Supporting information for Tettelin *et al.* (2002) *Proc. Natl. Acad. Sci. USA*, 10.1073/pnas.182380799

Table 6. Clusters of genes derived from phylogenetic profiling of *S. agalactiae* strains based on comparative genome hybridizations

Clusters of five or more genes (see Fig. 5) are listed.

| Cluster 1 | |
|---|--|
| SAG0230 | conserved hypothetical protein |
| SAG0231 | hypothetical protein |
| SAG0232 | hypothetical protein |
| SAG0233 | hypothetical protein |
| SAG0234 | hypothetical protein |
| SAG0235 | hypothetical protein |
| | |
| CI () | |
| Cluster 2 | |
| SAG0222 | conserved domain protein |
| C1015001 = | conserved domain protein conserved hypothetical protein, fusion |
| SAG0222 | 1 |
| SAG0222 SAG0223 | conserved hypothetical protein, fusion |
| SAG0222 SAG0223 SAG0225 | conserved hypothetical protein, fusion hypothetical protein |
| SAG0222 SAG0223 SAG0225 SAG0226 | conserved hypothetical protein, fusion hypothetical protein recombination protein |
| SAG0222 SAG0223 SAG0225 SAG0226 SAG0227 | conserved hypothetical protein, fusion hypothetical protein recombination protein hypothetical protein |

| Cluster 3 | |
|-----------------------|---|
| SAG0634 | hypothetical protein |
| SAG0635 | acid phosphatase, class B |
| SAG0636 | conserved hypothetical protein |
| SAG0638 | cell wall surface anchor family protein, interruption-N |
| SAG0640 | transposase OrfA, IS3 family |
| SAG0642 | hypothetical protein |
| SAG0643 | chaperonin, 33 kDa, degenerate |
| SAG0644 | transcriptional regulator, AraC family |
| SAG0645 | cell wall surface anchor family protein |
| SAG0646 | cell wall surface anchor family protein |
| SAG0647 | sortase family protein |
| SAG0648 | sortase family protein |
| SAG0649 | cell wall surface anchor family protein, putative |
| SAG0650 | sortase family protein |
| SAG0651 | protein of unknown function |
| | |
| Cluster 4 | |
| SAG1898 | PTS system, IID component |
| SAG1899 | PTS system, IIC component |
| SAG1900 | PTS system, IIB component |
| SAG1901 | glucuronyl hydrolase |
| SAG1902 | PTS system, IIA component |
| SAG1905 | conserved hypothetical protein |
| SAG1906 | carbohydrate kinase, PfkB family |
| 6 7 4 5 | |
| Cluster 5 | |
| SAG0247 | hypothetical protein |
| SAG0248 | hypothetical protein |
| SAG0249 | hypothetical protein |
| SAG0674 | hypothetical protein |
| SAG0675 | putative secreted protein |
| SAG0676 | proteinase, putative |
| SAG0677 | hypothetical protein |
| SAG0680 | protein of unknown function |
| SAG0681 | conserved domain protein |
| SAG0684 | ABC transporter, ATP-binding protein |
| SAG1698 | conserved hypothetical protein |
| Cluster 6 | |
| SAG0261 | IS1381, transposase OrfB |
| SAG0262 | IS1381, transposase OrfA |
| SAG0965 | IS1381, transposase OrfA |
| SAG0966 | IS1381, transposase OrfB |
| SAG2002 | IS1381, transposase OffB |
| 51102002 | 101001, transposase OHD |

| Cluster 7 | |
|------------|--|
| SAG1027 | conserved hypothetical protein |
| SAG1028 | hypothetical protein |
| SAG1029 | hypothetical protein |
| SAG1030 | protein of unknown function |
| SAG1031 | conserved domain protein |
| SAG1032 | conserved hypothetical protein |
| Cluster 8 | |
| SAG1253 | transposase, ISL3 family |
| SAG1254 | mercuric reductase |
| SAG1255 | mercuric resistance operon regulatory protein MerR |
| SAG2022 | transposase, ISL3 family |
| SAG2023 | mercuric reductase |
| SAG2024 | mercuric resistance operon regulatory protein MerR |
| Cluster 9 | |
| SAG1993 | site-specific recombinase, phage integrase family |
| SAG1994 | conserved hypothetical protein |
| SAG1995 | hypothetical protein |
| SAG1996 | cell wall surface anchor family protein, putative |
| SAG1997 | hypothetical protein |
| SAG1998 | hypothetical protein |
| SAG2000 | membrane protein, putative |
| SAG2001 | conjugal transfer protein, interruption-C |
| SAG2007 | conserved hypothetical protein |
| SAG2008 | conserved hypothetical protein |
| SAG2009 | conserved hypothetical protein |
| SAG2010 | hypothetical protein |
| SAG2011 | conserved hypothetical protein |
| SAG2012 | hypothetical protein |
| SAG2016 | hypothetical protein |
| SAG2017 | transcriptional regulator, Cro/CI family |
| SAG2025 | Mn2+/Fe2+ transporter, NRAMP family |
| Cluster 10 | |
| SAG1039 | conserved hypothetical protein |
| SAG1447 | conserved hypothetical protein |
| SAG1448 | glycosyl transferase, group 1 family protein |
| SAG1449 | preprotein translocase SecA subunit, putative |
| SAG1450 | conserved domain protein |
| SAG1452 | conserved hypothetical protein |
| SAG1453 | preprotein translocase SecY family protein |
| SAG1454 | glycosyl transferase, putative |
| SAG1455 | glycosyl transferase, group 2 family protein |
| SAG1456 | glycosyl transferase, family 8, degenerate |
| | |

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|--------------------|--|
| SAG1459 | glycosyl transferase family 8 |
| SAG1460 | glycosyl transferase, family 8 |
| SAG1461 | conserved hypothetical protein |
| SAG1462 | cell wall surface anchor family protein |
| SAG1463 | transcriptional regulator, RofA family, authentic point mutation |
| SAG1469 | conserved hypothetical protein |
| SAG1471 | conserved hypothetical protein |
| SAG1933 | PTS system, IIC component, putative |
| 21101900 | 1 12 by stem, 110 component, pulluri |
| Cluster 11 | |
| SAG0009 | hypothetical protein |
| SAG0120 | hypothetical protein |
| SAG0120 SAG0157 | deoxyribonuclease-related protein, degenerate |
| SAG0137 SAG0186 | · · · · · · · · · · · · · · · · · · · |
| | hypothetical protein |
| SAG0216 | hypothetical protein |
| SAG0236 | hypothetical protein |
| SAG0307 | hypothetical protein |
| SAG0308 | ABC transporter, ATP-binding protein |
| SAG0311 | DNA-binding response regulator, authentic point mutation |
| SAG0518 | peptide chain release factor 2, programmed frameshift |
| SAG0553 | hypothetical protein |
| SAG0555 | prophage LambdaSa1, antirepressor, putative |
| SAG0564 | conserved hypothetical protein |
| SAG0579 | conserved hypothetical protein |
| SAG0580 | conserved hypothetical protein, truncation |
| SAG0611 | transposase, degenerate |
| SAG0637 | transcriptional regulator, TetR family, putative, authentic frameshift |
| SAG0641 | Tn5252, Orf 10 protein, degenerate |
| SAG0652 | Tn5252, Orf 28 protein, degenerate |
| SAG0655 | conserved hypothetical protein |
| SAG0678 | endopeptidase O, degenerate |
| SAG0683 | transmembrane protein Vexp3, putative, degenerate |
| SAG0855 | glycogen biosynthesis protein GlgD, authentic frameshift |
| SAG0898 | hypothetical protein |
| SAG0899 | hypothetical protein |
| SAG0901 | hypothetical protein |
| SAG0902 | hypothetical protein |
| SAG0903 | hypothetical protein |
| SAG0917 | Tn916, hypothetical protein |
| SAG0920 | Tn916, hypothetical protein |
| SAG0922 | Tn916, hypothetical protein |
| SAG0924 | Tn916, tetM leader peptide |
| SAG0928 | Tn916, hypothetical protein, authentic frameshift |
| SAG0936 | Tn916, hypothetical protein |
| SAG0930 SAG0943 | hypothetical protein |
| SAG0972 | conserved hypothetical protein, authentic frameshift |
| 5/100//2 | conserved hypothetical protein, authoritic framesimit |

| SAG1023 | hypothetical protein |
|----------|---|
| SAG1080 | hypothetical protein |
| SAG1123 | hypothetical protein |
| SAG1129 | hypothetical protein |
| SAG1136 | conserved hypothetical protein |
| SAG1217 | conserved hypothetical protein, authentic frameshift |
| SAG1231 | transposase OrfB, IS3 family, degenerate |
| SAG1242 | transposase OrfB, IS3 family, truncation |
| SAG1309 | hypothetical protein |
| SAG1331 | R5 protein |
| SAG1437 | hypothetical protein |
| SAG1445 | MutT/nudix family protein, authentic frameshift |
| SAG1484 | ribosomal protein L33 |
| SAG1493 | hypothetical protein |
| SAG1539 | hypothetical protein |
| SAG1543 | conserved hypothetical protein, authentic frameshift |
| SAG1560 | hypothetical protein |
| SAG1568 | phosphoserine aminotransferase, authentic frameshift |
| SAG1570 | conserved hypothetical protein |
| SAG1601 | conserved hypothetical protein |
| SAG1644 | hypothetical protein |
| SAG1646 | hypothetical protein |
| SAG1699 | hypothetical protein |
| SAG1705 | peptidase, M24 family, authentic point mutation |
| SAG1708 | hypothetical protein |
| SAG1857 | prophage LambdaSa2, HNH endonuclease family protein |
| SAG1864 | hypothetical protein |
| SAG1868 | hypothetical protein |
| SAG1869 | prophage LambdaSa2, type II DNA modification methyltransferase, |
| putative | FF8,,, |
| SAG1872 | hypothetical protein |
| SAG1874 | hypothetical protein |
| SAG1876 | prophage LambdaSa2, HNH endonuclease family protein |
| SAG1878 | conserved domain protein |
| SAG1881 | hypothetical protein |
| SAG1883 | conserved hypothetical protein |
| SAG1886 | hypothetical protein |
| SAG1903 | hypothetical protein |
| SAG1937 | streptococcal histidine triad family protein, degenerate |
| SAG1971 | hypothetical protein |
| SAG1979 | membrane protein, putative |
| SAG1980 | ABC transporter, ATP-binding protein |
| SAG1981 | hypothetical protein |
| SAG1982 | transcriptional regulator, Cro/CI family |
| SAG1983 | conserved hypothetical protein |
| SAG1984 | conserved hypothetical protein TIGR00730 |
| 51131707 | constructing positional protein 110100/30 |

| SAG1985 | hypothetical protein |
|------------|--|
| SAG1991 | transcriptional regulator, Cro/CI family |
| SAG1992 | protein of unknown function |
| SAG1999 | hypothetical protein |
| SAG2004 | conjugal transfer protein, interruption-N |
| SAG2039 | conserved hypothetical protein |
| SAG2044 | hypothetical protein |
| SAG2052 | hypothetical protein |
| SAG2065 | ribosomal protein L33 |
| SAG2094 | competence/damage-inducible protein CinA, authentic frameshift |
| SAG2099 | hypothetical protein |
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| Cluster 12 | |
| SAG1164 | glycosyl transferase CpsJ(V) |
| SAG1165 | glycosyl transferase CpsO(V) |
| SAG1166 | glycosyl transferase CpsN(V) |
| SAG1167 | polysaccharide biosynthesis protein CpsM(V) |
| SAG1168 | polysaccharide biosynthesis protein cpsH(V) |
| | Forgonian coordinates broken specific () |
| Cluster 13 | |
| SAG0581 | conserved hypothetical protein |
| SAG0582 | conserved hypothetical protein |
| SAG0583 | conserved hypothetical protein |
| SAG0585 | conserved hypothetical protein |
| SAG0586 | conserved hypothetical protein |
| SAG0587 | prophage LambdaSa1, structural protein, putative |
| SAG0588 | conserved hypothetical protein |
| SAG0589 | conserved hypothetical protein |
| SAG0590 | conserved hypothetical protein |
| SAG0591 | conserved hypothetical protein |
| SAG0593 | prophage LambdaSa1, structural protein |
| SAG0594 | conserved hypothetical protein |
| SAG0595 | conserved hypothetical protein |
| SAG0596 | prophage LambdaSa1, pblA protein, internal deletion |
| 5110000 | propriage Zamoaasur, pon i protein, internar acteuen |
| Cluster 14 | |
| SAG0915 | Tn916, transposase |
| SAG0918 | Tn916, hypothetical protein |
| SAG0919 | Tn916, hypothetical protein |
| SAG0921 | Tn916, transcriptional regulator, putative |
| SAG0925 | Tn916, hypothetical protein |
| SAG0926 | Tn916, NLP/P60 family protein |
| SAG0927 | membrane protein, putative |
| SAG0929 | Tn916, hypothetical protein |
| SAG0930 | Tn916, hypothetical protein |
| SAG0931 | Tn916, hypothetical protein |
| | ,, pomewer proven |
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|----------------|---|
| SAG0932 | Tn916, transcriptional regulator, putative |
| SAG0933 | Tn916, FtsK/SpoIIIE family protein |
| SAG0934 | Tn916, hypothetical protein |
| SAG0935 | Tn916, hypothetical protein |
| SAG0937 | ABC transporter, ATP-binding protein, authentic frameshift |
| Cluster 15 | |
| SAG1835 | conserved hypothetical protein |
| SAG1837 | prophage LambdaSa2, lysin, putative |
| SAG1839 | conserved hypothetical protein |
| SAG1840 | hypothetical protein |
| SAG1842 | prophage LambdaSa2, PblB, putative |
| SAG1843 | conserved hypothetical protein |
| SAG1844 | conserved hypothetical protein |
| SAG1849 | hypothetical protein |
| SAG1851 | conserved domain protein |
| SAG1852 | conserved domain protein |
| SAG1853 | prophage LambdaSa2, protease, putative |
| SAG1854 | conserved hypothetical protein |
| SAG1855 | prophage LambdaSa2, terminase large subunit, putative |
| SAG1856 | hypothetical protein |
| SAG1858 | hypothetical protein |
| SAG1859 | prophage LambdaSa2, site-specific recombinase, phage integrase family |
| SAG1860 | conserved hypothetical protein |
| SAG1861 | prophage LambdaSa2, transcriptional regulator, Cro/CI family |
| SAG1862 | hypothetical protein |
| SAG1863 | prophage LambdaSa2, single-strand binding protein |
| SAG1865 | conserved hypothetical protein |
| SAG1866 | conserved hypothetical protein |
| SAG1867 | conserved hypothetical protein |
| SAG1870 | prophage LambdaSa2, DNA replication protein DnaC, putative |
| SAG1871 | prophage LambdaSa2, bacteriophage replication protein/hypothetical |
| protein, trunc | ration/fusion |
| SAG1873 | prophage LambdaSa2, replicative DNA helicase |
| SAG1877 | prophage LambdaSa2, antirepressor protein, putative |
| SAG1879 | hypothetical protein |
| SAG1882 | prophage LambdaSa2, repressor protein, putative |
| SAG1884 | hypothetical protein |
| SAG1885 | prophage LambdaSa2, site-specific recombinase, phage integrase family |
| Cluster 16 | |
| SAG1247 | site-specific recombinase, phage integrase family |
| SAG1250 | Tn5252, relaxase |
| SAG1251 | Tn5252, Orf 9 protein |
| SAG1252 | Tn5252, Orf 10 protein |
| SAG1256 | IS861, transposase OrfB, truncation |
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| SAG1257 | cation-transporting ATPase, E1-E2 family |
| SAG1258 | cadmium efflux system accessory protein |
| SAG1259 | conserved hypothetical protein |
| SAG1260 | hypothetical protein |
| SAG1261 | conserved hypothetical protein |
| SAG1262 | cation-transporting ATPase, E1-E2 family |
| SAG1263 | conserved domain protein, authentic frameshift |
| SAG1264 | transcriptional repressor CopY, putative |
| SAG1265 | cadmium resistance transporter, putative |
| SAG1266 | hypothetical protein |
| SAG1267 | hypothetical protein |
| SAG1268 | repressor protein, putative |
| SAG1270 | ImpB/MucB/SamB family protein |
| SAG1271 | conserved hypothetical protein |
| SAG1272 | conserved hypothetical protein |
| SAG1273 | conserved hypothetical protein |
| SAG1274 | conserved hypothetical protein |
| SAG1276 | conserved hypothetical protein |
| SAG1277 | hypothetical protein |
| SAG1278 | hypothetical protein |
| SAG1279 | conserved domain protein |
| SAG1280 | SNF2 family protein |
| SAG1281 | hypothetical protein |
| SAG1283 | agglutinin receptor |
| SAG1284 | abortive infection protein AbiGI |
| SAG1285 | abortive infection protein AbiGII |
| SAG1286 | Tn5252, Orf28 |
| SAG1287 | Tn5252, Orf26 |
| SAG1288 | Tn5252, Orf25, degenerate |
| SAG1289 | Tn5252, Orf23 |
| SAG1290 | hypothetical protein |
| SAG1291 | Tn5252, Orf 21 protein, internal deletion |
| SAG1292 | hypothetical protein |
| SAG1293 | protease, putative |
| SAG1294 | conserved hypothetical protein |
| SAG1295 | conserved hypothetical protein |
| SAG1296 | conserved hypothetical protein |
| SAG1297 | C-5 cytosine-specific DNA methylase |
| SAG1299 | conserved hypothetical protein |
| SAG1304 | hypothetical protein |
| | ** |