

Women's health is a team effort: probiogenomics supports the development of a multi-species vaginal probiotic

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Online Resource 5. Contigs in the genomes of newly isolated strains predicted as belonging to plasmids.

Genome ID	Plasmidic Contig ID
B5_ <i>L.crispatus</i>	NODE_54_length_12471_cov_1681.597168
C2_ <i>L.crispatus</i>	NODE_68_length_11124_cov_1744.180423
C3_ <i>L.crispatus</i>	NODE_68_length_11124_cov_1494.148643
J1_ <i>L.paragasseri</i>	NODE_6_length_26819_cov_3870.155721
ID686_ DSM 32277_ <i>L.fermentum</i>	NODE_7_length_18696_cov_137.345
ID1969_ DSM 32405_ <i>L.gasseri</i>	MB79-ID1969_contig_8
Q1_ <i>L.crispatus</i>	NODE_9_length_43888_cov_887.506156
Q2_ <i>L.gasseri</i>	NODE_15_length_2740_cov_1.284519
T1_ <i>L.crispatus</i>	NODE_9_length_46309_cov_1141.134811
U2_ <i>L.crispatus</i>	NODE_4_length_74800_cov_1045.453869
X2_ <i>L.crispatus</i>	NODE_85_length_6470_cov_2018.200975
Z1_ <i>L.crispatus</i>	NODE_84_length_6198_cov_1577.094627