

Bacillus cereus BC210(NZ_JADNQED10000085.1 WP_202338192.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Bacillus cereus J2(NZ_JA0022933.1 WP_236794102.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Ligilactobacillus salivarius L.S.05(NZ_CP101687.1 WP_250201211.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Ligilactobacillus salivarius SMX051(NZ_AICL01000010.1 WP_034984371.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Mammaliococcus sciri GDK8555(PNZ_CP065963.1 WP_198472373.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus arlettae SA-01(NZ_KK274135.1 WP_172687524.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus 21101007(NZ_JAHPK010000017.1 WP_248319807.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus 220.1(NZ_JAHHMQ010000023.1 WP_228571442.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus 580_537(NZ_JAKYYN010000037.1 WP_248314790.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus 625_547(NZ_JAKYV010000027.1 WP_248314363.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus AA35(NZ_JAEV010000073.1 WP_200661586.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus AG037(NZ_PTCG01000030.1 WP_031883967.1 60	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus BSAC1476(NZ_FGM001000040.1 WP_061837968.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus BSAC2445(NZ_FGW01000036.1 WP_061736134.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus BSAC2488(NZ_FYED1000037.1 WP_061821985.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus BSAC349(NZ_FFYU1000037.1 WP_061641947.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus BSAR745(NZ_CFVW01000053.1 WP_057520790.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus CFR8-178(NZ_RIYF01000042.1 WP_123117931.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus CFSA053(NZ_NDTU01000007.1 WP_001614676.1 102	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus CMRISA(NZ_OC07209.1 WP_011282421.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus DFSA04(NZ_JAQAAW010000023.1 WP_260354179.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus DY67-1(NZ_PTBM01000045.1 WP_014158488.1 60	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus FVRH6404(NZ_KK047747.1 WP_049886533.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus GD9M30A(NZ_IANVJN010000019.1 WP_181187314.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus GGM6053(NZ_KJF79226.1 WP_031862766.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus H41N2_VRQ001000081.1 WP_151129282.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus HP20814.072(NZ_JACOC001000028.1 WP_1910026755.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 1000(NZ_LKV010000228.1 WP_095300542.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 1002(NZ_LKVV01000095.1 WP_095231779.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 1007(NZ_LKVM01000120.1 WP_095286329.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 1008(NZ_LKVN01000041.1 WP_095280164.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 871(NZ_LKV01000091.1 WP_095257751.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 873(NZ_LKVV01000044.1 WP_095315204.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 877(NZ_LKV021000097.1 WP_095315729.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 877(NZ_LKV021000098.1 WP_095315731.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 878(NZ_LKWD01000067.1 WP_095341278.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 880(NZ_LKW001000116.1 WP_095303470.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 882(NZ_LKWE01000043.1 WP_095315968.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 883(NZ_LKWF01000183.1 WP_095292735.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 887(NZ_LKWH01000075.1 WP_095327504.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 978(NZ_LKYF01000096.1 WP_095316456.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 992(NZ_LKYL01000256.1 WP_154701212.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 994(NZ_LKYN01000146.1 WP_095241969.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus ISU 996(NZ_LKYP01000124.1 WP_095251374.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus JARB-OU1216(NZ_BQZV01000083.1 WP_251360741.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus MN 1(NZ_LLB001000233.1 WP_154702228.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus MN 2(NZ_LLBP01000055.1 WP_095321120.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus MN 38(NZ_LLBR01000098.1 WP_095324565.1 88	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus MN 3(NZ_LLBO01000082.1 WP_095256749.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus MN 4(NZ_LLBS01000056.1 WP_095257151.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus MN 5(NZ_LLBT01000053.1 WP_095257063.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus NC-3(NZ_JAUEM010000067.1 WP_233233975.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus SC0A6048(NZ_KJF76550.1 WP_031905348.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus SCPM-O-B-8379(NZ_PXVA01000057.1 WP_050443248.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus SJ55(NZ_PSYZ01000056.1 WP_113559904.1 60	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus SK1396(NZ_SK1396(NZ_OC13345.1 WP_002503928.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus SKLX41347(NZ_JAJAHZ010000035.1 WP_234115137.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus c46(NZ_NDWWH01000029.1 WP_001003264.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus aureus subsp. aureus 300-169(NZ_JASL01000037.1 WP_031929051.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus cohnii DE0431(NZ_VDSK01000038.1 WP_002457017.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus epidermidis C100(NZ_CP094871.1 WP_243997260.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus epidermidis JMF-2008-C-0018(NZ_JACKOQ010000034.1 WP_241526145.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus epidermidis M1310451(NZ_KY579373.1 WP_172689289.1 62	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus epidermidis UMBP183(NZ_VYVQ01000051.1 WP_150861853.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus haemolyticus 109P1F51(NZ_JAKVFE010000055.1 WP_237617870.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus haemolyticus B2P1N52(NZ_JAKVFC010000080.1 WP_210136564.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus haemolyticus 95671(NZ_CUIFA01000058.1 WP_001003263.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus haemolyticus CS8F5(NZ_JAKVFM010000049.1 WP_240566558.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus saprophyticus 10007621(NZ_OC07621.1 WP_011382144.1 85	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus sp. GDH8C106P(NZ_JAESM2010000021.1 WP_014125936.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus sp. GDQ8D205P(NZ_JAESMQ010000003.1 WP_210123703.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus sp. GDY8P44P(NZ_JAESL1010000032.1 WP_007410443.1 47	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus sp. HM5C65H10(NZ_KV840330.1 WP_070867177.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Staphylococcus xylosum SNUC 5423(NZ_PZEF01000072.1 WP_107531727.1 63	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K
Thermus scotodactes D5M 8553(NZ_KB905809.1 WP_081624258.1 62	M	G	I	F	S	I	F	V	L	N	T	V	D	Y	Q	P	N	K