

Table S1. Estimates of average evolutionary divergence over sequence pairs within lineages (shaded in gray) and between lineages.

genomic dataset					
	Outgroup	Lineage 1	Lineage 2	Lineage 3	Lineage 4
Outgroup	n.d.				
Lineage 1	0.357 ± 0.0005	n.d.			
Lineage 2	0.356 ± 0.0005	0.168 ± 0.0004	n.d.		
Lineage 3	0.363 ± 0.0005	0.181 ± 0.0004	0.171 ± 0.0004	0.070 ± 0.0002	
Lineage 4	0.353 ± 0.0005	0.185 ± 0.0004	0.175 ± 0.0004	0.183 ± 0.0004	0.121 ± 0.0003
<i>gltA, groEL, ribE, rpoB</i> dataset					
	Outgroup	Lineage 1	Lineage 2	Lineage 3	Lineage 4
Outgroup	n.d.				
Lineage 1	0.301 ± 0.0082	n.d.			
Lineage 2	0.292 ± 0.0077	0.139 ± 0.0058	0.031 ± 0.0020		
Lineage 3	0.301 ± 0.0081	0.145 ± 0.0064	0.131 ± 0.0056	0.061 ± 0.0033	
Lineage 4	0.291 ± 0.0068	0.157 ± 0.0055	0.139 ± 0.0042	0.145 ± 0.0049	0.107 ± 0.0032

n.d., not determined