

Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Percent of Identity	Accession Length	Accession
Streptococcus anginosus strain NCTC10713 genome assembly, chromosome: 1	Streptococcus anginosus	3751	3751	100%	0.0	97.29%	1953654	LR134283.1
Streptococcus anginosus subsp. anginosus strain FDAARGOS_1569 chromosome, complete genome	Streptococcus anginosus subsp. anginosus	3751	3751	100%	0.0	97.29%	1876334	CP085939.1
Streptococcus anginosus CCUG 27928 DNA, complete genome	Streptococcus anginosus	3751	3751	100%	0.0	97.29%	1955308	AP018548.1
Streptococcus constellatus strain FDAARGOS_1015 chromosome, complete genome	Streptococcus constellatus	3397	3397	100%	0.0	94.39%	2038583	CP066055.1
Streptococcus anginosus subsp. whileyi MA5624 DNA, complete genome	Streptococcus anginosus subsp. whileyi MA5624	3397	3397	100%	0.0	94.39%	2122284	AP013072.1
Streptococcus anginosus C238, complete genome	Streptococcus anginosus C238	3391	3391	100%	0.0	94.35%	2233640	CP003861.1
Streptococcus anginosus strain FDAARGOS_1155 chromosome, complete genome	Streptococcus anginosus	3391	3391	100%	0.0	94.35%	2284041	CP068060.1
Streptococcus constellatus strain FDAARGOS_1208 chromosome, complete genome	Streptococcus constellatus	3386	3386	100%	0.0	94.30%	1978680	CP069558.1
Streptococcus constellatus subsp. pharyngis C232, complete genome	Streptococcus constellatus subsp. pharyngis C232	3386	3386	100%	0.0	94.30%	1935414	CP003800.1
Streptococcus constellatus subsp. pharyngis C818, complete genome	Streptococcus constellatus subsp. pharyngis C818	3386	3386	100%	0.0	94.30%	1935662	CP003840.1
Streptococcus constellatus subsp. pharyngis C1050, complete genome	Streptococcus constellatus subsp. pharyngis C1050	3386	3386	100%	0.0	94.30%	1991156	CP003859.1
Streptococcus anginosus C1051, complete genome	Streptococcus anginosus C1051	3330	3330	100%	0.0	93.85%	1911706	CP003860.1
Streptococcus anginosus strain NCTC11064 genome assembly, chromosome: 1	Streptococcus anginosus	3330	3330	100%	0.0	93.85%	1904537	LR594037.1
Streptococcus anginosus strain J4211 chromosome, complete genome	Streptococcus anginosus	3325	3325	100%	0.0	93.81%	1924513	CP012805.1
Streptococcus anginosus strain VSI13 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	1926555	CP118081.1
Streptococcus anginosus strain VSI52 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	2081644	CP118046.1
Streptococcus anginosus strain VSI12 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	1926669	CP118082.1
Streptococcus anginosus strain VSI68 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	2010423	CP118029.1
Streptococcus anginosus strain VSI67 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	2091268	CP118031.1
Streptococcus anginosus strain VSI40 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	1958159	CP118053.1
Streptococcus anginosus strain UMB8616 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	2033097	CP136160.1
Streptococcus anginosus strain VSI37 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	2138327	CP118054.1
Streptococcus anginosus strain VSI16 chromosome, complete genome	Streptococcus anginosus	3070	3070	100%	0.0	91.72%	2085682	CP118078.1
Streptococcus anginosus strain VSI85 chromosome, complete genome	Streptococcus anginosus	3064	3064	100%	0.0	91.68%	2130825	CP118036.1
Streptococcus anginosus strain UMB4708 chromosome, complete genome	Streptococcus anginosus	3059	3059	100%	0.0	91.63%	2148204	CP126961.1
Streptococcus milleri strain NCTC11169 genome assembly, chromosome: 1	Streptococcus milleri	3042	3042	100%	0.0	91.50%	1856804	LR134288.1
Streptococcus periodonticum strain KCOM 2412 chromosome, complete genome	Streptococcus periodonticum	3042	3042	100%	0.0	91.50%	1903815	CP034543.1
Streptococcus anginosus strain FDAARGOS_1357 chromosome, complete genome	Streptococcus anginosus	2976	2976	100%	0.0	90.96%	1981414	CP069892.1
Streptococcus anginosus strain J4206 chromosome, complete genome	Streptococcus anginosus	2904	2904	100%	0.0	90.37%	2001352	CP012719.1
Streptococcus anginosus strain SA1, complete genome	Streptococcus anginosus	2898	2898	100%	0.0	90.33%	2036353	CP007573.1
Streptococcus anginosus strain 4751 chromosome	Streptococcus anginosus	2898	2898	100%	0.0	90.47%	1981612	CP068916.1
Streptococcus sp. KHUD_010 chromosome, complete genome	Streptococcus sp. KHUD_010	2887	2887	100%	0.0	90.24%	1883665	CP157941.1
Streptococcus sp. KS 6 chromosome, complete genome	Streptococcus sp. KS 6	2887	2887	100%	0.0	90.24%	1951018	CP041968.1
Streptococcus constellatus strain TCY107 chromosome, complete genome	Streptococcus constellatus	2815	2815	100%	0.0	89.73%	1908997	CP029207.1
Streptococcus constellatus subsp. constellatus DNA, complete genome, strain: CCUG 24889	Streptococcus constellatus subsp. constellatus	2719	2719	99%	0.0	89.07%	1901581	AP014647.1
Streptococcus constellatus strain FDAARGOS_1156 chromosome, complete genome	Streptococcus constellatus	2719	2719	99%	0.0	89.07%	1903262	CP068213.1
Streptococcus sp. NSJ-72 chromosome	Streptococcus sp. NSJ-72	2665	2665	100%	0.0	88.62%	2014404	CP060489.1
Streptococcus milleri strain NCTC10708 genome assembly, chromosome: 1	Streptococcus milleri	2658	2658	99%	0.0	88.58%	1890902	LR134307.1
Streptococcus intermedius B196, complete genome	Streptococcus intermedius B196	2045	2045	97%	0.0	84.02%	1996214	CP003857.1
Streptococcus intermedius strain XH2169 chromosome, complete genome	Streptococcus intermedius	2034	2034	97%	0.0	83.93%	1944282	CP136799.1
Streptococcus intermedius strain KCOM 1545 chromosome, complete genome	Streptococcus intermedius	2023	2023	91%	0.0	84.90%	1908201	CP012718.1
Streptococcus intermedius strain FDAARGOS_233 chromosome, complete genome	Streptococcus intermedius	2023	2023	91%	0.0	84.90%	1914006	CP020433.2
Streptococcus intermedius strain FDAARGOS_769 chromosome, complete genome	Streptococcus intermedius	2001	2001	91%	0.0	84.70%	1933377	CP053999.1
Streptococcus intermedius JTH08 DNA, complete genome	Streptococcus intermedius JTH08	1989	1989	91%	0.0	84.61%	1933610	AP010969.1
Streptococcus intermedius strain NCTC11324 genome assembly, chromosome: 1	Streptococcus intermedius	1989	1989	91%	0.0	84.61%	1932951	LS483436.1
Streptococcus intermedius C270, complete genome	Streptococcus intermedius C270	1973	1973	91%	0.0	84.46%	1960728	CP003858.1
Streptococcus intermedius TYG1620 DNA, complete genome	Streptococcus intermedius	1962	1962	91%	0.0	84.36%	2006877	AP014880.1
Gemella sanguinis strain FDAARGOS_742 chromosome, complete genome	Gemella sanguinis	424	424	52%	4,00E-113	73.54%	1796606	CP046313.1
Enterococcus cecorum strain 66 chromosome, complete genome	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	97.87%	2708228	CP144498.1
Enterococcus cecorum strain 85 chromosome, complete genome	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	97.87%	2601676	CP144494.1
Enterococcus cecorum isolate CIRMBP-1261 genome assembly, chromosome: 1	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	97.87%	2620135	OX346402.1
Enterococcus cecorum strain 65 chromosome, complete genome	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	97.87%	2569129	CP144500.1
Enterococcus cecorum strain 67 chromosome, complete genome	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	94.34%	2574331	CP144496.1
Enterococcus cecorum strain 128 chromosome, complete genome	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	97.87%	2542076	CP144489.1
Enterococcus cecorum strain 127 chromosome, complete genome	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	97.87%	2670571	CP144490.1
Enterococcus cecorum isolate CIRMBP-1228 genome assembly, chromosome: 1	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	94.34%	2799887	OX346405.1
Enterococcus cecorum strain NCTC12421 genome assembly, chromosome: 1	Enterococcus cecorum	82.4	82.4	2%	3,00E-10	94.34%	2421598	LS483306.1
Enterococcus cecorum strain 70 chromosome, complete genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2312828	CP144495.1
Enterococcus cecorum strain 117 chromosome, complete genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2427853	CP144492.1
Enterococcus cecorum isolate CIRMBP-1246 genome assembly, chromosome: 1	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2290161	OX346401.1
Enterococcus cecorum strain SA1 genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	95.74%	2261297	CP010060.1
Enterococcus cecorum isolate CIRMBP-1287 genome assembly, chromosome: 1	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2444274	OX346411.1
Enterococcus cecorum strain SA2 genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	95.74%	2281990	CP010061.1
Enterococcus cecorum isolate CIRMBP-1302 genome assembly, chromosome: 1	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2391303	OX346408.1
Enterococcus cecorum strain 89 chromosome, complete genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2427811	CP144493.1
Enterococcus cecorum isolate CIRMBP-1212 genome assembly, chromosome: 1	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2422110	OX346406.1
Enterococcus cecorum isolate CIRMBP-1274 genome assembly, chromosome: 1	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2479239	OX346407.1
Enterococcus cecorum SMUHEC01 DNA, complete genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2457796	AP035890.1
Enterococcus cecorum isolate CIRMBP-1281 genome assembly, chromosome: 1	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2542510	OX346404.1
Enterococcus cecorum isolate CIRMBP-1283 genome assembly, chromosome: 1	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2260504	OX346403.1
Enterococcus cecorum strain CE3 genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	95.74%	2372196	CP010063.1
Enterococcus cecorum strain SA3 genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	95.74%	2320599	CP010064.1
Enterococcus cecorum strain 45 chromosome, complete genome	Enterococcus cecorum	76.8	76.8	2%	1,00E-08	94.00%	2548069	CP144502.1
Streptococcus suis strain SRD478 chromosome, complete genome	Streptococcus suis	56.5	56.5	2%	0.017	85.45%	2062916	CP017088.1
Streptococcus suis strain ISU2912 chromosome, complete genome	Streptococcus suis	56.5	56.5	2%	0.017	85.45%	2720381	CP017785.1

The sequences clustering with *sanA* of *S. anginosus* 3792/10 are shaded.