K18179//c
101262252	1154	22.5	62.75	1.4796905	1.23E-201	Up	2.03E-203	LOC101262252	PREDICTED: Sol:NA NA NA K09841//
101246760	1085	0.69	1.91	1.4689044	1.16E-06	Up	3.07E-07	LOC101246760	PREDICTED: Sol:NA NA GO:200002NA
101245177	1115	18.55	51.32	1.468102	1.31E-157	Up	2.63E-159	LOC101245177	PREDICTED: Sol:NA NA NA K17911//
101255616	2593	0.66	1.82	1.4634005	5.70E-14	Up	9.03E-15	LOC101255616	PREDICTED: Sol:GO:0005622.GO:004657NA K17756//l
101055611	3554	1.22	3.36	1.4615801	1.95E-34	Up	1.47E-35	LOC101055611	Solanum lycopersiGO:0005886.GO:004792GO:000595K00696//s
101261981	1592	0.28	0.77	1.4594316	0.000257	Up	8.87E-05	LOC101261981	PREDICTED: Sol:NA GO:001675GO:000652NA
101257978	1242	69.05	189.77	1.4585386	0	Up	0	LOC101257978	PREDICTED: Sol:NA NA NA K22068//i
101267821	829	2.67	7.32	1.4550039	5.20E-17	Up	7.02E-18	LOC101267821	PREDICTED: Sol:GO:0005576.NA NA K20412//
104645668	1399	1.41	3.86	1.4529057	2.25E-14	Up	3.48E-15	LOC104645668	PREDICTED: Sol:NA GO:001676(NA K12639//c
104648136	1202	0.41	1.12	1.4498029	0.000148	Up	4.95E-05	LOC104648136	PREDICTED: Sol:NA GO:000382NA K17508//
101267261	838	1.67	4.55	1.4460184	3.92E-11	Up	7.34E-12	LOC101267261	PREDICTED: Sol:NA NA NA NA

104646621	1914	0.32	0.87	1.4429435	2.77E-05	Up	8.44E-06	LOC104646621	PREDICTED: Sol:GO:0005730.NA	NA	NA	K14855//r
101268608	1018	43.94	119.2	1.4397775	0	Up	0	LOC101268608	PREDICTED: Sol:NA	NA	NA	K13447//r
101262956	1442	15.27	41.28	1.4347429	7.35E-160	Up	1.46E-161	LOC101262956	PREDICTED: Sol:NA	GO:00167{	NA	NA
544237	1782	82.18	221.81	1.4324652	0	Up	0	LOC544237	Solanum lycopersiGO:0005623.GO:00090{GO:00454{K00382//			
101258983	1747	3.58	9.66	1.4320636	3.88E-46	Up	2.33E-47	LOC101258983	PREDICTED: Sol:GO:0016021.GO:000521NA			K00850//
101268660	756	4.67	12.6	1.4319293	2.61E-25	Up	2.53E-26	LOC101268660	PREDICTED: Sol:NA	NA	NA	NA
101243914	1627	9.52	25.67	1.4310498	1.66E-112	Up	4.60E-114	LOC101243914	Solanum lycopersiGO:0005886.NA	NA	NA	NA
101254402	1705	19.56	52.73	1.4307176	2.17E-241	Up	3.08E-243	LOC101254402	PREDICTED: Sol:GO:0043231.GO:00800{GO:00081{K13495//c			
101245976	1477	25.15	67.73	1.4292366	8.17E-267	Up	1.06E-268	LOC101245976	PREDICTED: Sol:NA	NA	NA	K13115//c
101259800	575	1.17	3.14	1.424256	1.68E-05	Up	4.98E-06	LOC101259800	PREDICTED: Sol:NA	NA	NA	K03860//
101247878	1089	0.79	2.11	1.4173184	6.14E-07	Up	1.59E-07	LOC101247878	PREDICTED: Sol:NA	NA	NA	NA
101260430	1455	27.84	74.29	1.4160088	5.77E-284	Up	6.97E-286	LOC101260430	PREDICTED: Sol:GO:0016021.GO:00081{NA			K12589//e
101251870	1070	0.59	1.57	1.4119777	2.77E-05	Up	8.44E-06	LOC101251870	PREDICTED: Sol:GO:0016021.NA	NA	NA	K19041//
101251239	716	52.85	140.51	1.4106974	1.07E-226	Up	1.63E-228	LOC101251239	PREDICTED: Sol:NA	NA	NA	NA
101262198	1496	2.12	5.63	1.4090707	8.44E-23	Up	9.00E-24	LOC101262198	Solanum lycopersi NA	GO:00469{	NA	K16190//
101268283	958	5.28	14.02	1.4088765	4.14E-35	Up	3.07E-36	LOC101268283	PREDICTED: Sol:NA	NA	GO:00069{	K10745//r
101258163	3255	2.49	6.59	1.4041327	2.59E-57	Up	1.30E-58	LOC101258163	PREDICTED: Sol:NA	GO:00302{	GO:00059{	K12309//
101250586	1623	41.78	110.39	1.4017251	0	Up	0	LOC101250586	PREDICTED: Sol:GO:0016021.GO:00164{GO:00066{K20416//			
101261193	1736	2.54	6.7	1.3993326	6.41E-31	Up	5.34E-32	LOC101261193	PREDICTED: Sol:GO:0043231.GO:00800{GO:00081{K12356//c			
101260610	1593	4.3	11.34	1.3990121	2.45E-47	Up	1.43E-48	LOC101260610	PREDICTED: Sol:NA	GO:00167{	NA	K13065//s
101254642	2166	6.11	15.99	1.3879257	6.65E-90	Up	2.22E-91	LOC101254642	PREDICTED: Sol:NA	NA	NA	NA
101249824	2602	11.32	29.41	1.3774328	3.24E-196	Up	5.54E-198	PAL2	Solanum lycopersiGO:0005737.GO:00455{GO:00098{K10775//			
101248819	1917	8.7	22.6	1.3772355	1.83E-110	Up	5.12E-112	LOC101248819	PREDICTED: Sol:GO:0005829.GO:00167{GO:00081{K13258//			
101268314	605	689.98	1789.73	1.3751155	0	Up	0	LOC101268314	PREDICTED: Sol:NA	NA	NA	NA
101268846	2662	0.9	2.33	1.372333	9.53E-17	Up	1.30E-17	LOC101268846	PREDICTED: Sol:NA	GO:00050{	NA	K10418//
101254974	1033	1.08	2.78	1.3640536	5.90E-08	Up	1.39E-08	LOC101254974	PREDICTED: Sol:GO:0005634.GO:00036{GO:00063{NA			
101267854	2519	0.54	1.39	1.3640536	7.45E-10	Up	1.53E-10	LOC101267854	PREDICTED: Sol:NA	NA	GO:00165{	K03063//
104646517	1315	0.79	2.03	1.3615552	1.71E-07	Up	4.21E-08	LOC104646517	PREDICTED: Sol:NA	NA	NA	NA
101264067	1651	2.52	6.45	1.3558754	4.17E-27	Up	3.83E-28	LOC101264067	PREDICTED: Sol:GO:0016021.NA	NA	NA	K04733//i
101251812	847	1.11	2.84	1.3553313	9.24E-07	Up	2.43E-07	LOC101251812	PREDICTED: Sol:NA	NA	NA	NA
101249327	664	32.03	81.88	1.3540873	1.65E-127	Up	4.18E-129	LOC101249327	PREDICTED: Sol:GO:0009507.NA	NA	NA	NA
104647914	2388	0.2	0.51	1.3504972	0.000413	Up	0.000147	LOC104647914	PREDICTED: Sol:NA	NA	NA	K19525//
101263714	1466	8.81	22.44	1.3488588	8.34E-81	Up	3.05E-82	LOC101263714	PREDICTED: Sol:GO:0005737.GO:00468{GO:00300{K17046//			
101256987	1010	2.16	5.5	1.3484003	3.32E-14	Up	5.18E-15	LOC101256987	PREDICTED: Sol:NA	NA	NA	NA
101260911	1839	8.68	22.07	1.3463197	6.24E-100	Up	1.90E-101	LOC101260911	PREDICTED: Sol:GO:0016021.GO:00468{GO:00454{K01728//			
100736527	1149	71.94	182.88	1.3460312	0	Up	0	LOC100736527	Solanum lycopersiGO:0005887.GO:00152{GO:00342{K09872//a			
101258159	832	22.96	58.27	1.3436307	2.12E-114	Up	5.80E-116	LOC101258159	PREDICTED: Sol:GO:0005634.GO:00036{GO:00458{NA			

101256655	1530	1.05	2.66	1.3410369	7.27E-11	Up	1.39E-11	LOC101256655	PREDICTED: Sol:NA	GO:000367	NA	K20557//t	
101246041	2533	1.82	4.59	1.3345557	3.47E-29	Up	3.02E-30	LOC101246041	PREDICTED: Sol:GO:0016021	GO:000467	NA	K04730//i	
101267289	2413	3.5	8.81	1.3317871	3.35E-49	Up	1.90E-50	LOC101267289	PREDICTED: Sol:NA	GO:004687	NA	NA	
101249287	1130	85.17	214.34	1.3314839	0	Up	0	LOC101249287	PREDICTED: Sol:GO:0005838	NA	GO:000656	K03031//	
101261631	2530	0.37	0.93	1.3297054	8.63E-07	Up	2.26E-07	LOC101261631	PREDICTED: Sol:NA	GO:000425	NA	K12619//	
109120511	1126	0.43	1.08	1.3286227	0.000675	Up	0.000246	LOC109120511	PREDICTED: Sol:NA	NA	NA	K03006//	
101261158	1945	9	22.58	1.3270486	1.22E-105	Up	3.57E-107	LOC101261158	PREDICTED: Sol:GO:0009507	GO:000816	GO:001935	K02303//	
101243919	909	3.1	7.76	1.3237884	3.44E-17	Up	4.60E-18	LOC101243919	PREDICTED: Sol:GO:0005634	GO:000552	GO:003256	K20496//l	
109118704	590	3.14	7.86	1.3237648	3.65E-11	Up	6.80E-12	LOC109118704	PREDICTED: Sol:NA	NA	GO:000975	K14488//	
101252601	6534	24.64	61.62	1.3223964	0	Up	0	LOC101252601	PREDICTED: Sol:NA	NA	GO:000975	K03255//	
101265617	1109	177.45	443.69	1.3221395	0	Up	0	LOC101265617	PREDICTED: Sol:GO:0010287	GO:003146	GO:000975	K08909//l	
104645933	1669	0.48	1.2	1.3219281	1.22E-05	Up	3.56E-06	LOC104645933	PREDICTED: Sol:GO:0016021	NA	NA	NA	
543582	1146	71.17	177.43	1.3179088	0	Up	0	EXP2	Solanum lycopersi	GO:0016020	NA	GO:000966	K20628//e
101254918	933	277.41	690.17	1.314932	0	Up	0	LOC101254918	PREDICTED: Sol:GO:0005618	GO:004575	NA	NA	
100125906	1425	43.58	108.37	1.3142273	0	Up	0	ODD	Solanum lycopersi	NA	GO:004687	NA	K06892//f
101247176	3257	0.33	0.82	1.3131579	1.60E-07	Up	3.92E-08	LOC101247176	PREDICTED: Sol:GO:0005634	GO:004356	GO:000635	K14494//	
101244240	2770	0.97	2.41	1.3129765	3.19E-16	Up	4.48E-17	LOC101244240	PREDICTED: Sol:NA	NA	NA	K12619//	
101248862	710	29.64	73.6	1.3121603	2.15E-117	Up	5.80E-119	LOC101248862	PREDICTED: Sol:GO:0016021	NA	NA	NA	
101252626	1178	25.59	63.53	1.3118579	4.22E-173	Up	7.91E-175	LOC101252626	PREDICTED: Sol:NA	NA	GO:008016	K04733//i	
101243657	1124	0.92	2.28	1.3093281	7.43E-07	Up	1.94E-07	LOC101243657	PREDICTED: Sol:NA	NA	NA	K13070//	
104645884	982	629.08	1557.18	1.3076203	0	Up	0	Cab-1A	Solanum lycopersi	GO:0010287	GO:003146	GO:000975	K08912//l
101248844	872	1.95	4.82	1.305559	1.75E-10	Up	3.43E-11	LOC101248844	PREDICTED: Sol:NA	NA	GO:000975	K14488//	
101252871	2766	0.51	1.26	1.3048546	2.82E-09	Up	6.04E-10	LOC101252871	PREDICTED: Sol:NA	GO:004355	GO:000695	K19613//l	
101258235	3895	0.17	0.42	1.3048546	4.25E-05	Up	1.32E-05	LOC101258235	PREDICTED: Sol:NA	GO:001678	GO:004687	K01099//i	
101260157	1584	0.62	1.53	1.3031915	1.29E-06	Up	3.43E-07	LOC101260157	PREDICTED: Sol:GO:0043231	GO:000372	GO:000945	K19589//r	
101251463	3475	1.12	2.76	1.3011695	2.01E-23	Up	2.09E-24	LOC101251463	PREDICTED: Sol:GO:0016021	GO:000521	NA	K14638//s	
101258284	885	24.76	60.86	1.297483	9.10E-121	Up	2.42E-122	LOC101258284	PREDICTED: Sol:GO:0016021	NA	NA	NA	
101248254	687	128.79	316.54	1.2973672	0	Up	0	LOC101248254	Solanum lycopersi	GO:0005576	NA	NA	K03860//
101247257	552	3.11	7.64	1.2966581	5.04E-10	Up	1.02E-10	LOC101247257	PREDICTED: Sol:GO:0005576	GO:000825	GO:000961	NA	
101260372	2271	10.63	26.04	1.2925879	7.92E-138	Up	1.87E-139	LOC101260372	PREDICTED: Sol:NA	NA	NA	K17535//s	
101250945	1927	1.25	3.06	1.2916036	1.82E-14	Up	2.80E-15	LOC101250945	PREDICTED: Sol:GO:0005634	GO:004356	GO:001021	K18835//	
101267047	1396	3.54	8.66	1.2906177	3.59E-28	Up	3.22E-29	LOC101267047	PREDICTED: Sol:GO:0005634	GO:000552	GO:003256	K15601//l	
101259886	1792	8.29	20.28	1.2906136	1.19E-83	Up	4.27E-85	LOC101259886	PREDICTED: Sol:GO:0016021	GO:000467	GO:000646	K04733//i	
101252930	727	2.04	4.99	1.2904707	7.51E-09	Up	1.66E-09	LOC101252930	PREDICTED: Sol:NA	NA	GO:000975	K14488//	
101247792	717	38.56	94.11	1.287243	1.15E-143	Up	2.57E-145	LOC101247792	PREDICTED: Sol:GO:0005685	GO:000372	GO:000025	K11088//s	
104648148	580	10.78	26.24	1.2834105	4.53E-33	Up	3.55E-34	LOC104648148	PREDICTED: Sol:NA	NA	NA	NA	
101257526	2017	2.52	6.12	1.2801079	8.92E-29	Up	7.86E-30	LOC101257526	PREDICTED: Sol:GO:0016021	GO:000842	GO:190165	K05350//	

101257576	2504	41.83	101.51	1.279012	0	Up	0	LOC101257576	PREDICTED: Sol:NA	NA	NA	K17275//	
101252446	1278	10.79	26.15	1.2771161	1.23E-73	Up	4.94E-75	LOC101252446	PREDICTED: Sol:NA	NA	GO:005126	K21919//	
544110	1752	16.08	38.93	1.2756149	2.14E-153	Up	4.43E-155	AMT1-2	Solanum lycopersi	GO:0016021	GO:000851	GO:000962	K03320//a
543609	3037	26.87	65.05	1.2755527	0	Up	0	TBG3	Solanum lycopersi	NA	GO:003022	GO:000597	K12309//
101244055	627	40.36	97.58	1.2736593	1.13E-129	Up	2.82E-131	LOC101244055	Solanum lycopersi	GO:0043231	GO:005153	GO:005507	NA
101249800	2904	1.71	4.12	1.268648	2.35E-27	Up	2.15E-28	LOC101249800	PREDICTED: Sol:GO:0005759	GO:004687	GO:000662	K01410//	
101255140	1565	18.13	43.65	1.2676027	1.22E-151	Up	2.55E-153	LOC101255140	PREDICTED: Sol:GO:0005634	GO:000377	GO:000701	K10364//c	
101254923	1233	22.68	54.54	1.2658941	1.66E-147	Up	3.61E-149	LOC101254923	PREDICTED: Sol:NA	NA	NA	K10268//	
101268070	3676	0.47	1.13	1.2655901	1.38E-10	Up	2.67E-11	LOC101268070	PREDICTED: Sol:GO:0016021	NA	GO:000681	K05740//	
101249141	3473	33.59	80.65	1.2636427	0	Up	0	AGO2A	Solanum lycopersi	NA	GO:000367	GO:003102	K11593//e
101264349	1215	0.63	1.51	1.2611248	5.56E-05	Up	1.76E-05	LOC101264349	PREDICTED: Sol:GO:0005634	GO:000098	GO:000635	K09422//t	
101247488	694	28.01	67.12	1.2608007	3.49E-98	Up	1.09E-99	LOC101247488	PREDICTED: Sol:NA	GO:000482	NA	K01051//	
101260536	1792	0.31	0.74	1.2552571	0.000869	Up	0.000322	LOC101260536	PREDICTED: Sol:GO:0016020	GO:001676	GO:004452	K20619//c	
101244290	2122	2.13	5.08	1.2539751	1.34E-24	Up	1.33E-25	LOC101244290	PREDICTED: Sol:NA	GO:000556	NA	K13448//c	
101254040	1820	13.69	32.65	1.2539605	2.83E-130	Up	7.02E-132	LOC101254040	PREDICTED: Sol:NA	GO:001649	NA	K10767//	
101244145	1493	26.98	64.24	1.2515815	6.90E-208	Up	1.12E-209	LOC101244145	PREDICTED: Sol:NA	NA	NA	NA	
101265927	1008	0.98	2.33	1.2494763	5.42E-06	Up	1.53E-06	LOC101265927	PREDICTED: Sol:GO:0048046	GO:001676	GO:001041	K08235//	
101246552	4534	0.24	0.57	1.2479275	9.55E-07	Up	2.51E-07	LOC101246552	PREDICTED: Sol:NA	GO:004353	GO:000716	K22038//	
101250847	803	1.2	2.85	1.2479275	9.39E-06	Up	2.71E-06	LOC101250847	PREDICTED: Sol:NA	NA	GO:000972	K14488//	
778300	488	5.02	11.91	1.2464141	1.85E-12	Up	3.18E-13	LOC778300	Solanum lycopersi	NA	GO:004687	NA	NA
101248037	1119	52.21	123.48	1.2418793	6.13E-292	Up	7.10E-294	LOC101248037	PREDICTED: Sol:GO:0016021	GO:000521	NA	K09872//a	
101055515	987	12.94	30.52	1.2379173	1.74E-63	Up	7.95E-65	LOC101055515	Solanum lycopersi	NA	GO:004698	NA	K18485//
101244203	824	3.71	8.75	1.2378638	8.84E-16	Up	1.26E-16	LOC101244203	PREDICTED: Sol:GO:0016021	NA	NA	NA	
101257365	1072	0.56	1.32	1.2370392	0.000524	Up	0.000189	LOC101257365	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	NA	
101249624	2095	36.39	85.7	1.2357532	0	Up	0	4CL	Solanum lycopersi	NA	GO:001626	GO:000812	K01904//
101249503	732	3.52	8.28	1.2340553	2.73E-13	Up	4.46E-14	LOC101249503	PREDICTED: Sol:NA	NA	GO:000972	K14488//	
101266576	1543	43.42	101.91	1.230864	0	Up	0	LOC101266576	PREDICTED: Sol:NA	GO:000372	GO:000635	K03124//t	
101264755	1047	0.75	1.76	1.2306129	8.36E-05	Up	2.70E-05	LOC101264755	PREDICTED: Sol:NA	NA	NA	NA	
101250276	2149	30.53	71.52	1.2281191	0	Up	0	LOC101250276	PREDICTED: Sol:NA	GO:000817	GO:000635	NA	
101249679	3218	2.17	5.08	1.2271335	4.49E-37	Up	3.22E-38	LOC101249679	PREDICTED: Sol:NA	NA	NA	K21842//	
101256557	1848	55.82	130.61	1.2264113	0	Up	0	LOC101256557	PREDICTED: Sol:NA	GO:000402	GO:000016	K13811//	
101263191	962	0.62	1.45	1.2257128	0.000826	Up	0.000305	LOC101263191	PREDICTED: Sol:NA	NA	NA	NA	
544018	610	1.16	2.71	1.224168	0.000274	Up	9.53E-05	PRO1	Solanum lycopersi	GO:0005856	GO:000377	GO:004298	K05759//
101259387	2146	7.07	16.5	1.2226839	1.43E-72	Up	5.85E-74	LOC101259387	PREDICTED: Sol:GO:0043231	GO:008002	GO:000812	K21373//	
101259032	884	1.7	3.96	1.2199657	4.27E-08	Up	9.97E-09	LOC101259032	PREDICTED: Sol:NA	NA	NA	K21994//	
101248942	1060	195.86	455.23	1.2167728	0	Up	0	LOC101248942	PREDICTED: Sol:GO:0009507	GO:004687	NA	K03940//	
101253752	2312	0.31	0.72	1.2157287	0.000152	Up	5.07E-05	LOC101253752	PREDICTED: Sol:NA	GO:000367	NA	K14794//r	

543794	1287	40.17	93.14	1.2132824	1.23E-245	Up	1.72E-247	exp11	Solanum lycopersi	GO:0016020	NA	GO:000966	K20628//e
101266110	1986	1.22	2.82	1.208814	1.63E-12	Up	2.78E-13	LOC101266110	PREDICTED: Sol:	NA	NA	NA	K13422//t
100134915	1523	4.66	10.77	1.2086164	1.96E-34	Up	1.48E-35	LOC100134915	Solanum lycopersi	GO:0016021	GO:001671	GO:000662	K10257//a
101251551	791	0.94	2.17	1.2069624	0.000236	Up	8.13E-05	LOC101251551	PREDICTED: Sol:	NA	GO:006162	GO:004316	K19045//
101263369	3279	0.26	0.6	1.2064509	6.14E-05	Up	1.95E-05	LOC101263369	PREDICTED: Sol:	NA	GO:004687	GO:000601	K01191//a
101252437	2799	1.2	2.76	1.2016339	1.00E-16	Up	1.37E-17	LOC101252437	PREDICTED: Sol:	GO:0016021	GO:000552	NA	K13420//
101244088	931	36.95	84.77	1.1979794	4.81E-156	Up	9.77E-158	LOC101244088	PREDICTED: Sol:	GO:0009507	NA	NA	NA
101256323	2600	11.61	26.63	1.1976845	4.20E-142	Up	9.54E-144	LOC101256323	PREDICTED: Sol:	GO:0005829	GO:005066	GO:000205	K03495//t
101265239	2824	0.41	0.94	1.1970368	1.30E-06	Up	3.46E-07	LOC101265239	PREDICTED: Sol:	NA	NA	NA	K14963//
101243687	2399	13.03	29.84	1.1954105	3.52E-145	Up	7.77E-147	LOC101243687	PREDICTED: Sol:	NA	GO:000367	NA	K05857//
101256329	1317	1.49	3.41	1.1944594	5.22E-10	Up	1.06E-10	LOC101256329	PREDICTED: Sol:	NA	NA	NA	NA
101256495	2283	0.28	0.64	1.1926451	0.000453	Up	0.000162	LOC101256495	PREDICTED: Sol:	GO:0016021	NA	NA	K15503//s
101248885	1240	152.16	347.43	1.1911332	0	Up	0	LOC101248885	PREDICTED: Sol:	NA	NA	NA	NA
101267437	1035	93.26	212.93	1.1910489	0	Up	0	LOC101267437	PREDICTED: Sol:	NA	NA	NA	K12603//
101258036	2334	1.13	2.58	1.1910483	3.66E-13	Up	6.02E-14	LOC101258036	PREDICTED: Sol:	GO:0016021	GO:001672	GO:000662	K15397//
101250944	1211	9.94	22.69	1.1907389	4.40E-55	Up	2.28E-56	LOC101250944	PREDICTED: Sol:	NA	NA	NA	K14496//a
101246621	757	6.8	15.52	1.1905219	1.89E-23	Up	1.97E-24	LOC101246621	PREDICTED: Sol:	NA	GO:004698	NA	K17693//
101260921	1300	1.24	2.83	1.1904619	2.84E-08	Up	6.56E-09	LOC101260921	PREDICTED: Sol:	GO:0016021	NA	NA	K11153//r
101251647	600	4.01	9.15	1.1901695	2.57E-11	Up	4.74E-12	LOC101251647	PREDICTED: Sol:	NA	NA	NA	NA
101264832	2292	4.01	9.14	1.1885919	5.12E-43	Up	3.29E-44	LOC101264832	PREDICTED: Sol:	NA	NA	NA	K13140//i
101252654	2044	15.03	34.22	1.1869948	1.93E-140	Up	4.48E-142	LOC101252654	PREDICTED: Sol:	GO:0031982	NA	GO:001615	NA
101251995	1158	1.78	4.05	1.1860447	4.02E-10	Up	8.08E-11	LOC101251995	PREDICTED: Sol:	NA	NA	NA	NA
109121043	467	1.61	3.66	1.184783	0.000409	Up	0.000145	LOC109121043	PREDICTED: Sol:	NA	NA	NA	NA
101248362	955	17.04	38.73	1.1845262	2.04E-72	Up	8.36E-74	LOC101248362	PREDICTED: Sol:	GO:0031977	NA	NA	NA
101250583	936	156.83	356.45	1.1844982	0	Up	0	LOC101250583	PREDICTED: Sol:	GO:0005576	GO:004572	NA	NA
101266372	1413	3.3	7.49	1.1824997	2.76E-21	Up	3.10E-22	LOC101266372	PREDICTED: Sol:	NA	GO:000367	GO:000632	K01872//a
101252466	1736	3.08	6.99	1.1823621	6.09E-25	Up	5.97E-26	LOC101252466	PREDICTED: Sol:	GO:0005623	GO:001667	GO:001941	K05907//a
104648440	745	1.86	4.22	1.1819404	5.54E-07	Up	1.43E-07	LOC104648440	PREDICTED: Sol:	NA	NA	NA	NA
101250755	774	1.42	3.22	1.1811698	1.03E-05	Up	3.00E-06	LOC101250755	PREDICTED: Sol:	NA	NA	NA	NA
101266496	1450	3.27	7.4	1.1782346	6.77E-22	Up	7.45E-23	LOC101266496	PREDICTED: Sol:	NA	NA	NA	NA
101250059	2801	9.13	20.56	1.1711535	1.55E-114	Up	4.24E-116	LOC101250059	PREDICTED: Sol:	NA	NA	NA	K10352//
101255525	348	7.61	17.13	1.1705568	3.77E-11	Up	7.04E-12	LOC101255525	PREDICTED: Sol:	GO:0016021	NA	NA	NA
101261645	574	53.04	119.32	1.1696832	5.47E-93	Up	1.77E-94	LOC101261645	PREDICTED: Sol:	GO:0000439	NA	GO:000632	K10845//
101245631	868	50.99	114.5	1.1670614	1.95E-187	Up	3.46E-189	LOC101245631	PREDICTED: Sol:	NA	NA	NA	NA
101246802	3530	6.95	15.57	1.1636841	1.19E-108	Up	3.37E-110	LOC101246802	PREDICTED: Sol:	GO:0005634	GO:000828	NA	K09338//
101252293	2266	6.86	15.33	1.1600772	7.97E-68	Up	3.45E-69	LOC101252293	PREDICTED: Sol:	GO:0016021	NA	GO:000962	K17085//t
104648350	801	2.48	5.54	1.1595459	3.29E-09	Up	7.08E-10	LOC104648350	PREDICTED: Sol:	NA	NA	NA	K10393//

100529103	1893	25.91	57.72	1.1555623	9.67E-210	Up	1.55E-211	LOC100529103	Solanum lycopersiGO:0005886.GO:010212GO:000597K12373//
101257650	1182	17.73	39.47	1.154564	9.26E-89	Up	3.11E-90	LOC101257650	PREDICTED: Sol:GO:0005737.GO:004687GO:00300(K17046//
101261316	1679	0.93	2.07	1.1543281	1.16E-07	Up	2.81E-08	LOC101261316	PREDICTED: Sol:GO:0005576.GO:000462GO:000597K01213//
101265639	1691	0.49	1.09	1.1534745	0.000171	Up	5.77E-05	LOC101265639	PREDICTED: Sol:NA GO:000552NA K08900//
101252151	1011	79.32	176.07	1.1503925	0	Up	0	LOC101252151	PREDICTED: Sol:GO:0010287.GO:00314(GO:00097(K08908//I
104648418	2253	1.07	2.37	1.1472763	5.11E-11	Up	9.66E-12	LOC104648418	PREDICTED: Sol:GO:0016021.NA NA K17046//
101262127	713	19.48	43.13	1.146698	3.84E-56	Up	1.96E-57	LOC101262127	PREDICTED: Sol:NA NA GO:000692NA
101258238	1820	39.23	86.84	1.1464024	6.42E-299	Up	7.21E-301	LOC101258238	PREDICTED: Sol:GO:0016020.GO:00167(GO:004452K07409//c
101244302	725	13.42	29.68	1.1451064	1.46E-39	Up	9.92E-41	LOC101244302	PREDICTED: Sol:GO:0016021.NA NA NA
101255499	1667	7.79	17.22	1.1443899	1.26E-54	Up	6.57E-56	LOC101255499	PREDICTED: Sol:NA GO:000367NA K20557//t
543991	2192	16.15	35.68	1.1435815	1.65E-148	Up	3.56E-150	GP1	Solanum lycopersiGO:0048046.NA GO:000982K06699//
101254978	626	6.52	14.39	1.1421227	7.64E-17	Up	1.04E-17	LOC101254978	PREDICTED: Sol:NA NA NA NA
104646222	2605	0.39	0.86	1.1408625	3.92E-05	Up	1.22E-05	LOC104646222	PREDICTED: Sol:NA GO:000367NA K21026//a
101268886	1622	1.67	3.68	1.1398577	4.31E-12	Up	7.56E-13	LOC101268886	PREDICTED: Sol:NA NA NA K19307//
101245640	1822	8.44	18.59	1.1392119	4.27E-64	Up	1.92E-65	LOC101245640	PREDICTED: Sol:NA NA NA NA
101262607	1808	0.8	1.76	1.1375035	5.79E-07	Up	1.50E-07	LOC101262607	PREDICTED: Sol:GO:0016021.GO:000848NA K16903//
101251000	1668	0.45	0.99	1.1375035	0.000503	Up	0.000181	LOC101251000	PREDICTED: Sol:GO:0005634.GO:000367NA K14794//r
101251196	573	84.22	184.9	1.1345104	1.45E-184	Up	2.59E-186	LOC101251196	PREDICTED: Sol:GO:0005665.GO:000385GO:00063(K03007//
101247764	1004	1.29	2.83	1.133431	5.24E-06	Up	1.48E-06	LOC101247764	PREDICTED: Sol:NA NA NA NA
101267980	1986	47.64	104.4	1.1318764	0	Up	0	LOC101267980	PREDICTED: Sol:NA NA NA K19613//I
101250846	1493	47.52	103.98	1.1296993	7.08E-285	Up	8.52E-287	LOC101250846	PREDICTED: Sol:GO:0005829.GO:00048(GO:00195(K01803//t
544152	1943	249.45	544.42	1.1259694	0	Up	0	LOC544152	Solanum lycopersiGO:0009507.GO:000382GO:000942K01626//
101248871	2256	6.94	15.13	1.1244044	1.59E-63	Up	7.22E-65	LOC101248871	PREDICTED: Sol:NA GO:00506(GO:000922K11752//
544158	1373	7.12	15.52	1.1241794	2.00E-39	Up	1.37E-40	LOC544158	PREDICTED: Sol:NA GO:000515GO:00096(K03006//
101266629	1605	117.62	256.01	1.1220668	0	Up	0	LOC101266629	PREDICTED: Sol:GO:0005737.GO:004687GO:000597K03841//f
101268821	2469	6.19	13.46	1.1206671	9.02E-62	Up	4.21E-63	LOC101268821	PREDICTED: Sol:GO:0043231.GO:000372GO:000942K19090//
101244999	507	1.79	3.89	1.1198106	0.000261	Up	9.03E-05	LOC101244999	PREDICTED: Sol:GO:0016021.NA GO:004852NA
101254427	1268	3.2	6.95	1.1189411	9.97E-17	Up	1.37E-17	LOC101254427	PREDICTED: Sol:GO:0005737.GO:00055(GO:001931K00469//i
104646795	435	4.81	10.44	1.1180129	2.01E-08	Up	4.58E-09	LOC104646795	PREDICTED: Sol:NA NA NA K01868//t
101258047	1574	6.4	13.88	1.1168638	4.10E-40	Up	2.76E-41	LOC101258047	PREDICTED: Sol:NA NA NA K15502//s
104645726	447	32.74	70.72	1.111064	2.41E-52	Up	1.30E-53	LOC104645726	PREDICTED: Sol:GO:0016021.NA NA NA
101265090	2729	0.63	1.36	1.1101829	9.93E-08	Up	2.39E-08	LOC101265090	PREDICTED: Sol:NA GO:000382GO:000812K00666//f
101254141	5090	0.89	1.92	1.1092291	1.18E-16	Up	1.62E-17	LOC101254141	PREDICTED: Sol:GO:0005643.NA GO:000681K14297//
101267742	1618	1.72	3.71	1.1090106	6.61E-11	Up	1.26E-11	LOC101267742	PREDICTED: Sol:GO:0016021.NA GO:005511K03189//
101254886	1609	7.28	15.69	1.107835	8.90E-46	Up	5.38E-47	LOC101254886	PREDICTED: Sol:NA NA NA K03063//
543809	1952	173.71	373.63	1.1049295	0	Up	0	LOC543809	Solanum lycopersiGO:0005737.GO:000451GO:000862K01858//
101260945	856	1.67	3.59	1.1041357	2.57E-06	Up	7.04E-07	LOC101260945	PREDICTED: Sol:GO:0016021.NA NA NA

101249313	1241	36.21	77.74	1.1022689	2.39E-169	Up	4.55E-171	LOC101249313	PREDICTED: Sol:GO:0005840.GO:000372GO:003252K02935//
100037489	1648	2.18	4.68	1.1021804	1.51E-14	Up	2.31E-15	CYP85A1	Solanum lycopersiGO:0016021.GO:00167(GO:001026K09590//
101251724	1182	11.54	24.77	1.1019506	9.46E-52	Up	5.13E-53	LOC101251724	PREDICTED: Sol:GO:0016021.NA GO:000962K09422//t
104649338	1033	7.11	15.24	1.0999414	4.75E-28	Up	4.28E-29	LOC104649338	PREDICTED: Sol:NA NA NA K01868//t
101268119	2017	11.47	24.58	1.0996195	2.28E-88	Up	7.71E-90	LOC101268119	PREDICTED: Sol:GO:0016021.GO:000521GO:000682K14638//s
101247340	2205	7.47	16	1.0988918	3.41E-63	Up	1.56E-64	LOC101247340	PREDICTED: Sol:NA GO:001616GO:000027K01177//
101260340	1123	37.55	80.24	1.0955087	9.31E-155	Up	1.92E-156	LOC101260340	PREDICTED: Sol:GO:0005623.GO:004712GO:000666K03671//t
101265433	1593	3.52	7.52	1.0951572	8.77E-22	Up	9.67E-23	LOC101265433	PREDICTED: Sol:NA NA NA NA
101255125	1673	22.79	48.67	1.0946319	9.80E-143	Up	2.21E-144	LOC101255125	PREDICTED: Sol:NA GO:006162GO:003012K15685//
101257036	1527	12.12	25.88	1.0944479	1.84E-69	Up	7.76E-71	LOC101257036	PREDICTED: Sol:NA NA NA K11247//e
101251848	670	5.54	11.82	1.0932722	4.60E-14	Up	7.24E-15	LOC101251848	PREDICTED: Sol:GO:0005623.GO:001502GO:004542K03676//
101252202	1305	7.49	15.98	1.0932298	7.07E-37	Up	5.08E-38	LOC101252202	PREDICTED: Sol:GO:0005783.GO:006162GO:003042K20417//
101244251	2796	0.6	1.28	1.0931094	1.79E-07	Up	4.42E-08	LOC101244251	PREDICTED: Sol:GO:0043231.GO:000372GO:000942K01176//a
101246107	1281	5.48	11.69	1.0930271	4.60E-26	Up	4.36E-27	LOC101246107	PREDICTED: Sol:NA GO:000028GO:001987K06133//
101264842	2160	34.16	72.83	1.0922249	3.04E-276	Up	3.78E-278	LOC101264842	PREDICTED: Sol:GO:0009941.NA GO:000642K10578//
101243647	2605	9.39	19.98	1.0893595	8.84E-92	Up	2.90E-93	LOC101243647	PREDICTED: Sol:GO:0005634.GO:000367GO:000632K16670//
101250679	880	14.79	31.47	1.0893551	2.47E-47	Up	1.44E-48	LOC101250679	PREDICTED: Sol:NA NA NA NA
101252854	812	6.45	13.72	1.0889094	1.49E-19	Up	1.80E-20	LOC101252854	PREDICTED: Sol:GO:0005623.GO:004712GO:000666K03671//t
101246156	717	1.58	3.36	1.0885367	5.27E-05	Up	1.66E-05	LOC101246156	PREDICTED: Sol:GO:0016021.NA NA NA
101266147	752	50.6	107.59	1.0883347	4.93E-135	Up	1.18E-136	LOC101266147	PREDICTED: Sol:NA NA NA K19363//I
543720	1892	91.65	194.8	1.0877869	0	Up	0	mek1	Solanum lycopersiGO:0005737.GO:000467GO:002301K20603//
101266182	1023	608.22	1291.99	1.0869297	0	Up	0	LOC101266182	PREDICTED: Sol:GO:0010287.GO:00314(GO:000976K08912//I
100736484	1875	0.68	1.44	1.0824622	1.20E-05	Up	3.49E-06	LOC100736484	Solanum lycopersiGO:0016021.GO:000467NA K13429//c
101261120	1131	5.56	11.73	1.0770462	4.15E-23	Up	4.37E-24	LOC101261120	PREDICTED: Sol:GO:0005737.GO:000472NA K18045//t
101261588	1099	281.83	594.43	1.0766817	0	Up	0	LOC101261588	Solanum lycopersi NA NA NA K00430//
101266036	901	0.93	1.96	1.075551	0.000636	Up	0.000232	LOC101266036	PREDICTED: Sol:NA NA NA K15697//
101253663	1875	16.05	33.79	1.0740231	2.72E-108	Up	7.77E-110	LOC101253663	PREDICTED: Sol:GO:0005743.GO:001652GO:000681K15119//s
544272	1281	33.8	71.06	1.0720144	1.16E-152	Up	2.43E-154	XTH1	Solanum lycopersiGO:0048046.GO:001676GO:001041K08235//
101245586	658	103.42	217.06	1.0695787	9.91E-229	Up	1.50E-230	LOC101245586	Solanum lycopersi NA NA NA K11855//
101264365	682	7.11	14.92	1.0693261	2.48E-17	Up	3.30E-18	LOC101264365	PREDICTED: Sol:NA NA NA K01180//e
101265605	986	791.45	1659.11	1.0678394	0	Up	0	LOC101265605	PREDICTED: Sol:GO:0010287.GO:00314(GO:000976K08912//I
104644351	1066	1.8	3.77	1.0665676	1.46E-07	Up	3.57E-08	LOC104644351	PREDICTED: Sol:NA NA NA K00799//
101266952	838	220.97	461.75	1.0632614	0	Up	0	LOC101266952	PREDICTED: Sol:GO:0005840.GO:000372GO:000641K02879//I
101257878	1373	0.68	1.42	1.0622843	0.000775	Up	0.000285	LOC101257878	PREDICTED: Sol:GO:0005576.GO:004572NA NA
101268276	1811	28.79	60.12	1.0622772	7.72E-182	Up	1.40E-183	LOC101268276	PREDICTED: Sol:GO:0016020.GO:001676GO:004452K14985//
101253234	730	2.09	4.36	1.0608252	4.56E-06	Up	1.28E-06	LOC101253234	PREDICTED: Sol:NA NA GO:000972K14488//
101261433	898	263.04	548.38	1.0598938	0	Up	0	LOC101261433	PREDICTED: Sol:GO:0005840.GO:000372GO:000641K02895//I

104648378	750	1.19	2.48	1.0593785	0.000561	Up	0.000203	LOC104648378	PREDICTED: Sol:NA	GO:004698	NA	NA	
101249376	1040	39.31	81.75	1.0563224	7.34E-138	Up	1.73E-139	LOC101249376	PREDICTED: Sol:GO:0005686	GO:000372	GO:000035	K11094//	
101248252	3252	4.65	9.67	1.0562852	2.29E-53	Up	1.22E-54	LOC101248252	PREDICTED: Sol:GO:0016021	GO:000467	NA	K00924//	
101249898	2078	0.63	1.31	1.0561431	1.58E-05	Up	4.66E-06	LOC101249898	PREDICTED: Sol:NA	GO:000552	NA	K08900//	
101258063	1586	1.54	3.2	1.0551416	3.03E-09	Up	6.48E-10	LOC101258063	PREDICTED: Sol:NA	GO:000372	NA	K12741//	
101265791	943	1.83	3.8	1.0541558	4.77E-06	Up	1.34E-06	LOC101265791	PREDICTED: Sol:GO:0005730	GO:004302	GO:000047	K03264//t	
101249132	2783	10.76	22.32	1.0526589	7.94E-103	Up	2.37E-104	LOC101249132	PREDICTED: Sol:GO:0005737	GO:000467	GO:002301	K04421//	
101266763	1303	0.69	1.43	1.0513469	0.000494	Up	0.000177	LOC101266763	PREDICTED: Sol:GO:0016021	NA	NA	K11649//	
544223	1783	43.37	89.87	1.0511422	1.21E-262	Up	1.59E-264	LOC544223	Solanum lycopersi	GO:0005777	GO:000418	GO:001972	K16296//s
101247889	617	68.45	141.82	1.0509386	1.47E-135	Up	3.52E-137	LOC101247889	PREDICTED: Sol:GO:0009523	NA	GO:001597	K02259//c	
101255295	1118	99.98	206.9	1.0492222	0	Up	0	LOC101255295	PREDICTED: Sol:GO:0015934	GO:000372	GO:000641	K02876//l	
101256723	992	157.12	324.79	1.0476404	0	Up	0	LOC101256723	PREDICTED: Sol:GO:0009535	GO:000551	GO:001597	K02717//	
543511	1280	80.87	167.17	1.0476395	0	Up	0	LOC543511	Solanum lycopersi	GO:0048046	GO:001676	GO:001041	K08235//
101262309	1450	2.75	5.68	1.0464593	3.25E-14	Up	5.07E-15	LOC101262309	PREDICTED: Sol:NA	NA	NA	NA	
101258880	1595	8.56	17.68	1.0464356	1.95E-46	Up	1.16E-47	LOC101258880	PREDICTED: Sol:GO:0005634	GO:000367	GO:000632	K09286//	
101266928	684	22.17	45.78	1.0461087	3.76E-49	Up	2.13E-50	LOC101266928	PREDICTED: Sol:NA	NA	GO:000692	K10745//r	
101268623	1094	82.94	171.14	1.045037	4.07E-298	Up	4.61E-300	aquaporin	Solanum lycopersi	GO:0005887	GO:001522	GO:003422	K09872//a
101245895	920	1.12	2.31	1.0443941	0.000226	Up	7.75E-05	LOC101245895	PREDICTED: Sol:NA	NA	GO:000972	K14488//	
101243986	740	2.22	4.57	1.0416345	3.11E-06	Up	8.59E-07	LOC101243986	PREDICTED: Sol:NA	NA	NA	NA	
101252082	1516	28.22	58.08	1.0413235	1.12E-141	Up	2.56E-143	LOC101252082	PREDICTED: Sol:NA	GO:004687	GO:004545	K01728//	
101266892	2058	1.21	2.49	1.0411387	3.47E-09	Up	7.46E-10	LOC101266892	PREDICTED: Sol:NA	GO:000552	NA	K08900//	
101243878	693	20.36	41.89	1.0408683	2.62E-45	Up	1.60E-46	LOC101243878	PREDICTED: Sol:NA	NA	NA	NA	
101263716	745	59.5	122.3	1.0394628	7.73E-141	Up	1.78E-142	LOC101263716	PREDICTED: Sol:NA	NA	NA	K11855//	
101055598	1719	7.83	16.09	1.0390801	1.22E-43	Up	7.67E-45	SICKX1	PREDICTED: Sol:NA	GO:001661	GO:000965	K00279//c	
101252569	1654	0.55	1.13	1.0388192	0.00033	Up	0.000116	LOC101252569	PREDICTED: Sol:GO:0005634	GO:000367	NA	K18490//	
104645946	291	96.53	198.07	1.0369611	4.88E-79	Up	1.83E-80	LOC104645946	PREDICTED: Sol:NA	NA	NA	NA	
100736446	1759	134.94	276.47	1.0348049	0	Up	0	CYP51	Solanum lycopersi	GO:0005783	GO:001676	GO:001612	K05917//s
101252642	1397	1.1	2.25	1.0324215	1.85E-05	Up	5.52E-06	LOC101252642	PREDICTED: Sol:NA	GO:001645	NA	K13070//	
101251933	972	3.6	7.36	1.0317089	1.51E-10	Up	2.95E-11	LOC101251933	PREDICTED: Sol:NA	GO:000382	NA	K08695//a	
101267670	3511	0.46	0.94	1.0310269	1.24E-06	Up	3.29E-07	LOC101267670	PREDICTED: Sol:GO:0005886	GO:004687	NA	K10615//	
101267965	2368	46.09	94.16	1.0306606	0	Up	0	LOC101267965	PREDICTED: Sol:GO:0016021	GO:004687	NA	K16075//	
101259820	954	14.84	30.29	1.0293505	3.05E-45	Up	1.86E-46	LOC101259820	PREDICTED: Sol:GO:0005779	GO:004281	GO:004437	K13352//	
101263424	1351	1.24	2.53	1.0287973	2.26E-06	Up	6.15E-07	LOC101263424	PREDICTED: Sol:GO:0016021	NA	GO:005511	NA	
101266010	4227	1	2.04	1.0285692	6.52E-15	Up	9.77E-16	LOC101266010	PREDICTED: Sol:NA	NA	NA	K19613//l	
101244634	942	11.38	23.19	1.0270023	2.70E-34	Up	2.04E-35	LOC101244634	PREDICTED: Sol:GO:0016021	GO:000902	NA	K03006//	
101257983	2042	5.83	11.87	1.0257521	1.20E-38	Up	8.35E-40	LOC101257983	PREDICTED: Sol:NA	NA	NA	NA	
104648605	633	3.68	7.49	1.02526	5.27E-08	Up	1.24E-08	LOC104648605	Solanum lycopersi	NA	NA	NA	NA

101257993	2300	18.14	36.92	1.0252281	9.59E-135	Up	2.31E-136	LOC101257993	PREDICTED: Sol:NA	GO:000827	NA	K03452//	
101247979	1721	1.72	3.5	1.0249464	2.30E-10	Up	4.53E-11	LOC101247979	PREDICTED: Sol:NA	GO:000897	GO:000662	K00145//	
101251036	1599	13.49	27.45	1.0249158	2.59E-69	Up	1.10E-70	LOC101251036	PREDICTED: Sol:NA	GO:001672	GO:000812	K21371//c	
101263383	1442	24.68	50.19	1.0240576	3.80E-113	Up	1.05E-114	LOC101263383	PREDICTED: Sol:GO:0010598	GO:003022	GO:000977	NA	
101249445	1605	311.22	632.7	1.0235868	0	Up	0	LOC101249445	PREDICTED: Sol:NA	GO:001662	GO:000600	K05298//	
101265489	3448	5.63	11.43	1.0216186	4.66E-55	Up	2.42E-56	LOC101265489	PREDICTED: Sol:NA	NA	NA	K18046//t	
100136876	3478	6.72	13.64	1.0213105	4.07E-76	Up	1.58E-77	LOC100136876	Solanum lycopersi	GO:0005737	GO:000452	GO:001022	NA
101257548	3424	6.11	12.4	1.0210958	5.45E-68	Up	2.35E-69	LOC101257548	PREDICTED: Sol:GO:0005634	GO:000828	NA	K09338//	
101263526	553	9.67	19.6	1.0192659	1.18E-16	Up	1.62E-17	LOC101263526	PREDICTED: Sol:NA	NA	NA	NA	
101247017	2313	8.21	16.63	1.018334	1.34E-60	Up	6.33E-62	LOC101247017	PREDICTED: Sol:GO:0005622	GO:001682	NA	K09835//	
101247929	2540	1.96	3.97	1.0182854	7.99E-17	Up	1.09E-17	LOC101247929	PREDICTED: Sol:GO:0016021	GO:000552	NA	K04733//i	
101267554	720	2.86	5.79	1.0175482	3.67E-07	Up	9.29E-08	LOC101267554	PREDICTED: Sol:NA	NA	NA	NA	
101246730	1906	18.36	37.15	1.0167962	1.78E-110	Up	4.98E-112	LOC101246730	PREDICTED: Sol:NA	GO:000022	GO:000662	K05929//	
101249062	2065	17.42	35.2	1.0148308	2.78E-113	Up	7.64E-115	LOC101249062	PREDICTED: Sol:NA	GO:004687	NA	K11975//	
104645444	968	2.56	5.17	1.0140205	1.64E-08	Up	3.70E-09	LOC104645444	PREDICTED: Sol:NA	NA	NA	NA	
101256217	3243	18.55	37.45	1.0135465	1.65E-202	Up	2.71E-204	LOC101256217	PREDICTED: Sol:GO:0005739	GO:000482	GO:000642	K14164//	
101264273	1448	196.88	397.26	1.012767	0	Up	0	LOC101264273	Solanum lycopersi	GO:0005829	GO:004212	GO:000592	K03841//f
101266880	1354	4.11	8.29	1.0122337	4.92E-18	Up	6.34E-19	LOC101266880	PREDICTED: Sol:NA	NA	NA	K08081//t	
101262881	1451	15.99	32.23	1.0112342	5.81E-72	Up	2.40E-73	LOC101262881	PREDICTED: Sol:GO:0000151	NA	GO:004312	NA	
101247302	2351	17.59	35.42	1.0098087	7.31E-129	Up	1.83E-130	LOC101247302	PREDICTED: Sol:NA	GO:003022	GO:003022	K01179//e	
101246648	2006	1.7	3.42	1.0084616	3.86E-11	Up	7.22E-12	LOC101246648	PREDICTED: Sol:GO:0016021	NA	NA	K08150//	
101254500	523	19.7	39.53	1.0047523	5.74E-30	Up	4.87E-31	LOC101254500	PREDICTED: Sol:NA	NA	NA	K03860//	
101244316	1732	3.71	7.44	1.0038834	1.98E-20	Up	2.30E-21	LOC101244316	PREDICTED: Sol:NA	GO:001672	GO:000812	K17193//a	
101252838	2302	9.85	19.72	1.0014639	9.83E-70	Up	4.13E-71	LOC101252838	PREDICTED: Sol:NA	GO:000550	NA	K17871//	
543759	612	2.31	4.62	1	5.07E-05	Up	1.59E-05	fsm1	Solanum lycopersi	GO:0005634	GO:000367	NA	NA
101267897	2550	0.48	0.96	1	9.67E-05	Up	3.15E-05	LOC101267897	PREDICTED: Sol:NA	NA	NA	K09527//	
101268614	4435	0.5	1	1	1.11E-07	Up	2.69E-08	LOC101268614	PREDICTED: Sol:GO:0005871	GO:000552	GO:000701	K10400//	
104645696	785	2.98	5.96	1	9.33E-08	Up	2.24E-08	LOC104645696	PREDICTED: Sol:NA	GO:001672	NA	K21026//a	
101266908	2306	19.75	0.01	-10.94764	0	Down	0	LOC101266908	PREDICTED: Sol:NA	NA	NA	NA	
101251387	592	11.97	0.01	-10.22521	6.65E-69	Down	2.83E-70	LOC101251387	PREDICTED: Sol:NA	NA	NA	NA	
101247133	932	8.1	0.01	-9.661778	1.27E-76	Down	4.91E-78	LOC101247133	PREDICTED: Sol:GO:0005576	NA	NA	K20628//e	
101259487	999	6.67	0.01	-9.381543	6.48E-68	Down	2.80E-69	LOC101259487	PREDICTED: Sol:GO:0005829	NA	GO:000941	NA	
101256409	1058	17.7	0.03	-9.204571	1.73E-195	Down	2.96E-197	USP	Solanum lycopersi	NA	NA	GO:000692	NA
104647185	435	5.79	0.01	-9.17742	2.53E-23	Down	2.64E-24	LOC104647185	PREDICTED: Sol:NA	NA	NA	NA	
101257497	1342	5.18	0.01	-9.016808	4.50E-72	Down	1.86E-73	LOC101257497	PREDICTED: Sol:GO:0005886	GO:000467	GO:000642	K00924//	
101261225	691	4.15	0.01	-8.696968	3.12E-28	Down	2.79E-29	LOC101261225	PREDICTED: Sol:NA	NA	NA	K13993//	
101253038	540	258.53	0.78	-8.372642	0	Down	0	LE25	Solanum lycopersi	NA	NA	GO:000972	NA

101267532	745	3.21	0.01	-8.326429	1.03E-23	Down	1.05E-24	LOC101267532	PREDICTED: Sol:GO:0016021.GO:005152	NA	K11000//c
101262752	982	3.11	0.01	-8.280771	5.51E-31	Down	4.57E-32	LOC101262752	PREDICTED: Sol:NA	NA	NA
101251603	534	2.96	0.01	-8.209453	4.05E-15	Down	6.01E-16	LOC101251603	PREDICTED: Sol:GO:0016021.NA	NA	NA
101245172	1022	8.43	0.03	-8.134426	6.67E-86	Down	2.34E-87	LOC101245172	PREDICTED: Sol:NA	GO:000382	NA
101250895	688	2.08	0.01	-7.70044	3.80E-14	Down	5.94E-15	LOC101250895	PREDICTED: Sol:GO:0005576.NA	GO:001992	NA
101249653	621	2.01	0.01	-7.651052	3.34E-12	Down	5.81E-13	LOC101249653	PREDICTED: Sol:GO:0005576.NA	GO:001992	NA
100750252	723	35.74	0.23	-7.279762	8.78E-257	Down	1.18E-258	LOC100750252	Solanum lycopersi	NA	NA
101244844	411	1.51	0.01	-7.238405	7.48E-06	Down	2.14E-06	LOC101244844	PREDICTED: Sol:NA	GO:004698	NA
101249875	559	1.17	0.01	-6.870365	2.00E-06	Down	5.42E-07	LOC101249875	PREDICTED: Sol:NA	NA	GO:000975
100037736	1205	1.1	0.01	-6.78136	1.46E-13	Down	2.35E-14	tbn1	Solanum lycopersi	NA	GO:000451
101248880	791	1.09	0.01	-6.768184	9.97E-09	Down	2.22E-09	LOC101248880	PREDICTED: Sol:GO:0005634.GO:000098	GO:000632	K09422//t
101250514	1032	57.57	0.61	-6.560364	0	Down	0	LOC101250514	Solanum lycopersi	GO:0009705.GO:001522	GO:003422
101246841	666	0.87	0.01	-6.442943	7.48E-06	Down	2.14E-06	LOC101246841	PREDICTED: Sol:GO:0005634.GO:000367	GO:000632	K09286//
101264804	957	0.81	0.01	-6.33985	5.88E-08	Down	1.39E-08	LOC101264804	PREDICTED: Sol:GO:0016021.NA	NA	NA
101251154	1149	27.43	0.34	-6.334076	1.08E-301	Down	1.20E-303	LOC101251154	Solanum lycopersi	GO:0009705.GO:001522	GO:003422
104647854	1772	0.77	0.01	-6.266787	2.42E-14	Down	3.76E-15	LOC104647854	PREDICTED: Sol:GO:0016021.GO:000552	NA	K05681//
101249857	1991	12.23	0.16	-6.256209	5.29E-238	Down	7.61E-240	vicilin	Solanum lycopersi	NA	GO:004572
101246412	838	2.85	0.04	-6.154818	2.83E-22	Down	3.08E-23	LOC101246412	PREDICTED: Sol:GO:0016021.NA	NA	NA
101257892	1172	0.7	0.01	-6.129283	1.55E-08	Down	3.51E-09	LOC101257892	PREDICTED: Sol:NA	GO:004687	GO:000941
104645830	1081	13.93	0.21	-6.051662	1.86E-141	Down	4.25E-143	LOC104645830	PREDICTED: Sol:NA	NA	NA
101267061	880	0.66	0.01	-6.044394	4.82E-06	Down	1.36E-06	LOC101267061	PREDICTED: Sol:GO:0016021.NA	NA	NA
101250974	1179	63.94	0.97	-6.04259	0	Down	0	LOC101250974	PREDICTED: Sol:NA	GO:001649	NA
101245363	854	0.64	0.01	-6	1.16E-05	Down	3.39E-06	LOC101245363	PREDICTED: Sol:GO:0016021.NA	NA	NA
101252753	699	0.63	0.01	-5.97728	0.000158	Down	5.30E-05	LOC101252753	PREDICTED: Sol:GO:0016021.NA	GO:001597	K03541//
101263785	1763	0.61	0.01	-5.930737	3.09E-11	Down	5.74E-12	LOC101263785	PREDICTED: Sol:GO:0005773.NA	NA	NA
101263037	729	0.6	0.01	-5.906891	0.000158	Down	5.30E-05	LOC101263037	PREDICTED: Sol:GO:0016021.NA	GO:001991	NA
544282	710	0.59	0.01	-5.882643	0.000243	Down	8.38E-05	LOC544282	Solanum lycopersi	NA	NA
101247056	513	31.02	0.62	-5.644787	1.29E-143	Down	2.89E-145	LOC101247056	PREDICTED: Sol:GO:0005829.NA	GO:004831	NA
101254720	1541	1	0.02	-5.643856	1.26E-14	Down	1.92E-15	LOC101254720	PREDICTED: Sol:NA	GO:001676	GO:001026
101267438	1249	1.45	0.03	-5.594947	4.25E-17	Down	5.71E-18	LOC101267438	PREDICTED: Sol:NA	GO:001649	NA
101255207	1474	0.92	0.02	-5.523562	9.81E-13	Down	1.65E-13	LOC101255207	PREDICTED: Sol:GO:0016021.NA	NA	NA
104648161	1468	0.89	0.02	-5.475733	3.61E-12	Down	6.28E-13	LOC104648161	PREDICTED: Sol:NA	GO:001672	NA
101258525	1640	4.3	0.1	-5.426265	3.01E-65	Down	1.34E-66	LOC101258525	PREDICTED: Sol:NA	NA	NA
101261928	2740	0.33	0.01	-5.044394	1.08E-09	Down	2.24E-10	LOC101261928	PREDICTED: Sol:GO:0016021.GO:000837	GO:000662	K00679//
101263695	516	2.2	0.07	-4.974005	1.48E-09	Down	3.10E-10	LOC101263695	PREDICTED: Sol:GO:0005737.GO:004687	GO:003006	K07213//c
101246023	1338	0.27	0.01	-4.754888	0.000579	Down	0.000209	LOC101246023	PREDICTED: Sol:GO:0009705.GO:000526	GO:003032	K05389//
101262930	495	1.88	0.07	-4.747234	1.04E-07	Down	2.51E-08	LOC101262930	Solanum lycopersi	NA	GO:000692

544056	746	1553.55	62.31	-4.639961	0	Down	0	TAS14	Solanum lycopersi	GO:0005829	NA	GO:000941	NA	
101246280	736	4.46	0.18	-4.630975	8.26E-27	Down	7.66E-28	LOC101246280	PREDICTED: Sol:	NA	NA	NA	NA	
101250900	2808	0.24	0.01	-4.584963	2.97E-06	Down	8.21E-07	LOC101250900	PREDICTED: Sol:GO:0016021	NA	NA	K21989	//c	
101266883	1689	16.6	0.71	-4.54722	3.01E-237	Down	4.34E-239	LOC101266883	PREDICTED: Sol:NA	GO:001672	NA	K19747	//	
101256004	1531	0.46	0.02	-4.523562	2.97E-06	Down	8.21E-07	LOC101256004	PREDICTED: Sol:NA	GO:001687	NA	K10260	//	
101256556	1610	0.21	0.01	-4.392317	0.000892	Down	0.000331	LOC101256556	PREDICTED: Sol:GO:0016021	NA	GO:001572	K03695	//	
104647879	1817	0.21	0.01	-4.392317	0.000375	Down	0.000132	LOC104647879	PREDICTED: Sol:GO:0010494	GO:00040	(GO:00339	K12614	//	
101268587	1053	0.61	0.03	-4.345775	2.36E-05	Down	7.12E-06	LOC101268587	PREDICTED: Sol:NA	NA	GO:000972	K14488	//	
100134906	1151	2.68	0.14	-4.258734	8.47E-25	Down	8.35E-26	LOC100134906	Solanum lycopersi	GO:0016021	NA	GO:001992	K20628	//e
101265833	715	42.07	2.23	-4.237676	1.80E-230	Down	2.68E-232	LOC101265833	PREDICTED: Sol:NA	NA	NA	NA		
101255307	513	18.16	0.98	-4.211839	9.84E-69	Down	4.20E-70	LOC101255307	PREDICTED: Sol:GO:0016021	NA	NA	NA		
101245668	1055	14.83	0.81	-4.194453	7.57E-123	Down	1.98E-124	LOC101245668	PREDICTED: Sol:GO:0048046	GO:00167	(GO:001041	K14504	//	
101262901	1219	2.83	0.16	-4.144658	1.96E-27	Down	1.79E-28	LOC101262901	PREDICTED: Sol:GO:0005634	GO:00167	NA	K07766	//	
101247602	1640	2.82	0.16	-4.139551	2.75E-37	Down	1.97E-38	LOC101247602	PREDICTED: Sol:GO:0016021	GO:00167	(NA	K20562	//	
101248072	782	6.52	0.39	-4.063326	1.21E-38	Down	8.42E-40	LOC101248072	PREDICTED: Sol:GO:0005829	NA	GO:000941	NA		
101257315	500	4.58	0.28	-4.031849	1.31E-17	Down	1.72E-18	LOC101257315	PREDICTED: Sol:NA	GO:000827	NA	K02995	//s	
101250492	906	7.15	0.44	-4.022368	3.97E-49	Down	2.25E-50	LOC101250492	PREDICTED: Sol:NA	NA	NA	K14611	//s	
104648547	966	3.85	0.24	-4.003752	1.94E-28	Down	1.73E-29	LOC104648547	PREDICTED: Sol:GO:0016021	NA	NA	NA		
101244961	1632	3.83	0.24	-3.996238	6.03E-49	Down	3.43E-50	LOC101244961	PREDICTED: Sol:NA	GO:001672	NA	K13065	//s	
101263222	1327	10.53	0.66	-3.995896	1.20E-107	Down	3.44E-109	LOC101263222	PREDICTED: Sol:GO:0005737	GO:00055	(GO:001931	K00469	//i	
101267002	2128	16.81	1.13	-3.894925	1.39E-274	Down	1.73E-276	LOC101267002	PREDICTED: Sol:GO:0005774	GO:001517	GO:000332	K15015	//s	
101254960	789	27.46	1.96	-3.808406	1.49E-155	Down	3.03E-157	LOC101254960	PREDICTED: Sol:NA	NA	NA	NA		
101258131	1961	0.41	0.03	-3.77259	1.96E-06	Down	5.30E-07	LOC101258131	PREDICTED: Sol:NA	GO:000447	NA	K00029	//	
101267582	366	2.63	0.2	-3.716991	1.31E-06	Down	3.49E-07	LOC101267582	PREDICTED: Sol:GO:0009507	GO:000028	GO:003421	K01255	//l	
101256744	2338	5.78	0.44	-3.715494	3.58E-101	Down	1.08E-102	LOC101256744	PREDICTED: Sol:GO:0016021	GO:004572	NA	K18626	//t	
101268729	1741	13.4	1.04	-3.687578	1.48E-171	Down	2.81E-173	LOC101268729	PREDICTED: Sol:NA	GO:000848	NA	K00827	//a	
101249086	857	2.01	0.16	-3.651052	1.32E-12	Down	2.24E-13	LOC101249086	PREDICTED: Sol:NA	NA	NA	K13944	//	
101249100	1839	2.38	0.19	-3.64689	3.08E-32	Down	2.47E-33	LOC101249100	PREDICTED: Sol:NA	GO:004687	NA	K15172	//t	
101252059	1875	7.5	0.6	-3.643856	4.54E-103	Down	1.35E-104	LOC101252059	PREDICTED: Sol:NA	NA	NA	K17987	//	
101264644	1927	2.5	0.22	-3.506353	1.01E-34	Down	7.54E-36	LOC101264644	PREDICTED: Sol:GO:0016021	GO:00167	(NA	K07428	//c	
101259229	2643	0.9	0.08	-3.491853	3.10E-17	Down	4.14E-18	LOC101259229	PREDICTED: Sol:NA	GO:001678	GO:00513	(K13525	//t	
101254463	3138	0.11	0.01	-3.459432	0.000892	Down	0.000331	LOC101254463	PREDICTED: Sol:GO:0016021	GO:001672	NA	K20887	//	
101262843	763	1.96	0.18	-3.444785	1.62E-10	Down	3.16E-11	LOC101262843	PREDICTED: Sol:NA	NA	NA	NA		
101251623	1755	1.18	0.11	-3.423211	6.15E-15	Down	9.21E-16	LOC101251623	PREDICTED: Sol:GO:0016021	GO:000412	GO:004501	K15406	//	
101244638	2110	0.21	0.02	-3.392317	0.000595	Down	0.000216	LOC101244638	PREDICTED: Sol:GO:0005840	GO:000372	GO:004222	NA		
101259407	784	3.91	0.39	-3.325623	2.80E-20	Down	3.28E-21	LOC101259407	PREDICTED: Sol:GO:0016021	NA	GO:001991	NA		
101259649	1938	5.3	0.53	-3.321928	7.44E-70	Down	3.12E-71	LOC101259649	PREDICTED: Sol:NA	NA	NA	K01835	//	

101267981	1407	31.8	3.19	-3.317398	2.15E-298	Down	2.43E-300	LOC101267981	PREDICTED: Sol:GO:0016021.NA	NA	NA	NA
101251576	1718	17.62	1.78	-3.307265	5.94E-203	Down	9.70E-205	LOC101251576	PREDICTED: Sol:GO:0016020.GO:00167(GO:00445:K20623//t			
101265270	1518	1.08	0.11	-3.295456	1.31E-11	Down	2.37E-12	LOC101265270	PREDICTED: Sol:NA	NA	NA	K20473//
101251122	1032	2.45	0.26	-3.236198	8.76E-17	Down	1.20E-17	LOC101251122	PREDICTED: Sol:NA	GO:00435(GO:00063:K14431//t		
101243665	951	0.65	0.07	-3.215013	0.000145	Down	4.82E-05	LOC101243665	PREDICTED: Sol:NA	NA	NA	NA
101246267	462	2.04	0.23	-3.148863	5.10E-06	Down	1.44E-06	LOC101246267	PREDICTED: Sol:NA	NA	NA	NA
101250584	1192	42.78	4.86	-3.137908	0	Down	0	LOC101250584	PREDICTED: Sol:GO:0016021.NA	NA	NA	K20254//
101259817	1030	1.93	0.22	-3.133025	8.32E-13	Down	1.40E-13	LOC101259817	PREDICTED: Sol:NA	GO:00167(GO:00098(K08233//		
101249495	1783	0.35	0.04	-3.129283	9.86E-05	Down	3.22E-05	LOC101249495	PREDICTED: Sol:GO:0005634.GO:00055(GO:00181(K13412//c			
101258357	817	11.6	1.35	-3.103093	2.30E-63	Down	1.05E-64	LOC101258357	PREDICTED: Sol:NA	NA	NA	NA
778339	2356	5.75	0.68	-3.079955	1.73E-86	Down	6.01E-88	LOC778339	PREDICTED: Sol:GO:0005887.GO:00228:NA			K08176//
101252649	1450	0.59	0.07	-3.075288	5.10E-06	Down	1.44E-06	LOC101252649	PREDICTED: Sol:GO:0016020.GO:00083:NA			K20891//
101254946	849	0.67	0.08	-3.066089	0.00045	Down	0.00016	LOC101254946	PREDICTED: Sol:NA	NA	NA	K13993//
101262066	2120	2.93	0.35	-3.065474	1.64E-39	Down	1.12E-40	LOC101262066	PREDICTED: Sol:GO:0016021.NA	NA	NA	NA
104647629	567	0.92	0.11	-3.06413	0.000399	Down	0.000141	LOC104647629	PREDICTED: Sol:NA	NA	NA	NA
101252588	1960	5.6	0.67	-3.063194	3.63E-67	Down	1.59E-68	LOC101252588	PREDICTED: Sol:NA	GO:00037(NA		K09060//
101248026	1418	31.29	3.78	-3.049244	5.10E-274	Down	6.41E-276	JA2L	Solanum lycopersiGO:0005634.GO:00036:GO:00063:K13126//			
544091	1212	1.53	0.19	-3.00946	8.30E-12	Down	1.48E-12	Q'a	Solanum lycopersi NA	GO:00045:GO:00059:K19893//		
778275	1287	6.58	0.84	-2.969626	3.53E-51	Down	1.93E-52	SNF4	Solanum lycopersi NA	GO:00163(NA		K07200//
101244308	3619	1.4	0.18	-2.959358	2.01E-31	Down	1.64E-32	LOC101244308	PREDICTED: Sol:GO:0005634.GO:00055:NA			K13254//s
101267951	490	2.25	0.29	-2.9558	1.19E-06	Down	3.16E-07	PSK2	Solanum lycopersiGO:0005576.GO:00080(GO:00082:NA			
101268303	726	2.16	0.28	-2.947533	1.52E-09	Down	3.19E-10	LOC101268303	PREDICTED: Sol:GO:0005576.NA	GO:00199:NA		
101250270	715	12.32	1.61	-2.93587	2.68E-50	Down	1.48E-51	SP2G	Solanum lycopersi NA	NA	NA	K16223//
101249811	2327	2.29	0.3	-2.932313	1.04E-32	Down	8.23E-34	LOC101249811	PREDICTED: Sol:GO:0005634.GO:00048:NA			K08332//
543808	2466	20.68	2.82	-2.874469	1.81E-303	Down	2.00E-305	cuAO	Solanum lycopersiGO:0016021.GO:00081:GO:00093(K00276//			
101264953	603	4.09	0.57	-2.843067	7.39E-14	Down	1.17E-14	LOC101264953	PREDICTED: Sol:NA	NA	NA	NA
544066	661	77.72	10.86	-2.839262	1.91E-279	Down	2.36E-281	TSW12	Solanum lycopersiGO:0016020.GO:00082(GO:00068(NA			
101249794	1505	88.28	12.34	-2.838744	0	Down	0	LOC101249794	PREDICTED: Sol:NA	GO:00468:NA		K14497//
101256853	904	5.98	0.85	-2.814611	1.51E-30	Down	1.27E-31	LOC101256853	PREDICTED: Sol:GO:0005634.GO:00036:GO:00063:K09286//			
101261478	1731	3.51	0.5	-2.811471	2.31E-35	Down	1.71E-36	LOC101261478	PREDICTED: Sol:GO:0016020.GO:00167(GO:00445:K00512//s			
544069	651	177.42	25.35	-2.807111	0	Down	0	PR-P2	Solanum lycopersiGO:0005618.NA	GO:00508:NA		
101262661	1167	172.53	24.97	-2.78858	0	Down	0	LOC101262661	PREDICTED: Sol:GO:0005634.GO:00435(NA			K09338//
101255592	1239	339.56	49.6	-2.775254	0	Down	0	LOC101255592	PREDICTED: Sol:GO:0005887.GO:00511(GO:00714:K15382//s			
101248335	671	1.7	0.25	-2.765535	1.19E-06	Down	3.15E-07	LOC101248335	PREDICTED: Sol:GO:0009507.GO:00090:GO:00229(K02639//f			
104646748	1341	13.14	1.94	-2.759837	2.88E-98	Down	8.95E-100	LOC104646748	PREDICTED: Sol:NA	NA	NA	K07466//r
101261974	1013	10.04	1.53	-2.714156	1.66E-53	Down	8.78E-55	LOC101261974	PREDICTED: Sol:NA	NA	NA	K08064//
101250467	1607	6.5	1.02	-2.671871	4.77E-57	Down	2.39E-58	LOC101250467	PREDICTED: Sol:GO:0043231.GO:00800:GO:00081:K21354//			

104645665	828	4.14	0.65	-2.671119	1.69E-18	Down	2.13E-19	LOC104645665	PREDICTED: Sol:NA	NA	NA	NA
101247830	1407	3.8	0.6	-2.662965	3.04E-29	Down	2.64E-30	LOC101247830	PREDICTED: Sol:GO:0005743.GO:000525	GO:000656	K15109//s	
101261456	1304	5.67	0.91	-2.63941	7.51E-40	Down	5.09E-41	LOC101261456	PREDICTED: Sol:NA	NA	NA	K15095//()
101256209	1682	8.06	1.3	-2.632268	3.69E-64	Down	1.66E-65	LOC101256209	PREDICTED: Sol:GO:0016021.NA	NA	NA	NA
101258342	576	7.97	1.32	-2.594042	2.60E-23	Down	2.71E-24	XSP10	Solanum lycopersi	NA	NA	NA
101251233	2488	0.36	0.06	-2.584963	1.19E-06	Down	3.15E-07	LOC101251233	PREDICTED: Sol:NA	NA	NA	K09458//
101267241	1561	19.35	3.23	-2.582727	2.71E-158	Down	5.42E-160	LOC101267241	PREDICTED: Sol:GO:0016021.GO:001517	GO:000333	K14993//s	
101247647	1068	99.02	16.93	-2.548138	0	Down	0	LOC101247647	PREDICTED: Sol:GO:0005576.NA	GO:001993	K20628//e	
101255514	747	2.88	0.5	-2.526069	3.51E-11	Down	6.54E-12	LOC101255514	PREDICTED: Sol:NA	NA	NA	NA
544034	613	78.85	13.94	-2.49988	1.29E-227	Down	1.96E-229	LE16	Solanum lycopersi	GO:0016020.GO:000828	GO:000686	NA
101267939	1416	2.54	0.46	-2.465123	1.90E-18	Down	2.40E-19	LOC101267939	PREDICTED: Sol:NA	GO:004698	NA	K13422//t
101268445	1919	0.55	0.1	-2.459432	6.45E-06	Down	1.84E-06	LOC101268445	PREDICTED: Sol:NA	GO:000462	GO:000656	K00318//
101262282	3069	1.53	0.28	-2.450033	2.06E-24	Down	2.05E-25	LOC101262282	PREDICTED: Sol:GO:0016021.GO:003022	GO:004852	K04733//i	
101268663	2138	2.45	0.45	-2.444785	1.09E-26	Down	1.01E-27	LOC101268663	PREDICTED: Sol:GO:0016021.GO:000556	GO:000662	K15404//a	
101245070	2449	2.99	0.55	-2.442642	2.44E-37	Down	1.74E-38	LOC101245070	PREDICTED: Sol:GO:0016021.GO:000521	GO:005508	K13783//	
101263398	753	3.64	0.67	-2.441705	1.71E-13	Down	2.77E-14	LOC101263398	PREDICTED: Sol:GO:0016021.NA	GO:001991	NA	
101256448	1741	1.52	0.28	-2.440573	8.48E-14	Down	1.35E-14	LOC101256448	PREDICTED: Sol:NA	GO:001661	NA	K21840//
101266084	872	1.45	0.27	-2.425022	9.99E-07	Down	2.63E-07	LOC101266084	PREDICTED: Sol:GO:0005737.GO:000436	GO:000672	K00799//	
101253409	3731	0.16	0.03	-2.415037	0.000881	Down	0.000327	LOC101253409	PREDICTED: Sol:NA	GO:001636	NA	K21026//a
109120650	2122	0.31	0.06	-2.369234	0.00021	Down	7.17E-05	LOC109120650	PREDICTED: Sol:NA	NA	NA	NA
101251993	1651	0.92	0.18	-2.353637	1.70E-08	Down	3.85E-09	LOC101251993	PREDICTED: Sol:NA	GO:004356	GO:000633	K14431//t
101265388	786	1.53	0.3	-2.350497	3.95E-06	Down	1.10E-06	LOC101265388	PREDICTED: Sol:NA	NA	NA	NA
101243675	811	10.28	2.03	-2.340289	5.47E-38	Down	3.85E-39	LOC101243675	PREDICTED: Sol:NA	NA	NA	K16223//
101250521	1991	5.97	1.18	-2.338944	1.07E-56	Down	5.39E-58	LOC101250521	PREDICTED: Sol:GO:0016021.NA	GO:005508	K13754//s	
101247301	1557	0.96	0.19	-2.337035	9.57E-08	Down	2.30E-08	LOC101247301	PREDICTED: Sol:GO:0043231.GO:000372	GO:000942	K20892//	
101257623	2617	5.06	1.01	-2.324782	3.77E-63	Down	1.73E-64	LOC101257623	PREDICTED: Sol:NA	GO:001672	NA	K06617//r
101055596	2189	14.25	2.86	-2.316875	1.30E-146	Down	2.85E-148	hsc70.3	PREDICTED: Sol:NA	GO:000552	NA	K03283//
104645532	834	0.99	0.2	-2.307429	0.000228	Down	7.83E-05	LOC104645532	PREDICTED: Sol:NA	GO:000367	GO:001507	NA
101244860	321	6.93	1.43	-2.27684	3.91E-09	Down	8.46E-10	LOC101244860	PREDICTED: Sol:NA	NA	NA	NA
101245417	2043	24.74	5.17	-2.258609	2.26E-229	Down	3.39E-231	LOC101245417	PREDICTED: Sol:GO:0016021.NA	NA	NA	K15015//s
101268307	837	14.75	3.12	-2.241097	3.93E-53	Down	2.09E-54	LOC101268307	PREDICTED: Sol:NA	NA	NA	K13993//
109119578	1172	8.12	1.73	-2.230708	1.73E-42	Down	1.12E-43	LOC109119578	PREDICTED: Sol:NA	GO:004356	GO:000633	K21867//
104646084	1578	0.75	0.16	-2.228819	6.47E-06	Down	1.84E-06	LOC104646084	PREDICTED: Sol:NA	NA	NA	K14709//s
101264531	1045	5.86	1.26	-2.217477	7.67E-27	Down	7.10E-28	LOC101264531	PREDICTED: Sol:NA	GO:000372	NA	K01166//r
543983	2070	0.41	0.09	-2.187627	0.000186	Down	6.32E-05	LOC543983	Solanum lycopersi	GO:0009507.GO:004286	GO:000905	K01754//t
101250924	2316	12.59	2.8	-2.16878	6.19E-127	Down	1.58E-128	LOC101250924	PREDICTED: Sol:GO:0016021.GO:000521	NA	NA	K14638//s
101267723	3286	2.01	0.45	-2.159199	1.98E-29	Down	1.72E-30	ppc2	PREDICTED: Sol:NA	GO:000896	GO:000605	K01595//

101254378	1737	27.5	6.17	-2.156089	6.58E-208	Down	1.06E-209	LOC101254378	PREDICTED: Sol:NA	GO:001662	NA	K00166//	
100316895	1783	50.81	11.4	-2.156079	0	Down	0	PP2C-2	Solanum lycopersi	NA	GO:004687	NA	K14497//
104648458	507	1.87	0.42	-2.154577	0.000186	Down	6.32E-05	LOC104648458	PREDICTED: Sol:NA	NA	NA	NA	
101263880	336	6.49	1.46	-2.15225	1.71E-08	Down	3.87E-09	LOC101263880	PREDICTED: Sol:GO:0016021	NA	NA	NA	K03363//c
101266080	659	587.03	132.69	-2.145375	0	Down	0	LOC101266080	Solanum lycopersi	GO:0016020	GO:000828	GO:000686	NA
101246666	785	6.06	1.37	-2.145142	7.82E-20	Down	9.31E-21	LOC101246666	Solanum lycopersi	NA	NA	GO:000695	NA
104644397	1716	0.57	0.13	-2.13245	7.85E-05	Down	2.53E-05	LOC104644397	PREDICTED: Sol:NA	NA	NA	NA	K13161//
101244167	698	91.8	21.15	-2.117836	1.35E-251	Down	1.87E-253	LOC101244167	PREDICTED: Sol:NA	NA	NA	NA	K12741//
101260986	837	5.38	1.24	-2.117266	1.33E-18	Down	1.66E-19	LOC101260986	PREDICTED: Sol:GO:0016021	GO:006163	GO:004316	GO:004316	K19040//
101255654	730	15.27	3.52	-2.117053	7.97E-45	Down	4.93E-46	LOC101255654	PREDICTED: Sol:NA	NA	NA	NA	
101258100	885	10.12	2.34	-2.112629	1.91E-36	Down	1.39E-37	LOC101258100	PREDICTED: Sol:NA	NA	NA	NA	
100529102	2783	2.4	0.56	-2.099536	1.36E-28	Down	1.20E-29	ITTS1	Solanum lycopersi	NA	GO:001686	NA	K15813//
101257495	792	1.07	0.25	-2.097611	0.000491	Down	0.000176	LOC101257495	PREDICTED: Sol:GO:0005739	NA	NA	NA	K02267//c
101252861	2027	0.47	0.11	-2.095157	0.000108	Down	3.55E-05	LOC101252861	PREDICTED: Sol:GO:0009505	GO:001672	NA	NA	K19791//i
101263926	1610	9.77	2.29	-2.093011	1.29E-67	Down	5.62E-69	LOC101263926	PREDICTED: Sol:NA	GO:001682	NA	NA	K01761//
101258285	354	24.29	5.7	-2.091329	1.83E-30	Down	1.54E-31	LOC101258285	PREDICTED: Sol:NA	NA	NA	NA	
101259698	1503	2.11	0.5	-2.077243	1.23E-13	Down	1.97E-14	LOC101259698	PREDICTED: Sol:NA	GO:004687	NA	NA	K05277//l
544180	1196	6.37	1.51	-2.076745	4.01E-31	Down	3.32E-32	abz1	Solanum lycopersi	NA	GO:000370	NA	K09060//
778310	1267	4.63	1.11	-2.060453	6.15E-24	Down	6.26E-25	GGPS1	Solanum lycopersi	NA	GO:001672	GO:000828	K13789//
101245699	826	4.91	1.18	-2.056936	2.64E-16	Down	3.69E-17	LOC101245699	PREDICTED: Sol:NA	NA	NA	NA	K03006//
101264793	1498	28.03	6.76	-2.051877	9.68E-172	Down	1.83E-173	LOC101264793	PREDICTED: Sol:GO:0016021	GO:001517	GO:000333	GO:000333	K15015//s
101252966	1781	16.24	3.92	-2.050626	2.38E-116	Down	6.45E-118	LOC101252966	PREDICTED: Sol:GO:0016021	NA	NA	NA	K14209//s
104648861	2380	0.33	0.08	-2.044394	0.000675	Down	0.000246	LOC104648861	PREDICTED: Sol:NA	NA	NA	NA	K11843//
101266471	1381	0.66	0.16	-2.044394	0.000279	Down	9.69E-05	LOC101266471	PREDICTED: Sol:GO:0005737	GO:000467	GO:002301	GO:002301	K04421//
104644865	1663	1.27	0.31	-2.034488	0.000107	Down	3.49E-05	LOC104644865	PREDICTED: Sol:NA	NA	NA	NA	
101244894	658	228.78	56.4	-2.020194	0	Down	0	LOC101244894	PREDICTED: Sol:NA	NA	NA	NA	K09874//a
101246264	1777	4.7	1.16	-2.018536	2.24E-33	Down	1.74E-34	LOC101246264	PREDICTED: Sol:GO:0005886	GO:000412	GO:001945	GO:001945	K15406//
101259110	964	0.85	0.21	-2.017074	0.000356	Down	0.000125	LOC101259110	PREDICTED: Sol:NA	NA	NA	NA	
109119902	542	7.26	1.8	-2.011973	3.12E-15	Down	4.61E-16	LOC109119902	PREDICTED: Sol:NA	NA	NA	NA	
101248124	762	10.59	2.66	-1.993204	3.36E-30	Down	2.85E-31	LOC101248124	PREDICTED: Sol:NA	NA	NA	NA	
101249967	714	3.42	0.86	-1.991588	1.13E-09	Down	2.34E-10	LOC101249967	PREDICTED: Sol:GO:0048046	GO:001685	NA	NA	K01870//i
101055585	1086	13.03	3.3	-1.981299	1.95E-53	Down	1.03E-54	ZF2	Solanum lycopersi	GO:0005634	GO:000367	NA	K05747//
104646047	678	1.77	0.45	-1.975752	4.85E-05	Down	1.52E-05	LOC104646047	PREDICTED: Sol:NA	NA	NA	NA	
101263275	1965	35.61	9.07	-1.973108	4.72E-268	Down	6.04E-270	LOC101263275	PREDICTED: Sol:GO:0005634	GO:000122	GO:000636	GO:000636	K12126//
101260767	724	2	0.51	-1.971431	6.19E-06	Down	1.76E-06	LOC101260767	PREDICTED: Sol:GO:0005634	GO:000367	GO:000636	GO:000636	K14516//e
101252918	1693	2.81	0.72	-1.964501	1.38E-18	Down	1.73E-19	LOC101252918	PREDICTED: Sol:NA	GO:004687	NA	NA	K06892//f
101268519	1592	1.01	0.26	-1.957772	1.07E-06	Down	2.84E-07	LOC101268519	PREDICTED: Sol:GO:0016021	NA	NA	NA	K19041//

101055556	729	3.26	0.84	-1.956411	2.89E-09	Down	6.18E-10	LOC101055556	Solanum lycopersi	GO:0005634	NA	GO:000635	K14484//a
101255889	1578	2.51	0.65	-1.949176	8.29E-16	Down	1.19E-16	LOC101255889	PREDICTED: Sol:	NA	GO:001672	NA	K19747//
101245338	1784	15.25	3.95	-1.948885	1.21E-92	Down	3.94E-94	LOC101245338	PREDICTED: Sol:	NA	NA	NA	NA
778214	777	1657.52	429.57	-1.948061		Down	0	ltpg1	Solanum lycopersi	GO:0016020	GO:000828	GO:000686	NA
544185	705	94.51	24.53	-1.94592	2.68E-235	Down	3.92E-237	P4	Solanum lycopersi	GO:0005576	NA	GO:005083	K13449//
101266923	667	90.36	23.61	-1.936286	4.79E-210	Down	7.65E-212	LOC101266923	PREDICTED: Sol:	NA	NA	NA	K12121//
101246214	1813	2.16	0.57	-1.921997	3.12E-15	Down	4.60E-16	LOC101246214	PREDICTED: Sol:	NA	GO:004572	NA	K17815//e
101263226	975	6.59	1.77	-1.896529	3.29E-23	Down	3.45E-24	LOC101263226	PREDICTED: Sol:	NA	NA	GO:000692	K20165//
544017	1929	5.33	1.45	-1.878083	1.41E-37	Down	1.00E-38	LAPA1	Solanum lycopersi	GO:0009507	GO:000028	GO:003421	K01255//l
543875	2259	11.08	3.04	-1.865815	4.81E-90	Down	1.60E-91	LEAS1	Solanum lycopersi	NA	GO:000552	GO:000652	K01953//a
101263372	2072	0.69	0.19	-1.860597	4.78E-06	Down	1.34E-06	LOC101263372	PREDICTED: Sol:	NA	GO:004687	GO:004612	K00422//
104645765	1301	1.08	0.3	-1.847997	1.19E-05	Down	3.48E-06	LOC104645765	PREDICTED: Sol:	NA	GO:000415	GO:000656	NA
101262581	1490	4.52	1.26	-1.842899	5.76E-23	Down	6.09E-24	LOC101262581	PREDICTED: Sol:	NA	GO:000552	NA	K07910//
101267932	1947	0.93	0.26	-1.838719	4.83E-07	Down	1.24E-07	LOC101267932	PREDICTED: Sol:	NA	NA	NA	K13963//s
101254668	1794	17.79	4.98	-1.836849	3.00E-111	Down	8.36E-113	LOC101254668	PREDICTED: Sol:	GO:0016020	GO:001676	GO:004453	K20562//
543853	1470	3.77	1.06	-1.8305	1.07E-19	Down	1.29E-20	Xegip	Solanum lycopersi	NA	GO:000415	GO:000656	K00924//
101245242	1937	1.1	0.31	-1.827163	4.75E-08	Down	1.11E-08	LOC101245242	PREDICTED: Sol:	NA	GO:000422	NA	K02563//
104648973	1090	3.1	0.88	-1.816693	4.24E-12	Down	7.43E-13	LOC104648973	PREDICTED: Sol:	NA	NA	NA	NA
544149	956	1279.55	365.29	-1.808522		Down	0	CHI3	Solanum lycopersi	GO:0005615	GO:000456	GO:000602	K20547//
101247100	1488	1.75	0.5	-1.807355	1.44E-09	Down	3.01E-10	LOC101247100	PREDICTED: Sol:	NA	GO:000482	NA	K10260//
101244796	1532	52.82	15.1	-1.806536	2.49E-273	Down	3.14E-275	LOC101244796	PREDICTED: Sol:	NA	GO:007192	NA	K09838//z
101246743	595	42.07	12.03	-1.806155	6.80E-79	Down	2.56E-80	LOC101246743	PREDICTED: Sol:	GO:0016020	GO:000828	GO:000686	NA
544163	2170	47.2	13.6	-1.79518		Down	0	LOC544163	Solanum lycopersi	NA	GO:001676	NA	K09840//
101245487	1917	2.39	0.69	-1.792342	9.74E-16	Down	1.40E-16	LOC101245487	PREDICTED: Sol:	GO:0005783	GO:000828	NA	K19907//s
101260726	1315	10.2	2.95	-1.789782	7.00E-48	Down	4.05E-49	LOC101260726	PREDICTED: Sol:	GO:0016021	GO:000827	NA	K10661//
101055548	1151	33.78	9.77	-1.789739	5.36E-128	Down	1.36E-129	IAA8	Solanum lycopersi	GO:0005634	NA	GO:000635	K14484//a
101245919	945	3.04	0.88	-1.788496	4.57E-10	Down	9.21E-11	LOC101245919	PREDICTED: Sol:	NA	NA	NA	K17506//
101266086	1515	5.38	1.56	-1.78606	9.50E-28	Down	8.59E-29	LOC101266086	PREDICTED: Sol:	GO:0005615	GO:000822	GO:005166	K01373//c
101264922	2239	2.68	0.78	-1.780687	1.58E-20	Down	1.83E-21	LOC101264922	PREDICTED: Sol:	NA	NA	NA	K17046//
101261619	677	16.62	4.84	-1.779841	1.40E-35	Down	1.03E-36	LOC101261619	PREDICTED: Sol:	GO:0005737	GO:004687	GO:003006	K07213//c
101261956	877	3.67	1.07	-1.778169	4.50E-11	Down	8.46E-12	LOC101261956	PREDICTED: Sol:	NA	NA	NA	NA
101245348	1368	1.77	0.52	-1.767166	1.13E-08	Down	2.53E-09	LOC101245348	PREDICTED: Sol:	NA	NA	NA	K03097//c
101256490	1744	1.32	0.39	-1.758992	2.02E-08	Down	4.59E-09	LOC101256490	PREDICTED: Sol:	NA	GO:001672	NA	K19747//
101245840	1716	39.66	11.75	-1.755024	3.05E-222	Down	4.72E-224	LOC101245840	PREDICTED: Sol:	GO:0005623	GO:000475	GO:004543	K17609//
101263090	1728	0.64	0.19	-1.752072	0.000167	Down	5.63E-05	LOC101263090	PREDICTED: Sol:	NA	GO:001672	GO:000812	K21373//
544236	2628	40.83	12.13	-1.75105		Down	0	SIP	Solanum lycopersi	NA	GO:001672	NA	K06617//r
109118734	1331	31.37	9.32	-1.750984	1.53E-134	Down	3.69E-136	LOC109118734	PREDICTED: Sol:	NA	GO:004687	NA	K04124//

778293	1401	6.33	1.9	-1.736206	7.60E-29	Down	6.67E-30	9612	Solanum lycopersi	NA	GO:004687	GO:004545	K01728//
101246954	861	15.19	4.58	-1.729702	9.93E-41	Down	6.63E-42	LOC101246954	PREDICTED: Sol:NA	NA	NA	NA	K20855//
101247761	609	1.69	0.51	-1.728454	0.000722	Down	0.000264	LOC101247761	PREDICTED: Sol:NA	NA	GO:000972	K14488//	
101255191	2218	13.07	3.95	-1.726335	1.87E-90	Down	6.20E-92	LOC101255191	PREDICTED: Sol:NA	NA	GO:000636	K17398//	
104648541	952	3.11	0.94	-1.726182	6.25E-10	Down	1.27E-10	LOC104648541	PREDICTED: Sol:GO:0016021	NA	GO:001991	NA	
101261459	3138	7.45	2.27	-1.714548	1.25E-74	Down	4.94E-76	LOC101261459	PREDICTED: Sol:GO:0005886	NA	NA	NA	
101250764	1501	2.26	0.69	-1.711655	1.94E-11	Down	3.55E-12	LOC101250764	PREDICTED: Sol:GO:0005634	GO:000098	GO:000635	K09422//t	
101250001	1603	8.25	2.52	-1.71097	1.77E-41	Down	1.16E-42	LOC101250001	PREDICTED: Sol:NA	GO:004698	NA	K16189//	
101246665	4046	4.84	1.48	-1.70941	1.28E-63	Down	5.81E-65	LOC101246665	PREDICTED: Sol:NA	GO:005108	NA	NA	
544134	738	2.55	0.78	-1.708951	2.12E-06	Down	5.76E-07	TSI-1	Solanum lycopersi	NA	GO:000695	K14496//a	
101249144	1192	27.62	8.46	-1.706984	5.49E-102	Down	1.65E-103	LOC101249144	PREDICTED: Sol:GO:0005634	GO:004356	NA	K09338//	
100037510	1610	50.89	15.6	-1.705836	1.23E-256	Down	1.67E-258	SRK2C	Solanum lycopersi	GO:0005634	GO:000467	GO:000646	K14498//s
101261337	3010	0.39	0.12	-1.70044	0.000125	Down	4.14E-05	LOC101261337	PREDICTED: Sol:GO:0005737	GO:000482	GO:000996	K01931//	
101247242	525	3.05	0.94	-1.698077	2.18E-05	Down	6.55E-06	LOC101247242	PREDICTED: Sol:GO:0016020	GO:000828	GO:000686	NA	
101249340	1368	0.94	0.29	-1.696608	6.99E-05	Down	2.23E-05	LOC101249340	PREDICTED: Sol:GO:0043231	GO:000372	GO:000945	K20892//	
101266493	812	3.19	0.99	-1.688056	2.02E-08	Down	4.60E-09	LOC101266493	PREDICTED: Sol:NA	NA	NA	NA	
101260277	5172	0.61	0.19	-1.68281	3.44E-11	Down	6.41E-12	LOC101260277	PREDICTED: Sol:GO:0005886	GO:004352	GO:000716	K07198//	
101252201	1696	0.61	0.19	-1.68281	0.000532	Down	0.000192	LOC101252201	PREDICTED: Sol:GO:0016021	GO:001676	NA	K07428//c	
101258143	776	3.2	1	-1.678072	4.82E-08	Down	1.13E-08	LOC101258143	PREDICTED: Sol:GO:0016021	NA	GO:001991	NA	
101256151	797	5.52	1.73	-1.673896	1.39E-13	Down	2.24E-14	LOC101256151	PREDICTED: Sol:GO:0016021	GO:000521	GO:005508	K15400//	
101265369	1533	2.07	0.65	-1.671119	2.58E-10	Down	5.11E-11	LOC101265369	PREDICTED: Sol:GO:0043231	GO:008002	GO:000815	K12938//a	
104648156	1033	56.94	17.92	-1.667872	3.07E-174	Down	5.71E-176	LOC104648156	PREDICTED: Sol:GO:0046658	GO:000905	NA	K01179//e	
544092	1250	79.27	24.98	-1.666001	1.80E-296	Down	2.06E-298	Q`b	Solanum lycopersi	GO:0046658	GO:004297	GO:000597	K19893//
101245973	1522	3.52	1.11	-1.665016	9.79E-18	Down	1.28E-18	LOC101245973	PREDICTED: Sol:NA	NA	NA	NA	
101251084	1787	1.14	0.36	-1.662965	4.90E-07	Down	1.25E-07	LOC101251084	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	K12900//	
101247078	1279	1.04	0.33	-1.656046	9.13E-05	Down	2.97E-05	LOC101247078	PREDICTED: Sol:NA	GO:001678	GO:000815	K13258//	
104648615	941	1.75	0.56	-1.643856	1.58E-05	Down	4.68E-06	LOC104648615	PREDICTED: Sol:GO:0016021	NA	NA	NA	
104645408	834	4.15	1.33	-1.641685	1.07E-10	Down	2.06E-11	LOC104645408	PREDICTED: Sol:NA	NA	NA	K15920//	
101264952	522	218.01	70.02	-1.638555	3.65E-306	Down	4.00E-308	LOC101264952	PREDICTED: Sol:NA	NA	NA	NA	
101262489	2466	3.6	1.16	-1.633872	3.17E-27	Down	2.91E-28	LOC101262489	Solanum lycopersi	NA	GO:000028	GO:000961	K17982//
101266963	1866	6.78	2.19	-1.630354	4.18E-38	Down	2.94E-39	LOC101266963	PREDICTED: Sol:GO:0016020	GO:001676	GO:004455	K14985//	
101259682	910	49.98	16.17	-1.628031	2.87E-129	Down	7.15E-131	LOC101259682	PREDICTED: Sol:GO:0016021	NA	GO:000972	K03251//t	
104647829	469	3.52	1.14	-1.626542	3.74E-05	Down	1.16E-05	LOC104647829	PREDICTED: Sol:NA	NA	NA	NA	
109119848	1817	0.92	0.3	-1.616671	1.16E-05	Down	3.39E-06	LOC109119848	PREDICTED: Sol:GO:0016020	GO:001676	GO:004455	K07408//c	
544028	1942	3.37	1.1	-1.615245	4.93E-20	Down	5.83E-21	ACS1A	Solanum lycopersi	NA	GO:000382	GO:000905	K20772//
104649633	1231	0.98	0.32	-1.61471	0.000287	Down	9.98E-05	LOC104649633	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	NA	
101249481	1291	5.41	1.77	-1.611879	1.11E-20	Down	1.28E-21	LOC101249481	PREDICTED: Sol:NA	GO:004687	NA	K06892//f	

101262567	1187	19.79	6.48	-1.610706	2.64E-67	Down	1.15E-68	LOC101262567	PREDICTED: Sol:GO:0009506.GO:000487GO:000692NA
101263538	1999	1.66	0.55	-1.59368	1.83E-10	Down	3.59E-11	LOC101263538	PREDICTED: Sol:GO:0016021.GO:000521NA K14638//s
101264425	1258	5.65	1.89	-1.579865	2.37E-20	Down	2.76E-21	LOC101264425	PREDICTED: Sol:GO:0005576.GO:004687GO:000697K00430//
101257896	2613	40.56	13.58	-1.578574	1.08E-302	Down	1.20E-304	sbt4b	Solanum lycopersi NA GO:000422NA K12619//
544035	1286	3.85	1.29	-1.577487	2.06E-14	Down	3.18E-15	EXP1	Solanum lycopersiGO:0016020.NA GO:000962K20628//e
101248657	1431	4.64	1.56	-1.572579	7.39E-19	Down	9.16E-20	LOC101248657	PREDICTED: Sol:GO:0016021.GO:001671GO:000662K10256//
101244010	518	3.42	1.15	-1.572362	2.65E-05	Down	8.05E-06	LOC101244010	PREDICTED: Sol:NA NA GO:000692NA
101256065	1414	2.73	0.92	-1.569195	1.68E-11	Down	3.06E-12	LOC101256065	PREDICTED: Sol:GO:0005634.GO:004352GO:000632K13425//
101243802	2190	2.61	0.88	-1.568474	6.21E-17	Down	8.41E-18	LOC101243802	PREDICTED: Sol:GO:0016021.NA GO:001022K16055//t
101245037	1124	13.52	4.56	-1.567989	4.48E-42	Down	2.92E-43	LOC101245037	PREDICTED: Sol:GO:0005634.GO:004352NA K09338//
101254104	1731	1.33	0.45	-1.563429	2.52E-07	Down	6.31E-08	LOC101254104	PREDICTED: Sol:GO:0016021.GO:006162GO:004312K19038//
101245684	758	7.24	2.45	-1.563208	3.46E-15	Down	5.11E-16	LOC101245684	PREDICTED: Sol:GO:0005737.GO:004687GO:003002K07213//c
544250	1572	95.4	32.38	-1.558886	0	Down	0	fdh	Solanum lycopersiGO:0005739.GO:001661GO:004212K00122//f
109119181	1416	7.3	2.48	-1.557556	3.10E-29	Down	2.69E-30	LOC109119181	PREDICTED: Sol:GO:0016021.GO:001671GO:000662K10256//
101244867	1445	2.17	0.74	-1.552098	1.31E-09	Down	2.73E-10	LOC101244867	PREDICTED: Sol:GO:0016021.NA GO:005502K07088//
101253938	944	2.46	0.84	-1.550197	5.73E-07	Down	1.48E-07	LOC101253938	PREDICTED: Sol:GO:0005634.GO:004342GO:001631K14490//
543953	536	116.59	39.95	-1.545177	2.11E-155	Down	4.32E-157	ER1	Solanum lycopersiGO:0005576.GO:000482GO:000961NA
101259007	2774	64.54	22.21	-1.538984	0	Down	0	LOC101259007	PREDICTED: Sol:NA NA NA K09060//
109118713	1368	1.8	0.62	-1.537657	2.33E-07	Down	5.81E-08	LOC109118713	PREDICTED: Sol:GO:0016021.NA NA K13195//c
101258079	1845	2.67	0.92	-1.537134	4.34E-14	Down	6.82E-15	LOC101258079	PREDICTED: Sol:NA GO:004691GO:000605K01647//c
101267030	870	1.77	0.61	-1.536868	8.15E-05	Down	2.63E-05	LOC101267030	PREDICTED: Sol:GO:0016021.GO:005152NA NA
101244327	1998	4.06	1.4	-1.536053	7.77E-23	Down	8.26E-24	LOC101244327	PREDICTED: Sol:GO:0016021.GO:001522NA K03327//
109120086	745	3.55	1.23	-1.529161	1.30E-07	Down	3.17E-08	LOC109120086	PREDICTED: Sol:GO:0016021.NA NA NA
101265741	2419	0.49	0.17	-1.527247	0.000474	Down	0.00017	LOC101265741	PREDICTED: Sol:NA GO:000552NA K08835//s
101245149	3463	2.88	1	-1.526069	3.42E-28	Down	3.06E-29	LOC101245149	PREDICTED: Sol:GO:0005829.GO:000892GO:000605K01595//
101268052	880	59.41	20.66	-1.523866	9.53E-135	Down	2.29E-136	LOC101268052	PREDICTED: Sol:NA NA GO:000941NA
544083	917	8.41	2.94	-1.51629	1.74E-20	Down	2.02E-21	PR-1a1	Solanum lycopersiGO:0005576.NA NA K13449//
101252766	1587	2.8	0.98	-1.514573	1.98E-11	Down	3.61E-12	LOC101252766	PREDICTED: Sol:NA NA NA NA
544123	807	12527.42	4385.59	-1.514246	0	Down	0	PR1b1	Solanum lycopersiGO:0005576.NA GO:005082K13449//
544061	696	142.6	50.06	-1.510244	4.07E-249	Down	5.65E-251	LOC544061	PREDICTED: Sol:NA NA NA K10747//
101249257	1261	36.92	12.97	-1.509224	2.16E-121	Down	5.69E-123	LOC101249257	PREDICTED: Sol:GO:0005634.GO:000467GO:001812K04515//c
544297	978	5.69	2	-1.508429	5.04E-15	Down	7.52E-16	CrtR-b2	Solanum lycopersiGO:0016021.GO:000552GO:000861K15746//
101256789	2113	1.65	0.58	-1.508341	3.61E-10	Down	7.22E-11	LOC101256789	PREDICTED: Sol:NA GO:004692NA K13422//t
101266759	784	10.26	3.61	-1.50696	6.73E-21	Down	7.71E-22	LOC101266759	PREDICTED: Sol:NA NA NA NA
101247334	2078	1.79	0.63	-1.506536	3.11E-10	Down	6.19E-11	LOC101247334	PREDICTED: Sol:NA GO:001672NA K01426//a
101256979	2517	39.87	14.09	-1.500632	2.14E-266	Down	2.80E-268	EDS1	Solanum lycopersi NA GO:001672GO:000662K18875//e
101250069	513	5.03	1.78	-1.498681	6.80E-07	Down	1.76E-07	LOC101250069	PREDICTED: Sol:GO:0009506.GO:005102GO:000642NA

101265862	1010	6.3	2.23	-1.498308	5.91E-17	Down	7.99E-18	LOC101265862	PREDICTED: Sol:GO:0005618.GO:004572	NA	K00384//t
544286	512	125.27	44.41	-1.496084	3.28E-151	Down	6.91E-153	mcpi	Solanum lycopersiGO:0016021.GO:000418	NA	NA
543816	892	16.21	5.75	-1.49525	8.04E-37	Down	5.79E-38	GST-T4	Solanum lycopersiGO:0005737.GO:000436	GO:000672	K00799//
101267006	851	20.69	7.35	-1.493117	2.10E-44	Down	1.30E-45	LOC101267006	PREDICTED: Sol:NA	NA	NA
101267866	1673	2.7	0.96	-1.491853	2.15E-12	Down	3.70E-13	LOC101267866	PREDICTED: Sol:NA	NA	NA
101254501	656	38.81	13.82	-1.489671	4.82E-62	Down	2.24E-63	LOC101254501	PREDICTED: Sol:NA	NA	NA
101260320	1897	2.73	0.98	-1.478047	4.47E-14	Down	7.02E-15	LOC101260320	PREDICTED: Sol:NA	NA	NA
101262200	1627	115.92	41.84	-1.470175	0	Down	0	LOC101262200	PREDICTED: Sol:GO:0009505.GO:001988	NA	K07200//
101246420	894	251.94	91.03	-1.468666	0	Down	0	Asr4	Solanum lycopersi	NA	GO:000692
101268427	1516	2.6	0.94	-1.467779	9.19E-11	Down	1.76E-11	LOC101268427	PREDICTED: Sol:NA	NA	NA
101247540	603	2.68	0.97	-1.466176	0.000131	Down	4.33E-05	LOC101247540	PREDICTED: Sol:GO:0005622.GO:000827	NA	K19939//
101262533	1752	2.48	0.9	-1.462343	8.51E-12	Down	1.52E-12	LOC101262533	PREDICTED: Sol:GO:0016021.GO:001670	NA	K07428//c
101268860	1756	13.03	4.73	-1.461925	2.69E-58	Down	1.33E-59	LOC101268860	PREDICTED: Sol:NA	NA	GO:000716
101244426	1047	18.73	6.8	-1.461744	1.19E-48	Down	6.78E-50	LOC101244426	PREDICTED: Sol:GO:0016021	NA	NA
101251889	927	1.57	0.57	-1.461731	0.000234	Down	8.03E-05	LOC101251889	PREDICTED: Sol:NA	NA	NA
101257792	1554	13.63	4.95	-1.461285	1.14E-53	Down	6.01E-55	LOC101257792	PREDICTED: Sol:GO:0016021.GO:000521	NA	K00850//
101256765	1504	5.09	1.85	-1.46014	4.16E-20	Down	4.90E-21	LOC101256765	PREDICTED: Sol:GO:0043231.GO:008002	GO:000815	K13495//c
104646584	2267	1.32	0.48	-1.459432	2.39E-08	Down	5.47E-09	LOC104646584	PREDICTED: Sol:NA	NA	NA
101262352	1667	2.28	0.83	-1.457851	2.77E-10	Down	5.51E-11	LOC101262352	PREDICTED: Sol:GO:0016020.GO:001505	GO:001565	NA
104647880	465	4.01	1.46	-1.457634	5.27E-05	Down	1.66E-05	LOC104647880	PREDICTED: Sol:GO:0016021	NA	NA
101251222	863	11.15	4.06	-1.457492	8.49E-24	Down	8.68E-25	LOC101251222	PREDICTED: Sol:GO:0005795.GO:002002	GO:000677	K05770//t
101055579	1535	1.73	0.63	-1.457348	3.47E-10	Down	6.94E-11	LOC101055579	PREDICTED: Sol:GO:0016021.GO:001661	GO:000965	K00279//c
101259076	1144	1.81	0.66	-1.455452	6.64E-06	Down	1.89E-06	LOC101259076	PREDICTED: Sol:GO:0016021.GO:005111	GO:000862	K15382//s
101253759	1811	50.75	18.51	-1.455103	3.46E-231	Down	5.12E-233	LOC101253759	PREDICTED: Sol:GO:0016021.GO:000551	GO:000692	K08472//
101257588	1908	13.15	4.81	-1.450954	8.42E-64	Down	3.81E-65	LOC101257588	PREDICTED: Sol:GO:0005618.GO:001678	GO:007152	K19882//
101255907	573	9.26	3.39	-1.449727	6.95E-13	Down	1.16E-13	LOC101255907	PREDICTED: Sol:GO:0016020.GO:000828	GO:000686	NA
101253729	966	1.96	0.72	-1.444785	2.14E-05	Down	6.42E-06	LOC101253729	PREDICTED: Sol:GO:0005618.GO:004572	NA	K00384//t
101055581	1067	3.62	1.33	-1.444563	3.35E-11	Down	6.23E-12	LOC101055581	PREDICTED: Sol:NA	NA	NA
101249954	965	12.7	4.67	-1.443334	6.05E-30	Down	5.14E-31	LOC101249954	PREDICTED: Sol:NA	NA	NA
101244562	1041	2.66	0.98	-1.440573	1.77E-07	Down	4.37E-08	LOC101244562	PREDICTED: Sol:NA	GO:000486	NA
101260225	1524	1.49	0.55	-1.437809	2.51E-06	Down	6.88E-07	LOC101260225	PREDICTED: Sol:GO:0016020.GO:000462	GO:001602	K16815//c
101268129	1761	46.38	17.13	-1.436978	4.21E-201	Down	6.98E-203	NAOD	Solanum lycopersi	NA	NA
101265482	1006	13.2	4.89	-1.432632	1.21E-30	Down	1.01E-31	LOC101265482	PREDICTED: Sol:GO:0090575.GO:000367	GO:000635	NA
101262509	942	45.63	16.92	-1.431253	8.56E-102	Down	2.58E-103	LOC101262509	PREDICTED: Sol:GO:0016021	NA	NA
104649135	480	35.1	13.02	-1.430742	2.31E-37	Down	1.65E-38	LOC104649135	PREDICTED: Sol:NA	NA	NA
101258004	1703	2.55	0.95	-1.424498	0.000992	Down	0.00037	LOC101258004	PREDICTED: Sol:NA	GO:000372	GO:004422
101252465	834	321.13	120.2	-1.417721	0	Down	0	LOC101252465	PREDICTED: Sol:NA	NA	NA

101266879	790	764.35	286.51	-1.415648	0	Down	0	LOC101266879	PREDICTED: Sol:GO:0016020,GO:000828,GO:000686 NA
101245625	1209	1.36	0.51	-1.415037	0.000114	Down	3.75E-05	LOC101245625	PREDICTED: Sol:NA GO:000828 NA NA
101254922	802	6.93	2.6	-1.414344	1.68E-13	Down	2.72E-14	LOC101254922	PREDICTED: Sol:NA NA NA K11804//
101246932	2814	1.57	0.59	-1.411978	8.73E-13	Down	1.47E-13	LOC101246932	PREDICTED: Sol:GO:0000783,GO:000365,GO:003162, K00430//
101261722	3632	19.69	7.41	-1.409918	6.29E-175	Down	1.16E-176	LOC101261722	PREDICTED: Sol:GO:0016021,GO:000472,GO:000605, K14157//a
101255051	1012	16.39	6.17	-1.409473	4.45E-39	Down	3.06E-40	LOC101255051	PREDICTED: Sol:GO:0016021,GO:001662,GO:000662, K10258//
101250569	1429	1.46	0.55	-1.408465	8.05E-06	Down	2.32E-06	LOC101250569	PREDICTED: Sol:GO:0005737,GO:000467,GO:002301, K04421//
101258706	2099	40.23	15.19	-1.40515	1.74E-202	Down	2.86E-204	LOC101258706	PREDICTED: Sol:GO:0016021,GO:001522,GO:000972, K20456//
101261765	1745	3.01	1.14	-1.40073	3.63E-13	Down	5.98E-14	LOC101261765	PREDICTED: Sol:NA GO:001671 NA K09754//c
101245153	1741	7.52	2.85	-1.399771	1.10E-31	Down	8.98E-33	LOC101245153	PREDICTED: Sol:GO:0016020,GO:001671,GO:004452, K20562//
101267355	3159	13.66	5.18	-1.398933	1.05E-103	Down	3.10E-105	LOC101267355	PREDICTED: Sol:NA GO:004698 NA K12126//
101246025	2172	8.78	3.34	-1.394373	3.16E-44	Down	1.97E-45	LOC101246025	PREDICTED: Sol:GO:0005634,GO:000367, NA NA
101246381	848	147.06	56.02	-1.392391	6.24E-280	Down	7.69E-282	LOC101246381	Solanum lycopersi NA NA GO:000692, K14496//a
101256767	1359	6.66	2.54	-1.390694	1.09E-21	Down	1.21E-22	LOC101256767	PREDICTED: Sol:GO:0016021,GO:001671,GO:000662, K10256//
101250515	1945	65.9	25.15	-1.38972	1.82E-305	Down	2.00E-307	SIADH12A1	Solanum lycopersi NA GO:001645, NA K00294//
101260284	1631	9.94	3.8	-1.387246	2.20E-38	Down	1.53E-39	LOC101260284	PREDICTED: Sol:GO:0016021, NA NA K01180//e
101267816	954	2.17	0.83	-1.386512	1.73E-05	Down	5.14E-06	LOC101267816	PREDICTED: Sol:NA NA NA NA
101248076	494	32.36	12.39	-1.385035	4.95E-34	Down	3.78E-35	LOC101248076	PREDICTED: Sol:GO:0016021, NA NA NA
101266903	1499	6.12	2.35	-1.380871	3.17E-22	Down	3.46E-23	LOC101266903	PREDICTED: Sol:NA GO:004687, NA K06892//f
101246328	2025	8.15	3.13	-1.380637	4.02E-39	Down	2.76E-40	LOC101246328	PREDICTED: Sol:NA GO:000848,GO:000902, K00814//a
101245159	759	5.33	2.05	-1.378512	1.06E-09	Down	2.19E-10	LOC101245159	PREDICTED: Sol:NA NA NA NA
101255829	941	6.7	2.58	-1.37679	7.90E-15	Down	1.19E-15	LOC101255829	PREDICTED: Sol:NA GO:001671,GO:000812, K07025//
100134881	1613	2.49	0.96	-1.375039	5.52E-10	Down	1.12E-10	LOC100134881	Solanum lycopersi NA NA NA K18081//
101255474	1366	21.6	8.33	-1.374643	1.24E-67	Down	5.39E-69	LOC101255474	PREDICTED: Sol:GO:0016021,GO:001671,GO:000662, K10256//
101247196	1378	99.36	38.33	-1.374191	0	Down	0	LOC101247196	PREDICTED: Sol:NA GO:000451, NA K08999//
101244900	2736	0.57	0.22	-1.373458	0.000188	Down	6.37E-05	LOC101244900	PREDICTED: Sol:GO:0016021, NA NA K01240//
101262640	1161	4.89	1.89	-1.371448	1.53E-13	Down	2.46E-14	LOC101262640	PREDICTED: Sol:NA GO:001671, NA NA
101246396	2154	12.62	4.88	-1.370759	3.25E-63	Down	1.49E-64	LOC101246396	Solanum lycopersi NA GO:002002,GO:000692, K10529//a
101258987	950	3.98	1.54	-1.369838	1.57E-09	Down	3.30E-10	LOC101258987	Solanum lycopersi GO:0009507,GO:002002,GO:000692, K00434//
101268143	704	52.81	20.47	-1.3673	3.14E-80	Down	1.16E-81	LOC101268143	PREDICTED: Sol:NA NA NA NA
101249960	1658	40.32	15.63	-1.367178	2.67E-155	Down	5.47E-157	LOC101249960	PREDICTED: Sol:NA GO:003022,GO:000592, K01792//
101251368	2471	10.57	4.1	-1.36628	9.02E-61	Down	4.26E-62	LOC101251368	PREDICTED: Sol:GO:0016021,GO:000551,GO:000861, K15404//a
101264731	1798	16.44	6.39	-1.363322	7.45E-68	Down	3.23E-69	LOC101264731	PREDICTED: Sol:GO:0005634,GO:004351, NA K09338//
101266146	923	33.83	13.15	-1.36324	7.01E-69	Down	2.99E-70	LOC101266146	PREDICTED: Sol:NA NA NA K04733//i
101265980	1037	21.45	8.37	-1.357678	4.23E-50	Down	2.35E-51	LOC101265980	PREDICTED: Sol:NA GO:000371, NA K08064//
101244665	1559	5.53	2.16	-1.356248	2.49E-19	Down	3.03E-20	LOC101244665	PREDICTED: Sol:NA NA NA K15631//
101249528	1168	1.51	0.59	-1.355762	9.75E-05	Down	3.18E-05	LOC101249528	PREDICTED: Sol:GO:0005615,GO:000822,GO:005161, K13441//c

101250559	1690	8.87	3.47	-1.353998	2.55E-34	Down	1.93E-35	LOC101250559	PREDICTED: Sol:GO:0016020,GO:00167(GO:00985:K07418//c
101254322	2448	38	14.93	-1.347785	1.69E-210	Down	2.68E-212	LOC101254322	PREDICTED: Sol:NA GO:00042:NA K12619//
101251255	1263	3.93	1.55	-1.342261	1.11E-11	Down	1.99E-12	LOC101251255	Solanum lycopersi NA GO:00468:GO:00096:K05933//a
101263630	770	89.61	35.6	-1.331782	6.97E-144	Down	1.55E-145	LOC101263630	PREDICTED: Sol:GO:0005634,GO:00435:NA K09338//
101262584	864	16.92	6.73	-1.330051	2.42E-31	Down	1.99E-32	LOC101262584	PREDICTED: Sol:GO:0016021,GO:00515:NA NA
104649031	1266	2.01	0.8	-1.329124	1.21E-05	Down	3.53E-06	LOC104649031	PREDICTED: Sol:NA NA NA NA
101245158	1083	9.39	3.74	-1.328087	1.43E-22	Down	1.54E-23	LOC101245158	PREDICTED: Sol:GO:0016021,NA NA NA
101266972	902	8.69	3.47	-1.324421	3.70E-17	Down	4.96E-18	LOC101266972	PREDICTED: Sol:GO:0005737,GO:00043(GO:00067:K00799//
101263489	1637	23.72	9.48	-1.323145	1.51E-84	Down	5.34E-86	LOC101263489	PREDICTED: Sol:NA GO:00468:NA K15172//t
101263945	1478	1.1	0.44	-1.321928	0.000221	Down	7.58E-05	LOC101263945	PREDICTED: Sol:GO:0016021,GO:00087:GO:00322:NA
101262912	1841	181.86	72.87	-1.319431	0	Down	0	LOC101262912	PREDICTED: Sol:GO:0005829,GO:00168:GO:00197:K01590//
543986	1205	202.05	80.97	-1.319253	0	Down	0	LOC543986	Solanum lycopersiGO:0005615,GO:00429:GO:00069:K19891//
101257330	1941	5.8	2.33	-1.315723	5.03E-25	Down	4.93E-26	LOC101257330	PREDICTED: Sol:GO:0043231,GO:00800:GO:00081:K18823//s
101255324	1070	66.88	26.9	-1.313969	3.96E-150	Down	8.41E-152	LOC101255324	PREDICTED: Sol:GO:0016021,NA NA NA
101255164	2189	5.19	2.09	-1.312232	2.61E-25	Down	2.53E-26	LOC101255164	PREDICTED: Sol:NA GO:00055:NA K03283//
101260130	1258	2.18	0.88	-1.308753	1.35E-06	Down	3.60E-07	LOC101260130	PREDICTED: Sol:NA NA NA K14299//
101245781	671	89.45	36.12	-1.308284	3.99E-120	Down	1.07E-121	LOC101245781	PREDICTED: Sol:NA NA GO:00069:NA
101261835	2011	55.47	22.4	-1.308209	2.85E-230	Down	4.25E-232	LOC101261835	PREDICTED: Sol:NA GO:00468:NA K14497//
543645	1261	10.85	4.4	-1.30212	1.89E-29	Down	1.63E-30	LOC543645	Solanum lycopersi NA GO:00090:NA NA
543815	838	61.25	24.87	-1.300303	2.64E-104	Down	7.78E-106	LOC543815	Solanum lycopersiGO:0005737,GO:00043(GO:00067:K00799//
101266721	2134	20.42	8.3	-1.2988	4.57E-93	Down	1.48E-94	LOC101266721	PREDICTED: Sol:NA NA GO:00071:K05747//
101260687	1011	2.16	0.88	-1.295456	2.81E-05	Down	8.56E-06	LOC101260687	PREDICTED: Sol:GO:0016021,GO:00616:GO:00431:K19040//
543731	2853	27.64	11.27	-1.29427	1.05E-167	Down	2.01E-169	sus3	PREDICTED: Sol:NA GO:00161:GO:00059:K00695//s
100134914	2589	4.19	1.71	-1.292954	8.73E-24	Down	8.94E-25	HSP70	Solanum lycopersi NA GO:00055:GO:00064:K03283//
101259953	614	8.93	3.65	-1.290764	1.94E-11	Down	3.54E-12	LOC101259953	PREDICTED: Sol:NA NA NA NA
101254669	1686	4.45	1.82	-1.289867	1.88E-16	Down	2.60E-17	LOC101254669	PREDICTED: Sol:GO:0016020,GO:00167(GO:00445:K20562//
101248118	1371	20.65	8.46	-1.287412	8.56E-50	Down	4.78E-51	LOC101248118	PREDICTED: Sol:NA NA NA K07305//
101265425	794	5.07	2.08	-1.285402	7.61E-09	Down	1.68E-09	LOC101265425	PREDICTED: Sol:GO:0005886,GO:00436:GO:00069:NA
101246536	1893	0.9	0.37	-1.2824	0.000235	Down	8.09E-05	LOC101246536	PREDICTED: Sol:NA NA NA K15271//
108511945	856	45.82	18.86	-1.280648	2.66E-78	Down	1.01E-79	ERF5	Solanum lycopersiGO:0005634,GO:00036:GO:00063:K09286//
544204	1849	74.56	30.7	-1.280163	5.82E-287	Down	6.90E-289	AREB	PREDICTED: Sol:NA GO:00036:NA K14432//
101254813	520	29.93	12.36	-1.275914	1.48E-29	Down	1.27E-30	LOC101254813	PREDICTED: Sol:NA NA NA NA
101267720	2695	0.75	0.31	-1.274622	3.90E-05	Down	1.21E-05	LOC101267720	PREDICTED: Sol:NA GO:00161:GO:00059:K00695//s
101250923	1335	3	1.24	-1.274622	6.36E-09	Down	1.40E-09	LOC101250923	PREDICTED: Sol:NA NA NA NA
101251700	1193	12.29	5.08	-1.274585	2.92E-30	Down	2.46E-31	LOC101251700	PREDICTED: Sol:NA GO:00167:GO:00081:K14493//
101249565	2082	4.77	1.98	-1.268489	4.23E-21	Down	4.79E-22	LOC101249565	PREDICTED: Sol:NA GO:00167(GO:00102:K09843//
101268013	1862	29.33	12.18	-1.267863	6.28E-112	Down	1.74E-113	LOC101268013	PREDICTED: Sol:GO:0016021,GO:00167(NA K07428//c

101268294	1669	21.25	8.83	-1.266977	1.63E-72	Down	6.66E-74	LOC101268294	PREDICTED: Sol:GO:0005737.GO:003162GO:00431K10523//s
101262624	1370	1.13	0.47	-1.26559	0.000581	Down	0.00021	LOC101262624	PREDICTED: Sol:NA GO:001682GO:000902K21407//a
101260177	4464	1.13	0.47	-1.26559	3.30E-11	Down	6.15E-12	LOC101260177	PREDICTED: Sol:GO:0016021.GO:000552GO:000681K05681//
101243932	733	12.21	5.08	-1.265163	7.24E-18	Down	9.41E-19	LOC101243932	PREDICTED: Sol:NA NA NA NA
101255139	1714	30.79	12.82	-1.264066	2.49E-107	Down	7.16E-109	LOC101255139	PREDICTED: Sol:GO:0016021.NA NA K08145//
104644746	920	5.21	2.17	-1.263588	2.24E-10	Down	4.41E-11	LOC104644746	PREDICTED: Sol:NA NA NA NA
101260643	4547	7.31	3.05	-1.261062	1.91E-69	Down	8.06E-71	LOC101260643	PREDICTED: Sol:GO:0016021.GO:000552GO:000681K05681//
101250797	1076	3.66	1.53	-1.258312	1.42E-08	Down	3.20E-09	LOC101250797	PREDICTED: Sol:GO:0005634.GO:000367NA K10773//e
101254884	755	6.3	2.64	-1.254814	7.34E-10	Down	1.50E-10	LOC101254884	PREDICTED: Sol:GO:0005634.GO:000367NA K05747//
101245324	1627	1.19	0.5	-1.250962	8.73E-05	Down	2.83E-05	LOC101245324	PREDICTED: Sol:GO:0016021.NA NA K14209//s
101268010	1409	3.4	1.43	-1.24952	2.74E-10	Down	5.44E-11	LOC101268010	PREDICTED: Sol:NA NA NA K13126//
104647874	580	12.63	5.32	-1.247356	3.80E-14	Down	5.95E-15	LOC104647874	PREDICTED: Sol:GO:0016021.NA NA NA
101247545	4612	3.48	1.47	-1.243271	2.59E-33	Down	2.01E-34	LOC101247545	PREDICTED: Sol:GO:0016021.GO:000552GO:000681K05681//
101258887	1257	25.06	10.62	-1.238603	2.28E-61	Down	1.07E-62	LOC101258887	PREDICTED: Sol:GO:0005576.NA NA K20628//e
101249973	951	46.53	19.77	-1.234848	2.44E-84	Down	8.65E-86	LOC101249973	PREDICTED: Sol:GO:0009506.GO:000487GO:000692NA
101266533	733	37.18	15.8	-1.234602	8.01E-51	Down	4.39E-52	LOC101266533	PREDICTED: Sol:NA GO:000552NA K16465//c
101262714	1249	1.6	0.68	-1.234465	0.0001	Down	3.28E-05	LOC101262714	PREDICTED: Sol:NA NA NA NA
101265748	1745	3.08	1.31	-1.233364	2.87E-11	Down	5.32E-12	LOC101265748	PREDICTED: Sol:NA NA NA K01214//i
101246876	1745	9.49	4.04	-1.232053	5.79E-33	Down	4.55E-34	LOC101246876	PREDICTED: Sol:GO:0016021.GO:000851NA K03320//a
101260481	2158	49.95	21.34	-1.226925	1.79E-210	Down	2.85E-212	LOC101260481	PREDICTED: Sol:GO:0016021.NA GO:000962NA
101268009	910	11.97	5.12	-1.225207	1.59E-25	Down	1.53E-26	LOC101268009	PREDICTED: Sol:GO:0016021.NA NA K06890//
101262527	1202	47.39	20.28	-1.224525	2.46E-108	Down	7.00E-110	LOC101262527	PREDICTED: Sol:GO:0045263.GO:001507GO:001598K13153//
101252269	1092	13.15	5.63	-1.223856	1.16E-27	Down	1.05E-28	LOC101252269	PREDICTED: Sol:GO:0046658.GO:000902NA K03006//
101266245	938	84.18	36.06	-1.223078	5.41E-147	Down	1.18E-148	LOC101266245	PREDICTED: Sol:GO:0009507.GO:005032GO:000652K22246//t
101249488	775	11.4	4.89	-1.221127	6.31E-17	Down	8.56E-18	LOC101249488	PREDICTED: Sol:NA NA NA NA
101257929	744	120.61	51.77	-1.220161	4.11E-163	Down	8.04E-165	LOC101257929	PREDICTED: Sol:NA NA NA NA
101244335	1010	1.91	0.82	-1.219877	0.000176	Down	5.94E-05	LOC101244335	PREDICTED: Sol:NA NA NA K13945//
101256273	1953	7.38	3.17	-1.219138	6.26E-29	Down	5.48E-30	LOC101256273	PREDICTED: Sol:NA NA NA K16865//
101266149	896	2	0.86	-1.217591	0.000341	Down	0.00012	LOC101266149	PREDICTED: Sol:GO:0005737.GO:000432GO:000672K00799//
101243709	2240	2.09	0.9	-1.215506	6.34E-10	Down	1.29E-10	LOC101243709	PREDICTED: Sol:GO:0016021.GO:000552NA K04733//i
101257491	2571	15.6	6.72	-1.215013	7.19E-78	Down	2.74E-79	LOC101257491	PREDICTED: Sol:GO:0016021.NA NA K15503//s
101244959	2087	21.8	9.4	-1.213595	1.27E-87	Down	4.31E-89	LOC101244959	PREDICTED: Sol:NA NA NA K01599//
101265813	989	5.35	2.31	-1.211646	1.40E-10	Down	2.72E-11	LOC101265813	PREDICTED: Sol:NA NA NA NA
101263832	1799	0.9	0.39	-1.206451	0.000755	Down	0.000277	LOC101263832	PREDICTED: Sol:GO:0005886.GO:000552GO:000642K04733//i
101249525	1359	4.52	1.96	-1.205469	3.10E-12	Down	5.38E-13	LOC101249525	PREDICTED: Sol:NA GO:001672NA K19747//
101249743	1531	4.77	2.07	-1.204358	2.01E-14	Down	3.09E-15	LOC101249743	PREDICTED: Sol:GO:0016021.NA GO:005511NA
543895	2164	0.99	0.43	-1.203092	5.91E-05	Down	1.88E-05	LOC543895	Solanum lycopersi NA GO:002002GO:000692K10529//a

101265590	1068	108.68	47.37	-1.198041	6.70E-211	Down	1.06E-212	LOC101265590	PREDICTED: Sol:GO:0005737.GO:000892GO:000676NA
101268197	1268	1.53	0.67	-1.191299	0.0002	Down	6.80E-05	LOC101268197	PREDICTED: Sol:GO:0016021.GO:000837GO:000648K20843//
101265690	986	25.78	11.29	-1.191207	3.97E-46	Down	2.39E-47	LOC101265690	PREDICTED: Sol:NA GO:004687GO:00098(K00588//c
101268222	2093	10.04	4.4	-1.190184	1.22E-39	Down	8.32E-41	LOC101268222	PREDICTED: Sol:NA NA NA NA
100736441	1361	203.2	89.06	-1.190051	0	Down	0	LOC100736441	Solanum lycopersiGO:0005768.NA GO:00096:K06699//
101256484	833	2.21	0.97	-1.18799	0.000387	Down	0.000137	LOC101256484	PREDICTED: Sol:NA NA NA K21373//
101251778	700	3.98	1.75	-1.185414	1.17E-05	Down	3.40E-06	LOC101251778	Solanum lycopersiGO:0005634.GO:00435(GO:00063:K18835//
101261506	2383	3.16	1.39	-1.18484	8.32E-14	Down	1.32E-14	LOC101261506	PREDICTED: Sol:NA NA GO:00063:NA
101265775	1049	49.41	21.75	-1.183788	1.19E-92	Down	3.85E-94	LOC101265775	PREDICTED: Sol:GO:0016021.GO:00090:NA K03006//
101246190	1233	5.29	2.33	-1.182938	1.69E-12	Down	2.89E-13	LOC101246190	PREDICTED: Sol:NA GO:004687NA K18054//
104648182	3157	1.18	0.52	-1.182203	5.06E-05	Down	1.59E-05	LOC104648182	PREDICTED: Sol:NA GO:000367GO:001507K12447//
101261429	1607	8.94	3.94	-1.182079	3.82E-27	Down	3.51E-28	LOC101261429	PREDICTED: Sol:NA GO:001678GO:00092:K20884//r
101256961	2797	4.17	1.84	-1.180342	5.22E-21	Down	5.94E-22	LOC101256961	PREDICTED: Sol:GO:0005622.GO:000414GO:00072(K00901//
543963	714	14.98	6.61	-1.180315	4.31E-19	Down	5.30E-20	RSI-1	Solanum lycopersiGO:0009506.NA GO:00097:K03860//
101251958	1529	1.11	0.49	-1.179706	0.000541	Down	0.000195	LOC101251958	PREDICTED: Sol:GO:0016021.GO:006162GO:00431(K15692//
101261914	2238	1.97	0.87	-1.179108	4.84E-09	Down	1.05E-09	LOC101261914	PREDICTED: Sol:NA GO:001682GO:000597K18195//r
101258189	2334	3.26	1.44	-1.178803	7.92E-15	Down	1.19E-15	sbt4e	Solanum lycopersi NA GO:00042:NA K12619//
101250113	1406	2.71	1.2	-1.175258	1.17E-07	Down	2.85E-08	LOC101250113	PREDICTED: Sol:NA GO:001678NA K21026//a
101260072	1639	12.67	5.63	-1.17021	7.26E-38	Down	5.12E-39	LOC101260072	PREDICTED: Sol:NA GO:00200:GO:000697K14454//a
101265851	868	17.21	7.66	-1.167831	2.17E-26	Down	2.04E-27	LOC101265851	PREDICTED: Sol:NA GO:00163(NA K13418//s
101267225	785	5.59	2.49	-1.166703	2.56E-08	Down	5.88E-09	LOC101267225	PREDICTED: Sol:NA NA NA NA
101268896	1262	1.39	0.62	-1.164745	0.000609	Down	0.000221	LOC101268896	PREDICTED: Sol:GO:0005622.GO:000482GO:000651K05609//
101244740	1566	7.3	3.26	-1.163024	5.07E-21	Down	5.76E-22	LOC101244740	Solanum lycopersiGO:0016021.NA NA K20854//
778337	2213	3	1.34	-1.16273	1.01E-12	Down	1.70E-13	ftsH6	Solanum lycopersiGO:0016020.GO:00082:GO:00103(K03798//c
101257549	1062	2.28	1.02	-1.160465	4.89E-05	Down	1.54E-05	LOC101257549	PREDICTED: Sol:NA NA GO:00069:NA
101268787	1893	21.21	9.5	-1.158745	7.72E-72	Down	3.19E-73	LOC101268787	PREDICTED: Sol:GO:0005634.GO:00435(GO:00508:K13424//
101246306	1351	20.45	9.17	-1.157107	9.95E-49	Down	5.69E-50	LOC101246306	PREDICTED: Sol:NA NA NA K13783//
101251627	1362	19.33	8.67	-1.156738	1.53E-46	Down	9.09E-48	LOC101251627	PREDICTED: Sol:GO:0005829.GO:001858GO:00466:NA
101261104	1493	343.34	154.1	-1.155771	0	Down	0	LOC101261104	PREDICTED: Sol:GO:0016021.GO:005111GO:00086:K15382//s
101247272	1110	208.6	93.65	-1.155388	0	Down	0	LOC101247272	PREDICTED: Sol:GO:0016021.NA NA NA
101250935	1949	75	33.71	-1.153714	3.12E-258	Down	4.16E-260	LOC101250935	PREDICTED: Sol:GO:0016021.GO:001517GO:00033:K19476//
544074	1684	2.38	1.07	-1.153351	7.72E-08	Down	1.84E-08	ADH2	Solanum lycopersi NA GO:000402NA K18857//a
101254095	1203	11.3	5.09	-1.150585	4.34E-24	Down	4.38E-25	LOC101254095	PREDICTED: Sol:GO:0016021.NA NA NA
101258849	1085	1.95	0.88	-1.147899	0.000182	Down	6.16E-05	LOC101258849	PREDICTED: Sol:NA GO:00469:NA K04733//i
101258039	2460	5.27	2.38	-1.146841	1.81E-23	Down	1.87E-24	LOC101258039	PREDICTED: Sol:NA NA NA K19613//l
544082	895	2044.89	925.74	-1.143344	0	Down	0	TPM-1	Solanum lycopersi NA NA NA K04733//i
101265726	888	3.07	1.39	-1.143154	2.05E-05	Down	6.14E-06	LOC101265726	PREDICTED: Sol:NA NA NA NA

101247130	895	2.87	1.3	-1.142539	3.94E-05	Down	1.22E-05	LOC101247130	PREDICTED: Sol:NA	GO:000485	NA	K01051//	
101264592	2616	0.75	0.34	-1.141356	0.000253	Down	8.73E-05	LOC101264592	PREDICTED: Sol:NA	GO:004355	GO:000695	K13457//	
778266	840	9.38	4.26	-1.138734	9.41E-14	Down	1.50E-14	LOC778266	Solanum lycopersi	GO:0016021	NA	NA	
101257269	1737	15.67	7.12	-1.138056	2.07E-47	Down	1.21E-48	LOC101257269	PREDICTED: Sol:GO:0009505	GO:005265	GO:000595	K07407//a	
101251145	1276	72.47	32.93	-1.137981	6.60E-157	Down	1.34E-158	Rcr3	Solanum lycopersi	GO:0005615	GO:000415	GO:005160	K13441//c
101266204	1225	12.84	5.84	-1.136605	3.29E-27	Down	3.02E-28	LOC101266204	PREDICTED: Sol:GO:0009506	NA	NA	K15382//s	
101251753	1365	3.45	1.57	-1.135832	4.50E-09	Down	9.79E-10	LOC101251753	PREDICTED: Sol:NA	NA	NA	NA	
101244694	1697	5.14	2.34	-1.13526	9.67E-16	Down	1.39E-16	LOC101244694	Solanum lycopersi	NA	GO:001645	NA	K11153//r
109120085	1123	10.14	4.62	-1.134093	3.78E-20	Down	4.45E-21	LOC109120085	PREDICTED: Sol:NA	GO:001678	NA	K21026//a	
101265869	885	6.21	2.83	-1.133791	7.12E-10	Down	1.46E-10	LOC101265869	PREDICTED: Sol:GO:0016021	NA	NA	NA	
101255309	2084	6.03	2.75	-1.132726	3.50E-22	Down	3.82E-23	LOC101255309	PREDICTED: Sol:GO:0005737	GO:000372	NA	K03265//	
101257682	1422	9.56	4.36	-1.132682	1.07E-23	Down	1.10E-24	LOC101257682	PREDICTED: Sol:GO:0005618	GO:004535	GO:004545	K01051//	
104649684	740	4.51	2.06	-1.130483	0.000119	Down	3.93E-05	LOC104649684	PREDICTED: Sol:NA	GO:000367	NA	K05666//	
101259283	1553	11.28	5.16	-1.128324	3.38E-30	Down	2.86E-31	LOC101259283	PREDICTED: Sol:NA	GO:000382	NA	K08726//s	
101259198	791	838.75	383.92	-1.127435	0	Down	0	LOC101259198	PREDICTED: Sol:NA	NA	NA	NA	
101266243	1063	4.87	2.23	-1.126878	2.36E-09	Down	5.03E-10	LOC101266243	PREDICTED: Sol:NA	NA	NA	NA	
778321	807	7.73	3.54	-1.126719	6.50E-11	Down	1.24E-11	LOC778321	Solanum lycopersi	NA	NA	GO:000695	NA
101266856	1588	7.58	3.48	-1.123111	6.01E-21	Down	6.87E-22	LOC101266856	PREDICTED: Sol:GO:0043231	GO:008002	GO:000815	K21374//	
101248665	1239	22.36	10.28	-1.12108	2.39E-46	Down	1.43E-47	LOC101248665	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	K12862//	
101258824	1987	6.24	2.87	-1.120495	1.17E-21	Down	1.30E-22	LOC101258824	PREDICTED: Sol:NA	NA	NA	NA	
101252505	1388	273.52	125.83	-1.12017	0	Down	0	Cyp-3	Solanum lycopersi	GO:0005615	GO:000825	GO:005160	K01366//c
101267829	1523	16.3	7.5	-1.119909	5.01E-42	Down	3.27E-43	LOC101267829	PREDICTED: Sol:GO:0005886	GO:000395	GO:190100	NA	
101255714	2301	0.91	0.42	-1.115477	0.000163	Down	5.49E-05	LOC101255714	PREDICTED: Sol:NA	NA	NA	K10278//	
101244186	2257	22.55	10.41	-1.115157	5.22E-86	Down	1.81E-87	LOC101244186	PREDICTED: Sol:NA	NA	NA	NA	
101243979	958	2.62	1.21	-1.11456	7.56E-05	Down	2.43E-05	LOC101243979	Solanum lycopersi	NA	NA	NA	K04733//i
543990	1804	1661.58	767.53	-1.114261	0	Down	0	cat1	Solanum lycopersi	GO:0009514	GO:004687	GO:000695	K03781//c
101256047	1575	2.12	0.98	-1.113211	2.54E-06	Down	6.97E-07	LOC101256047	PREDICTED: Sol:NA	GO:000810	GO:000665	K00559//s	
101267548	1164	5.06	2.34	-1.112629	1.93E-10	Down	3.79E-11	LOC101267548	Solanum lycopersi	GO:0005634	GO:000367	GO:000635	K09286//
101261854	1691	13.11	6.08	-1.108524	3.54E-37	Down	2.54E-38	LOC101261854	PREDICTED: Sol:NA	NA	NA	K21440//a	
101263788	780	6.23	2.89	-1.108163	1.87E-08	Down	4.25E-09	LOC101263788	PREDICTED: Sol:NA	GO:004687	NA	K13811//	
101244582	1354	19.7	9.15	-1.106352	6.30E-44	Down	3.94E-45	LOC101244582	Solanum lycopersi	GO:0005634	GO:000367	GO:000635	K03094//
101268757	1419	1.57	0.73	-1.104796	0.000163	Down	5.46E-05	LOC101268757	PREDICTED: Sol:NA	GO:000552	NA	K20043//c	
101267987	1962	6.75	3.14	-1.104123	1.32E-22	Down	1.41E-23	LOC101267987	PREDICTED: Sol:GO:0016020	GO:001670	GO:004455	K14985//	
101253953	583	28.95	13.49	-1.101673	5.62E-26	Down	5.35E-27	LOC101253953	PREDICTED: Sol:NA	NA	NA	NA	
101261098	1682	2.51	1.17	-1.101179	8.63E-08	Down	2.07E-08	LOC101261098	PREDICTED: Sol:GO:0005743	GO:001511	GO:000965	K15102//s	
101250024	2277	6.65	3.1	-1.101086	2.21E-24	Down	2.21E-25	LOC101250024	PREDICTED: Sol:GO:0016021	NA	GO:000862	K08150//	
101247199	562	5.04	2.35	-1.100763	4.35E-05	Down	1.36E-05	LOC101247199	PREDICTED: Sol:NA	NA	NA	NA	

101247807	1488	23.24	10.84	-1.100245	3.02E-56	Down	1.53E-57	LOC101247807	PREDICTED: Sol:NA	GO:001682	GO:000652	K01620//t	
101257043	1048	3.51	1.64	-1.097775	1.06E-06	Down	2.80E-07	LOC101257043	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	K09286//	
101261451	1078	63.25	29.56	-1.097419	3.24E-108	Down	9.27E-110	LOC101261451	PREDICTED: Sol:NA	GO:001678	NA	K01061//c	
101249360	2435	0.92	0.43	-1.097297	0.000145	Down	4.84E-05	LOC101249360	PREDICTED: Sol:GO:0005634	GO:000482	NA	K08332//	
101261650	1284	1650.73	772.08	-1.096282	0	Down	0	LOC101261650	Solanum lycopersi	GO:0046658	GO:004297	GO:000695	K19891//
101259182	917	1410.05	659.59	-1.096105	0	Down	0	LOC101259182	PREDICTED: Sol:NA	NA	NA	K13344//	
101260164	1347	125.35	58.65	-1.095759	4.20E-270	Down	5.34E-272	LOC101260164	PREDICTED: Sol:NA	NA	NA	NA	
778364	1247	3.74	1.75	-1.095683	1.87E-08	Down	4.25E-09	Asc-1	Solanum lycopersi	GO:0005783	GO:005025	GO:004651	K04710//c
101245588	806	30.68	14.37	-1.094238	8.34E-39	Down	5.77E-40	LOC101245588	Solanum lycopersi	NA	NA	GO:000695	K06674//s
101255009	1506	8.39	3.93	-1.094141	5.90E-21	Down	6.73E-22	LOC101255009	PREDICTED: Sol:GO:0019898	GO:008002	GO:003445	K17908//a	
100191111	832	1615.3	758.09	-1.091361	0	Down	0	LOC100191111	Solanum lycopersi	GO:0005576	NA	NA	K13449//
101252347	902	12.03	5.65	-1.090314	1.51E-17	Down	1.99E-18	LOC101252347	PREDICTED: Sol:GO:0005840	GO:007018	GO:000641	K02990//s	
101261400	1867	27.34	12.85	-1.089245	1.43E-82	Down	5.16E-84	LOC101261400	PREDICTED: Sol:NA	GO:000482	NA	K06001//t	
101253070	1077	4.01	1.89	-1.085216	6.90E-06	Down	1.97E-06	LOC101253070	PREDICTED: Sol:NA	NA	NA	NA	
104647150	3789	0.7	0.33	-1.084889	4.28E-05	Down	1.34E-05	LOC104647150	PREDICTED: Sol:NA	NA	NA	K01528//	
544148	1154	1306.5	616.67	-1.083137	0	Down	0	CHI9	Solanum lycopersi	GO:0005618	GO:000806	GO:000605	K20547//
101247551	2554	7.09	3.35	-1.081625	4.54E-30	Down	3.85E-31	LOC101247551	PREDICTED: Sol:NA	GO:000367	NA	K14007//	
544193	974	5.1	2.41	-1.081464	1.83E-08	Down	4.16E-09	rnalx	Solanum lycopersi	GO:0005737	GO:000372	NA	K01166//r
101245962	1066	10.37	4.91	-1.078621	9.54E-18	Down	1.24E-18	LOC101245962	PREDICTED: Sol:GO:0048046	GO:001676	GO:001041	K08235//	
543871	2389	34.25	16.22	-1.07833	5.05E-131	Down	1.25E-132	FRO1	Solanum lycopersi	GO:0016021	GO:001645	NA	K00521//f
101253408	2746	13.67	6.48	-1.076948	1.43E-60	Down	6.77E-62	LOC101253408	PREDICTED: Sol:GO:0016021	NA	NA	NA	
101266225	777	7.32	3.47	-1.076908	7.90E-10	Down	1.62E-10	LOC101266225	PREDICTED: Sol:NA	NA	NA	K01513//e	
101262849	1047	2.13	1.01	-1.076498	0.000249	Down	8.57E-05	LOC101262849	PREDICTED: Sol:NA	NA	NA	NA	
101266039	2296	3.1	1.47	-1.076452	4.14E-07	Down	1.06E-07	LOC101266039	PREDICTED: Sol:NA	NA	NA	K11267//s	
101250708	637	26.51	12.61	-1.071968	3.31E-25	Down	3.22E-26	LOC101250708	PREDICTED: Sol:GO:0005737	GO:004687	GO:003006	K04733//i	
101263056	1132	6.57	3.13	-1.069731	4.07E-12	Down	7.12E-13	LOC101263056	PREDICTED: Sol:GO:0005634	GO:000367	GO:000635	K09286//	
101255440	1508	17.07	8.14	-1.068362	1.80E-40	Down	1.20E-41	LOC101255440	PREDICTED: Sol:GO:0000326	NA	GO:004831	K12472//e	
104647325	1746	1.74	0.83	-1.067904	1.29E-05	Down	3.77E-06	LOC104647325	PREDICTED: Sol:NA	GO:000812	NA	K01025//s	
101243985	1886	3.96	1.89	-1.067114	1.90E-12	Down	3.26E-13	LOC101243985	PREDICTED: Sol:NA	GO:000552	NA	K04420//	
101249493	1919	24.19	11.55	-1.066518	9.12E-73	Down	3.70E-74	LOC101249493	PREDICTED: Sol:NA	GO:000376	NA	K09060//	
101262056	2239	1.34	0.64	-1.066089	1.58E-05	Down	4.70E-06	LOC101262056	PREDICTED: Sol:NA	NA	NA	K20254//	
544142	874	104.64	49.99	-1.065723	4.90E-136	Down	1.17E-137	LOC544142	Solanum lycopersi	GO:0005634	GO:000367	GO:000941	K11275//
101055553	684	14.45	6.92	-1.062226	5.06E-15	Down	7.55E-16	LOC101055553	Solanum lycopersi	GO:0005634	NA	GO:000635	K14484//a
101258071	1641	8.99	4.31	-1.060633	2.29E-23	Down	2.38E-24	LOC101258071	Solanum lycopersi	NA	GO:004687	NA	K14497//
109119847	803	4.75	2.28	-1.058894	1.60E-06	Down	4.28E-07	LOC109119847	PREDICTED: Sol:NA	GO:000486	NA	NA	
100271890	1256	34.86	16.74	-1.058273	1.74E-66	Down	7.69E-68	ZIP	Solanum lycopersi	NA	GO:000376	NA	K09060//
101250791	2212	4.06	1.95	-1.058006	1.22E-14	Down	1.85E-15	LOC101250791	PREDICTED: Sol:GO:0016021	GO:000521	NA	K14638//s	

101261885	694	4.39	2.11	-1.056978	3.03E-05	Down	9.27E-06	LOC101261885	PREDICTED: Sol:NA	GO:000367	NA	NA	
101250444	1756	12.27	5.9	-1.056348	1.33E-33	Down	1.02E-34	LOC101250444	PREDICTED: Sol:GO:0016021	GO:001670	NA	K20661//c	
543944	1500	13.68	6.58	-1.055909	9.38E-32	Down	7.61E-33	ARG1	Solanum lycopersi	NA	GO:004687	NA	K01476//a
104649084	524	7.1	3.42	-1.053823	5.21E-06	Down	1.47E-06	LOC104649084	PREDICTED: Sol:GO:0016021	NA	NA	K00030//i	
109120423	1138	11.02	5.31	-1.05334	1.93E-19	Down	2.33E-20	LOC109120423	PREDICTED: Sol:NA	NA	NA	NA	
101249999	1876	4.69	2.26	-1.053265	3.20E-14	Down	4.99E-15	LOC101249999	PREDICTED: Sol:GO:0016021	GO:001670	NA	K20661//c	
101243997	4996	1.39	0.67	-1.052852	1.05E-11	Down	1.89E-12	LOC101243997	PREDICTED: Sol:GO:0016021	GO:000552	GO:000681	K05681//	
101257056	1107	59.15	28.54	-1.051393	2.58E-97	Down	8.10E-99	LOC101257056	PREDICTED: Sol:NA	NA	NA	NA	
101265728	1705	121	58.4	-1.050967		Down	0	LOC101265728	PREDICTED: Sol:NA	NA	NA	K19475//	
543878	984	877.56	424.13	-1.048991		Down	0	LOC543878	Solanum lycopersi	NA	NA	NA	K04733//i
101257730	1712	2.42	1.17	-1.048499	3.39E-07	Down	8.56E-08	LOC101257730	PREDICTED: Sol:GO:0016021	NA	GO:005508	K02575//	
543574	711	237.4	114.79	-1.048323	4.79E-239	Down	6.83E-241	ASR1	Solanum lycopersi	NA	GO:000692	NA	
101246437	2253	7.59	3.67	-1.04832	1.00E-26	Down	9.31E-28	LOC101246437	PREDICTED: Sol:GO:0016021	GO:001670	NA	K13680//	
104648101	1122	4.77	2.31	-1.046096	1.11E-08	Down	2.48E-09	LOC104648101	PREDICTED: Sol:GO:0009506	GO:005108	GO:000645	NA	
101246295	1578	49.33	23.96	-1.041837	4.18E-116	Down	1.14E-117	LOC101246295	PREDICTED: Sol:GO:0043231	GO:008002	GO:000815	K13495//c	
101255252	1802	9.95	4.84	-1.039689	1.64E-27	Down	1.50E-28	LOC101255252	PREDICTED: Sol:GO:0016021	GO:001522	NA	K03327//	
101259463	1955	3.7	1.8	-1.039528	1.38E-11	Down	2.50E-12	LOC101259463	PREDICTED: Sol:GO:0016021	NA	GO:005120	K03217//	
101249609	1271	2.01	0.98	-1.036342	0.000124	Down	4.10E-05	LOC101249609	PREDICTED: Sol:NA	GO:000875	GO:001945	K16040//t	
101245191	1052	7.7	3.76	-1.034126	1.90E-12	Down	3.26E-13	LOC101245191	PREDICTED: Sol:NA	NA	NA	NA	
101259361	1512	7.31	3.58	-1.029912	5.74E-17	Down	7.75E-18	LOC101259361	PREDICTED: Sol:GO:0009507	GO:000897	GO:000657	K15227//a	
104645573	1022	20.99	10.28	-1.029862	3.09E-31	Down	2.54E-32	LOC104645573	PREDICTED: Sol:GO:0016021	NA	NA	NA	
101246812	1434	15.94	7.81	-1.029257	7.38E-34	Down	5.65E-35	LOC101246812	Solanum lycopersi	GO:0005634	GO:004350	GO:000961	K13424//
101251574	1374	130.16	63.86	-1.027302	3.62E-258	Down	4.84E-260	LOC101251574	PREDICTED: Sol:NA	NA	NA	K18342//	
101251704	1399	2.14	1.05	-1.027221	2.95E-05	Down	9.00E-06	LOC101251704	PREDICTED: Sol:GO:0016020	NA	GO:000710	K15503//s	
101245084	819	20.93	10.27	-1.027136	1.59E-24	Down	1.58E-25	LOC101245084	PREDICTED: Sol:GO:0016021	NA	NA	NA	
104647159	1229	4.87	2.39	-1.026911	3.29E-08	Down	7.61E-09	LOC104647159	PREDICTED: Sol:NA	NA	NA	NA	
101244669	2594	11.69	5.74	-1.026152	2.49E-44	Down	1.55E-45	LOC101244669	PREDICTED: Sol:GO:0005887	GO:000522	GO:004235	K05391//c	
101259235	1788	3.93	1.93	-1.025928	3.62E-11	Down	6.75E-12	LOC101259235	PREDICTED: Sol:NA	GO:000435	GO:000655	K01580//	
101262338	1658	9.74	4.79	-1.023896	3.50E-24	Down	3.52E-25	LOC101262338	PREDICTED: Sol:GO:0012505	GO:000012	GO:000688	K08486//s	
101267118	1632	13.54	6.66	-1.023634	1.20E-32	Down	9.45E-34	LOC101267118	PREDICTED: Sol:NA	GO:001662	NA	K00166//	
100736519	1734	209.49	103.07	-1.023257		Down	0	TCP12	Solanum lycopersi	GO:0005634	GO:000367	GO:000635	K13035//
104648432	1594	2.09	1.03	-1.020859	1.18E-05	Down	3.46E-06	LOC104648432	PREDICTED: Sol:NA	NA	NA	NA	
101259340	744	316.99	156.25	-1.020581		Down	0	LOC101259340	PREDICTED: Sol:NA	NA	GO:000692	NA	
101268692	1654	11.82	5.83	-1.019662	7.63E-29	Down	6.70E-30	LOC101268692	PREDICTED: Sol:NA	NA	NA	K03283//	
101266611	2662	4.63	2.29	-1.015665	1.35E-18	Down	1.70E-19	LOC101266611	PREDICTED: Sol:GO:0016021	NA	GO:000662	K01052//l	
101250644	1430	1.88	0.93	-1.01543	0.000106	Down	3.49E-05	LOC101250644	PREDICTED: Sol:GO:0016021	GO:001670	NA	K01988//l	
101247777	1731	2.83	1.4	-1.015375	6.07E-08	Down	1.43E-08	LOC101247777	PREDICTED: Sol:GO:0016020	GO:001670	GO:004455	K07418//c	

101252574	1048	96.36	47.7	-1.014445	1.17E-140	Down	2.71E-142	LOC101252574	PREDICTED: Sol:NA	GO:00080	NA	K22246//t
101247504	1482	3.53	1.75	-1.012313	2.74E-08	Down	6.30E-09	LOC101247504	PREDICTED: Sol:GO:0005634.GO:00009	GO:00063	NA	K12135//z
778249	1259	179.51	89	-1.012187		Down	0	CIP2b	Solanum lycopersi	GO:0005634.GO:00469	NA	K08066//
101261342	1426	18.51	9.19	-1.010168	1.02E-36	Down	7.40E-38	LOC101261342	PREDICTED: Sol:GO:0005634.GO:00036	GO:00063	NA	K13126//
101244173	2308	1.75	0.87	-1.008268	1.54E-06	Down	4.14E-07	LOC101244173	PREDICTED: Sol:GO:0005739.NA	GO:00083	NA	K17964//l
101246456	369	7.68	3.82	-1.007534	0.000278	Down	9.65E-05	LOC101246456	PREDICTED: Sol:GO:0016020.GO:00082	GO:00068	NA	
543736	2390	13.71	6.82	-1.007385	1.97E-43	Down	1.25E-44	tEG3	PREDICTED: Sol:GO:0005618.GO:00045	GO:00059	NA	K12309//
101252968	1980	109.15	54.32	-1.006757		Down	0	LOC101252968	PREDICTED: Sol:NA	GO:00164	NA	K14085//a
101253864	1809	6.34	3.16	-1.004558	6.51E-17	Down	8.83E-18	LOC101253864	PREDICTED: Sol:GO:0043231.GO:00800	GO:00081	NA	K03809//
101247407	627	4.39	2.19	-1.00329	0.000179	Down	6.05E-05	LOC101247407	PREDICTED: Sol:GO:0043231.GO:00800	GO:00081	NA	K13495//c
101261035	593	13.21	6.59	-1.00328	5.98E-11	Down	1.13E-11	LOC101261035	PREDICTED: Sol:NA	NA	NA	NA
101257639	993	12.94	6.46	-1.002232	3.56E-18	Down	4.57E-19	LOC101257639	PREDICTED: Sol:NA	GO:00469	NA	K04733//i
101252947	1171	60.93	30.42	-1.002133	2.21E-98	Down	6.86E-100	LOC101252947	PREDICTED: Sol:NA	GO:00301	NA	K15631//
101258448	751	13.96	6.97	-1.002068	1.11E-14	Down	1.68E-15	LOC101258448	PREDICTED: Sol:NA	NA	NA	NA
101263123	1429	53.11	26.52	-1.001903	9.71E-106	Down	2.83E-107	LOC101263123	PREDICTED: Sol:GO:0005634.GO:00044	GO:00431	NA	K10523//s
101253557	6043	8.99	4.49	-1.001606	1.24E-78	Down	4.67E-80	LOC101253557	PREDICTED: Sol:NA	NA	NA	K04733//i
101254816	1688	4.44	2.22	-1	2.31E-11	Down	4.24E-12	LOC101254816	PREDICTED: Sol:GO:0016021.NA	GO:00157	NA	K03695//

Supplementary Table S3. Expression data of the differentially expressed genes (DEGs) identified by RNA-seq analysis. (B) Expression data of the 84 differentially expressed transcription factors (DETFs) identified by RNA-seq analysis.

GeneID	Length	Control FPKM	PepMV FPKM	log2FC (PepMV/Control)	Pvalue	Transcripts	TF_family	Symbol	Description	Cellular Component	Molecular Function	Biological Process	Kegg Orthology
101263989 (Solyc12g096070)	1836	2.14	13.07	2.610576439	5.95E-150	XM_010316518.2	GRF	LOC101263989	PREDICTED: ScNA	NA	NA	NA	K20496//i
101261180 (Solyc02g093420)	2060	5.9	23.46	1.991416154	2.86E-213	XM_004231794.3	NAC	LOC101261180	PREDICTED: ScGO:0005634GO:00036:GO:00063:NA	NA	NA	NA	NA
101254394 (Solyc06g075780)	1145	2.26	8.11	1.843379142	1.34E-37	XM_004241199.3	C2H2	LOC101254394	PREDICTED: ScNA	GO:00036:NA	NA	NA	NA
101264084 (Solyc02g077610)	1271	3.53	12.37	1.809105412	4.84E-61	XM_010318477.2	NAC	LOC101264084	PREDICTED: ScGO:0005634GO:00036:GO:00063:NA	NA	NA	NA	NA
101248376 (Solyc04g015500)	1199	11.9	35.83	1.59020647	1.19E-135	BGI_novel_T0019	ABI3VP1	LOC101248376	PREDICTED: ScGO:0005634GO:00036:GO:00063:NA	NA	NA	NA	NA
101248351	1087	0.01	0.24	4.584962501	0.00024506	XM_004232564.3	MYB	LOC101248351	PREDICTED: ScGO:0005634GO:00009:GO:00063:K09422//t	NA	NA	NA	NA
101252786	1384	0.05	0.38	2.925999419	8.93E-05	XM_004242184.3	NAC	LOC101252786	PREDICTED: ScGO:0005634GO:00036:GO:00063:K03094//	NA	NA	NA	NA
101245410	1412	0.25	1.4	2.485426827	6.84E-13	XM_010329417.2	AP2-EREBP	LOC101245410	PREDICTED: ScGO:0005634GO:00036:GO:00063:K09286//	NA	NA	NA	NA
101261864	1569	0.36	1.79	2.313890776	1.43E-16	XM_004249405.2	GRF	LOC101261864	PREDICTED: ScGO:0005634GO:00055:GO:00325(K15601//i	NA	NA	NA	NA
101262811	850	0.28	1.1	1.974004791	2.07E-05	XM_010321849.2	MADS	LOC101262811	PREDICTED: ScGO:0005634GO:00036:GO:00063:K09260//	NA	NA	NA	NA
101246590	749	0.5	1.94	1.956056652	2.01E-07	XM_004250498.2	AP2-EREBP	LOC101246590	PREDICTED: ScGO:0005634GO:00036:GO:00063:K14516//	NA	NA	NA	NA
101259870	1082	0.18	0.67	1.896164189	0.00024879	XM_004246133.3	C2H2	LOC101259870	PREDICTED: ScGO:0005634GO:00036:NA	NA	NA	NA	K01870//i
104647464	699	0.63	2.14	1.764187063	8.61E-07	XM_010322879.1	MADS	LOC104647464	PREDICTED: ScGO:0005634GO:00036:GO:00063:K09260//	NA	NA	NA	NA
101246259	1247	0.41	1.36	1.729910837	1.68E-07	XM_004242235.3	C2H2	LOC101246259	PREDICTED: ScGO:0005634GO:00036:NA	NA	NA	NA	NA
101246748	1304	0.44	1.35	1.617383978	3.21E-07	XM_004228325.3	MYB	LOC101246748	PREDICTED: ScGO:0005634GO:00009:GO:00063:K09422//t	NA	NA	NA	NA
101267169	1192	1.47	4.48	1.607682577	1.59E-18	XM_004231151.3	AP2-EREBP	LOC101267169	PREDICTED: ScGO:0005634GO:00036:GO:00063:K13434//	NA	NA	NA	NA
101248663	3693	0.47	1.38	1.553935605	2.52E-17	BGI_novel_T0018	FHA	LOC101248663	PREDICTED: ScGO:0005634GO:00080:GO:00062:K10862//t	NA	NA	NA	NA
101246760	1085	0.69	1.91	1.468904371	3.07E-07	XM_004229745.3	C2C2-YABBY	LOC101246760	PREDICTED: ScNA	NA	GO:20000:NA	NA	NA
101262198	1496	2.12	5.63	1.409070658	9.00E-24	BGI_novel_T0052	bHLH	LOC101262198	Solanum lycoper NA	GO:00469:NA	NA	NA	K16190//
101254974	1033	1.08	2.78	1.364053571	1.39E-08	XM_004251574.3	ABI3VP1	LOC101254974	PREDICTED: ScGO:0005634GO:00036:GO:00063:NA	NA	NA	NA	NA
101256655	1530	1.05	2.66	1.341036918	1.39E-11	XM_004251750.3	bZIP	LOC101256655	PREDICTED: ScNA	GO:00036:NA	NA	NA	K20557//t
101243919	909	3.1	7.76	1.323788437	4.60E-18	XM_004246087.3	GRF	LOC101243919	PREDICTED: ScGO:0005634GO:00055:GO:00325(K20496//i	NA	NA	NA	NA
101247176	3257	0.33	0.82	1.313157885	3.92E-08	XM_010314051.2	GRAS	LOC101247176	PREDICTED: ScGO:0005634GO:00435(GO:00063:K14494//	NA	NA	NA	NA
101250945	1927	1.25	3.06	1.291603558	2.80E-15	XM_010313966.2	FAR1	LOC101250945	PREDICTED: ScGO:0005634GO:00435(GO:00102:K18835//	NA	NA	NA	NA
101267047	1396	3.54	8.66	1.290617665	3.22E-29	XM_019215420.1	GRF	LOC101267047	PREDICTED: ScGO:0005634GO:00055:GO:00325(K15601//i	NA	NA	NA	NA
101264349	1215	0.63	1.51	1.261124816	1.76E-05	XM_004244680.3	MYB	LOC101264349	PREDICTED: ScGO:0005634GO:00009:GO:00063:K09422//t	NA	NA	NA	NA
101055515	987	12.94	30.52	1.237917345	7.95E-65	NM_001279002.1	bHLH	LOC101055515	Solanum lycoper NA	GO:00469:NA	NA	NA	K18485//
101257365	1072	0.56	1.32	1.237039197	0.00018879	XM_004238118.3	ABI3VP1	LOC101257365	PREDICTED: ScGO:0005634GO:00036:GO:00063:NA	NA	NA	NA	NA
101259032	884	1.7	3.96	1.219965684	9.97E-09	XM_004235189.3	LOB	LOC101259032	PREDICTED: ScNA	NA	NA	NA	K21994//
101253752	2312	0.31	0.72	1.215728691	5.07E-05	XM_004251070.2	C2H2	LOC101253752	PREDICTED: ScNA	GO:00036:NA	NA	NA	K14794//r
101266110	1986	1.22	2.82	1.208814015	2.78E-13	XM_004235794.3	bHLH	LOC101266110	PREDICTED: ScNA	NA	NA	NA	K13422//t
101246621	757	6.8	15.52	1.190521906	1.97E-24	XM_004236988.2	bHLH	LOC101246621	PREDICTED: ScNA	GO:00469:NA	NA	NA	K17693//
101266372	1413	3.3	7.49	1.182499694	3.10E-22	XM_010326422.2	Sigma70-like	LOC101266372	PREDICTED: ScNA	GO:00036:GO:00063:K01872//	NA	NA	NA
101255499	1667	7.79	17.22	1.144389909	6.57E-56	XM_004240817.3	bZIP	LOC101255499	PREDICTED: ScNA	GO:00036:NA	NA	NA	K20557//t
101251000	1668	0.45	0.99	1.137503524	0.00018074	XM_004241269.3	C2H2	LOC101251000	PREDICTED: ScGO:0005634GO:00036:NA	NA	NA	NA	K14794//r
101258880	1595	8.56	17.68	1.046435573	1.16E-47	XM_004244583.3	AP2-EREBP	LOC101258880	PREDICTED: ScGO:0005634GO:00036:GO:00063:K09286//	NA	NA	NA	NA
101265489	3448	5.63	11.43	1.021618576	2.42E-56	XM_019215477.1	Alfin-like	LOC101265489	PREDICTED: ScNA	NA	NA	NA	K18046//t
104645444	968	2.56	5.17	1.01402047	3.70E-09	XM_010317128.2	OFP	LOC104645444	PREDICTED: ScNA	NA	NA	NA	NA
101252588 (Solyc05g050220)	1960	5.6	0.67	-3.063193826	1.59E-68	XM_010323221.2	bZIP	LOC101252588	PREDICTED: ScNA	GO:00037:NA	NA	NA	K09060//
101248026 (Solyc07g063410)	1418	31.29	3.78	-3.049243519	6.41E-276	NM_001319178.1	NAC	JA2L	Solanum lycoper GO:0005634GO:00036:GO:00063:K13126//	NA	NA	NA	NA
101256853 (Solyc12g044390)	904	5.98	0.85	-2.814610738	1.27E-31	XM_004252453.3	AP2-EREBP	LOC101256853	PREDICTED: ScGO:0005634GO:00036:GO:00063:K09286//	NA	NA	NA	NA
101055585 (Solyc01g107170)	1086	13.03	3.3	-1.981299154	1.03E-54	NM_001279257.1	C2H2	ZF2	Solanum lycoper GO:0005634GO:00036:NA	NA	NA	NA	K05747//

101263275 (Solyc11g056650)	1965	35.61	9.07	-1.97310798	6.04E-270	XM_004250809.3	bHLH	LOC101263275	PREDICTED: S	GO:0005634	GO:00012	GO:00063	K12126//
101250001 (Solyc03g121240)	1603	8.25	2.52	-1.710970386	1.16E-42	XM_010320219.2	bHLH	LOC101250001	PREDICTED: S	NA	GO:00469	NA	K16189//
101244844	411	1.51	0.01	-7.238404739	2.14E-06	XM_004234674.1	bHLH	LOC101244844	PREDICTED: S	NA	GO:00469	NA	K10863//
101248880	791	1.09	0.01	-6.768184325	2.22E-09	XM_004240173.3	MYB	LOC101248880	PREDICTED: S	GO:0005634	GO:00009	GO:00063	K09422//t
101246841	666	0.87	0.01	-6.442943496	2.14E-06	XM_004243645.3	AP2-EREBP	LOC101246841	PREDICTED: S	GO:0005634	GO:00036	GO:00063	K09286//
101249086	857	2.01	0.16	-3.651051691	2.24E-13	XM_004240230.3	LOB	LOC101249086	PREDICTED: S	NA	NA	NA	K13944//
101249100	1839	2.38	0.19	-3.64689025	2.47E-33	XM_004243322.3	C3H	LOC101249100	PREDICTED: S	NA	GO:00468	NA	K15172//t
101267939	1416	2.54	0.46	-2.465122731	2.40E-19	XM_004248047.3	bHLH	LOC101267939	PREDICTED: S	NA	GO:00469	NA	K13422//t
101251993	1651	0.92	0.18	-2.353636955	3.85E-09	XM_010324134.2	bZIP	LOC101251993	PREDICTED: S	NA	GO:00435	GO:00063	K14431//t
101260767	724	2	0.51	-1.971430848	1.76E-06	XM_004247876.3	AP2-EREBP	LOC101260767	PREDICTED: S	GO:0005634	GO:00036	GO:00063	K14516//
101250764	1501	2.26	0.69	-1.711654506	3.55E-12	XM_004232741.3	MYB	LOC101250764	PREDICTED: S	GO:0005634	GO:00009	GO:00063	K09422//t
101251084	1787	1.14	0.36	-1.662965013	1.25E-07	XM_004237769.3	AP2-EREBP	LOC101251084	PREDICTED: S	GO:0005634	GO:00036	GO:00063	K12900//
101256065	1414	2.73	0.92	-1.569195185	3.06E-12	XM_004234227.3	WRKY	LOC101256065	PREDICTED: S	GO:0005634	GO:00435	GO:00063	K13425//
101256789	2113	1.65	0.58	-1.508341219	7.22E-11	XM_010325751.2	bHLH	LOC101256789	PREDICTED: S	NA	GO:00469	NA	K13422//t
101251889	927	1.57	0.57	-1.461730735	8.03E-05	XM_004240553.2	LOB	LOC101251889	PREDICTED: S	NA	NA	NA	K13944//
101265482	1006	13.2	4.89	-1.432631559	1.01E-31	BGI_novel_T0042	bHLH	LOC101265482	PREDICTED: S	GO:0090575	GO:00036	GO:00063	NA
101246932	2814	1.57	0.59	-1.4119777	1.47E-13	XM_010324049.2	MYB	LOC101246932	PREDICTED: S	GO:0000783	GO:00036	GO:00316	K00430//
101267355	3159	13.66	5.18	-1.398933481	3.10E-105	XM_010315661.1	bHLH	LOC101267355	PREDICTED: S	NA	GO:00469	NA	K12126//
101263489	1637	23.72	9.48	-1.323145046	5.34E-86	XM_004251602.3	C3H	LOC101263489	PREDICTED: S	NA	GO:00468	NA	K15172//t
543645	1261	10.85	4.4	-1.302119614	1.63E-30	NM_001247500.1	WRKY	LOC543645	Solanum lycoper	NA	GO:00090	NA	NA
108511945	856	45.82	18.86	-1.280647783	1.01E-79	NM_001330445.1	AP2-EREBP	ERF5	Solanum lycoper	GO:0005634	GO:00036	GO:00063	K09286//
544204	1849	74.56	30.7	-1.280163204	6.90E-289	XM_010321757.2	bZIP	AREB	PREDICTED: S	NA	GO:00036	NA	K14432//
101250797	1076	3.66	1.53	-1.258311996	3.20E-09	XM_004238868.3	C2H2	LOC101250797	PREDICTED: S	GO:0005634	GO:00036	NA	K10773//
101254884	755	6.3	2.64	-1.254813899	1.50E-10	XM_004252770.3	C2H2	LOC101254884	PREDICTED: S	GO:0005634	GO:00036	NA	K05747//
101268010	1409	3.4	1.43	-1.249519599	5.44E-11	XM_004243835.3	NAC	LOC101268010	PREDICTED: S	NA	NA	NA	K13126//
101244335	1010	1.91	0.82	-1.219876823	5.94E-05	XM_004232313.3	LOB	LOC101244335	PREDICTED: S	NA	NA	NA	K13945//
101251778	700	3.98	1.75	-1.185413509	3.40E-06	NM_001323315.1	WRKY	LOC101251778	Solanum lycoper	GO:0005634	GO:00435	GO:00063	K18835//
101268787	1893	21.21	9.5	-1.158745202	3.19E-73	XM_004246260.3	FAR1	LOC101268787	PREDICTED: S	GO:0005634	GO:00435	GO:00508	K13424//
101258849	1085	1.95	0.88	-1.147898695	6.16E-05	XM_004236949.3	bHLH	LOC101258849	PREDICTED: S	NA	GO:00469	NA	K04733//i
101248665	1239	22.36	10.28	-1.121079924	1.43E-47	XM_004236996.3	NAC	LOC101248665	PREDICTED: S	GO:0005634	GO:00036	GO:00063	K12862//
101267548	1164	5.06	2.34	-1.112628855	3.79E-11	NM_001328614.1	AP2-EREBP	LOC101267548	Solanum lycoper	GO:0005634	GO:00036	GO:00063	K09286//
101244582	1354	19.7	9.15	-1.106351981	3.94E-45	NM_001329523.1	NAC	LOC101244582	Solanum lycoper	GO:0005634	GO:00036	GO:00063	K03094//
101257043	1048	3.51	1.64	-1.097775216	2.80E-07	XM_004234061.3	AP2-EREBP	LOC101257043	PREDICTED: S	GO:0005634	GO:00036	GO:00063	K09286//
101260164	1347	125.35	58.65	-1.095758983	5.34E-272	XM_010327256.1	PLATZ	LOC101260164	PREDICTED: S	NA	NA	NA	
101263056	1132	6.57	3.13	-1.069730713	7.12E-13	XM_004229716.3	AP2-EREBP	LOC101263056	PREDICTED: S	GO:0005634	GO:00036	GO:00063	K09286//
101261885	694	4.39	2.11	-1.056977941	9.27E-06	XM_010315247.2	C2H2	LOC101261885	PREDICTED: S	NA	GO:00036	NA	NA
101257056	1107	59.15	28.54	-1.051392834	8.10E-99	XM_004235767.3	LOB	LOC101257056	PREDICTED: S	NA	NA	NA	NA
101246812	1434	15.94	7.81	-1.029257176	5.65E-35	NM_001316915.1	WRKY	LOC101246812	Solanum lycoper	GO:0005634	GO:00435	GO:00096	K13424//
100736519	1734	209.49	103.07	-1.023256902	0	NM_001246877.1	TCP	TCP12	Solanum lycoper	GO:0005634	GO:00036	GO:00063	K13035//
101261342	1426	18.51	9.19	-1.010168129	7.40E-38	XM_019214651.1	NAC	LOC101261342	PREDICTED: S	GO:0005634	GO:00036	GO:00063	K13126//
101257639	993	12.94	6.46	-1.002231547	4.57E-19	XM_004252186.3	bHLH	LOC101257639	PREDICTED: S	NA	GO:00469	NA	K04733//i
101263123	1429	53.11	26.52	-1.001902754	2.83E-107	XM_004241479.2	TAZ	LOC101263123	PREDICTED: S	GO:0005634	GO:00044	GO:00431	K10523//

Supplementary Table S4. The rice (*Oryza sativa*) and *Arabidopsis thaliana* homologs of the highly responsive tomato TFs against *PepMV* infection

	Tomato Gene ID	Gene symbol	length (aa)	TF family	Homologs	Description	length (aa)	Query Cover	Per. Ident	E value
1	<i>Solyc04g015500</i>	Solyc04g015500	245	B3	At3g18990	AP2/B3-like transcriptional factor family protein	341	40%	45.54%	2.00E-23
					Os03g0621900	Transcriptional factor B3 domain containing protein	143	39%	34.38%	3.00E-14
2	<i>Solyc02g093420</i>	Solyc02g093420	393	NAC	At3g01600	NAC domain containing protein 44	366	100%	43.04%	3.00E-96
					Os02g0594800	No apical meristem (NAM) family protein, Awn development, Regulation of boundary formation, lateral organ separation and floral organ identity, NAC50	393	100%	49.01%	4.00E-121
3	<i>Solyc12g096070</i>	GRF4	349	GRF	At3g13960	growth-regulating factor 5	397	33%	60.16%	6.00E-47
					Os06g0116200	growth-regulating factor 5	349	93%	45.10%	9.00E-86
4	<i>Solyc02g077610</i>	NAC073	292	NAC	At4g28500	NAC domain containing protein 73	305	76%	72.49%	2.00E-117
					Os05g0563000	NAC domain-containing protein 62	315	85%	57.24%	2.00E-115
5	<i>Solyc06g075780</i>	ZAT1-like	323	C2H2	At5g59820	zinc finger protein Zat12	162	81%	45.75%	8.00E-30
					Os03g0279700	Similar to ZPT2-12	192	43%	63.01%	2.00E-23
					Os01g0838600	Zinc finger, C2H2-like domain containing protein	217	61%	39.85%	1.00E-20
6	<i>Solyc11g056650</i>	bHLH81	441	bHLH	At1g51140	basic helix-loop-helix (bHLH) DNA-binding superfamily protein, FBH3	379	54%	42.64%	6.00E-50
					Os08g0506700	Homolog of Arabidopsis FBH, Control of flowering	400	67%	39.32%	2.00E-46
7	<i>Solyc07g063410</i>	JA2L	356	NAC	At4g27410	NAC (No Apical Meristem) domain transcriptional regulator superfamily protein, ANAC072, RD26	297	72%	69.62%	1.00E-128
					At1g52890	NAC domain containing protein 19	317	71%	63.97%	7.00E-118
					Os01g0816100	Plant-specific transcription factor, Regulation of hypersensitive response (HR) cell death, OsNAC4	319	51%	60.21%	2.00E-79
8	<i>Solyc01g107170</i>	ZF2	310	C2H2	At1g27730	salt tolerance zinc finger, STZ, ZAT10	227	99%	35.76%	1.00E-33
					At5g04340	salt tolerance zinc finger, ZAT6	238	99%	34.29%	2.00E-30
					Os03g0764100	Zinc finger transcription factor ZF1	270	56%	39.06%	2.00E-26
					Os12g0583700	TFIIIA-type zinc finger protein, Tolerance to salt and drought stresses	248	89%	32.41%	6.00E-26
9	<i>Solyc03g121240</i>	bHLH87	380	bHLH	At3g21330	basic helix-loop-helix (bHLH) DNA-binding superfamily protein	373	78%	49.53%	2.00E-79
					Os01g0707500	Similar to Transcription factor LAX PANICLE	394	31%	62.10%	1.00E-45
					Os01g0566800	Hypothetical conserved gene, bHLH117	467	50%	48.47%	7.00E-45
10	<i>Solyc12g044390</i>	DREB3	143	ERF	At5g11590	Integrase-type DNA-binding superfamily protein, TINY2	236	65%	76.84%	1.00E-49
					Os04g0549800	ethylene-responsive transcription factor ERF037	289	64%	71.74%	3.00E-42
11	<i>Solyc05g050220</i>	TAF-1-like	416	bZIP	At2g46270	G-box binding factor 3	359	99%	44.63%	1.00E-94
					At4g01120	G-box binding factor 2	360	96%	39.46%	5.00E-67
					Os05g0569300	Similar to G-box binding factor 1	381	96%	35.68%	2.00E-62
					Os01g0658900	bZIP class abscisic acid responsive element (ABRE)-binding factor, Salt tolerance	361	100%	32.38%	8.00E-48

Supplementary Table S5. Identification of distinct signature motifs within the members of each TF family

TF family	motif	Sequence	length (aa)
B3	1	CFCKCGRQKFMQ	12
	2	DGHVWVKVGL	9
	3	GKVWLQDGWQEFVDRYSIKIGYFLIFRYEGNSQFDVFIFDLSGSEIEY	48
GRF	1	YYAAYGKKFDPEPGRCRRTDGKKWRCSKEAYPDSKYCERHMHRGRNRSRK	50
	2	PFTAVQWQELEHQALIYKYL VAGVPVPPDLILPIR	35
	3	VKQENRPLRPFDEWPGRKESWSELDEEGSNKNFSTTQLSISIPM	46
NAC	1	KVLYFYYGKAPKPEKTNWVMHZYHLG	26
	2	PGLPPGVKFDPTDEELLEHLECKKVAGNSKPHPLIDEFILT	41
	3	ICYTHPENLPGVNDGTSRHHFHRPSKAYTTGTRKRRKIHT	41
C2H2	1	KVHECSICGKEFPTGQALGGHKRRHYE	27
	2	FECKTCNKQFSSYQALGGHKASHKKP	26
	3	TEDEYLALCLMLLAR	15
bHLH	1	RKNVGISTDPQTVAARVRRERISERIRVLQELVPGGDKMDT	41
	2	ASMLDEAABYJKFLKSQVKALEELRAKCT	29
	3	YEPDPEAIAQVKEMIYRAAAFVPVNFVGAE	29
bZIP	1	BERELKREKRKQSNRESARRSRLRKQAETEELAVKVEALTAENTSLRSEI	50
	2	YPDWAAMQAYYGPRVAPPYYSSAIAPGH	29
	3	MPPPYGAPYAAVYPHGG	17
ERF	1	HPVYRGVVRMRSWGKWWSEIREPRKKSRIWLGTFPTPEMAARAHDVAALSI	50
	2	KGNAAILNFPELADSLRPASJSPRDVQ	28
	3	EDNLGQGIEPNFGKG	15

Supplementary Table S6. The *cis*-elements identified within the promoter regions of the 11 highly differentiated DETFs

Gene	Site name	Organism	Position	Strand	Matrix score	Sequence	Function
Solyc04g015500	ABRE	<i>Arabidopsis thaliana</i>	-705	+	5	ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-16	-	5	CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-16	+	5	TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	W box	<i>Arabidopsis thaliana</i>	-632	+	6	TTGACC	
Solyc02g093420	ABRE	<i>Triticum aestivum</i>	-578	+	9	GACACGTGGC	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-432	+	5	ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-1487	-	5	CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	ERE	<i>Nicotiana glutinos</i>	-1986	+	8	ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-526	-	8	ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-768	-	8	ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-205	-	8	ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-1800	+	8	ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-746	-	8	ATTTTAAA	
	TC-rich repeats	<i>Nicotiana tabacum</i>	-963	+	9	ATTCTCTAAC	cis-acting element involved in defense and stress responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-1487	+	5	TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	W box	<i>Arabidopsis thaliana</i>	-14	-	6	TTGACC	
	WUN-motif	<i>Nicotiana glutinosa</i>	-1675	+	9	AAATTACTA	
	WUN-motif	<i>Nicotiana glutinosa</i>	-251	+	8	AAATTACT	
WUN-motif	<i>Nicotiana glutinosa</i>	-1466	-	8	AAATTACT		
GRF4	ABRE	<i>Arabidopsis thaliana</i>	-1720	+	5	ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-213	+	5	ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-235	-	5	CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-132	+	5	CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	ERE	<i>Nicotiana glutinos</i>	-993	+	8	ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-716	-	8	ATTTTAAA	
	TGACG-motif	<i>Hordeum vulgare</i>	-132	-	5	TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-235	+	5	TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	WUN-motif	<i>Nicotiana glutinosa</i>	-396	+	9	TAATTACTC	
	WUN-motif	<i>Nicotiana glutinosa</i>	-1644	+	8	AAATTACT	
	WUN-motif	<i>Nicotiana glutinosa</i>	-177	-	9	TTATTACAT	
	WUN-motif	<i>Nicotiana glutinosa</i>	-221	+	9	AAATTACTA	
NAC073	ABRE	<i>Arabidopsis thaliana</i>	-1759	+	5	ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-622	+	5	ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-623	-	6	CACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-38	+	5	ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-516	-	5	CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-308	-	5	CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-516	+	5	TGACG	cis-acting regulatory element involved in the MeJA-responsiveness

	TGACG-motif	<i>Hordeum vulgare</i>	-308	+	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	W box	<i>Arabidopsis thaliana</i>	-1068	-	6 TTGACC	
	W box	<i>Arabidopsis thaliana</i>	-484	+	6 TTGACC	
	WUN-motif	<i>Nicotiana glutinosa</i>	-1817	+	9 CCATTTCAA	
	WUN-motif	<i>Nicotiana glutinosa</i>	-935	-	8 AAATTACT	
	WUN-motif	<i>Nicotiana glutinosa</i>	-936	-	9 AAATTACTA	
ZAT1-like	ABRE	<i>Arabidopsis thaliana</i>	-238	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-86	-	5 CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-86	+	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	WUN-motif	<i>Nicotiana glutinosa</i>	-1485	+	9 AAATTTCTT	
bHLH81	ABRE	<i>Arabidopsis thaliana</i>	-1714	-	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ERE	<i>Nicotiana glutinos</i>	-1998	-	8 ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-165	+	8 ATTTTAAA	
	TC-rich repeats	<i>Nicotiana tabacum</i>	-1743	+	9 GTTTTCTTAC	cis-acting element involved in defense and stress responsiveness
	TC-rich repeats	<i>Nicotiana tabacum</i>	-1254	-	9 GTTTTCTTAC	cis-acting element involved in defense and stress responsiveness
JA2L	ABRE	<i>Arabidopsis thaliana</i>	-837	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-1180	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Oryza sativa</i>	-838	+	8 TACGTGTC	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-573	-	7 TACGGTC	cis-acting element involved in the abscisic acid responsiveness
	ERE	<i>Nicotiana glutinos</i>	-1108	+	8 ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-348	-	8 ATTTCATA	
	WUN-motif	<i>Nicotiana glutinosa</i>	-1654	+	9 TTATTACAT	
	WUN-motif	<i>Nicotiana glutinosa</i>	-72	-	9 AAATTTCTT	
ZF2	ABRE	<i>Arabidopsis thaliana</i>	-239	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-1667	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-927	-	6 CACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-1259	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-926	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-1562	+	5 CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	ERE	<i>Nicotiana glutinos</i>	-1862	-	8 ATTTTAAA	
	TGACG-motif	<i>Hordeum vulgare</i>	-1562	-	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	W box	<i>Arabidopsis thaliana</i>	-547	-	6 TTGACC	
	W box	<i>Arabidopsis thaliana</i>	-269	+	6 TTGACC	
	W box	<i>Arabidopsis thaliana</i>	-362	+	6 TTGACC	
	WUN-motif	<i>Brassica oleracea</i>	-1474	+	9 AAATTTCCCT	wound-responsive element
	WUN-motif	<i>Nicotiana glutinosa</i>	-406	-	8 AAATTACT	
	ABRE	<i>Arabidopsis thaliana</i>	-655	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-510	-	5 CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	ERE	<i>Nicotiana glutinos</i>	-1930	-	8 ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-1678	-	8 ATTTCATA	

bHLH87	ERE	<i>Nicotiana glutinos</i>	-1857	-	8 ATTTTAAA	
	ERE	<i>Nicotiana glutinos</i>	-1102	-	8 ATTTTAAA	
	TC-rich repeats	<i>Nicotiana tabacum</i>	-1988	+	9 ATTCTCTAAC	cis-acting element involved in defense and stress responsiveness
	TC-rich repeats	<i>Nicotiana tabacum</i>	-790	-	9 ATTCTCTAAC	cis-acting element involved in defense and stress responsiveness
	TCA-element	<i>Nicotiana tabacum</i>	-1307	-	9 CCATCTTTTT	cis-acting element involved in salicylic acid responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-510	+	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	W box	<i>Arabidopsis thaliana</i>	-1968	+	6 TTGACC	
	W box	<i>Arabidopsis thaliana</i>	-1341	+	6 TTGACC	
DREB3	ABRE	<i>Arabidopsis thaliana</i>	-1498	-	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-592	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-1419	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-1040	-	5 CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	ERE	<i>Nicotiana glutinos</i>	-1217	+	8 ATTTTAAA	
	TC-rich repeats	<i>Nicotiana tabacum</i>	-1447	+	9 ATTCTCTAAC	cis-acting element involved in defense and stress responsiveness
	TC-rich repeats	<i>Nicotiana tabacum</i>	-530	+	9 ATTCTCTAAC	cis-acting element involved in defense and stress responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-1040	+	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	W box	<i>Arabidopsis thaliana</i>	-643	+	6 TTGACC	
	WUN-motif	<i>Nicotiana glutinosa</i>	-1319	-	8 AAATTACT	
WUN-motif	<i>Brassica oleracea</i>	-1554	+	9 AAATTCCT	wound-responsive element	
WUN-motif	<i>Nicotiana glutinosa</i>	-1626	+	9 CCATTTCAA		
TAF-1-like	ABRE	<i>Arabidopsis thaliana</i>	-1274	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-158	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Oryza sativa</i>	-263	-	8 TACGTGTC	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-171	-	6 CACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-648	-	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-170	+	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	ABRE	<i>Arabidopsis thaliana</i>	-261	-	5 ACGTG	cis-acting element involved in the abscisic acid responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-646	+	5 CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-160	-	5 CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	CGTCA-motif	<i>Hordeum vulgare</i>	-219	-	5 CGTCA	cis-acting regulatory element involved in the MeJA-responsiveness
	ERE	<i>Nicotiana glutinos</i>	-203	-	8 ATTTTAAA	
	TCA-element	<i>Nicotiana tabacum</i>	-148	+	9 CCATCTTTTT	cis-acting element involved in salicylic acid responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-646	-	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-160	+	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	TGACG-motif	<i>Hordeum vulgare</i>	-219	+	5 TGACG	cis-acting regulatory element involved in the MeJA-responsiveness
	W box	<i>Arabidopsis thaliana</i>	-771	-	6 TTGACC	
WUN-motif	<i>Nicotiana glutinosa</i>	-904	+	8 AAATTACT		
WUN-motif	<i>Nicotiana glutinosa</i>	-669	-	8 AAATTACT		