

Table S5 Ancestral haplotype diversity in the CC. The table provides the fraction of the genome with different number of haplotypes and whether these haplotypes are from one, two or three subspecies for the five classical founder strains (5) and all eight CC founders (8).

# of haplotypes	<u># of subspecies</u>					
	1 (5)	1 (8)	2 (5)	2 (8)	3 (5)	3 (8)
1	0.018	0.000	0.000	0.000	0.000	0.000
2	0.083	0.000	0.005	0.000	0.000	0.000
3	0.378	0.000	0.055	0.000	0.000	0.000
4	0.284	0.000	0.083	0.009	0.000	0.012
5	0.044	0.001	0.047	0.033	0.002	0.069
6	0.000	0.001	0.000	0.067	0.000	0.377
7	0.000	0.001	0.000	0.051	0.000	0.296
8	0.000	0.000	0.000	0.014	0.000	0.068