

Table S1. Sequence assembly statistics for newly sequenced *E. faecium* strains

Strain	k	Assembly size (Mbp)	Largest scaffold (bp)	N50 (bp)	Nr of scaffolds	Average Nt coverage
E155	35	3.09	153,272	46,336	242	79
E525	37	2.93	143,241	43,370	241	69
E1165	31	2.81	93,227	40,594	185	117

Assembly statistics are based on scaffolds of size ≥ 200 bp.