

Original citation:

Archbold, J. and Griffiths, Nathan (2015) Maximising influence in non-blocking cascades of interacting concepts. In: Multi-Agent-Based Simulation (MABS), Istanbul, Turkey, 5 May 2015

Permanent WRAP url:

<http://wrap.warwick.ac.uk/67163>

Copyright and reuse:

The Warwick Research Archive Portal (WRAP) makes this work by researchers of the University of Warwick available open access under the following conditions. Copyright © and all moral rights to the version of the paper presented here belong to the individual author(s) and/or other copyright owners. To the extent reasonable and practicable the material made available in WRAP has been checked for eligibility before being made available.

Copies of full items can be used for personal research or study, educational, or not-for profit purposes without prior permission or charge. Provided that the authors, title and full bibliographic details are credited, a hyperlink and/or URL is given for the original metadata page and the content is not changed in any way.

A note on versions:

The version presented here may differ from the published version or, version of record, if you wish to cite this item you are advised to consult the publisher's version. Please see the 'permanent WRAP url' above for details on accessing the published version and note that access may require a subscription.

For more information, please contact the WRAP Team at: publications@warwick.ac.uk



<http://wrap.warwick.ac.uk>

Maximising Influence in Non-blocking Cascades of Interacting Concepts

James Archbold¹ and Nathan Griffiths²

¹ University of Warwick, UK archbold@dcs.warwick.ac.uk

² University of Warwick, UK nathan@dcs.warwick.ac.uk

Abstract. In large populations of autonomous individuals, the propagation of ideas, strategies or infections is determined by the composite effect of interactions between individuals. The propagation of concepts in a population is a form of influence spread and can be modelled as a cascade from a set of initial individuals through the population. Understanding influence spread and information cascades has many applications, from informing epidemic control and viral marketing strategies to understanding the emergence of conventions in multi-agent systems. Existing work on influence spread has mainly considered single concepts, or small numbers of blocking (exclusive) concepts. In this paper we focus on non-blocking cascades, and propose a new model for characterising concept interaction in an independent cascade. Furthermore, we propose two heuristics, Concept Aware Single Discount and Expected Infected, for identifying the individuals that will maximise the spread of a particular concept, and show that in the non-blocking multi-concept setting our heuristics out-perform existing methods.

1 Introduction

When autonomous individuals interact, as part of a large population, the propagation of ideas, strategies or infections throughout the population is determined by the composite effect of interactions between individuals. Populations can be viewed as complex systems, with net effects that are hard to predict or influence despite being due to individual behaviour. The propagation of concepts, strategies or infections is a form of influence spread and can be modelled as a cascade from a set of initial individuals through the population.

Understanding how to limit or increase the spread of cascades through a population provides valuable insight into how to influence populations towards a particular state. Such insight has many applications, from informing epidemic control and viral marketing strategies to understanding the emergence of conventions in multi-agent systems. For example, characterising the spread of disease aids in identifying groups of individuals who are at risk, enabling containment efforts to be focused intelligently to help avoid wider spread. Understanding how ideas and their adoption propagates can also be used to inform viral marketing strategies, or calculate the network value of individuals in a population. In these cases the key is being able to identify the set of individuals who can help to

spread an idea or product, or who can restrict future spreading (e.g. through their vaccination).

Several models have been developed to simulate how influence spreads in a network, and much attention has been focused on the *influence maximisation problem*: finding a set of k nodes (individuals) whose activation will maximise the spread of a particular concept. This problem has been shown to be NP-hard, which has led to the development of heuristics to approximate optimal solutions. Many cascade models assume that cascades are *blocking*, in that a node that has been infected/activated by an idea or concept cannot be activated by any others. However, in many application domains individuals can hold multiple opinions, adopt multiple strategies, or have multiple infections that interact with each other and can further influence other individuals. The concepts held by an individual will affect those that they are likely to adopt later, and those that they are likely to propagate to others. This informs the idea of cascades or concepts interacting, however, most existing work on influence spread has considered single concepts, or small numbers of blocking (exclusive) concepts.

There has been relatively little consideration of cascades with multiple concepts, and such work has made simplifying assumptions. For example in the domain of epidemic spread Sanz *et al.* developed a model that allows two concepts to interact. The concepts active on a node affect its ability to activate other nodes, and so the spread of a concept is affected by the other concepts within the network. Concept interaction could be applied in other cascade models, requiring re-evaluation of existing influence maximisation heuristics. There is also the opportunity to develop heuristics that leverage concept interaction to improve concept spread.

In this paper we focus on non-blocking cascades, and propose a new model for characterising concept interaction in an independent cascade, specifically we propose a modification to the independent cascade model of influence spread that incorporates interacting cascades for an arbitrary number of concepts. Furthermore, we propose two heuristics, Concept Aware Single Discount and Expected Infected, for identifying individuals that will maximise the spread of a given concept, and show that in the non-blocking multi-concept setting our heuristics out-perform existing methods.

2 Related Work

In many application areas it would be valuable to leverage influential nodes within a population to maximise the spread of a concept throughout the population. This is referred to as the *influence maximisation problem* where we aim to pick a (minimal) set of nodes that would maximise the spread of information through the population. Several influence propagation models have been proposed in social network analysis literature [8, 14]. The target set of nodes is activated at the start of influence propagation, and in subsequent cycles, neighbours of active nodes are activated according some model of influence propaga-

tion. Such models can be classified into two types: those that use node-specific thresholds and those based on interacting particle systems [14].

In the *linear threshold model*, discussed by Kempe *et al.* [14], a node is influenced by each of its neighbours to varying degrees, as defined by the edge weights. Each node v has a threshold θ_v . When the sum of the weights of the active neighbours of v exceeds θ_v , v becomes active. Methods have been proposed for maximising influence spread within this model [7], but for now our focus is the independent cascade model.

In the *independent cascade model* (ICM) [10], when a node v becomes active it gets one chance to activate each of its inactive neighbours w , with some probability p_{vw} . Kempe *et al.* showed that a hill-climbing approach can be guaranteed to find a set of target nodes that has a performance slightly better than 63% of the optimal set [14]. One of the key issues with the greedy approach is the need to estimate target set quality. Numerous heuristics have been proposed to improve the speed of estimating the influence spread of a node [1, 6], but it remains problematic in large networks. Building on the greedy approach, Chen *et al.* proposed a degree discount heuristic that accounts for the existing activations in the network and attempts to reduce the impact of ‘double counting’ [6]. The degree discount heuristic has been shown to have similar effectiveness to the greedy approaches, while remaining computationally tractable.

The problem of influence maximisation has been studied in many contexts. For example in viral marketing, knowing the influential individuals in a network facilitates designing effective marketing strategies. Early studies into influence spread and maximisation focused on the network worth of users [8, 18]. Influence cascades have also been studied in relation to epidemic spread [16, 15, 20]. The two most commonly applied models when characterising epidemics are the Susceptible Infected Susceptible (SIS) and Susceptible Infected Recovered (SIR) models [9, 5]. These both take a probabilistic approach to the independent cascade model, allowing nodes to become deactivated.

Many of these studies have used single cascade models. In many real-world environments, there may be many concepts vying for the attention of an individual. As such, the effect of multiple influence cascades within a single network has been the focus of more recent work on influence spread [11, 12], with consideration of competing cascades that model competing products [2], epidemics [13] or general influence spread [3]. Existing work in this area, has typically assumed that the cascades are blocking, meaning that nodes activated/infected by one cascade cannot be activated/infected by another. Additionally, most existing work assumes only two concepts, while in reality there could be many interacting concepts. It is also often assumed that once activated a node remains active, though there are exceptions to this [17].

Sanz *et al.* developed a multi-layer network model in which concepts may only spread on a given layer but nodes can be activated by more than one concept at a time. Other work on the spread of epidemics also limits their travel to a single layer [19]. Existing research typically either assumes blocking concepts on a network layer, or non-blocking concepts that are each limited to a single

layer [12]. There has been little consideration of non-blocking concepts in a single layered network. Much of the work in epidemics focuses on the SIS model and the survival thresholds of viruses, with little exploration of multiple concepts interacting within other models [4].

3 Concept Interaction

To model concept interaction, we extend the work of Sanz *et al.* for two interacting diseases [20]. To represent these interactions, two cases must be considered. When attempting to infect a susceptible *receiver*, the infectiousness will change if the receiver is infected with the other disease. Conversely, the infectiousness of a disease is affected by the state of the *infector* spreading it. If the infector has both diseases, their infectiousness will change. These attributes form the basis of this model, which was originally intended for use with SIS and SIR cascade models, with modification, we can extend the formulation for use with the independent cascade model.

We must allow for both positive and negative relationships when concepts can interact. If a concept c affects c' in a positive way, we call it *boosting*, while if c is *inhibiting* c' then the effect is negative. How concepts spread in a given cascade model will change the exact effect of boosting and inhibiting. In general, boosting a concept makes it more likely to activate on a node and inhibiting makes it less likely. These relationships can be asymmetric: a concept could boost another concept that inhibits it and vice versa.

The relationship between two concepts, c and c' is defined by two *concept interaction factor* (CIF) functions, which describe the effect on the interaction of the infector and receiver respectively. Concepts active on an infector will affect each other's spread from that node. We refer to this as the *internal* effect of the infector on a given concept. The function $CIF_{int}(c, c')$ represents how concept c' affects c when an infector with both active attempts to spread c . For the receiver, we consider the concepts it has active and the *external* concept that attempts activation. The concepts already active on a node affect how willing it is to adopt new concepts. The function $CIF_{ext}(c, c')$ represents how concept c' affects the attempt by an infector to activate concept c on a receiver with c' active. These functions are both bounded in the range of $[-1, 1]$. If c' inhibits c , these functions will return a value below 0, while above 0 indicates a boosting relationship. If c' does not affect c the functions will return 0.

Since real-world environments may have more than 2 concepts we must evaluate the effect on the currently spreading concept of the infector's *internal* environment and the receiver's *external* environment. Two *concept interaction environment* functions characterise these effects, $CIE_{int}(C_n, c)$ and $CIE_{ext}(C_n, c)$ describe the internal and external environment respectively for a spreading concept c and set of concepts active on node n , C_n .

The notion of concept interaction is independent of the cascade model considered. In this paper we focus on the independent cascade model [14], as it has been the focus for much influence maximisation research, and extend it to ac-

count for multiple interacting concepts. In the standard independent cascade, an infector has chance p of making a neighbour active. With multiple concepts this probability is affected by the CIE_{int} function of the infector and the CIE_{ext} function of the receiver. When node n attempts to activate concept c on node m , the probability of success in the interactive independent cascade becomes:

$$p_c^s = p_c * (1 + CIE_{int}(C_n, c) + CIE_{ext}(C_m, c))$$

Where p_c is the baseline probability for that concept. CIE_{int} and CIE_{ext} are bounded to prevent unbalanced boosting compared to inhibiting. Boosting and inhibiting should have similar impact, rather than one offering more significant change. Therefore, we define the *concept interaction environment* functions as:

$$CIE_{int}(C_n, c) = \max(-1/2, \min(1/2, \sum_{c' \in C_n} CIF_{int}(c, c')))$$

$$CIE_{ext}(C_n, c) = \max(-1/2, \min(1/2, \sum_{c' \in C_n} CIF_{ext}(c, c')))$$

This means that p_c^s can range between $[0, p_c * 2]$. Since we must consider each node's environment, this value will be calculated for each interaction.

Cascades proceed in rounds, with an initial set of active nodes for each concept. Nodes can be in more than one initial set. Each node in the initial set for a concept will attempt to active that concept on each neighbour that is inactive for that concept. Each successfully activated neighbour will attempt to activate its neighbours in the next round. Nodes make one attempt on each neighbour for each concept they have active, and when no concept activates new nodes the cascade ends. For simplicity in this paper, we adopt the assumption that nodes will never deactivate a concept.

4 Heuristics for Node Selection

Several heuristics have been proposed for influence maximisation, as discussed in Section 2. In this section we introduce the main existing heuristics and propose two new methods: Concept Aware Single Discount and Expected Infected, which aim to take advantage of concept interaction.

Degree based selection. Degree based selection is the simplest heuristic, and has the advantage of only using attributes of the network, meaning that it is cheap to compute. With the degree heuristic we simply select the k nodes with the highest degree, an approach that has previously been shown to be effective [14].

Single Discount. When a node is added to the selection set, each of its neighbours has a chance to be activated in the first subsequent round of a cascade. However, if it is known that a node will become activated, adding it to the selection set provides no additional value, since that node will be activated regardless of

whether it is added to the selection set. This is the motivation behind the single discount heuristic. When a node n is placed into the selection set, the degree of all neighbouring nodes is lowered by 1 to represent this reduced network value (i.e. the number of potential activations they can create has reduced since n is already known to be active). Selection using the single discount heuristic proceeds in rounds, selecting the highest degree node and discounting its neighbours until the desired selection size is reached [6].

Concept Aware Single Discount Heuristic. Introducing concept interaction into the environment requires us to reconsider how concepts spread through a network. Each node can now affect the reach of a targeted concept’s spread based on the other concepts they have active. Node degree is typically a good indicator of influence, however in a concept interactive environment, this is not always the case. A node with many inhibiting concepts will be less desirable than a node with many boosting concepts if they have the same degree. Similarly, if a node has many neighbours who have active inhibiting concepts, its influence is likely to be low.

We propose a new heuristic, Concept Aware Single Discount (CASD), that weights the degree of a node based on its own concept environment and that of its neighbours, with the aim of providing a more accurate value of node desirability. Specifically, for CASD we define node utility as:

$$U_c(v) = CIE_{int}(C_v, c) + \sum_{n \in N(v)} 1 + CIE_{ext}(C_n, c)$$

where $N(v)$ is v ’s set of neighbours. Since we are attempting to select nodes that would help to maximise the spread of the targeted concept, the internal environment of a node is a good indicator of node value along with a its weighted degree. The external environment of a neighbour of v affects the likelihood of v activating it. The aim of the heuristic is to target nodes with many boosting neighbours and avoid those surrounded by inhibiting nodes. Therefore, $CIE_{ext}(C_n, c)$ is used to increase or decrease the contribution a neighbour makes to the degree of a node. This allows for the concept environment of a node and its neighbours to be considered when evaluating it’s worth to the selection set.

Selection proceeds in rounds, with the highest valued node selected each round. When a node n is added to the seed set, neighbour v has its utility updated accordingly:

$$U_c(v) = U_c(v) - (1 + CIE_{ext}(C_n, c))$$

In the same way as Single Discount, we remove the value contributed by the neighbour as it can no longer be activated. Once all neighbours have been updated, the next selection is made, until the required number of nodes is selected.

Degree Discount. Degree discount has been shown to be effective in approaching the optimal solution with reasonable computational overheads [6]. It relies on

calculating the expected nodes gained from adding a given node to the selection set. When a node is added, the expected gain of adding its neighbours is lowered. Additionally, those neighbours now have a chance to be activated in the first round of a cascade. The heuristic therefore weights the degree of a node based on these factors, updating the value for any neighbours when a node is added to the selection set. Nodes are initially ranked by degree, and when a node is added to the seed set neighbours have their degree set to $d_v - 2t_v - (d_v - t_v) * t_v * p$, where d_v is the original degree, t_v is the number of neighbours in the seed set and p is the probability of infection. This calculation is based on the expected benefit of such nodes (details of its derivation can be found in [6]).

Expected Infected Heuristic. It is important to consider the environment of a node and its neighbours when selecting nodes. The expected payoff from a node will change if it is surrounded by inhibiting concepts compared to boosting ones. Degree Discount is successful because it considers the expected number of activations for a node to decide its value. However, since it is intended for a single cascade model, it requires updating to consider concept interaction. We propose a new heuristic, Expected Infected, with the aim of accounting for these effects.

For each node, v , we consider the set of neighbours with chosen concept c active, $AN_c(v)$. Each of these neighbours will have a chance to activate v , which if successful removes any additional value v would have. The probability of v having c activated by one of these neighbours, $p_a(c, v)$, is:

$$p_a(c, v) = \sum_{n \in AN_c(v)} OP_c(n, v)$$

The sum of the individual chances of each neighbour to activate concept c on v , known as $OP_c(n, v)$, can be defined as:

$$OP_c(n, v) = p_c * (1 + CIE_{int}(C_n, c) + CIE_{ext}(C_v, c))$$

We can now determine the number of activations from $N(v)$ that can be expected as a result of activating v , as follows:

$$EA_c(v) = 1 + \sum_{n \in N(v) \setminus AN_c(v)} OP_c(v, n)$$

In addition to v itself, for each non-active neighbour, we have a $OP_c(v, n)$ chance to activate concept c . Summing the probabilities for each neighbour gives the expected number of neighbours v will activate. However, the chance that v will be activated by a neighbour must also be considered, and so the expected utility for adding v , to the seed set is given by:

$$U_c(v) = (1 - p_a(c, v)) * EA_c(v)$$

where $1 - p_a(c, v)$ is the chance of v not being activated. If activated anyway, v will give no additional value. Accounting for this requires scaling $EA_c(v)$ by

Table 1: Experimental Parameters

Parameter	Values
Graph Type	Small-world, Scale-free
Graph Size (nodes)	1000, 5000
Boost Proportion	0, 0.1, 0.2, 0.3, 0.4
Inhibit Proportion	0, 0.1, 0.2, 0.3, 0.4
Initial set size	1%, 2.5% or 5% of graph size
Intervention set size	1%, 2.5%, 5%, 7.5% or 10% of graph size
Rounds before intervention	5, 10, 25

the probability that v does not get activated. Initially nodes have a value of $EA_c(v)$, since there will be no active neighbours.

In each selection round, we add the node with the highest value for this heuristic and update its neighbours accordingly, continuing until the selection set is the desired size.

5 Experimental Approach

To evaluate the effectiveness of our proposed heuristics in the context of multiple interacting concepts, we perform simulations using the interactive independent cascade model introduced in Section 3. Each simulation has 10 concepts, with an activation probability for any concept of 0.05. We use the heuristics introduced in Section 4, along with random selection to provide a baseline for comparison. The network topologies listed in Table 1 were used, since they exhibit characteristics found in real-world social networks.

For each simulation, we determine the number of concepts boosted and inhibited by a given concept by selecting from a Gaussian distribution, with a mean of $boost_proportion * 10$ or $inhibit_proportion * 10$ respectively, and a standard deviation of 2.5. This, with the proportions defined in Table 1, prevents concepts being too similar and allows for more realistic environments. Though randomly selected, the final number of concepts boosted or inhibited by a single concept is restricted to be in the range $[0, 5)$.

The initial set of nodes for each concept is selected uniformly at random, and is the same for all concepts. The cascade proceeds for a fixed number of iterations (a burn-in period) until an intervention occurs, during which the targeted concept will activate an additional set of nodes selected using a chosen heuristic. The burn-in period before intervention is necessary since the concept aware heuristics require nodes to have concepts activated prior to selection. When selecting, the initial value of a node incorporates discounts from active neighbours as dictated by the chosen heuristic. This helps to compensate for heuristics that assume no nodes are active at the start. Each heuristic is used for interventions in 100 runs for each combination of parameters in Table 1.

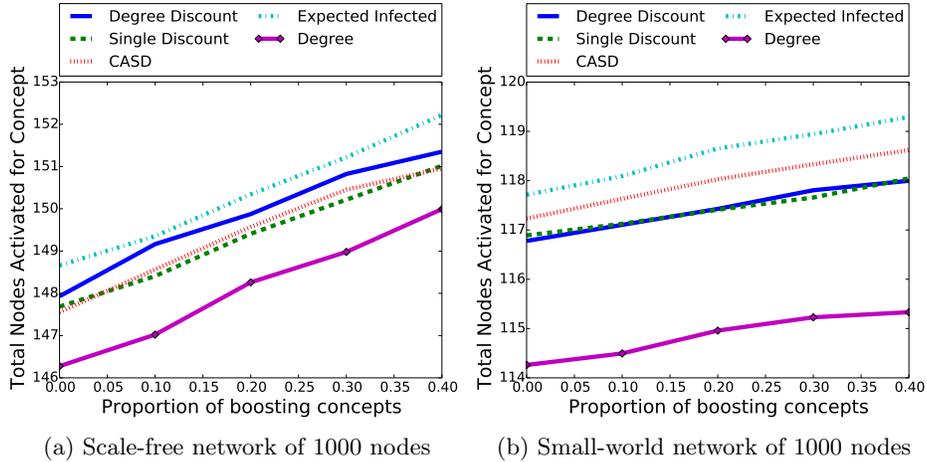


Fig. 1: Total activations against proportion of boosting concepts

6 Results

We initially compare the performance of each of the heuristics introduced above for a range of parameter settings. Random selection performed significantly worse than other heuristics in all cases, and while Degree was less effective than the other heuristics it was by a much smaller margin. Therefore, for simplicity of presentation, we do not consider Random further. Fig 1a shows the performance of the heuristics as the proportion of boosting concepts increases. We can see that Expected Infected performs best and outperforms our other proposed heuristic, CASD, with results for other topologies and populations mirroring this result. Therefore the remainder of our analysis focuses on comparing Expected Infected to the best performing of the existing heuristics, namely Degree Discount.

Expected Infected consistently outperforms Degree Discount, although in some situations the improvement is small. The difference in performance is larger in small-world networks than in scale-free, as shown by Fig 1. As the proportion of boosting concepts rises, all heuristics improve their total activations, demonstrating the impact of concepts interacting. The advantage of Expected Infected is stable for smaller populations, but is more varied in larger populations. It seems then that other network aspects counteract the benefit of targeting boosting concepts. For instance, in larger graphs encountering other concepts may be rarer, making smaller populations more sensitive to concept interaction.

As their proportion within the network increases, targeting boosting concepts becomes less effective. As Fig 2a shows, the performance difference between Expected Infected and Degree Discount decreases as boosting proportion increases. This demonstrates that as boosting concepts become more numerous, targeting them explicitly is less advantageous. The difference between performance declines earlier in scale-free graphs, likely owing to their construction through preferential attachment resulting in a group of nodes with high degree. Both Expected

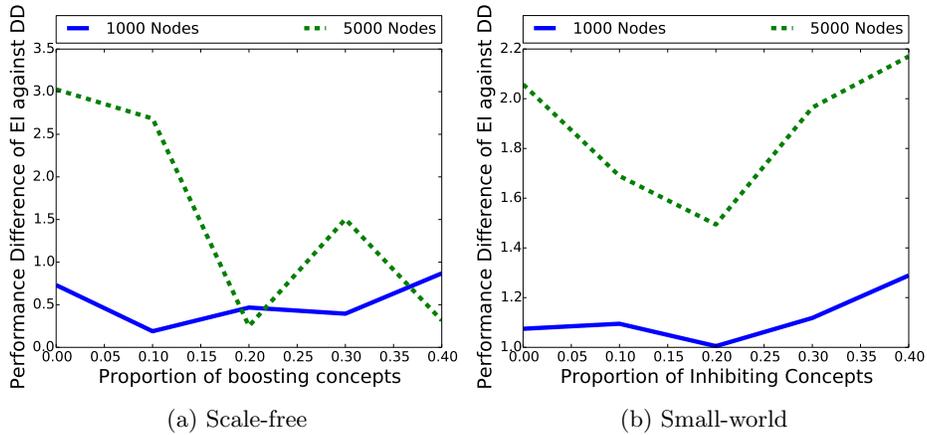


Fig. 2: Difference in activations of Expected Infected (EI) and Degree Discount (DD) against boosting and inhibiting proportions for 1000 and 5000 node graphs

Infected and Degree Discount will target nodes of high degree, such nodes will be more capable of utilising nearby boosting concepts without explicitly targeting them as the proportion of boosting concepts increases.

Conversely, as shown by Fig 2b, when the proportion of inhibiting concepts increases, so does the performance of Expected Infected compared to Degree Discount. As the number of inhibiting concepts increases it becomes harder to avoid them by chance. In a heavily inhibiting environment, high degree nodes have a higher chance to encounter inhibiting nodes and consequently have their influence diminished. The effectiveness of Expected Infected appears to be in avoiding inhibiting concepts, rather than in taking advantage of boosting ones.

The size of the initial and intervention sets also impacts performance as shown by Fig 3. At the largest initial set size, either 50 for Fig 3a or 250 for Fig 3b, performance improves as the intervention set size increases. In general, there is a slight upward trend in performance as the size of the initial set increases. The increased coverage results in concepts being more likely to interact, resulting in the consideration of other concepts being more advantageous. Additionally, as the initial and intervention sets increase in size, Degree Discount will find it harder to avoid inhibiting concepts, again demonstrating the advantage of avoiding them. It is possible that being able to target more boosting concepts is a factor here, but Fig 2a suggests this contribution is likely to be minor.

A relationship also appears to exist between graph density and performance Expected Infected, since as density increases Expected Infected performs better. The denser a graph, the more edges each node has, and avoiding inhibiting nodes by chance becomes unlikely as node degree increases, potentially impacting performance. The effect of network properties will be a key focus in future work.

Overall the Expected Infected heuristic performs slightly, but consistently, better than Degree Discount in a non-blocking multi-concept environment, and

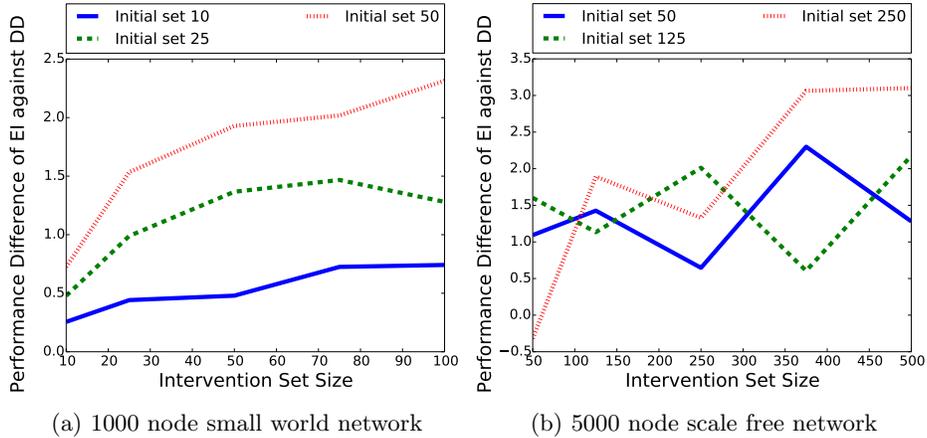


Fig. 3: Difference in activations of Expected Infected (EI) against Degree Discount (DD) for different initial sets against intervention set size

out-performs all other heuristics considered. Further, one of the core reasons for this performance seems to be in Expected Infected attempting to avoid inhibiting environments and thus not having the chosen concept’s spread hindered.

7 Conclusion

The study of how ideas, strategies or concepts propagate through a network has many applications. For example, simulations of disease and their infection characteristics can help identify areas at risk of an epidemic that should be the focus of containment and detecting the influential individuals in a social network allows for the improvement and refining of marketing strategies. This paper introduced an extension of the concept interaction model by Sanz *et al.* [20] to allow for n concepts within the independent cascade. We also proposed two new heuristics, including Expected Infected which made use of concept relationships to find the expected value of activating a node. Expected Infected was found to out-perform Degree Discount consistently in a concept interactive environment, specifically the avoidance of inhibiting factors seems to provide most of this advantage. Further work to quantify the effect of network properties on concept interactions is needed, to give a better understanding of when best to utilise concept interaction based heuristics. Observing how our results scale with the increase of concepts within the network would also be of interest, to see if the consideration of inhibiting concepts remains important.

References

1. A. Anagnostopoulos, R. Kumar, and M. Mahdian. Influence and correlation in social networks. In *Proc. of the 14th ACM SIGKDD Int. Conf. on Knowledge*

- discovery and data mining*, pages 7–15, 2008.
2. K. R. Apt and E. Markakis. Diffusion in social networks with competing products. In *Algorithmic Game Theory*, pages 212–223. 2011.
 3. A. Borodin, Y. Filmus, and J. Oren. Threshold models for competitive influence in social networks. In *Internet and network economics*, pages 539–550. 2010.
 4. C. D. Brummitt, K.-M. Lee, and K.-I. Goh. Multiplexity-facilitated cascades in networks. *Physical Review E*, 85(4):45–102, 2012.
 5. D. Chakrabarti, Y. Wang, C. Wang, J. Leskovec, and C. Faloutsos. Epidemic thresholds in real networks. *ACM TISSEC*, 10(4):1, 2008.
 6. W. Chen, Y. Wang, and S. Yang. Efficient influence maximization in social networks. In *Proc. of the 15th ACM SIGKDD Int. Conf. on Knowledge discovery and data mining*, pages 199–208, 2009.
 7. W. Chen, Y. Yuan, and L. Zhang. Scalable influence maximization in social networks under the linear threshold model. In *IEEE 10th ICDM*, pages 88–97, 2010.
 8. P. Domingos and M. Richardson. Mining the network value of customers. In *Proc. of the 7th ACM SIGKDD Int. Conf. on Knowledge discovery and data mining*, pages 57–66, 2001.
 9. S. C. Ferreira, C. Castellano, and R. Pastor-Satorras. Epidemic thresholds of the susceptible-infected-susceptible model on networks: A comparison of numerical and theoretical results. *Physical Review E*, 86(4):41–125, 2012.
 10. J. Goldenberg, B. Libai, and E. Muller. Using complex systems analysis to advance marketing theory development. *Academy of Marketing Science Review*, 2001(9):1–20, 2001.
 11. S. Goyal and M. Kearns. Competitive contagion in networks. In *Proc. of the 44th annual ACM symposium on Theory of computing*, pages 759–774, 2012.
 12. X. He, G. Song, W. Chen, and Q. Jiang. Influence blocking maximization in social networks under the competitive linear threshold model. In *SDM*, pages 463–474, 2012.
 13. B. Karrer and M. Newman. Competing epidemics on complex networks. *Physical Review E*, 84(3):36–106, 2011.
 14. D. Kempe, J. Kleinberg, and É. Tardos. Maximizing the spread of influence through a social network. In *Proc. of the 9th ACM SIGKDD Int. Conf. on Knowledge discovery and data mining*, pages 137–146, 2003.
 15. J. Leskovec, A. Krause, C. Guestrin, C. Faloutsos, J. VanBriesen, and N. Glance. Cost-effective outbreak detection in networks. In *Proc. of the 13th ACM SIGKDD Int. Conf. on Knowledge discovery and data mining*, pages 420–429, 2007.
 16. R. Pastor-Satorras and A. Vespignani. Epidemic spreading in scale-free networks. *Physical review letters*, 86(14):3200, 2001.
 17. N. Pathak, A. Banerjee, and J. Srivastava. A generalized linear threshold model for multiple cascades. In *IEEE 10th ICDM*, pages 965–970, 2010.
 18. M. Richardson and P. Domingos. Mining knowledge-sharing sites for viral marketing. In *Proc. of the 8th ACM SIGKDD Int. Conf. on Knowledge discovery and data mining*, pages 61–70, 2002.
 19. F. D. Sahneh and C. Scoglio. May the best meme win!: New exploration of competitive epidemic spreading over arbitrary multi-layer networks. *arXiv preprint arXiv:1308.4880*, 2013.
 20. J. Sanz, C.-Y. Xia, S. Meloni, and Y. Moreno. Dynamics of interacting diseases. *arXiv preprint arXiv:1402.4523*, 2014.