

ParaC genome SNPs 1. Numbers of mutational and recombinational SNPs per clade within the ParaC Lineage plus Birkenhead

Clade	No. Core SNPs ¹		
	Mutation	Recombination	Total
PC-1	4,001	57	4,058
PC-2	1,651	72	1,723
Modern Paratyphi C (w/o Ragna)	5,652	129	5,781
Paratyphi C (including Ragna)	6,327	132	6,459
Ragna	263	5	268
Paratyphi C & Typhisuis	11,891	219	12,110
Choleraesuis var. Kunzendorf	2,799	934	3,733
Choleraesuis <i>sensu stricto</i>	1,207	90	1,297
Choleraesuis	7,246	1,373	8,619
Paratyphi C, Typhisuis & Choleraesuis	20,247	1,602	21,849
Para C Lineage	-	-	37,971
Para C Lineage plus Birkenhead	-	-	62,868
<i>S. enterica</i> subsp. I	-	-	711,009

¹SNPs (ParaC genome SNPs 2) were extracted from core genomic regions in the Para C Lineage except for *S. enterica* subsp. I, where they correspond to all SNPs within the concatenated core genes. SNPs that were brought in by recombination after the MRCA of Paratyphi C, Typhisuis & Choleraesuis were identified by RECHMM and excluded for downstream analyses.

Note that this table counts the total number of events, and thus homoplastic SNPs were counted multiple times. Number of events per SNP sites that are leading to Paratyphi C, Choleraesuis and Typhisuis are summarized in “Para C genome SNPs 1”.