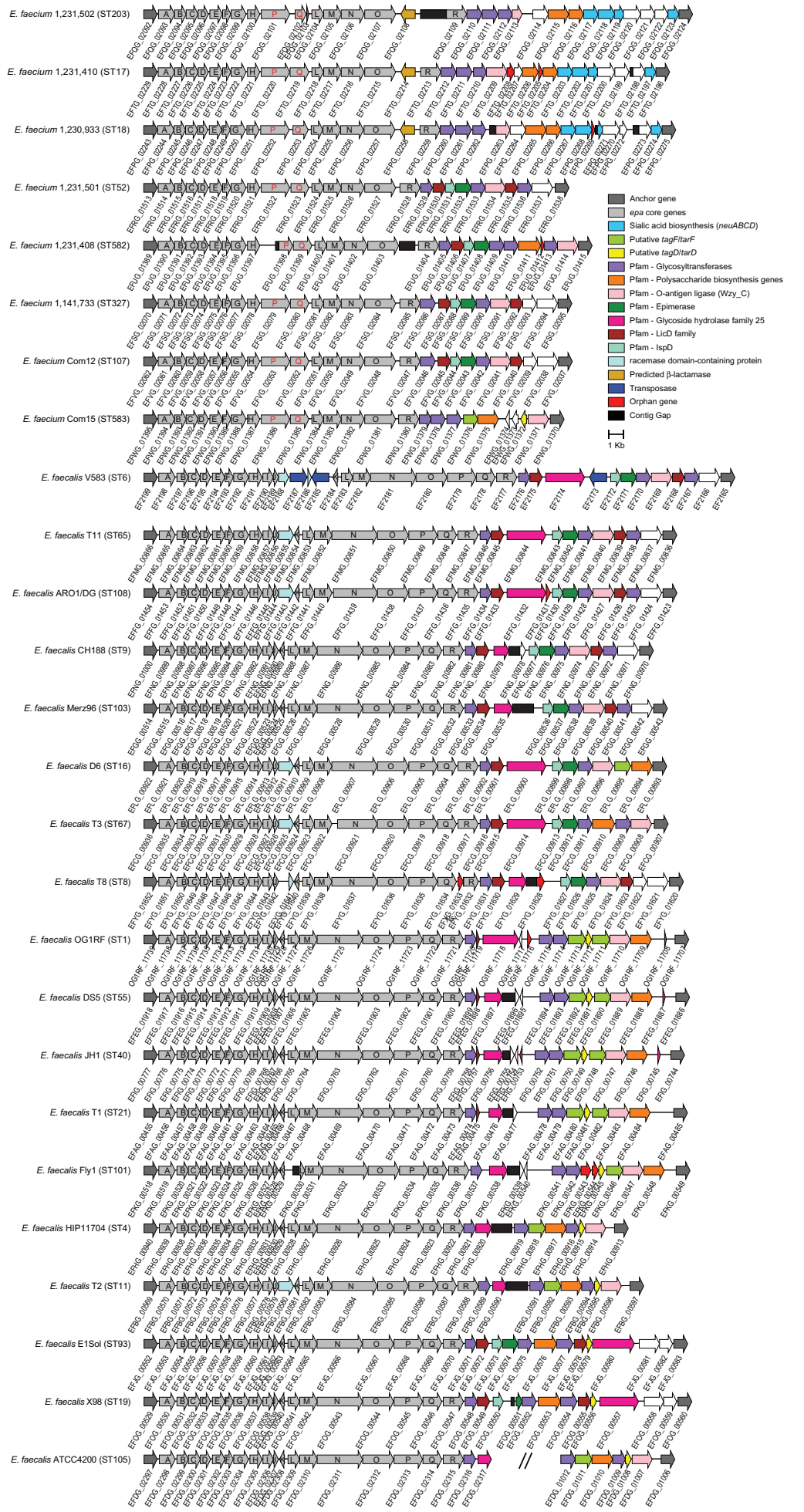
E. faecium 1,231,502 (ST203)
E. faecium 1,231,410 (ST17)
E. faecium 1,230,933 (ST18)
E. faecium 1,231,501 (ST52)
E. faecium 1,231,408 (ST582)
E. faecium 1,141,733 (ST327)
E. faecium Com12 (ST107)
E. faecium Com15 (ST583)
E. faecalis V583 (ST6)
E. faecalis T11 (ST65)
E. faecalis ARO1/DG (ST108)
E. faecalis CH188 (ST9)
E. faecalis Merz96 (ST103)
E. faecalis D6 (ST16)
E. faecalis T3 (ST67)
E. faecalis T8 (ST8)
E. faecalis OG1RF (ST1)
E. faecalis DS5 (ST55)
E. faecalis JH1 (ST40)
E. faecalis T1 (ST21)
E. faecalis Fly1 (ST101)
E. faecalis HIP11704 (ST4)
E. faecalis T2 (ST11)
E. faecalis E1Sol (ST93)
E. faecalis X98 (ST19)
E. faecalis ATCC4200 (ST105)
Anchor gene
epa core genes
Sialic acid biosynthesis (neuABCD)
Putative tagF/tarF
Putative tagD/tarD
Pfam - Glycosyltransferases
Pfam - Polysaccharide biosynthesis genes
Pfam - O-antigen ligase (Wzy_C)
Pfam - Epimerase
Pfam - Glycoside hydrolase family 25
Pfam - LicD family
Pfam - IspD
racemase domain-containing protein
Predicted β-lactamase
Transposase
Orphan gene
Contig Gap
1 Kb