A New Method Based on Coding Sequence Density to Cluster Bacteria

Supplementary materials

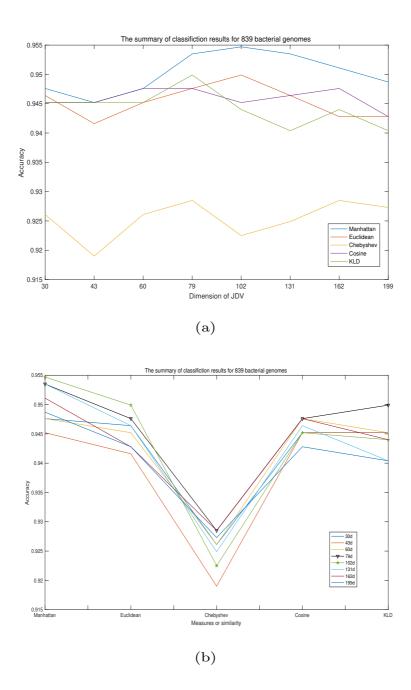
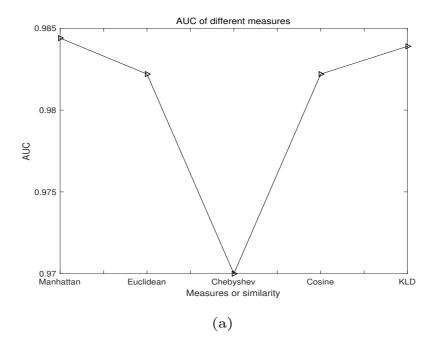


Figure S1: these two figures are consistent with table 2. (a)For different dimensional JDV, Manhattan distance (blue line) performs best. (b) The accuracy rates corresponding to 102 (green line) and 79 (black line) dimensional JDVs are higher than the other dimensions, while the result corresponding to 79 dimensional JDV is more stable, and calculating 102 dimensional JDV takes more time.



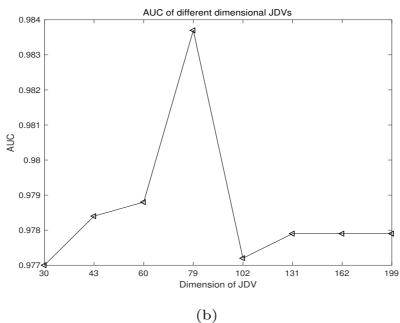


Figure S2: the AUC results of 10-Fold Cross-Validation.

$$\begin{split} \text{(a)} \text{AUC}_{\text{M}} &= 0.9844, \text{AUC}_{\text{E}} = 0.9822, \text{AUC}_{\text{Co}} = 0.9822, \text{AUC}_{\text{KLD}} = 0.9839, \\ \text{error} &= \frac{1}{10} \sum_{i=1}^{10} \text{error}_{\text{i}} = 0 \text{ ; (b)} \text{AUC}_{30} = 0.9770, \text{AUC}_{43} = 0.9784, \text{AUC}_{60} = 0.9788, \\ \text{AUC}_{79} &= 0.9837, \text{AUC}_{102} = 0.9772, \text{AUC}_{131} = 0.9779, \text{AUC}_{162} = 0.9779, \text{AUC}_{199} = 0.9779, \\ \text{error} &= \frac{1}{10} \sum_{i=1}^{10} \text{error}_{\text{i}} = 0. \end{split}$$

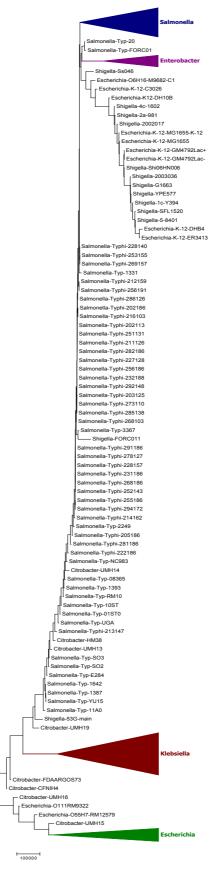


Figure S3: phylogenetic tree for the first dataset by NV method. Bacteria can't be separated.

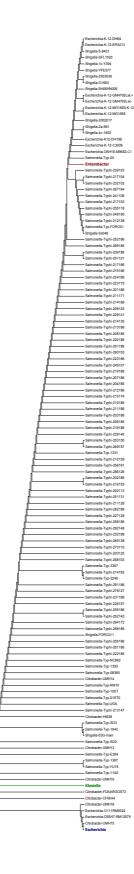


Figure S4: phylogenetic tree for the first dataset by FFT method. Bacteria from different groups can't be separated.

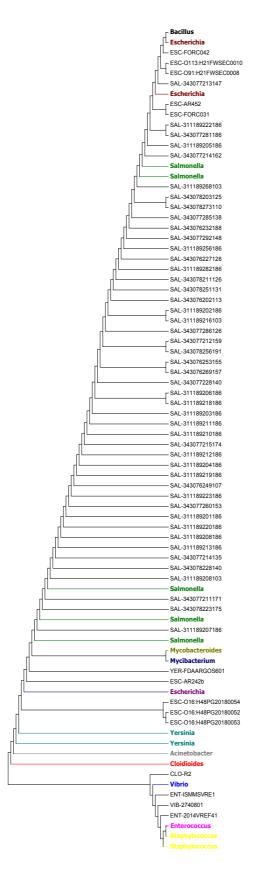


Figure S5: phylogenetic tree for the second dataset by NV method. Bacteria from different genera can't be separated.

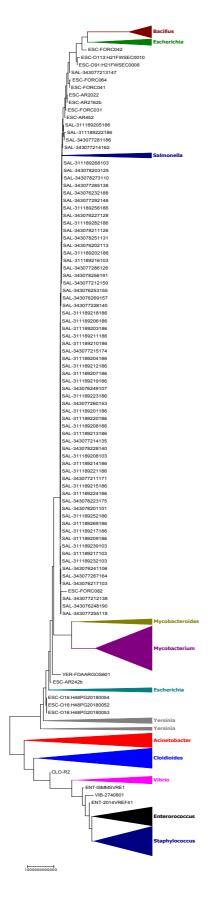


Figure S6: phylogenetic tree for the second dataset by FFT method. Bacteria from different genera can't be separated.

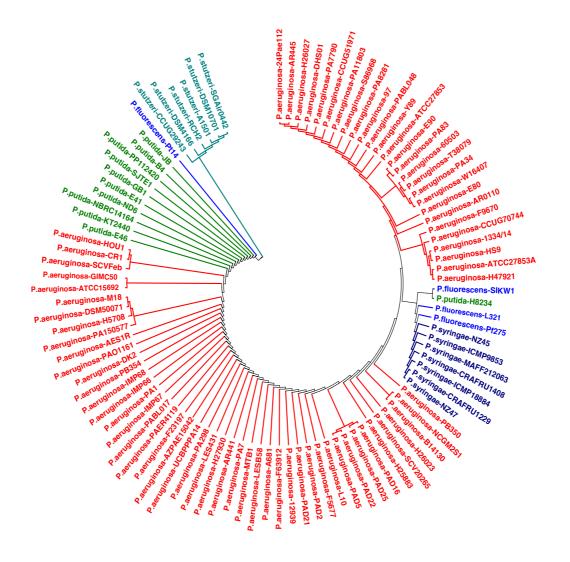


Figure S7: phylogenetic tree for the third dataset by NV method. Bacteria from same species are in the same color

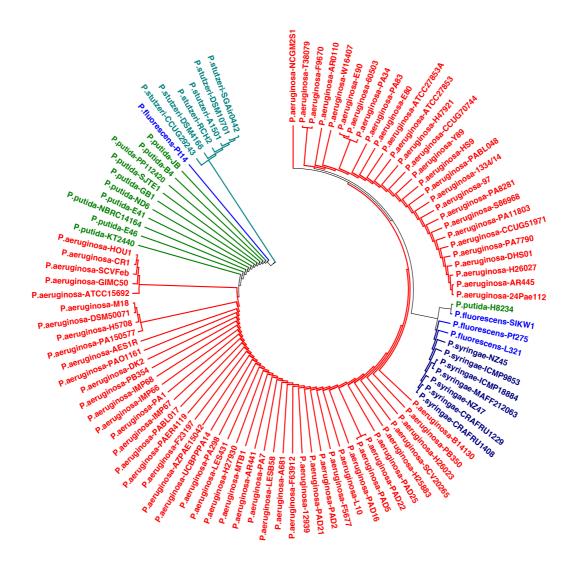


Figure S8: phylogenetic tree for the third dataset by FFT method. The blue species has two branches in this tree.

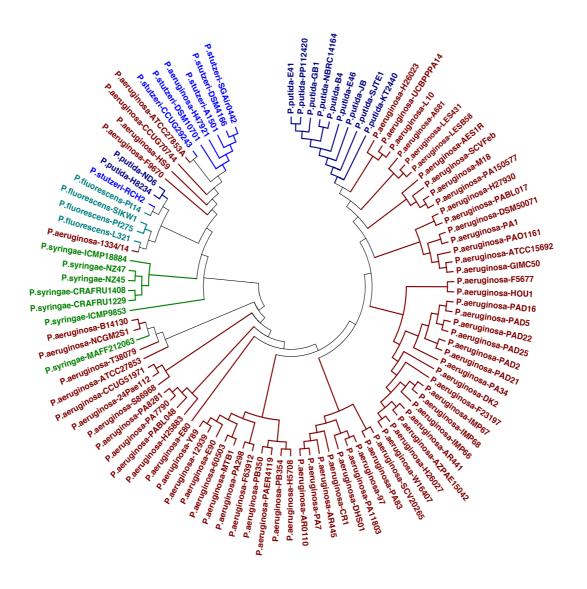


Figure S9: phylogenetic tree for the third dataset by 60d JDV. Bacteria from the same species are in the same color. Bacteria from different species can't be separated. The tree is drawn by NJ in Mega7.

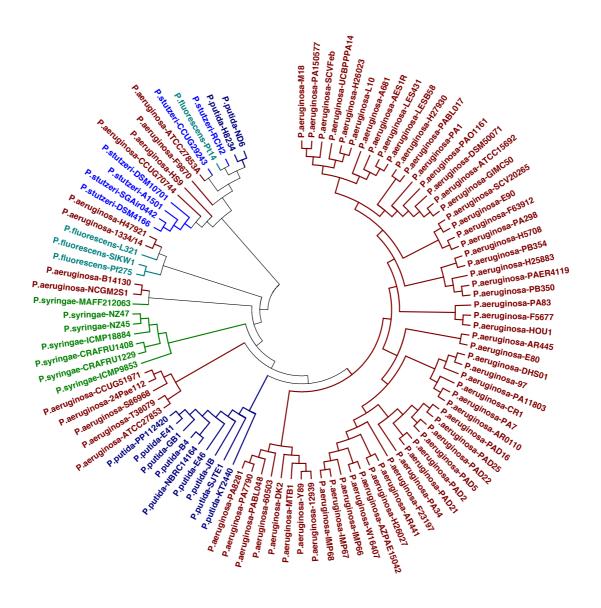


Figure S10: phylogenetic tree for the third dataset by 79d JDV, but the distance matrix is calculated by Euclidean distance. Bacteria from different species can't be separated.

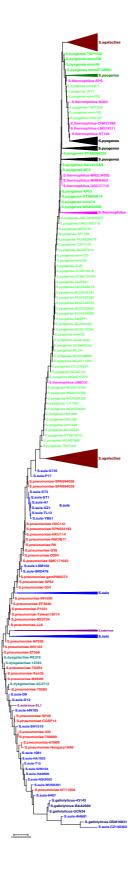


Figure S11: phylogenetic tree for the fourth dataset is constructed by FFT method. Bacteria from different species can't be separated. Bacteria from same species are in the same color.

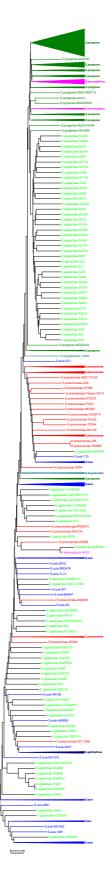


Figure S12: phylogenetic tree for the fourth dataset is constructed by CGR method. Bacteria from different species can't be separated. Bacteria from same species are in the same color.

Table S1: The dataset to determine K

| Name | Accession Num. | Sequence Length | CDS number |
|---|------------------|-----------------|------------|
| Clostridioides difficile 08ACD0030 | NZ_CP010888 | 4167076 | 3779 |
| Clostridioides difficile BR81 | $NZ_CP019870$ | 4124384 | 3684 |
| Clostridioides difficile DH/NAP11/106/ST-42 | $NZ_CP022524$ | 4087127 | 3615 |
| Clostridioides difficile 630 delta erm | NZ_CP016318 | 4293048 | 3944 |
| Clostridioides difficile W0023a | $NZ_CP025045$ | 4110080 | 3675 |
| Clostridioides difficile W0022a | $NZ_CP025046$ | 4188456 | 3812 |
| Clostridioides difficile R0104a | $NZ_CP025044$ | 4190038 | 3793 |
| Clostridioides difficile W0003a | $NZ_CP025047$ | 4075361 | 3635 |
| Clostridioides difficile CD-10-00484 | $NZ_CP026592$ | 4119484 | 3715 |
| Clostridioides difficile 08-00495 | $NZ_CP026594$ | 4177956 | 3774 |
| Clostridioides difficile CD-17-01474 | $NZ_CP026591$ | 4201330 | 3812 |
| Clostridioides difficile 12-00008 | $NZ_CP026593$ | 4119604 | 3714 |
| Clostridioides difficile 12-00011 | $NZ_CP026595$ | 4119932 | 3717 |
| Clostridioides difficile 10-00253 | $NZ_CP026598$ | 4120065 | 3715 |
| Clostridioides difficile 10-00071 | $NZ_CP026596$ | 4119720 | 3715 |
| Clostridioides difficile 10-00078 | $NZ_CP026597$ | 4119749 | 3717 |
| Clostridioides difficile 09-00072 | $NZ_{-}CP026599$ | 4119929 | 3714 |
| Clostridioides difficile FDAARGOS-267 | $NZ_{-}CP020424$ | 4109635 | 3694 |
| Clostridioides difficile CDT4 | $NZ_{-}CP029152$ | 4240954 | 3836 |
| Clostridioides difficile CD161 | NZ_CP029154 | 4295210 | 3925 |
| Clostridioides difficile DSM 27638 | NZ_CP011846 | 4229698 | 3838 |
| Clostridioides difficile DSM 27640 | NZ_CP011848 | 4229629 | 3836 |
| Clostridioides difficile DSM 27639 | NZ_CP011847 | 4263997 | 3899 |
| Clostridioides difficile R2 | NZ_CP026614 | 4093145 | 3707 |
| Clostridioides difficile R3 | $NZ_CP026615$ | 4093147 | 3707 |
| Clostridioides difficile R1 | NZ_CP026613 | 4093143 | 3707 |
| Clostridioides difficile DSM 102859 | $NZ_{-}CP020378$ | 4247015 | 3922 |
| Clostridioides difficile DSM 102978 | NZ_CP020380 | 4253783 | 3943 |
| Clostridioides difficile DSM 102860 | $NZ_{-}CP020379$ | 4253785 | 3943 |
| Clostridioides difficile DSM 29688 | $NZ_{-}CP019858$ | 4228964 | 3913 |
| Clostridioides difficile DSM 29632 | NZ_CP019860 | 4175826 | 3763 |
| Clostridioides difficile DSM 28669 | NZ_CP012323 | 4137693 | 3780 |
| Clostridioides difficile DSM 29637 | NZ_CP016106 | 4114585 | 3716 |
| Clostridioides difficile DSM 29020 | $NZ_CP012325$ | 4139418 | 3776 |
| Clostridioides difficile DSM 28666 | $NZ_{-}CP012321$ | 4122585 | 3701 |
| Clostridioides difficile DSM 29747 | NZ_CP019864 | 4071596 | 3681 |
| Clostridioides difficile DSM 29745 | NZ_CP019857 | 4240646 | 3847 |
| Clostridioides difficile DSM 29629 | NZ_CP016104 | 4111235 | 3674 |
| Clostridioides difficile DSM 29627 | NZ_CP016102 | 4209547 | 3788 |
| Clostridioides difficile DSM 28670 | NZ_CP012312 | 4191653 | 3833 |
| Clostridioides difficile DSM 28196 | NZ_CP012320 | 4205365 | 3820 |
| Clostridioides difficile DSM 28668 | NZ_CP012309 | 4214812 | 3772 |
| Clostridioides difficile 12038 | $NZ_{-}CP033214$ | 4066982 | 3661 |
| Clostridioides difficile CD21062 | NZ_CP033216 | 3996135 | 3610 |
| Clostridioides difficile CD10010 | NZ_CP033213 | 4049580 | 3657 |
| Clostridioides difficile Mta-79 | $NZ_{-}CP042267$ | 4123685 | 3719 |
| Clostridioides difficile 630 | NC_009089 | 4290252 | 3756 |
| Clostridioides difficile 630 | NZ_CP010905 | 4274782 | 3925 |
| Clostridioides difficile ATCC 9689 | NZ_CP011968 | 4109692 | 3697 |
| Clostridium difficile CD196 | $NC_{-}013315$ | 4110554 | 3691 |

| continued table S1 | | | |
|--|------------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Clostridium difficile M120. | NC_017174 | 4047729 | 3634 |
| Enterococcus faecalis LD33 | $NZ_{-}CP014949$ | 2803429 | 2790 |
| Enterococcus faecalis L9 | NZ_CP018004 | 2693039 | 2647 |
| Enterococcus faecalis L12 | $NZ_{-}CP018102$ | 2668255 | 2586 |
| Enterococcus faecalis CLB21560 | $NZ_{-}CP019512$ | 3111017 | 3147 |
| Enterococcus faecalis KB1 | $NZ_{-}CP015410$ | 3026009 | 2939 |
| Enterococcus faecalis sorialis | $NZ_{-}CP015883$ | 2873100 | 2778 |
| Enterococcus faecalis FDAARGOS-338 | $NZ_{-}CP022059$ | 2861022 | 2858 |
| Enterococcus faecalis FDAARGOS-324 | $NZ_{-}CP028285$ | 2928142 | 2896 |
| Enterococcus faecalis H25 | NZ_CP029612 | 3132715 | 3214 |
| Enterococcus faecalis TY1 | $NZ_{-}CP031027$ | 3065679 | 3106 |
| Enterococcus faecalis CVM N48037F | NZ_CP028720 | 2997828 | 2971 |
| Enterococcus faecalis CVM N60443F | NZ_CP028724 | 2758818 | 2678 |
| Enterococcus faecalis FC | NZ_CP028835 | 2934454 | 2918 |
| Enterococcus faecalis SGAir0397 | NZ_CP039434 | 2696714 | 2650 |
| Enterococcus faecalis C25 | NZ_CP030042 | 2825273 | 2793 |
| Enterococcus faecalis C54 | NZ_CP030045 | 2813589 | 2787 |
| Enterococcus faecalis FDAARGOS-611 | NZ_CP041012 | 2845446 | 2815 |
| Enterococcus faecalis HA-1 | NZ_CP040898 | 2881081 | 2879 |
| Enterococcus faecalis VE18379 | NZ_CP039548 | 3218315 | 3317 |
| Enterococcus faecalis VE18395 | NZ_CP039549 | 3201501 | 3304 |
| Enterococcus faecalis VE14089 | NZ_CP039296 | 3238854 | 3346 |
| Enterococcus faecalis DM01 | NZ_CP041344 | 2785968 | 2750 |
| Enterococcus faecalis ARO1/DG | NZ_CP022488 | 2717680 | 2670 |
| Enterococcus faecalis ATCC 29212 | NZ_CP008816 | 2939973 | 2940 |
| Enterococcus faecalis D32 | NC_018221 | 2987450 | 3006 |
| Enterococcus faecalis DENG1 | NZ_CP004081 | 2961043 | 2981 |
| Enterococcus faecalis EnGen0107 B594 | NZ_CP041738 | 3152103 | 3229 |
| Enterococcus faecalis OG1RF | NC_017316 | 2739625 | 2636 |
| Enterococcus faecalis Symbioflor 1 | NC_019770 | 2810675 | 2806 |
| Enterococcus faecalis V583 | NC_004668 | 3218031 | 3112 |
| Escherichia coli BL21-Gold(DE3)pLysS AG | NC_012947 | 4570938 | 4356 |
| Escherichia coli 042 | NC_017626 | 5241977 | 5053 |
| Escherichia coli 536 | NC_008253 | 4938920 | 4663 |
| Escherichia coli 55989 | NC_011748 | 5154862 | 4985 |
| Escherichia coli ABU 83972 | NC_01743 NC_017631 | 5131397 | 4928 |
| Escherichia coli APEC O1 | NC_008563 | 5082025 | 4956 |
| Escherichia coli APEC O78 | NC_020163 | 4798435 | 4596 |
| Escherichia coli ATCC 8739 | NC_010468 | 4746218 | 4448 |
| Escherichia coli B REL606 | NC_010408 NC_012967 | 4629812 | 4372 |
| Escherichia coli BL21(DE3) | NC_012892 | 4558947 | 4324 |
| Escherichia coli BL21(DE3) Escherichia coli BL21(DE3) | NC_012892 NC_012971 | 4558953 | 4324 |
| Escherichia coli BW2952 | | | |
| | NC_012759 | 4578159 | 4333 |
| Escherichia coli CFT073 | NC_004431 | 5231428 | 5055 |
| Escherichia coli DH1 | NC_017625 | 4630707 | 4358 |
| Escherichia coli DH1 (ME8569) | NC 017638 | 4621430 | 4347 |
| Escherichia coli ETEC H10407 | NC_017633 | 5153435 | 5278 |
| Escherichia coli HS | NC_009800 | 4643538 | 4387 |
| Escherichia coli IAI1 | NC_011741 | 4700560 | 4408 |
| Escherichia coli IAI39 | NC_011750 | 5132068 | 4725 |
| Mycobacterium tuberculosis I0002353-6 | NZ_CP018300 | 4385578 | 4208 |
| Mycobacterium tuberculosis M0002959-6 | NZ_CP018304 | 4386447 | 4208 |

continued table S1

| continued table S1 | | | |
|---|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Mycobacterium tuberculosis I0002801-4 | NZ_CP018301 | 4376067 | 4203 |
| Mycobacterium tuberculosis DK9897 | NZ_CP018778 | 4411511 | 4232 |
| Mycobacterium tuberculosis MTB1 | NZ_CP020381 | 4433542 | 4255 |
| Mycobacterium tuberculosis Beijing-like/36918 | $NZ_{-}CP017594$ | 4441591 | 4260 |
| Mycobacterium tuberculosis Beijing-like/50148 | NZ_CP017597 | 4444417 | 4266 |
| Mycobacterium tuberculosis Beijing-like/35049 | NZ_CP017593 | 4427062 | 4250 |
| Mycobacterium tuberculosis Beijing-like/1104 | NZ_CP017598 | 4380156 | 4220 |
| Mycobacterium tuberculosis Beijing-like/38774 | $NZ_{-}CP017595$ | 4431885 | 4259 |
| Mycobacterium tuberculosis Beijing/391 | NZ_CP017596 | 4406925 | 4246 |
| Mycobacterium tuberculosis MTB2 | NZ_CP022014 | 4417716 | 4238 |
| Mycobacterium tuberculosis CSV4644 | $NZ_CP023574$ | 4411271 | 4238 |
| Mycobacterium tuberculosis LE492 | $NZ_{-}CP023579$ | 4411208 | 4236 |
| Mycobacterium tuberculosis LN3588 | NZ_CP023614 | 4411379 | 4237 |
| Mycobacterium tuberculosis MDRDM627 | $NZ_{-}CP023584$ | 4411215 | 4236 |
| Mycobacterium tuberculosis LN3589 | $NZ_CP023615$ | 4411392 | 4234 |
| Mycobacterium tuberculosis TBV5000 | $NZ_{-}CP023589$ | 4411318 | 4234 |
| Mycobacterium tuberculosis LN1100 | NZ_CP023619 | 4411392 | 4235 |
| Mycobacterium tuberculosis SLM056 | $NZ_{-}CP023594$ | 4411306 | 4232 |
| Mycobacterium tuberculosis LN1856 | $NZ_CP023620$ | 4411299 | 4236 |
| Mycobacterium tuberculosis MDRMA701 | $NZ_{-}CP023624$ | 4411338 | 4235 |
| Mycobacterium tuberculosis CSV383 | $NZ_{-}CP023599$ | 4411115 | 4235 |
| Mycobacterium tuberculosis LE76 | NZ_CP023604 | 4411211 | 4238 |
| Mycobacterium tuberculosis MDRMA863 | $NZ_CP023625$ | 4411432 | 4236 |
| Mycobacterium tuberculosis LN55 | $NZ_{-}CP023609$ | 4411186 | 4233 |
| Mycobacterium tuberculosis MDRMA2441 | $NZ_{-}CP023630$ | 4411454 | 4238 |
| Mycobacterium tuberculosis TBDM2489 | $NZ_CP023635$ | 4411369 | 4238 |
| Mycobacterium tuberculosis TBV4952 | $NZ_{-}CP023640$ | 4411414 | 4234 |
| Mycobacterium tuberculosis CSV10399 | $NZ_{-}CP023576$ | 4411180 | 4232 |
| Mycobacterium tuberculosis LN2358 | $NZ_{-}CP023581$ | 4411353 | 4239 |
| Mycobacterium tuberculosis LN3756 | $NZ_CP023582$ | 4411315 | 4234 |
| Mycobacterium tuberculosis MDRMA2491 | $NZ_{-}CP023586$ | 4411121 | 4229 |
| Mycobacterium tuberculosis ME1473 | $NZ_{-}CP023587$ | 4411217 | 4238 |
| Mycobacterium tuberculosis TBV5365 | $NZ_{-}CP023591$ | 4411398 | 4234 |
| Mycobacterium tuberculosis SLM036 | $NZ_CP023592$ | 4411342 | 4236 |
| Mycobacterium tuberculosis CSV4519 | $NZ_CP023573$ | 4411288 | 4239 |
| Mycobacterium tuberculosis SLM088 | $NZ_{-}CP023597$ | 4411385 | 4236 |
| Mycobacterium tuberculosis CSV11678 | $NZ_{-}CP023577$ | 4411382 | 4237 |
| Mycobacterium tuberculosis LE13 | $NZ_CP023602$ | 4411412 | 4234 |
| Mycobacterium tuberculosis LE486 | $NZ_{-}CP023578$ | 4411180 | 4236 |
| Mycobacterium tuberculosis LE371 | $NZ_{-}CP023607$ | 4411137 | 4234 |
| Mycobacterium tuberculosis MDRDM260 | $NZ_{-}CP023583$ | 4411280 | 4237 |
| Mycobacterium tuberculosis LN2978 | $NZ_{-}CP023612$ | 4411331 | 4233 |
| Mycobacterium tuberculosis TBDM425 | $NZ_{-}CP023588$ | 4411143 | 4235 |
| Mycobacterium tuberculosis LN3672 | $NZ_{-}CP023617$ | 4411335 | 4236 |
| Mycobacterium tuberculosis SLM040 | $NZ_{-}CP023593$ | 4411408 | 4236 |
| Mycobacterium tuberculosis MDRDM827 | $NZ_{-}CP023622$ | 4411315 | 4238 |
| Mycobacterium tuberculosis SLM100 | $NZ_{-}CP023598$ | 4411394 | 4234 |
| Mycobacterium tuberculosis MDRMA2019 | $NZ_{-}CP023627$ | 4411229 | 4236 |
| Escherichia coli IHE3034 | $NC_{-}017628$ | 5108383 | 4966 |
| Escherichia coli JJ1886 | $NC_{-}022648$ | 5129938 | 5010 |
| Escherichia coli KO11 | $NC_{-}016902$ | 4920168 | 4952 |
| Escherichia coli KO11FL | $NC_{-}017660$ | 5021812 | 4834 |

| Name | Accession Num. | Sequence Length | CDS number |
|--|------------------|-----------------|------------|
| Escherichia coli LF82 | NC_011993 | 4773108 | 453 |
| Escherichia coli LY180 | $NC_{-}022364$ | 4835601 | 460 |
| Escherichia coli O103:H2 12009 | $NC_{-}013353$ | 5449314 | 542 |
| Escherichia coli O104:H4 2009EL-2050 | $NC_{-}018650$ | 5253138 | 512 |
| Escherichia coli O104:H4 2009EL-2071 | $NC_{-}018661$ | 5312586 | 519 |
| Escherichia coli O104:H4 2011C-3493 | $NC_{-}018658$ | 5273097 | 496 |
| Escherichia coli O111:H- 11128 | $NC_{-}013364$ | 5371077 | 533 |
| Escherichia coli 0127:H6 E2348/69 | NC_011601 | 4965553 | 513 |
| Escherichia coli E24377A | NC_009801 | 4979619 | 504 |
| Escherichia coli O157:H7 EC4115 | $NC_{-}011353$ | 5572075 | 557 |
| Escherichia coli O157:H7 Sakai | $NC_{-}002695$ | 5498578 | 520 |
| Escherichia coli O157:H7 TW14359 | $NC_{-}013008$ | 5528136 | 551 |
| Escherichia coli O26:H11 11368 | $NC_{-}013361$ | 5697240 | 577 |
| Escherichia coli O55:H7 CB9615 | $NC_{-}013941$ | 5386352 | 523 |
| Escherichia coli O55:H7 RM12579 | $NC_{-}017656$ | 5263980 | 509 |
| Escherichia coli O7:K1 CE10 | $NC_{-}017646$ | 5313531 | 514 |
| Escherichia coli O83:H1 NRG 857C | $NC_{-}017634$ | 4747819 | 442 |
| Escherichia coli P12b | $NC_{-}017663$ | 4935294 | 478 |
| Escherichia coli S88 | $NC_{-}011742$ | 5032268 | 514 |
| Escherichia coli SE11 | $NC_{-}011415$ | 4887515 | 465 |
| Escherichia coli SE15 | $NC_{-}013654$ | 4717338 | 444 |
| Escherichia coli SMS-3-5 | NC_010498 | 5068389 | 476 |
| Escherichia coli clone D i14 | $NC_{-}017652$ | 5038386 | 480 |
| Escherichia coli clone D i2 | $NC_{-}017651$ | 5038386 | 480 |
| Escherichia coli K12 subDH10B | $NC_{-}010473$ | 4686137 | 473 |
| Escherichia coli K-12 subMDS42 | $NC_{-}020518$ | 3976195 | 365 |
| Escherichia coli K-12 subMG1655 | NC_000913 | 4641652 | 435 |
| Escherichia coli K-12 subW3110 | $NC_{-}007779$ | 4646332 | 467 |
| Mycobacterium tuberculosis 96121 | $NZ_{-}CP009427$ | 4410945 | 425 |
| Mycobacterium tuberculosis 96075 | $NZ_{-}CP009426$ | 4379376 | 423 |
| Mycobacterium tuberculosis KIT87190 | NZ_CP007809 | 4410788 | 425 |
| Mycobacterium tuberculosis ZMC13-88 | NZ_CP009101 | 4411515 | 423 |
| Mycobacterium tuberculosis ZMC13-264 | NZ_CP009100 | 4411507 | 424 |
| Mycobacterium tuberculosis 22103 | NZ_CP010339 | 4399422 | 424 |
| Mycobacterium tuberculosis 22115 | NZ_CP010337 | 4401829 | 420 |
| Mycobacterium tuberculosis F28 | NZ_CP010330 | 4421903 | 420 |
| Mycobacterium tuberculosis 26105 | NZ_CP010340 | 4426489 | 420 |
| Mycobacterium tuberculosis SCAID 320.0 | NZ_CP016794 | 4406628 | 425 |
| Mycobacterium tuberculosis SCAID 252.0 | NZ_CP016888 | 4439387 | 425 |
| Mycobacterium tuberculosis SCAID 187.0 | $NZ_{-}CP012506$ | 4411829 | 424 |
| Mycobacterium tuberculosis Beijing | NZ_CP011510 | 4378588 | 419 |
| Mycobacterium tuberculosis 1458 | NZ_CP013475 | 4402033 | 422 |
| Mycobacterium tuberculosis TB282 | NZ_CP017920 | 4425860 | 425 |
| Mycobacterium tuberculosis I0004000-1 | $NZ_{-}CP018302$ | 4365724 | 420 |
| Mycobacterium tuberculosis I0004241-1 | NZ_CP018303 | 4386132 | 420 |
| Mycobacterium tuberculosis M0018684-2 | NZ_CP018305 | 4359825 | 418 |
| Mycobacterium tuberculosis LE63 | NZ_CP023603 | 4411415 | 423 |
| Mycobacterium tuberculosis TBDM2189 | $NZ_CP023632$ | 4411316 | 423 |
| Mycobacterium tuberculosis LE410 | $NZ_{-}CP023608$ | 4411365 | 425 |
| Mycobacterium tuberculosis LN3584 | NZ_CP023613 | 4411326 | 423 |
| Mycobacterium tuberculosis TBDM2717 | NZ_CP023637 | 4411442 | 423 |
| Mycobacterium tuberculosis LN3695 | $NZ_{-}CP023618$ | 4411449 | 423 |

continued table S1

| continued table S1 | | | |
|--|----------------------------|-----------------|---------------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Mycobacterium tuberculosis MDRMA203 | NZ_CP023623 | 4411290 | 4234 |
| Mycobacterium tuberculosis MDRMA2082 | $NZ_{-}CP023628$ | 4411479 | 4234 |
| Mycobacterium tuberculosis MDRMA2260 | NZ_CP023629 | 4411352 | 4237 |
| Mycobacterium tuberculosis TBDM2444 | NZ_CP023633 | 4411284 | 4237 |
| Mycobacterium tuberculosis TBDM2487 | $NZ_{CP023634}$ | 4411314 | 4237 |
| Mycobacterium tuberculosis CSV5769 | $NZ_{-}CP023575$ | 4411312 | 4235 |
| Mycobacterium tuberculosis TBV4766 | NZ_CP023638 | 4411310 | 4234 |
| Mycobacterium tuberculosis LN180 | $NZ_{-}CP023580$ | 4411436 | 4234 |
| Mycobacterium tuberculosis TBV4768 | NZ_CP023639 | 4411173 | 4229 |
| Mycobacterium tuberculosis MDRDM1098 | NZ_CP023585 | 4411148 | 4235 |
| Mycobacterium tuberculosis TBV5362 | NZ_CP023590 | 4411331 | 4235 |
| Mycobacterium tuberculosis SLM060 | $NZ_{-}CP023595$ | 4411134 | 4231 |
| Mycobacterium tuberculosis SLM063 | NZ_CP023596 | 4411337 | 4234 |
| Mycobacterium tuberculosis CSV3611 | NZ_CP023600 | 4411439 | 4236 |
| Mycobacterium tuberculosis CSV9577 | NZ_CP023601 | 4411230 | 4237 |
| Mycobacterium tuberculosis LE79 | NZ_CP023605 | 4411475 | 4237 |
| Mycobacterium tuberculosis LE103 | NZ_CP023606 | 4411243 | 4233 |
| Mycobacterium tuberculosis LN317 | NZ_CP023610 | 4411340 | 4234 |
| Mycobacterium tuberculosis LN763 | NZ_CP023611 | 4411321 | 4235 |
| Mycobacterium tuberculosis LN3668 | NZ_CP023616 | 4411494 | 4234 |
| Mycobacterium tuberculosis LN2900 | NZ_CP023621 | 4411327 | 4235 |
| Mycobacterium tuberculosis MDRMA1565 | NZ_CP023626 | 4411159 | 4233 |
| Mycobacterium tuberculosis TBDM1506 | NZ_CP023631 | 4411157 | 4230 |
| Mycobacterium tuberculosis TBDM2699 | NZ_CP023636 | 4411457 | 4235 |
| Mycobacterium tuberculosis GG-90-10 | NZ_CP025601 | 4411602 | 4250 |
| Mycobacterium tuberculosis GG-5-10 | NZ_CP025594 | 4411442 | 4246 |
| Mycobacterium tuberculosis GG-129-11 | NZ_CP025604 | 4411413 | 4242 |
| Mycobacterium tuberculosis GG-123-11 Mycobacterium tuberculosis GG-111-10 | NZ_CP025593 | 4411563 | 4245 |
| Mycobacterium tuberculosis GG-20-11 | NZ_CP025595 | 4411504 | 4242 |
| Mycobacterium tuberculosis GG-20-11 Mycobacterium tuberculosis GG-134-11 | NZ_CP025605 | 4411399 | 4244 |
| Mycobacterium tuberculosis GG-134-11 Mycobacterium tuberculosis GG-229-10 | NZ_CP025608 | 4411519 | 4244 |
| Mycobacterium tuberculosis GG-229-10 Mycobacterium tuberculosis GG-37-11 | NZ_CP025598 | 4411519 | 4244 4247 |
| Mycobacterium tuberculosis GG-186-10 | NZ_CP025607 | 4411478 | 4245 |
| | NZ_CP025603 | 4411510 | 4243 |
| Mycobacterium tuberculosis GG-121-10 Mycobacterium tuberculosis GG-27-11 | | | $\frac{4241}{4247}$ |
| Mycobacterium tuberculosis GG-27-11 Mycobacterium tuberculosis GG-137-10 | NZ_CP025596 | 4411443 | |
| | NZ_CP025606 NZ_CP025602 | 4411446 | 4243 |
| Mycobacterium tuberculosis GG-109-10 | | 4411463 | 4243 |
| Mycobacterium tuberculosis GG-45-11 | NZ_CP025599 | 4411469 | 4243 |
| Mycobacterium tuberculosis GG-36-11 | NZ_CP025597 | 4411469 | 4245 |
| Mycobacterium tuberculosis GG-77-11 | NZ_CP025600 | 4411508 | 4243 |
| Mycobacterium tuberculosis WC059 | NZ_CP022578 | 4413669 | 4254 |
| Mycobacterium tuberculosis WC078 | NZ_CP022577 | 4413712 | 4254 |
| Mycobacterium tuberculosis H107 | NZ_CP019612 | 4418796 | 4257 |
| Mycobacterium tuberculosis RUS-B0 | NZ_CP030093 | 4418559 | 4253 |
| Mycobacterium tuberculosis H54 | NZ_CP019610 | 4416938 | 4263 |
| Mycobacterium tuberculosis H83 | NZ_CP019611 | 4413214 | 4242 |
| Mycobacterium tuberculosis H112 | NZ_CP019613 | 4406346 | 4249 |
| Mycobacterium tuberculosis TBMENG-03 | NZ_CP029065 | 4359659 | 4216 |
| Mycobacterium tuberculosis MT-0080 | NZ_CP041207 | 4426525 | 4267 |
| Mycobacterium tuberculosis L | NZ_CP044345 | 4416671 | 4260 |
| Mycobacterium tuberculosis Beijing2014PNGD | NZ_CP022704 | 4404064 | 4264 |
| Mycobacterium tuberculosis 7199-99 | NC_020089 | 4421197 | 4249 |

| continued table S1 | | | |
|--|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Mycobacterium tuberculosis BT2 | NZ_CP002882 | 4401899 | 4239 |
| Mycobacterium tuberculosis BT1 | NZ_CP002883 | 4399405 | 4239 |
| Mycobacterium tuberculosis CCDC5079 | $NC_{-}021251$ | 4414325 | 4249 |
| Mycobacterium tuberculosis CCDC5180 | $NC_{-}017522$ | 4405981 | 4251 |
| Mycobacterium tuberculosis CCDC5180 | $NZ_{-}CP002885$ | 4414346 | 4248 |
| Mycobacterium tuberculosis CDC1551 | $NC_{-}002755$ | 4403837 | 4242 |
| Mycobacterium tuberculosis CTRI-2 | $NC_{-}017524$ | 4398525 | 4241 |
| Mycobacterium tuberculosis EAI5 | $NC_{-}021740$ | 4391174 | 4223 |
| Mycobacterium tuberculosis EAI5/NITR206 | NC_021194 | 4390306 | 4226 |
| Mycobacterium tuberculosis F11 | $NC_{-}009565$ | 4424435 | 4237 |
| Mycobacterium tuberculosis H37Ra | $NC_{-}009525$ | 4419977 | 4257 |
| Mycobacterium tuberculosis H37Ra | $NZ_{-}CP016972$ | 4426109 | 4248 |
| Mycobacterium tuberculosis H37Rv | $NC_{-}000962$ | 4411532 | 3906 |
| Mycobacterium tuberculosis H37Rv | $NC_{-}018143$ | 4411709 | 4243 |
| Mycobacterium tuberculosis H37RvSiena | $NZ_CP007027$ | 4410911 | 4242 |
| Mycobacterium tuberculosis HKBS1 | NZ_CP002871 | 4407929 | 4246 |
| Mycobacterium tuberculosis K | NZ_CP007803 | 4385518 | 4235 |
| Mycobacterium tuberculosis KZN 1435 | $NC_{-}012943$ | 4398250 | 4226 |
| Mycobacterium tuberculosis KZN 4207 | NC_016768 | 4394985 | 4220 |
| Mycobacterium tuberculosis KZN 605 | $NC_{-}018078$ | 4399120 | 4225 |
| Mycobacterium tuberculosis Erdman | $NC_{-}020559$ | 4392353 | 4251 |
| Mycobacterium tuberculosis Haarlem | $NC_{-}022350$ | 4408224 | 4233 |
| Mycobacterium africanum GM041182 | NC_015758 | 4389314 | 4192 |
| Mycobacterium bovis BCG 3281 | NZ_CP008744 | 4410431 | 4217 |
| Mycobacterium bovis BCG Russia 368 | NZ_CP009243 | 4370138 | 4182 |
| Mycobacterium bovis BCG ATCC 35743 | NZ_CP003494 | 4334064 | 4175 |
| Mycobacterium bovis BCG Korea 1168P | $NC_{-}020245$ | 4376711 | 4190 |
| Mycobacterium bovis BCG Mexico | $NC_{-}016804$ | 4350386 | 4162 |
| Mycobacterium bovis BCG Pasteur 1173P2 | $NC_{-}008769$ | 4374522 | 4186 |
| Mycobacterium bovis BCG Tokyo 172 | $NC_{-}012207$ | 4371711 | 4183 |
| Mycobacterium tuberculosis variant bovis BCG Tokyo 172 | NZ_CP014566 | 4371707 | 4224 |
| Mycobacterium tuberculosis W-148 | $NZ_{-}CP012090$ | 4418548 | 4237 |
| Pseudomonas aeruginosa FRD1 | $NZ_{-}CP010555$ | 6712339 | 6290 |
| Pseudomonas aeruginosa Carb01 63 | NZ_CP011317 | 7497593 | 7228 |
| Pseudomonas aeruginosa F9676 | NZ_CP012066 | 6368008 | 5927 |
| Pseudomonas aeruginosa F22031 | NZ_CP007399 | 6603099 | 6182 |
| Pseudomonas aeruginosa PA1RG | NZ_CP012679 | 6500439 | 6128 |
| Pseudomonas aeruginosa 12-4-4(59) | NZ_CP013696 | 6431911 | 6056 |
| Pseudomonas aeruginosa H47921 | NZ_CP008861 | 6836415 | 6468 |
| Pseudomonas aeruginosa T52373 | NZ_CP008867 | 6322459 | 5869 |
| Pseudomonas aeruginosa T63266 | NZ_CP008868 | 6456866 | 6066 |
| Pseudomonas aeruginosa M37351 | NZ_CP008863 | 6897231 | 6552 |
| Pseudomonas aeruginosa F30658 | NZ_CP008857 | 7273258 | 6942 |
| Pseudomonas aeruginosa USDA-ARS-USMARC-41639 | NZ_CP013989 | 6364583 | 5933 |
| Pseudomonas aeruginosa VA-134 | NZ_CP013245 | 6400418 | 5955 |
| Pseudomonas aeruginosa F9670 | NZ_CP008873 | 6794354 | 6388 |
| Pseudomonas aeruginosa N17-1 | NZ_CP014948 | 6370730 | 5959 |
| Pseudomonas aeruginosa ATCC 27853 | NZ_CP015117 | 6827737 | 6473 |
| Pseudomonas aeruginosa BAMCPA07-48 | NZ_CP015377 | 7021552 | 6716 |
| Pseudomonas aeruginosa PA121617 | NZ_CP016214 | 6430493 | 5982 |
| Pseudomonas aeruginosa ATCC 27853 | NZ_CP011857 | 6833187 | 6463 |
| Pseudomonas aeruginosa PA-D22 | NZ_CP012583 | 6681981 | 6248 |

| Name | Accession Num. | Sequence Length | CDS numbe |
|-------------------------------------|------------------|-----------------|-----------|
| Pseudomonas aeruginosa DN1 | NZ_CP017099 | 6641902 | 626 |
| Pseudomonas aeruginosa PA-D1 | $NZ_CP012585$ | 6643823 | 620 |
| Pseudomonas aeruginosa PA-D16 | $NZ_CP012581$ | 6681975 | 624 |
| Pseudomonas aeruginosa PA-D5 | $NZ_{-}CP012579$ | 6681992 | 624 |
| Pseudomonas aeruginosa PA-D21 | $NZ_{-}CP012582$ | 6639108 | 620 |
| Pseudomonas aeruginosa PA-D25 | $NZ_CP012584$ | 6683204 | 625 |
| Pseudomonas aeruginosa PA-D9 | NZ_CP012580 | 6645477 | 621 |
| Pseudomonas aeruginosa PA-D2 | $NZ_{-}CP012578$ | 6642996 | 620 |
| Pseudomonas aeruginosa ATCC 15692 | NZ_CP017149 | 6276434 | 584 |
| Pseudomonas aeruginosa FA-HZ1 | $NZ_{-}CP017353$ | 6866790 | 651 |
| Pseudomonas aeruginosa PA11803 | NZ_CP015003 | 7006578 | 673 |
| Pseudomonas aeruginosa PA8281 | NZ_CP015002 | 6928736 | 656 |
| Pseudomonas aeruginosa PA1088 | NZ_CP015001 | 6721480 | 633 |
| Pseudomonas aeruginosa PA7790 | NZ_CP014999 | 7018690 | 668 |
| Pseudomonas aeruginosa isolate B10W | NZ_CP017969 | 6723378 | 635 |
| Pseudomonas aeruginosa PAER4-119 | NZ_CP013113 | 6504659 | 605 |
| Pseudomonas aeruginosa SCVJan | NZ_CP013478 | 6213029 | 57 |
| Pseudomonas aeruginosa NHmuc | NZ_CP013479 | 6213276 | 578 |
| Pseudomonas aeruginosa SCVFeb | NZ_CP013477 | 6213026 | 57' |
| Pseudomonas aeruginosa PA-154197 | NZ_CP014866 | 6445239 | 603 |
| Pseudomonas aeruginosa RIVM-EMC2982 | NZ_CP016955 | 7380063 | 708 |
| Pseudomonas aeruginosa E6130952 | NZ_CP020603 | 7040952 | 673 |
| Pseudomonas aeruginosa T38079 | NZ_CP008866 | 6795741 | 639 |
| Pseudomonas aeruginosa F23197 | NZ_CP008856 | 6517340 | 60 |
| Pseudomonas aeruginosa W60856 | NZ_CP008864 | 6896700 | 649 |
| Pseudomonas aeruginosa W45909 | NZ_CP008871 | 6777566 | 64 |
| Pseudomonas aeruginosa M1608 | NZ_CP008862 | 6460023 | 608 |
| Pseudomonas aeruginosa X78812 | NZ_CP008872 | 6348761 | 590 |
| Pseudomonas aeruginosa H5708 | NZ_CP008859 | 6334378 | 590 |
| Pseudomonas aeruginosa W16407 | NZ_CP008869 | 6808844 | 649 |
| Pseudomonas aeruginosa W36662 | NZ_CP008870 | 6791731 | 63 |
| Pseudomonas aeruginosa H27930 | NZ_CP008860 | 6568228 | 61 |
| Pseudomonas aeruginosa S86968 | NZ_CP008865 | 6934277 | 65 |
| 9 | | | 66 |
| Pseudomonas aeruginosa PASGNDM699 | NZ_CP020704 | 6985102 | |
| Pseudomonas aeruginosa PASGNDM345 | NZ_CP020703 | 6893164 | 65 |
| Pseudomonas aeruginosa F63912 | NZ_CP008858 | 6618768 | 62 |
| Pseudomonas aeruginosa Pa58 | NZ_CP021775 | 7241575 | 69 |
| Pseudomonas aeruginosa Pa124 | NZ_CP021774 | 7008516 | 66 |
| Pseudomonas aeruginosa Pa84 | NZ_CP021999 | 6566724 | 62 |
| Pseudomonas aeruginosa Pa1242 | NZ_CP022002 | 7050510 | 66 |
| Pseudomonas aeruginosa NCGM2.S1 | NC_017549 | 6764661 | 64 |
| Pseudomonas aeruginosa PA1 | NC_022808 | 6498072 | 61 |
| Pseudomonas aeruginosa PA1R | NC_022806 | 6309305 | 593 |
| Pseudomonas aeruginosa PA7 | NC_009656 | 6588339 | 623 |
| Pseudomonas aeruginosa PAK | NZ_CP020659 | 6283316 | 59 |
| Pseudomonas aeruginosa PAO1 | NC_002516 | 6264404 | 55' |
| Pseudomonas aeruginosa RP73 | NC_021577 | 6342034 | 59. |
| Pseudomonas aeruginosa SCV20265 | NC_023149 | 6725183 | 63 |
| Pseudomonas aeruginosa SJTD-1 | $NZ_{-}CP015877$ | 6243825 | 580 |
| Pseudomonas aeruginosa UCBPP-PA14 | $NC_{-}008463$ | 6537648 | 61 |
| Pseudomonas aeruginosa VRFPA04 | NZ_CP008739 | 6818030 | 64 |
| Pseudomonas aeruginosa YL84 | NZ_CP007147 | 6433441 | 60 |

continued table S1

| continued table S1 | | | |
|---------------------------------|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Salmonella enterica CFSAN000189 | NC_021844 | 4730612 | 4479 |
| Salmonella enterica CFSAN000191 | $NZ_{-}CP032622$ | 4727462 | 4473 |
| Salmonella enterica CFSAN000211 | NZ_CP039500 | 4727567 | 4477 |
| Salmonella enterica CFSAN000212 | NZ_CP039499 | 4727570 | 4473 |
| Salmonella enterica CFSAN000228 | NZ_CP039498 | 4774876 | 4554 |
| Salmonella enterica CFSAN000661 | NZ_CP039496 | 4726555 | 4473 |
| Salmonella enterica CFSAN000669 | NZ_CP039493 | 4726500 | 4475 |
| Salmonella enterica CFSAN000700 | NZ_CP039491 | 4726555 | 4473 |
| Salmonella enterica CFSAN000752 | NZ_CP039489 | 4726556 | 4474 |
| Salmonella enterica CFSAN000753 | $NZ_{-}CP039487$ | 4726557 | 4473 |
| Salmonella enterica CFSAN000951 | NZ_CP039486 | 4726556 | 4472 |
| Salmonella enterica CFSAN000952 | NZ_CP039484 | 4726550 | 4473 |
| Salmonella enterica CFSAN000954 | NZ_CP039482 | 4726558 | 4475 |
| Salmonella enterica CFSAN000958 | $NZ_{-}CP039480$ | 4726543 | 4473 |
| Salmonella enterica CFSAN000960 | $NZ_{-}CP039478$ | 4726552 | 4473 |
| Salmonella enterica CFSAN000961 | $NZ_{-}CP039477$ | 4726558 | 4475 |
| Salmonella enterica CFSAN000963 | $NZ_{-}CP039476$ | 4726558 | 4473 |
| Salmonella enterica CFSAN000968 | $NZ_{-}CP039473$ | 4726548 | 4473 |
| Salmonella enterica CFSAN000970 | $NZ_{-}CP039471$ | 4726554 | 4473 |
| Salmonella enterica CFSAN001112 | $NZ_{-}CP039469$ | 4726545 | 4473 |
| Salmonella enterica CFSAN001115 | $NZ_{-}CP039467$ | 4726542 | 4472 |
| Salmonella enterica CFSAN001118 | $NZ_{-}CP039465$ | 4726540 | 4473 |
| Salmonella enterica CFSAN001140 | $NZ_{-}CP039463$ | 4726559 | 4473 |
| Salmonella enterica EC20090641 | $NZ_{-}CP007249$ | 4712309 | 4492 |
| Salmonella enterica EC20090698 | $NZ_{-}CP007248$ | 4680996 | 4454 |
| Salmonella enterica EC20100101 | $NZ_{-}CP007246$ | 4654156 | 4428 |
| Salmonella enterica EC20100325. | NZ_CP007360 | 4685849 | 4467 |
| Salmonella enterica EC20110221 | $NZ_{-}CP007247$ | 4709033 | 4489 |
| Salmonella enterica EC20110354 | NZ_CP007175 | 4711160 | 4491 |
| Salmonella enterica EC20110355 | $NZ_{-}CP007250$ | 4681839 | 4455 |
| Salmonella enterica EC20110356 | $NZ_{-}CP007262$ | 4682805 | 4454 |
| Salmonella enterica EC20110357 | NZ_CP007261 | 4678418 | 4453 |
| Salmonella enterica EC20110358 | $NZ_{-}CP007260$ | 4678687 | 4454 |
| Salmonella enterica EC20110359 | $NZ_{-}CP007259$ | 4711474 | 4492 |
| Salmonella enterica EC20110360 | $NZ_{-}CP007258$ | 4678618 | 4453 |
| Salmonella enterica EC20110361 | NZ_CP007263 | 4682911 | 4456 |
| Salmonella enterica EC20111095 | $NZ_{-}CP007254$ | 4679953 | 4452 |
| Salmonella enterica EC20111175 | $NZ_{-}CP007252$ | 4679808 | 4453 |
| Salmonella enterica EC20120002. | NZ_CP007329 | 4679418 | 4452 |
| Salmonella enterica EC20120008 | $NZ_{-}CP007245$ | 4711924 | 4492 |
| Salmonella enterica EC20120200 | $NZ_{-}CP007434$ | 4685848 | 4466 |
| Salmonella enterica EC20121175 | $NZ_{-}CP007269$ | 4679953 | 4452 |
| Salmonella enterica EC20121177 | NZ_CP007333 | 4685848 | 4466 |
| Salmonella enterica EC20121178 | $NZ_{-}CP007271$ | 4685848 | 4466 |
| Salmonella enterica EC20121179. | $NZ_{-}CP007272$ | 4685848 | 4467 |
| Salmonella enterica EC20121180. | $NZ_{-}CP007273$ | 4685848 | 4466 |
| Salmonella enterica SA19960848 | $NZ_CP007280$ | 4685848 | 4467 |
| Salmonella enterica SA19970769 | $NZ_{-}CP007283$ | 4685848 | 4466 |
| Salmonella enterica SA19980677 | $NZ_{-}CP007285$ | 4685849 | 4466 |
| Salmonella enterica SA19981522 | $NZ_{-}CP007286$ | 4685848 | 4467 |
| Salmonella enterica SA19981857 | $NZ_{-}CP007288$ | 4685848 | 4467 |
| Salmonella enterica SA19992322. | $NZ_{-}CP007291$ | 4685849 | 4467 |

continued table S1

| continued table S1 | | | |
|------------------------------------|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Salmonella enterica SA19994216 | NZ_CP007292 | 4685847 | 4466 |
| Salmonella enterica SA20082034 | $NZ_{-}CP007425$ | 4685659 | 4466 |
| Salmonella enterica SA20094352 | $NZ_{-}CP007312$ | 4685849 | 4467 |
| Salmonella enterica PM016/13 | NZ_CP012091 | 4793553 | 4669 |
| Salmonella enterica B/SF/13/03/195 | $NZ_{-}CP012151$ | 4798993 | 4682 |
| Salmonella enterica Ty21a | $NZ_{-}CP023975$ | 4791341 | 4874 |
| Salmonella enterica 311189-215186 | NZ_CP029930 | 4782608 | 4774 |
| Salmonella enterica 311189-213186 | NZ_CP029936 | 4783243 | 4777 |
| Salmonella enterica 311189-211186 | $NZ_{-}CP029945$ | 4786194 | 4782 |
| Salmonella enterica 311189-220186 | NZ_CP029886 | 4783528 | 4775 |
| Salmonella enterica 311189-208186 | NZ_CP029949 | 4783337 | 4775 |
| Salmonella enterica 311189-207186 | NZ_CP029938 | 4784615 | 4778 |
| Salmonella enterica 311189-205186 | $NZ_{-}CP029950$ | 4821321 | 4846 |
| Salmonella enterica 343077-228140 | NZ_CP029866 | 4790593 | 4799 |
| Salmonella enterica 343077-214162 | $NZ_{-}CP029862$ | 4814028 | 4816 |
| Salmonella enterica 343076-249107 | NZ_CP029913 | 4783891 | 4776 |
| Salmonella enterica 343076-232188 | NZ_CP029900 | 4800621 | 4803 |
| Salmonella enterica 311189-282186 | NZ_CP029920 | 4799956 | 4800 |
| Salmonella enterica 311189-268103 | $NZ_{-}CP029902$ | 4804086 | 4815 |
| Salmonella enterica 311189-232103 | NZ_CP029918 | 4778368 | 4765 |
| Salmonella enterica 311189-223186 | NZ_CP029880 | 4783762 | 4778 |
| Salmonella enterica 343078-228140 | $NZ_{-}CP029962$ | 4783156 | 4775 |
| Salmonella enterica 343077-267164 | NZ_CP029906 | 4775386 | 4775 |
| Salmonella enterica 311189-201186 | $NZ_{-}CP029958$ | 4783542 | 4778 |
| Salmonella enterica 311189-203186 | $NZ_{-}CP029942$ | 4786232 | 4783 |
| Salmonella enterica 311189-209186 | $NZ_CP029952$ | 4782252 | 4775 |
| Salmonella enterica 311189-219186 | NZ_CP029911 | 4784370 | 4779 |
| Salmonella enterica 311189-208103 | $NZ_{-}CP029932$ | 4783092 | 4776 |
| Salmonella enterica 343077-214135 | $NZ_{-}CP029872$ | 4783204 | 4774 |
| Salmonella enterica 343077-213147 | $NZ_{-}CP029897$ | 4897593 | 4940 |
| Salmonella enterica 343077-212138 | NZ_CP029919 | 4774558 | 4773 |
| Salmonella enterica 343077-211171 | $NZ_{-}CP029873$ | 4782825 | 4777 |
| Salmonella enterica 343076-294172 | $NZ_{-}CP029888$ | 4809171 | 4820 |
| Salmonella enterica 343076-253155 | $NZ_{-}CP029890$ | 4791587 | 4800 |
| Salmonella enterica 343076-248190 | $NZ_{-}CP029882$ | 4773823 | 4777 |
| Salmonella enterica 311189-291186 | $NZ_{-}CP029894$ | 4806999 | 4815 |
| Pseudomonas aeruginosa Pa127 | $NZ_{-}CP022000$ | 7148302 | 6783 |
| Pseudomonas aeruginosa Pa1207 | $NZ_{-}CP022001$ | 7411863 | 7080 |
| Pseudomonas aeruginosa PA83 | $NZ_{-}CP017293$ | 6816227 | 6385 |
| Pseudomonas aeruginosa L10 | $NZ_{-}CP019338$ | 6661962 | 6217 |
| Pseudomonas aeruginosa Ocean-1155 | $NZ_{-}CP022526$ | 6952237 | 6601 |
| Pseudomonas aeruginosa Ocean-1175 | $NZ_{-}CP022525$ | 6943220 | 6592 |
| Pseudomonas aeruginosa PA-150577 | $NZ_{-}CP017306$ | 6334472 | 5928 |
| Pseudomonas aeruginosa 12939 | $NZ_{-}CP024477$ | 6621378 | 6270 |
| Pseudomonas aeruginosa PB369 | $NZ_{-}CP025049$ | 6526814 | 6216 |
| Pseudomonas aeruginosa PB367 | $NZ_{-}CP025056$ | 6752906 | 6340 |
| Pseudomonas aeruginosa PB350 | $NZ_CP025055$ | 6752870 | 6340 |
| Pseudomonas aeruginosa PB368 | $NZ_CP025050$ | 6638559 | 6320 |
| Pseudomonas aeruginosa PB353 | $NZ_CP025051$ | 6437515 | 6075 |
| Pseudomonas aeruginosa PB354 | $NZ_CP025053$ | 6434352 | 6066 |
| Pseudomonas aeruginosa F5677 | $NZ_CP026680$ | 6645227 | 6301 |
| Pseudomonas aeruginosa AR-0360 | $NZ_CP027165$ | 6463575 | 6052 |

| continued table S1 Name | Accession Num. | Sequence Length | CDS numbe |
|------------------------------------|------------------|-----------------|-----------|
| Pseudomonas aeruginosa AR-0354 | NZ_CP027171 | 6747010 | 633 |
| Pseudomonas aeruginosa AR-0230 | $NZ_{-}CP027174$ | 7012922 | 660 |
| Pseudomonas aeruginosa AR-0357 | $NZ_{-}CP027166$ | 7162784 | 685 |
| Pseudomonas aeruginosa AR-0356 | $NZ_{-}CP027169$ | 6586916 | 622 |
| Pseudomonas aeruginosa AR-0353 | $NZ_{-}CP027172$ | 7240677 | 700 |
| Pseudomonas aeruginosa AR-0095 | $NZ_{-}CP027538$ | 6822666 | 639 |
| Pseudomonas aeruginosa CR1 | $NZ_{-}CP020560$ | 6118054 | 571 |
| Pseudomonas aeruginosa MRSN12280 | $NZ_{-}CP028162$ | 7050928 | 670 |
| Pseudomonas aeruginosa PPF-1 | NZ_CP023316 | 6930893 | 651 |
| Pseudomonas aeruginosa JB2 | $NZ_{-}CP028917$ | 6867314 | 650 |
| Pseudomonas aeruginosa AR445 | NZ_CP029088 | 7125975 | 679 |
| Pseudomonas aeruginosa AR444 | NZ_CP029089 | 6853499 | 642 |
| Pseudomonas aeruginosa AR441 | NZ_CP029093 | 6584825 | 620 |
| Pseudomonas aeruginosa AR442 | NZ_CP029090 | 7267567 | 702 |
| Pseudomonas aeruginosa AR439 | $NZ_{-}CP029097$ | 7139518 | 675 |
| Pseudomonas aeruginosa CCUG 70744 | $NZ_{-}CP023255$ | 6859232 | 653 |
| Pseudomonas aeruginosa AR-0446 | NZ_CP029660 | 6475581 | 612 |
| Pseudomonas aeruginosa AR-0110 | $NZ_{-}CP029745$ | 6799785 | 643 |
| Pseudomonas aeruginosa K34-7 | $NZ_{-}CP029707$ | 7038012 | 670 |
| Pseudomonas aeruginosa AR-460 | NZ_CP030351 | 6303875 | 585 |
| Pseudomonas aeruginosa AR-458 | $NZ_{-}CP030327$ | 6685102 | 628 |
| Pseudomonas aeruginosa AR-455 | $NZ_{-}CP030328$ | 6540996 | 626 |
| Pseudomonas aeruginosa HS9 | NZ_CP030861 | 6876988 | 653 |
| Pseudomonas aeruginosa Y89 | NZ_CP030913 | 6868832 | 649 |
| Pseudomonas aeruginosa Y82 | NZ_CP030912 | 7106857 | 679 |
| Pseudomonas aeruginosa Y31 | NZ_CP030910 | 6831076 | 641 |
| Pseudomonas aeruginosa Y71 | NZ_CP030911 | 6940949 | 660 |
| Pseudomonas aeruginosa PABL017 | $NZ_{-}CP031660$ | 6503460 | 609 |
| Pseudomonas aeruginosa PABL012 | $NZ_{-}CP031659$ | 6546467 | 616 |
| Pseudomonas aeruginosa 24Pae112 | $NZ_{-}CP029605$ | 7097241 | 67' |
| Pseudomonas aeruginosa N15-01092 | NZ_CP012901 | 6972899 | 661 |
| Pseudomonas aeruginosa AR-0111 | $NZ_{-}CP032257$ | 6946231 | 657 |
| Pseudomonas aeruginosa WCHPA075019 | $NZ_{-}CP028584$ | 6886080 | 640 |
| Pseudomonas aeruginosa 268 | $NZ_{-}CP032761$ | 7030474 | 669 |
| Salmonella enterica 311189-269186 | $NZ_{-}CP029922$ | 4782083 | 477 |
| Salmonella enterica 311189-256186 | $NZ_{-}CP029917$ | 4800376 | 480 |
| Salmonella enterica 311189-231186 | $NZ_{-}CP029904$ | 4807430 | 483 |
| Salmonella enterica 343078-203125 | $NZ_{-}CP029850$ | 4801819 | 480 |
| Salmonella enterica 343078-201101 | $NZ_{-}CP029852$ | 4782235 | 477 |
| Salmonella enterica 311189-221186 | $NZ_{-}CP029923$ | 4782756 | 47 |
| Salmonella enterica 311189-217103 | $NZ_{-}CP029927$ | 4778473 | 477 |
| Salmonella enterica 311189-214186 | $NZ_{-}CP029933$ | 4782854 | 477 |
| Salmonella enterica 311189-212186 | $NZ_{-}CP029944$ | 4784815 | 47 |
| Salmonella enterica 311189-204186 | $NZ_{-}CP029954$ | 4784767 | 478 |
| Salmonella enterica 311189-202186 | $NZ_{-}CP029956$ | 4799946 | 480 |
| Salmonella enterica 343077-260153 | $NZ_{-}CP029861$ | 4783691 | 477 |
| Salmonella enterica 343077-215174 | $NZ_CP029868$ | 4785665 | 477 |
| Salmonella enterica 343076-269157 | $NZ_CP029881$ | 4791108 | 479 |
| Salmonella enterica 343076-227128 | $NZ_CP029875$ | 4800199 | 480 |
| Salmonella enterica 343076-217103 | $NZ_{-}CP029914$ | 4775784 | 477 |
| Salmonella enterica 311189-268186 | $NZ_CP029883$ | 4807681 | 481 |
| Salmonella enterica 311189-255186 | $NZ_{-}CP029885$ | 4808675 | 485 |

continued table S1

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|---------------------------------------|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Salmonella enterica 311189-222186 | NZ_CP029909 | 4842715 | 4867 |
| Salmonella enterica 343078-251131 | $NZ_CP029960$ | 4799747 | 4804 |
| Salmonella enterica 343078-211126 | $NZ_CP029848$ | 4799748 | 4803 |
| Salmonella enterica 343077-292148 | $NZ_CP029855$ | 4800586 | 4802 |
| Salmonella enterica 343077-286126 | $NZ_CP029856$ | 4797628 | 4804 |
| Salmonella enterica 311189-218186 | $NZ_{-}CP029925$ | 4786711 | 4783 |
| Salmonella enterica 311189-216103 | $NZ_{-}CP029928$ | 4799426 | 4803 |
| Salmonella enterica 311189-210186 | $NZ_{-}CP029946$ | 4785705 | 4784 |
| Salmonella enterica 343077-228157 | $NZ_CP029864$ | 4807408 | 4818 |
| Salmonella enterica 343076-252143 | $NZ_{-}CP029892$ | 4808285 | 4817 |
| Salmonella enterica 343076-241106 | $NZ_CP029899$ | 4775702 | 4775 |
| Salmonella enterica 343076-202113 | $NZ_CP029915$ | 4799269 | 4801 |
| Salmonella enterica 311189-252186 | $NZ_{-}CP029896$ | 4781801 | 4773 |
| Salmonella enterica 311189-224186 | $NZ_CP029878$ | 4782665 | 4776 |
| Salmonella enterica 343078-273110 | $NZ_CP029846$ | 4801146 | 4804 |
| Salmonella enterica 343078-223175 | $NZ_CP029964$ | 4782707 | 4780 |
| Salmonella enterica 343077-285138 | $NZ_CP029858$ | 4800928 | 4805 |
| Salmonella enterica 343077-278127 | $NZ_CP029863$ | 4807245 | 4816 |
| Salmonella enterica 311189-217186 | $NZ_CP029646$ | 4782348 | 4781 |
| Salmonella enterica 311189-206186 | $NZ_{-}CP029940$ | 4786487 | 4782 |
| Salmonella enterica 343077-255118 | $NZ_{-}CP029907$ | 4774500 | 4776 |
| Salmonella enterica 311189-239103 | NZ_CP029908 | 4778864 | 4768 |
| Salmonella enterica LXYSH | NZ_CP030936 | 4780894 | 4772 |
| Salmonella enterica 343077-212159 | $NZ_CP029870$ | 4795465 | 4804 |
| Salmonella enterica 343078-256191 | $NZ_CP029959$ | 4795890 | 4803 |
| Salmonella enterica 343077-281186 | $NZ_CP029853$ | 4829489 | 4852 |
| Salmonella enterica WGS1146 | $NZ_CP040575$ | 4813117 | 4712 |
| Salmonella enterica CT18 | $NC_{-}003198$ | 4809037 | 4110 |
| Salmonella enterica P-stx-12 | $NC_{-}016832$ | 4768352 | 4640 |
| Salmonella enterica Ty2 | NC_004631 | 4791961 | 4858 |
| Salmonella enterica Ty21a | $NC_{-}021176$ | 4791958 | 4666 |
| Salmonella entericamurium ATCC 13311 | $NZ_{-}CP009102$ | 4793299 | 4551 |
| Pseudomonas aeruginosa PA34 | $NZ_{-}CP032552$ | 6810079 | 6462 |
| Pseudomonas aeruginosa SP4528 | $NZ_{-}CP033439$ | 6877287 | 6421 |
| Pseudomonas aeruginosa FDAARGOS-501 | NZ_CP033843 | 6865838 | 6494 |
| Pseudomonas aeruginosa FDAARGOS-505 | $NZ_{-}CP033832$ | 7029824 | 6620 |
| Pseudomonas aeruginosa H25883 | $NZ_{-}CP033686$ | 6706800 | 6337 |
| Pseudomonas aeruginosa FDAARGOS-570 | $NZ_{-}CP033835$ | 7119105 | 6855 |
| Pseudomonas aeruginosa H26023 | $NZ_CP033685$ | 6729216 | 6347 |
| Pseudomonas aeruginosa FDAARGOS-532 | $NZ_{-}CP033771$ | 6928914 | 6486 |
| Pseudomonas aeruginosa FDAARGOS-571 | $NZ_{-}CP033833$ | 6999770 | 6660 |
| Pseudomonas aeruginosa H26027 | $NZ_CP033684$ | 7079598 | 6747 |
| Pseudomonas aeruginosa IMP-13 | $NZ_CP034354$ | 7047704 | 6630 |
| Pseudomonas aeruginosa B17932 | NZ_CP034436 | 6744658 | 6341 |
| Pseudomonas aeruginosa GIMC5015:PAKB6 | NZ_CP034429 | 6258491 | 5824 |
| Pseudomonas aeruginosa SP4527 | NZ_CP034409 | 7005215 | 6567 |
| Pseudomonas aeruginosa LW | NZ_CP022478 | 6824837 | 6408 |
| Pseudomonas aeruginosa PAO1161 | NZ_CP032126 | 6383803 | 5962 |
| Pseudomonas aeruginosa SP4371 | NZ_CP034369 | 6937609 | 6520 |
| Pseudomonas aeruginosa E80 | NZ_CP031677 | 6819140 | 6462 |
| Pseudomonas aeruginosa AES1M | NZ_CP037925 | 6373139 | 6003 |
| Pseudomonas aeruginosa AES1R | $NZ_CP037926$ | 6373893 | 6015 |

| Name | Accession Num. | Sequence Length | CDS number |
|--------------------------------------|------------------------|-----------------|------------|
| Pseudomonas aeruginosa B41226 | NZ_CP034368 | 6814295 | 641 |
| Pseudomonas aeruginosa BA15561 | $NZ_{-}CP033432$ | 6793961 | 635 |
| Pseudomonas aeruginosa B14130 | $NZ_{-}CP034435$ | 6759594 | 634 |
| Pseudomonas aeruginosa SP2230 | $NZ_{-}CP034434$ | 6976603 | 656 |
| Pseudomonas aeruginosa PABL048 | $NZ_{-}CP039293$ | 6879622 | 652 |
| Pseudomonas aeruginosa PA298 | $NZ_{-}CP040127$ | 6539894 | 614 |
| Pseudomonas aeruginosa 97 | NZ_CP031449 | 6925889 | 661 |
| Pseudomonas aeruginosa FDAARGOS-610 | NZ_CP041013 | 6792215 | 649 |
| Pseudomonas aeruginosa FDAARGOS-767 | NZ_CP041008 | 6263676 | 583 |
| Pseudomonas aeruginosa BA7823 | $NZ_{-}CP032569$ | 6876113 | 648 |
| Pseudomonas aeruginosa AZPAE15042 | NZ_CP041354 | 6527298 | 616 |
| Pseudomonas aeruginosa 243931 | $NZ_{-}CP041772$ | 6514208 | 608 |
| Pseudomonas aeruginosa A681 | NZ_CP041771 | 6616247 | 627 |
| Pseudomonas aeruginosa 519119 | NZ_CP041773 | 6714255 | 629 |
| Pseudomonas aeruginosa 60503 | NZ_CP041774 | 6809062 | 640 |
| Pseudomonas aeruginosa C79 | NZ_CP040684 | 6756154 | 638 |
| Pseudomonas aeruginosa IMP68 | NZ_CP028849 | 6481697 | 608 |
| Pseudomonas aeruginosa IMP67 | NZ_CP028848 | 6501414 | 61 |
| Pseudomonas aeruginosa IMP66 | NZ_CP028959 | 6486336 | 609 |
| Pseudomonas aeruginosa CCUG 51971 | NZ_CP043328 | 7012798 | 66 |
| Pseudomonas aeruginosa HOU1 | NZ_CP042269 | 6123373 | 57 |
| Pseudomonas aeruginosa E90 | NZ_CP044006 | 6813135 | 64 |
| Pseudomonas aeruginosa B136-33 | NC_020912 | 6421010 | 60 |
| Pseudomonas aeruginosa DHS01 | NZ_CP013993 | 7055752 | 68: |
| Pseudomonas aeruginosa DK2 | NC_018080 | 6402658 | 60 |
| Pseudomonas aeruginosa DSM 50071 | NZ_CP012001 | 6317050 | 59 |
| Pseudomonas aeruginosa LES431 | NC_023066 | 6550070 | 61- |
| Pseudomonas aeruginosa LESB58 | NC_023000 NC_011770 | 6601757 | 62 |
| Pseudomonas aeruginosa M18 | NC_011770 NC_017548 | 6327754 | 59 |
| | | | |
| Pseudomonas aeruginosa MTB-1 | NC_023019 | 6580038 | 62 |
| Staphylococcus aureus ST228 18341. | NC_020536 | 2759473 | 26 |
| Staphylococcus aureus ST228 18412. | NC_020537 | 2759263 | 26 |
| Staphylococcus aureus ST228 10497. | NC_020564 | 2759512 | 27 |
| Staphylococcus aureus ST228 16125. | NC_020566 | 2759457 | 26 |
| Staphylococcus aureus ST228 18583. | NC_020568 | 2759328 | 26 |
| Staphylococcus aureus ST398 | NC_017333 | 2872582 | 28 |
| Staphylococcus aureus JKD6008 | NC_017341 | 2924344 | 29 |
| Staphylococcus aureus Newman | NC_009641 | 2878897 | 29 |
| Staphylococcus aureus T0131 | NC_017347 | 2913900 | 29 |
| Staphylococcus aureus TCH60 | $NC_{-}017342$ | 2802675 | 27 |
| Staphylococcus aureus TW20 | $NC_{-}017331$ | 3043210 | 30 |
| Staphylococcus aureus USA300-FPR3757 | $NC_{-}007793$ | 2872769 | 28 |
| Staphylococcus aureus USA300-TCH1516 | $NC_{-}010079$ | 2872915 | 28 |
| Staphylococcus aureus Z172 | $NC_{-}022604$ | 2987966 | 30 |
| Streptococcus pyogenes 1E1 | $NZ_{-}CP007241$ | 1796152 | 17 |
| Streptococcus pyogenes ATCC 19615 | $NZ_{-}CP008926$ | 1844804 | 18 |
| Streptococcus pyogenes M23ND | $NZ_{-}CP008695$ | 1846477 | 18- |
| Streptococcus pyogenes HKU360 | $NZ_{-}CP009612$ | 1944537 | 18 |
| Streptococcus pyogenes 7F7 | $NZ_{-}CP007240$ | 1709790 | 16 |
| Streptococcus pyogenes AP1 | $NZ_{-}CP007537$ | 1908294 | 19 |
| Streptococcus pyogenes D471 | NZ_CP011415 | 1811968 | 17 |
| Streptococcus pyogenes NGAS743 | NZ_CP007560 | 1915554 | 18 |

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|--|------------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Streptococcus pyogenes NGAS596 | NZ_CP007561 | 1791306 | 1711 |
| Streptococcus pyogenes NGAS327 | $NZ_CP007562$ | 1702054 | 1611 |
| Streptococcus pyogenes 5448 | NZ_CP008776 | 1829516 | 1800 |
| Streptococcus pyogenes STAB10015 | NZ_CP011068 | 1950454 | 1902 |
| Streptococcus pyogenes HKU488 | $NZ_CP012045$ | 1943415 | 1924 |
| Streptococcus pyogenes NGAS638 | NZ_CP010450 | 1791401 | 1740 |
| Streptococcus pyogenes NGAS322 | NZ_CP010449 | 1950469 | 1898 |
| Streptococcus pyogenes MEW123 | NZ_CP014139 | 1878699 | 1794 |
| Streptococcus pyogenes AP53 | $NZ_{-}CP013672$ | 1860554 | 1848 |
| Streptococcus pyogenes MGAS23530 | NZ_CP013839 | 1709394 | 1623 |
| Streptococcus pyogenes MGAS27061 | NZ_CP013840 | 1741348 | 1660 |
| Streptococcus pyogenes MGAS11027 | $NZ_{-}CP013838$ | 1786874 | 1718 |
| Streptococcus pyogenes STAB09014 | NZ_CP011069 | 1862487 | 1801 |
| Streptococcus pyogenes M28PF1 | NZ_CP011535 | 1896976 | 1852 |
| Streptococcus pyogenes NS53 | NZ_CP015238 | 1765123 | 1696 |
| Streptococcus pyogenes STAB14018 | NZ_CP014542 | 1890465 | 1877 |
| Streptococcus pyogenes GURSA1 | $NZ_{-}CP022206$ | 1893171 | 1883 |
| Streptococcus pyogenes GUR | $NZ_{-}CP022354$ | 1890204 | 1870 |
| Streptococcus pyogenes STAB090229 | NZ_CP020027 | 1846347 | 1811 |
| Streptococcus pyogenes STAB120304 | NZ_CP020082 | 1890354 | 1875 |
| Streptococcus pyogenes HarveyGAS | $NZ_{-}CP023769$ | 1852980 | 1788 |
| Streptococcus pyogenes JS12 | NZ_CP021640 | 1810283 | 1722 |
| Streptococcus pyogenes FDAARGOS-149 | NZ_CP014027 | 1839644 | 1801 |
| Streptococcus pyogenes TJ11-001 | NZ_CP028148 | 1939844 | 1899 |
| Streptococcus pyogenes NGAS979 | NZ_CP028140 | 1790938 | 1725 |
| Streptococcus pyogenes DMG1800716 | NZ_CP027771 | 1869673 | 1794 |
| Streptococcus pyogenes SASM4-Duke | NZ_CP031770 | 1903997 | 1875 |
| Streptococcus pyogenes ABC020055975 | NZ_CP029694 | 1866513 | 1859 |
| Salmonella entericamurium CDC 2011K-0870 | NZ_CP007523 | 4769471 | 4511 |
| Salmonella entericamurium CDC 2011K-1702 | NZ_CP014967 | 4906321 | 4702 |
| Salmonella entericamurium CDC H2662 | NZ_CP014979 | 4884395 | 4695 |
| Salmonella entericamurium D23580 | NC_016854 | 4879400 | 4671 |
| Salmonella entericamurium DT104 main | NC_022569 | 4933631 | 4949 |
| Salmonella entericamurium DT2 | NC_022544 | 4814801 | 4582 |
| Salmonella entericamurium LT2 | NC_003197 | 4857450 | 4446 |
| Salmonella entericamurium SARA13 | NZ_CP017728 | 4819807 | 4770 |
| Salmonella entericamurium SL1344 | NC_016810 | 4878012 | 4886 |
| Salmonella entericamurium ST4/74 | NC_016857 | 4878013 | 4660 |
| Salmonella entericamurium T000240 | NC_016860 | 4954814 | 4770 |
| Salmonella entericamurium U288 | NC_021151 | 4852606 | 4631 |
| Salmonella entericamurium UK-1 | NC_016863 | 4817868 | 4576 |
| Salmonella entericamurium USDA-1808 | NZ_CP014969 | 4936894 | 4977 |
| Salmonella entericamurium USDA-1810 | NZ_CP014982 | 4927145 | 4734 |
| Salmonella entericamurium USDA-1880 | NZ_CP014981 | 4815208 | 4793 |
| Salmonella entericamurium USDA-1896 | NZ_CP014977 | 4856440 | 4880 |
| Salmonella entericamurium USDA-1898 | NZ_CP014971 | 4784385 | 4715 |
| Salmonella entericamurium USDA-1899 | NZ_CP007235 | 4856438 | 4824 |
| Staphylococcus aureus | NC_021670 | 2980548 | 3004 |
| Staphylococcus aureus 04-02981 | NC_021070 NC_017340 | 2821452 | 2780 |
| Staphylococcus aureus 08BA02176 | NC_017540 NC_018608 | 2782313 | 2704 |
| Staphylococcus aureus CA-347 | NC_018008 NC_021554 | 2850503 | 2819 |
| Staphylococcus aureus RF122 | NC_021534 NC_007622 | 2742531 | 2714 |
| Diaphyrococcus aureus fui 122 | 110_001022 | 2142001 | 2114 |

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|--|----------------------------|-----------------|----------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Staphylococcus aureus 11819-97 | $NC_{-}017351$ | 2846546 | 2825 |
| Staphylococcus aureus 55/2053 | $NC_{-}022113$ | 2756919 | 2711 |
| Staphylococcus aureus 6850 | $NC_{-}022222$ | 2736560 | 2662 |
| Staphylococcus aureus CN1 | $NC_{-}022226$ | 2751266 | 2699 |
| Staphylococcus aureus COL | $NC_{-}002951$ | 2809422 | 2784 |
| Staphylococcus aureus ECT-R 2 | $NC_{-}017343$ | 2729540 | 2668 |
| Staphylococcus aureus ED133 | $NC_{-}017337$ | 2832478 | 2808 |
| Staphylococcus aureus ED98 | $NC_{-}013450$ | 2824404 | 2793 |
| Staphylococcus aureus HO 5096 0412 | $NC_{-}017763$ | 2832299 | 2781 |
| Staphylococcus aureus JH1 | $NC_{-}009632$ | 2906507 | 2906 |
| Staphylococcus aureus JH9 | NC_009487 | 2906700 | 2906 |
| Staphylococcus aureus JKD6159 | $NC_{-}017338$ | 2811435 | 2753 |
| Staphylococcus aureus LGA251 | $NC_{-}017349$ | 2750834 | 2654 |
| Staphylococcus aureus M013 | $NC_{-}016928$ | 2795015 | 2715 |
| Staphylococcus aureus MRSA252 | NC_002952 | 2902619 | 2937 |
| Staphylococcus aureus MSSA476 | NC_002953 | 2799802 | 2827 |
| Staphylococcus aureus Mu3 | NC_009782 | 2880168 | 2863 |
| Staphylococcus aureus Mu50 | NC_002758 | 2878529 | 2862 |
| Staphylococcus aureus MW2 | NC_003923 | 2820462 | 2771 |
| Staphylococcus aureus N315 | NC_002745 | 2814816 | 2760 |
| Staphylococcus aureus NCTC 8325 | NC_007795 | 2821361 | 2767 |
| Staphylococcus aureus SA40 | NC_022443 | 2728308 | 2631 |
| Staphylococcus aureus SA40 Staphylococcus aureus SA957 | NC_022443 NC_022442 | 2789538 | 2716 |
| Staphylococcus aureus ST228 10388. | NC_020529 | 2759510 | 2700 |
| Staphylococcus aureus ST228 10588. Staphylococcus aureus ST228 15532. | NC_020529 NC_020532 | 2759883 | 2700 |
| Staphylococcus aureus ST228 16035. | NC_020532 NC_020533 | 2759835 | 2701 |
| Salmonella entericamurium VNP20009 | NZ_CP007804 | 4763557 | 4547 |
| Salmonella entericamurium FORC-015 | NZ_CP011365 | 4759372 | 4547 |
| Salmonella entericamurium 33676 | NZ_CP012681 | 4809574 | 4581 |
| Salmonella entericamurium FORC-020 | | 4799793 | |
| Salmonella entericamurium FORC-020 Salmonella entericamurium SO2 | NZ_CP012144 | | 4584 4716 |
| | NZ_CP014356 NZ_CP014536 | 4916170 | |
| Salmonella entericamurium SO3 Salmonella entericamurium NC983 | | 4912959 | $4708 \\ 4625$ |
| | NZ_CP015157 | 4846304 | |
| Salmonella entericamurium 13-931 | NZ_CP016385 | 4852920 | 4625 |
| Salmonella entericamurium 22792 | NZ_CP017621 | 4806964 | 4586 |
| Salmonella entericamurium RM10961 | NZ_CP013702 | 4856460 | 4651 |
| Salmonella entericamurium 81741 | NZ_CP019442 | 4974856 | 4801 |
| Salmonella entericamurium 22495 | NZ_CP017617 | 4817416 | 4724 |
| Salmonella entericamurium TW-Stm6 | NZ_CP019649 | 4999862 | 5041 |
| Salmonella entericamurium WW012 | NZ_CP022168 | 4991167 | 5015 |
| Salmonella entericamurium BL10 | NZ_CP024619 | 4835517 | 4613 |
| Salmonella entericamurium FORC50 | NZ_CP019383 | 4673894 | 4651 |
| Salmonella entericamurium AR-0031 | NZ_CP026700 | 4870262 | 4878 |
| Salmonella entericamurium CFSAN018746 | NZ_CP028199 | 4857558 | 4849 |
| Salmonella entericamurium PIR00538 | $NZ_{-}CP025555$ | 4870262 | 4882 |
| Salmonella entericamurium FORC58 | $NZ_{-}CP020565$ | 4698556 | 4653 |
| Salmonella entericamurium FDAARGOS-321 | $NZ_{-}CP022070$ | 4884508 | 4679 |
| Salmonella entericamurium UGA14 | $NZ_{-}CP021462$ | 4879285 | 4915 |
| Salmonella entericamurium ATCC 14028 | $NZ_{-}CP034230$ | 4869644 | 4761 |
| Salmonella entericamurium RSE04 | $NZ_{-}CP034719$ | 4766197 | 4636 |
| Salmonella entericamurium B3589 | $NZ_{-}CP034968$ | 4913154 | 4832 |
| Salmonella entericamurium sg-wt7 | NZ_CP036168 | 4930312 | 4821 |

| continued table S1 | | | |
|--|----------------------------|--------------------|--------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Salmonella entericamurium 08-365-0802 | NZ_CP038847 | 4854979 | 4663 |
| Salmonella entericamurium 16425992 | $NZ_CP039576$ | 4931770 | 4771 |
| Salmonella entericamurium 11A0003616 | NZ_CP039591 | 4942800 | 4785 |
| Salmonella entericamurium FORC88 | $NZ_{-}CP029029$ | 4634278 | 4395 |
| Salmonella entericamurium E2840153550 | $NZ_{-}CP040321$ | 4923481 | 4746 |
| Salmonella entericamurium TJWQ005 | NZ_CP040458 | 4982919 | 4821 |
| Salmonella entericamurium SAP17-8290 | $NZ_{-}CP040568$ | 4965236 | 4796 |
| Salmonella entericamurium SAP17-7399 | $NZ_{-}CP040562$ | 4902004 | 4707 |
| Salmonella entericamurium SAP17-7699 | $NZ_{-}CP040564$ | 4883741 | 4671 |
| Salmonella entericamurium SAP17-7299 | $NZ_{-}CP040566$ | 4857789 | 4651 |
| Salmonella entericamurium 14028 | $NZ_{-}CP034479$ | 4870261 | 4647 |
| Salmonella entericamurium SAP18-6199 | NZ_CP040900 | 4813658 | 4606 |
| Salmonella entericamurium 01ST04081 | $NZ_{-}CP029840$ | 4870227 | 4648 |
| Salmonella entericamurium 10ST07093 | $NZ_{-}CP029839$ | 4865212 | 4639 |
| Salmonella entericamurium FORC098 | $NZ_{-}CP030029$ | 4855239 | 4629 |
| Salmonella entericamurium 14028S | $NC_{-}016856$ | 4870265 | 4649 |
| Salmonella entericamurium 14028S subGXS259 | $NZ_{-}CP043401$ | 4869261 | 4648 |
| Salmonella entericamurium 14028S subGXS254 | $NZ_{-}CP043402$ | 4869252 | 4651 |
| Salmonella entericamurium 14028S subJY996 | NZ_CP043400 | 4870263 | 4646 |
| Salmonella entericamurium 14028S subGXS275 | $NZ_{-}CP043399$ | 4870694 | 4649 |
| Salmonella entericamurium 798 | $NC_{-}017046$ | 4876219 | 4662 |
| Salmonella entericamurium CDC 2009K-1640 | $NZ_{-}CP014975$ | 4933707 | 4756 |
| Salmonella entericamurium CDC 2009K-2059 | NZ_CP014983 | 4823793 | 4596 |
| Salmonella entericamurium CDC 2010K-1587 | $NZ_{-}CP014965$ | 4799398 | 4567 |
| Streptococcus pyogenes FDAARGOS-534 | $NZ_{-}CP033767$ | 1837737 | 1786 |
| Streptococcus pyogenes Duke-Large | $NZ_{-}CP033907$ | 1903852 | 1874 |
| Streptococcus pyogenes RLGH | NZ_CP033908 | 1903972 | 1875 |
| Streptococcus pyogenes MGAS28085 | NZ_CP032666 | 1851471 | 1783 |
| Streptococcus pyogenes MGAS27961 | NZ_CP032665 | 1853912 | 1790 |
| Streptococcus pyogenes CCUG 4207 | NZ_CP028841 | 1914862 | 1920 |
| Streptococcus pyogenes TSPY453 | NZ_CP033337 | 1976011 | 1930 |
| Streptococcus pyogenes TSPY165 | NZ_CP033336 | 1812036 | 1761 |
| Streptococcus pyogenes TSPY556 | NZ_CP032700 | 1828132 | 1755 |
| Streptococcus pyogenes TSPY208 | NZ_CP033335 | 1801696 | 1732 |
| Streptococcus pyogenes M75 | NZ_CP033621 | 1852894 | 1807 |
| Streptococcus pyogenes MGAS29409 | NZ_CP031617 | 1894040 | 1840 |
| Streptococcus pyogenes MGAS28669 | NZ_CP031623 | 1905059 | 1857 |
| Streptococcus pyogenes MGAS28278 | NZ_CP031629 | 1893446 | 1832 |
| Streptococcus pyogenes MGAS29064 | NZ_CP031620 | 1839848 | 1761 |
| Streptococcus pyogenes SP1336 | NZ_CP031738 | 1878827 | 1842 |
| Streptococcus pyogenes MGAS8347 | NZ_CP031638 | 1833822 | 1742 |
| Streptococcus pyogenes MGAS29326 | NZ_CP031618 | 1940684 | 1884 |
| Streptococcus pyogenes MGAS28650 | NZ_CP031624 | 1851876 | 1784 |
| Streptococcus pyogenes MGAS28386 | NZ_CP031626 | 1837594 | 1759 |
| Streptococcus pyogenes MGAS10786 | NZ_CP031637 | 1937069 | 1895 |
| Streptococcus pyogenes MGAS29284 | NZ_CP031619 | 1946265 | 1919 |
| Streptococcus pyogenes MGAS28533 | NZ_CP031625 | 1893263 | 1851 |
| Streptococcus pyogenes MGAS28191 Streptococcus pyogenes MGAS11115 | NZ_CP031631 NZ_CP031633 | 1892473 1851403 | 1832 1788 |
| Streptococcus pyogenes MGAS11115 Streptococcus pyogenes MGAS28078 | NZ_CP031633 NZ_CP031632 | 1880780 | 1812 |
| Streptococcus pyogenes MGAS28078 Streptococcus pyogenes MGAS11052 | NZ_CP031632 NZ_CP031635 | 1907750 | 1812 1833 |
| Streptococcus pyogenes MGAS11052 Streptococcus pyogenes MGAS7914 | NZ_CP031639 NZ_CP031639 | 1885126 | 1832 |
| pareproceeds pyogenes mGA51314 | 1477CL 091098 | 1000120 | 1002 |

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|--|------------------|-----------------|----------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Streptococcus pyogenes MGAS7888 | NZ_CP031640 | 1962429 | 1898 |
| Streptococcus pyogenes MGAS28746 | $NZ_{-}CP031621$ | 1853620 | 1785 |
| Streptococcus pyogenes MGAS28686 | $NZ_{-}CP031622$ | 1894087 | 1851 |
| Streptococcus pyogenes MGAS28360 | $NZ_{-}CP031627$ | 1893041 | 1833 |
| Streptococcus pyogenes MGAS28330 | $NZ_{-}CP031628$ | 1851619 | 1776 |
| Streptococcus pyogenes MGAS11108 | $NZ_{-}CP031634$ | 1853812 | 1783 |
| Streptococcus pyogenes MGAS10826 | NZ_CP031636 | 1852263 | 1783 |
| Streptococcus pyogenes emm78.3 | $NZ_{-}CP035437$ | 1806521 | 1724 |
| Streptococcus pyogenes emm97.1 | $NZ_{-}CP035447$ | 1812090 | 1744 |
| Streptococcus pyogenes emm22.8 | NZ_CP035438 | 1950616 | 1920 |
| Streptococcus pyogenes emm123 | $NZ_{-}CP035452$ | 1768816 | 1662 |
| Streptococcus pyogenes emm75.1 | NZ_CP035434 | 1887700 | 1815 |
| Streptococcus pyogenes emm90.5 | NZ_CP035444 | 1764969 | 1667 |
| Streptococcus pyogenes emm58 | NZ_CP035443 | 1888806 | 1846 |
| Streptococcus pyogenes emm11 | NZ_CP035432 | 1809631 | 1738 |
| Streptococcus pyogenes emm25 | NZ_CP035442 | 1835714 | 1786 |
| Streptococcus pyogenes emm197 | NZ_CP035455 | 1797172 | 1764 |
| Streptococcus pyogenes emm55 | NZ_CP035430 | 1899479 | 1849 |
| Streptococcus pyogenes emm77 | NZ_CP035439 | 1806314 | 1740 |
| Streptococcus pyogenes TSPY155 | NZ_CP032699 | 1882327 | 1817 |
| Streptococcus pyogenes emm230 | NZ_CP035451 | 1826832 | 1775 |
| Streptococcus pyogenes emm1 | NZ_CP035441 | 1834762 | 1801 |
| Streptococcus pyogenes emini Streptococcus pyogenes MGAS28271 | NZ_CP031630 | 1842830 | 1771 |
| Streptococcus pyogenes mGAS26277 Streptococcus pyogenes emm93.4 | NZ_CP035450 | 1701466 | 1613 |
| Streptococcus pyogenes emmiso.4 Streptococcus pyogenes emm89.14 | NZ_CP035436 | 1806344 | 1746 |
| Streptococcus pyogenes emm100 Streptococcus pyogenes emm100 | NZ_CP035453 | 1839531 | 1800 |
| Streptococcus pyogenes emm100 Streptococcus pyogenes emm92 | NZ_CP035445 | 1758778 | 1683 |
| Streptococcus pyogenes emm52 Streptococcus pyogenes emm54 | NZ_CP035454 | 1742565 | 1664 |
| Streptococcus pyogenes emm64.3 | NZ_CP035435 | 1779766 | 1743 |
| Streptococcus pyogenes emmSTG866.1 | NZ_CP035428 | 1816007 | 1745 |
| | NZ_CP035431 | 1800116 | 1736 1716 |
| Streptococcus pyogenes emm105 | | | |
| Streptococcus pyogenes emm124 | NZ_CP035440 | 1790668 | $1706 \\ 1827$ |
| Streptococcus pyogenes emm74 | NZ_CP035427 | 1861037 | |
| Streptococcus pyogenes emm57 | NZ_CP035426 | 1756622 | 1676 |
| Streptococcus pyogenes emm56 | NZ_CP035449 | 1825427 | 1785 |
| Streptococcus pyogenes emm65 | NZ_CP035433 | 1789855 | 1709 |
| Streptococcus pyogenes emmNA | NZ_CP035429 | 1788166 | 1749 |
| Streptococcus pyogenes emm68.2 | NZ_CP035446 | 1747924 | 1659 |
| Streptococcus pyogenes FDAARGOS-774 | NZ_CP040997 | 1914862 | 1919 |
| Streptococcus pyogenes FDAARGOS-514 | NZ_CP033815 | 1899746 | 1903 |
| Streptococcus pyogenes FDAARGOS-668 | NZ_CP044093 | 1837758 | 1787 |
| Streptococcus pyogenes A20 | NC_018936 | 1837281 | 1805 |
| Streptococcus pyogenes Alab49 | NC_017596 | 1827308 | 1799 |
| Streptococcus pyogenes HSC5 | $NC_{-}021807$ | 1818351 | 1795 |
| Streptococcus pyogenes JRS4 | NZ_CP011414 | 1811968 | 1763 |
| Streptococcus pyogenes M1 476 | $NC_{-}020540$ | 1831128 | 1803 |
| Streptococcus pyogenes M1 GAS | $NC_{-}002737$ | 1852433 | 1693 |
| Streptococcus pyogenes MGAS10270 | $NC_{-}008022$ | 1928252 | 1899 |
| Streptococcus pyogenes MGAS10394 | $NC_{-}006086$ | 1899877 | 1859 |
| Streptococcus pyogenes MGAS10750 | $NC_{-}008024$ | 1937111 | 1901 |
| Streptococcus pyogenes MGAS15252 | $NC_{-}017040$ | 1750832 | 1672 |
| Streptococcus pyogenes MGAS1882 | NC_017053 | 1781029 | 1711 |

| Name | Accession Num. | Sequence Length | CDS number |
|---------------------------------|----------------|-----------------|------------|
| Streptococcus pyogenes MGAS315 | NC_004070 | 1900521 | 1902 |
| Streptococcus pyogenes MGAS5005 | $NC_{-}007297$ | 1838562 | 1806 |
| Streptococcus pyogenes MGAS6180 | $NC_{-}007296$ | 1897578 | 1852 |
| Streptococcus pyogenes MGAS8232 | $NC_{-}003485$ | 1895017 | 1907 |
| Streptococcus pyogenes MGAS9429 | $NC_{-}008021$ | 1836467 | 1785 |
| Streptococcus pyogenes NZ131 | $NC_{-}011375$ | 1815785 | 1764 |
| Streptococcus pyogenes SSI-1 | $NC_{-}004606$ | 1894275 | 1898 |
| Streptococcus pyogenes STAB902 | NZ_CP007041 | 1892124 | 1896 |
| Streptococcus pyogenes Manfredo | $NC_{-}009332$ | 1841271 | 1824 |

Table S2: The first dataset

| Name | Accession Num. | Sequence Length | CDS number |
|---|------------------|-----------------|------------|
| Citrobacter freundii CFNIH4 | NZ_CP026231 | 4528955 | 5057 |
| Citrobacter freundii FDAARGOS73 | NZ_CP026056 | 4494773 | 4791 |
| Citrobacter freundii HM38 | NZ_CP024672 | 4373270 | 4703 |
| Citrobacter freundii UMH13 | NZ_CP024683 | 4379931 | 4685 |
| Citrobacter freundii UMH14 | NZ_CP024680 | 4328381 | 4621 |
| Citrobacter freundii UMH15 | NZ_CP024679 | 4765600 | 5215 |
| Citrobacter freundii UMH16 | NZ_CP024677 | 4631234 | 5033 |
| Citrobacter freundii UMH19 | NZ_CP024673 | 4457564 | 4860 |
| Enterobacter cloacae ECNIH3 | NZ_CP008897 | 4158155 | 4378 |
| Enterobacter cloacae ECR091 | NZ_CP008905 | 4146824 | 4357 |
| Enterobacter hormaechei 234 | NZ_CP021162 | 4188393 | 4375 |
| Enterobacter hormaechei 388 | $NZ_{-}CP021167$ | 4187638 | 4375 |
| Enterobacter hormaechei C15117 | $NZ_{-}CP032841$ | 4251211 | 4470 |
| Enterobacter hormaechei CAV1176 | NZ_CP011662 | 4381416 | 4676 |
| Enterobacter hormaechei PG20180049 | $NZ_{-}CP043226$ | 4154531 | 4326 |
| Enterobacter hormaechei PG20180056 | $NZ_{-}CP043225$ | 4154327 | 4325 |
| Enterobacter hormaechei S1116 | $NZ_{-}CP035385$ | 4342669 | 4632 |
| Enterobacter hormaechei SCEH020042 | $NZ_{-}CP028538$ | 4271727 | 4537 |
| Enterobacter hormaechei WCHEH090011 | NZ_CP036310 | 4174171 | 4440 |
| Enterobacter hormaechei oharae DSM 16687 | NZ_CP017180 | 4228448 | 4479 |
| Enterobacter hormaechei steigerwaltii 34977 | $NZ_{-}CP010376$ | 4389914 | 4615 |
| Enterobacter hormaechei steigerwaltii 34998 | $NZ_{-}CP012167$ | 4209967 | 4415 |
| Enterobacter hormaechei steigerwaltii DSM 16691 | NZ_CP017179 | 4295233 | 4494 |
| Enterobacter hormaechei steigerwaltii ME-1 | NZ_CP041733 | 4317058 | 4556 |
| Escherichia coli K-12 GM4792 Lac+ | $NZ_{-}CP011342$ | 4074170 | 4346 |
| Escherichia coli K-12 GM4792 Lac- | NZ_CP011343 | 4073312 | 4345 |
| Escherichia coli K-12 C3026 | $NZ_{-}CP014272$ | 4178806 | 4565 |
| Escherichia coli K-12 DHB4 | $NZ_{-}CP014270$ | 4003293 | 4285 |
| Escherichia coli K-12 ER3413 | NZ_CP009789 | 4019277 | 4314 |
| Escherichia coli O111 RM9322 | NZ_CP028117 | 4633699 | 5371 |
| Escherichia coli O111:H- 11128 DNA | $NC_{-}013364$ | 4736848 | 5331 |
| Escherichia coli O157:H7 EC4115 | $NC_{-}011353$ | 4931236 | 5575 |
| Escherichia coli O157:H7 EDL933 | $NZ_{-}CP008957$ | 4904375 | 5533 |
| Escherichia coli O157:H7 SS17 | $NZ_{-}CP008805$ | 4887496 | 5509 |
| Escherichia coli O157:H7 SS52 | $NZ_CP010304$ | 4861233 | 5459 |
| Escherichia coli O157:H7 Sakai DNA | $NC_{-}002695$ | 4825468 | 5203 |
| Escherichia coli O157:H7 TW14359 | NC_013008 | 4892906 | 5518 |
| Escherichia coli O157:H7 C1-057 | NZ_CP035366 | 4832316 | 5423 |

| continued table S2 | | | |
|---|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Escherichia coli O157:H7 FRIK2069 | NZ_CP015846 | 5055812 | 5950 |
| Escherichia coli O157:H7 FRIK2455 | NZ_CP015843 | 5054062 | 5945 |
| Escherichia coli O157:H7 FRIK2533 | NZ_CP015842 | 5049678 | 5942 |
| Escherichia coli O157:H7 FRIK944 | $NZ_{-}CP016625$ | 5104800 | 5842 |
| Escherichia coli O157:H7 TR01 | $NZ_{-}CP033605$ | 4931755 | 5574 |
| Escherichia coli O157:H7 WS4202 | $NZ_{-}CP012802$ | 4722882 | 5254 |
| Escherichia coli O26 RM10386 | NZ_CP028126 | 5181115 | 6183 |
| Escherichia coli O26 RM8426 | NZ_CP028116 | 5051758 | 6001 |
| Escherichia coli O26:H11 11368 DNA | $NC_{-}013361$ | 5039508 | 5776 |
| Escherichia coli O55:H7 CB9615 | NC_013941 | 4782409 | 5236 |
| Escherichia coli O55:H7 RM12579 | $NC_{-}017656$ | 4672671 | 5052 |
| Escherichia coli O6:H16 M9682-C1 | $NZ_{-}CP024275$ | 4273639 | 4867 |
| Escherichia coli K-12 subMG1655 K-12 | $NZ_{-}CP025268$ | 4143348 | 4687 |
| Escherichia coli K-12 subMG1655 | NC_000913 | 4079027 | 4357 |
| Escherichia coli K12 subDH10B | $NC_{-}010473$ | 4197082 | 4738 |
| Klebsiella pneumoniae 121 | NZ_CP031849 | 4587213 | 5015 |
| Klebsiella pneumoniae 2-1 | $NZ_{-}CP031562$ | 4651813 | 5071 |
| Klebsiella pneumoniae 2e | NZ_CP040175 | 4623234 | 4970 |
| Klebsiella pneumoniae 4743 | $NZ_{-}CP033625$ | 4796918 | 5260 |
| Klebsiella pneumoniae AP8555 | $NZ_{-}CP035383$ | 4865381 | 5239 |
| Klebsiella pneumoniae B31 | $NZ_{-}CP035929$ | 4699303 | 5045 |
| Klebsiella pneumoniae BJCFK909 | $NZ_{-}CP034123$ | 4871883 | 5405 |
| Klebsiella pneumoniae CR-HvKP1 | $NZ_{-}CP040533$ | 4787304 | 5240 |
| Klebsiella pneumoniae CR-HvKP5 | $NZ_{-}CP040545$ | 4802269 | 5258 |
| Klebsiella pneumoniae I72 | NZ_CP034281 | 4845718 | 5283 |
| Klebsiella pneumoniae INF078 | $NZ_{-}CP032831$ | 4710656 | 5126 |
| Klebsiella pneumoniae INF116-sc-2279924 | $NZ_{-}CP031792$ | 4701024 | 5118 |
| Klebsiella pneumoniae INF125-sc-2279943 | $NZ_{-}CP031795$ | 4714641 | 5138 |
| Klebsiella pneumoniae INF206-sc-2280074 | $NZ_{-}CP031808$ | 4779695 | 5269 |
| Klebsiella pneumoniae INF235-sc-2280127 | $NZ_{-}CP031817$ | 4571468 | 4991 |
| Klebsiella pneumoniae KP14003 | NZ_CP041934 | 4702032 | 5041 |
| Klebsiella pneumoniae KP17-15 | $NZ_{-}CP034076$ | 4788639 | 5172 |
| Klebsiella pneumoniae KP17-16 | $NZ_{-}CP034077$ | 4788639 | 5172 |
| Klebsiella pneumoniae KP18-29 | $NZ_{-}CP034249$ | 4772769 | 5261 |
| Klebsiella pneumoniae KP58 | NZ_CP041373 | 4868529 | 5348 |
| Klebsiella pneumoniae KP65 | $NZ_{-}CP044258$ | 4829714 | 5299 |
| Klebsiella pneumoniae KPNORMBLD2014104014 | $NZ_{-}CP034045$ | 4743890 | 5100 |
| Klebsiella pneumoniae KPNORMBLD2015112126 | $NZ_{-}CP034053$ | 4710060 | 5052 |
| Klebsiella pneumoniae KSB11I-sc-2280289 | NZ_CP031789 | 4617932 | 5061 |
| Klebsiella pneumoniae KSB17F-sc-2280268 | NZ_CP031814 | 4696494 | 5186 |
| Klebsiella pneumoniae Kp202 | $NZ_{-}CP041082$ | 4839245 | 5266 |
| Klebsiella pneumoniae L392 | $NZ_{-}CP033954$ | 4860428 | 5334 |
| Klebsiella pneumoniae L482 | NZ_CP033960 | 4849398 | 5330 |
| Klebsiella pneumoniae NB5306 | NZ_CP034760 | 4685452 | 5161 |
| Klebsiella pneumoniae P094-1 | $NZ_{-}CP043562$ | 4477472 | 4848 |
| Klebsiella pneumoniae PIMB15ND2KP27 | NZ_CP041639 | 4665800 | 5085 |
| Klebsiella pneumoniae QMP B2-170 | NZ_CP031798 | 4705293 | 5181 |
| Klebsiella pneumoniae SCKP020135 | NZ_CP037963 | 4639860 | 5034 |
| Klebsiella pneumoniae ST23 | NZ_CP037742 | 4799363 | 5119 |
| Klebsiella pneumoniae WCHKP015625 | NZ_CP033396 | 4851409 | 5361 |
| Klebsiella pneumoniae WCHKP020098 | NZ_CP036305 | 4853482 | 5356 |
| Klebsiella pneumoniae WCHKP095845 | NZ_CP031885 | 4679182 | 5122 |

continued table S2

| continued table S2 | | | |
|---|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Klebsiella pneumoniae WCHKP115069 | NZ_CP033405 | 4841832 | 5339 |
| Klebsiella pneumoniae WCHKP2080 | NZ_CP036361 | 4850078 | 5361 |
| Klebsiella pneumoniae WCHKP3 | NZ_CP031721 | 4817973 | 5268 |
| Klebsiella pneumoniae XJ-K1 | NZ_CP032163 | 4842187 | 5322 |
| Klebsiella pneumoniae ZYST1 | NZ_CP031613 | 4682297 | 5118 |
| Klebsiella pneumoniae pneumoniae Kp13 | NZ_CP003999 | 4711070 | 5056 |
| Klebsiella pneumoniae pneumoniae 12208 | NZ_CP030172 | 4649576 | 5065 |
| Klebsiella pneumoniae pneumoniae 234-12 | NZ_CP011313 | 4675867 | 5065 |
| Klebsiella pneumoniae pneumoniae ARLG-3135 | NZ_CP033946 | 4737882 | 5152 |
| Klebsiella pneumoniae pneumoniae AUSMDU00008079 | NZ_CP022691 | 4859140 | 5384 |
| Klebsiella pneumoniae pneumoniae CCRI-22199 | $NZ_CP035540$ | 4706222 | 5125 |
| Klebsiella pneumoniae pneumoniae CRK0298 | NZ_CP034039 | 4892634 | 5395 |
| Klebsiella pneumoniae pneumoniae KPCTRSRTH01 | NZ_CP041092 | 4626990 | 4940 |
| Klebsiella pneumoniae pneumoniae KPCTRSRTH07 | NZ_CP041099 | 4673423 | 5059 |
| Klebsiella pneumoniae pneumoniae M5 | NZ_CP031734 | 4801265 | 5212 |
| Klebsiella pneumoniae pneumoniae R210-2 | NZ_CP034082 | 4867663 | 5251 |
| Klebsiella pneumoniae pneumoniae RJA166 | NZ_CP019047 | 4764728 | 5161 |
| Klebsiella pneumoniae pneumoniae SCKP020046 | NZ_CP028783 | 4833417 | 5366 |
| Klebsiella pneumoniae pneumoniae SCKP020079 | NZ_CP029384 | 4878258 | 5455 |
| Klebsiella pneumoniae pneumoniae SCKP020143 | NZ_CP028548 | 4904595 | 5451 |
| Klebsiella pneumoniae pneumoniae WCHKP015093 | NZ_CP036300 | 4865017 | 5380 |
| Salmonella enterica Typhi 311189201186 | NZ_CP029958 | 4236973 | 4778 |
| Salmonella enterica Typhi 311189202186 | NZ_CP029956 | 4255507 | 4809 |
| Salmonella enterica Typhi 311189203186 | NZ_CP029942 | 4240151 | 4783 |
| Salmonella enterica Typhi 311189204186 | $NZ_{CP029954}$ | 4238692 | 4784 |
| Salmonella enterica Typhi 311189205186 | $NZ_{-}CP029950$ | 4275035 | 4846 |
| Salmonella enterica Typhi 311189206186 | NZ_CP029940 | 4239953 | 4782 |
| Salmonella enterica Typhi 311189207186 | NZ_CP029938 | 4237518 | 4778 |
| Salmonella enterica Typhi 311189208103 | $NZ_{-}CP029932$ | 4236430 | 4776 |
| Salmonella enterica Typhi 311189208186 | NZ_CP029949 | 4236277 | 4775 |
| Salmonella enterica Typhi 311189209186 | $NZ_CP029952$ | 4236302 | 4775 |
| Salmonella enterica Typhi 311189210186 | NZ_CP029946 | 4239943 | 4784 |
| Salmonella enterica Typhi 311189211186 | $NZ_{-}CP029945$ | 4239689 | 4782 |
| Salmonella enterica Typhi 311189212186 | NZ_CP029944 | 4237757 | 4777 |
| Salmonella enterica Typhi 311189213186 | NZ_CP029936 | 4236722 | 4777 |
| Salmonella enterica Typhi 311189214186 | $NZ_{-}CP029933$ | 4236583 | 4777 |
| Salmonella enterica Typhi 311189215186 | NZ_CP029930 | 4235774 | 4774 |
| Salmonella enterica Typhi 311189216103 | $NZ_{-}CP029928$ | 4254957 | 4803 |
| Salmonella enterica Typhi 311189217103 | $NZ_{-}CP029927$ | 4237323 | 4771 |
| Salmonella enterica Typhi 311189217186 | NZ_CP029646 | 4237112 | 4781 |
| Salmonella enterica Typhi 311189218186 | $NZ_{-}CP029925$ | 4239999 | 4783 |
| Salmonella enterica Typhi 311189219186 | NZ_CP029911 | 4238210 | 4779 |
| Salmonella enterica Typhi 311189220186 | NZ_CP029886 | 4236051 | 4775 |
| Salmonella enterica Typhi 311189221186 | NZ_CP029923 | 4236820 | 4779 |
| Salmonella enterica Typhi 311189222186 | $NZ_{-}CP029909$ | 4295377 | 4867 |
| Salmonella enterica Typhi 311189223186 | $NZ_{-}CP029880$ | 4237502 | 4778 |
| Salmonella enterica Typhi 311189224186 | NZ_CP029878 | 4236309 | 4776 |
| Salmonella enterica Typhi 311189231186 | NZ_CP029904 | 4259692 | 4815 |
| Salmonella enterica Typhi 311189232103 | $NZ_{-}CP029918$ | 4235827 | 4765 |
| Salmonella enterica Typhi 311189239103 | NZ_CP029908 | 4236639 | 4768 |
| Salmonella enterica Typhi 311189252186 | NZ_CP029896 | 4235511 | 4773 |
| Salmonella enterica Typhi 311189255186 | $NZ_{-}CP029885$ | 4260953 | 4824 |

continued table S2

| continued table S2 | | | |
|--|----------------------------|----------------------|--------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Salmonella enterica Typhi 311189256186 | NZ_CP029917 | 4252129 | 4800 |
| Salmonella enterica Typhi 311189268103 | NZ_CP029902 | 4259998 | 4815 |
| Salmonella enterica Typhi 311189268186 | NZ_CP029883 | 4259669 | 4815 |
| Salmonella enterica Typhi 311189269186 | $NZ_{-}CP029922$ | 4235855 | 4773 |
| Salmonella enterica Typhi 311189282186 | NZ_CP029920 | 4252366 | 4800 |
| Salmonella enterica Typhi 311189291186 | NZ_CP029894 | 4259514 | 4815 |
| Salmonella enterica Typhi 343076202113 | NZ_CP029915 | 4252474 | 4801 |
| Salmonella enterica Typhi 343076217103 | NZ_CP029914 | 4228169 | 4776 |
| Salmonella enterica Typhi 343076227128 | $NZ_{-}CP029875$ | 4252463 | 4800 |
| Salmonella enterica Typhi 343076232188 | NZ_CP029900 | 4252705 | 4803 |
| Salmonella enterica Typhi 343076241106 | NZ_CP029899 | 4228113 | 4775 |
| Salmonella enterica Typhi 343076248190 | NZ_CP029882 | 4228096 | 4777 |
| Salmonella enterica Typhi 343076249107 | NZ_CP029913 | 4236671 | 4776 |
| Salmonella enterica Typhi 343076252143 | NZ_CP029892 | 4260260 | 4817 |
| Salmonella enterica Typhi 343076253155 | NZ_CP029890 | 4248231 | 4800 |
| Salmonella enterica Typhi 343076269157 | NZ_CP029881 | 4247848 | 4798 |
| Salmonella enterica Typhi 343076294172 | NZ_CP029888 | 4261080 | 4820 |
| Salmonella enterica Typhi 343077211171 | NZ_CP029873 | 4237127 | 4777 |
| Salmonella enterica Typhi 343077212138 | NZ_CP029919 | 4227481 | 4773 |
| Salmonella enterica Typhi 343077212159 | NZ_CP029870 | 4251075 | 4804 |
| Salmonella enterica Typhi 343077213147 | NZ_CP029897 | 4344563 | 4940 |
| Salmonella enterica Typhi 343077214135 | NZ_CP029872 | 4236615 | 4774 |
| Salmonella enterica Typhi 343077214162 | NZ_CP029862 | 4266433 | 4816 |
| Salmonella enterica Typhi 343077215174 | NZ_CP029868 | 4238813 | 4779 |
| Salmonella enterica Typhi 343077228140 | NZ_CP029866 | 4247745 | 4799 |
| Salmonella enterica Typhi 343077228157 | NZ_CP029864 | 4259824 | 4818 |
| Salmonella enterica Typhi 343077255118 | NZ_CP029907 | 4228259 | 4776 |
| Salmonella enterica Typhi 343077260153 | NZ_CP029861 | 4237347 | 4778 |
| Salmonella enterica Typhi 343077267164 | NZ_CP029906 | 4227890 | 4775 |
| Salmonella enterica Typhi 343077278127 | NZ_CP029863 | 4259524 | 4816 |
| Salmonella enterica Typhi 343077281186 | NZ_CP029853 | 4281576 | 4852 |
| Salmonella enterica Typhi 343077285138 | NZ_CP029858 | 4252961 | 4805 |
| Salmonella enterica Typhi 343077286126 | NZ_CP029856 | 4250781 | 4804 |
| Salmonella enterica Typhi 343077292148 | NZ_CP029855 | 4253231 | 4802 |
| Salmonella enterica Typhi 343078201101 | NZ_CP029852 | 4235689 | 4774 |
| Salmonella enterica Typhi 343078203125 | NZ_CP029850 | 4253684 | 4803 |
| Salmonella enterica Typhi 343078211126 | NZ_CP029848 | 4252644 | 4803 |
| Salmonella enterica Typhi 343078223175 | NZ_CP029964 | 4237694 | 4780 |
| Salmonella enterica Typhi 343078228140 | NZ_CP029964 NZ_CP029962 | 4236329 | 4780 4775 |
| Salmonella enterica Typhi 343078251131 | NZ_CP029962 NZ_CP029960 | | 4804 |
| v <u>-</u> | | 4252907 | |
| Salmonella enterica Typhi 343078256191 | NZ_CP029959 | 4250926 | 4803 |
| Salmonella enterica Typhi 343078273110 | NZ_CP029846 NZ_CP029840 | 4253735 | 4804 |
| Salmonella enterica Typhimurium 01ST04081 | | 4291358 | 4648 |
| Salmonella enterica Typhimurium 08-365-0802 | NZ_CP038847 | 4285306 | 4663 |
| Salmonella enterica Typhimurium 10ST07093 | NZ_CP029839 | $4285734 \\ 4360344$ | 4639 |
| Salmonella enterica Typhimurium 11A0003616 | NZ_CP039591 | | 4785 4625 |
| Salmonella enterica Typhimurium 13-931 Salmonella enterica Typhimurium 138736 | NZ_CP016385 | 4274602 | 4625 |
| 0 2 | NZ_CP007581 | 4347161 | 4755 4771 |
| Salmonella enterica Typhimurium 16425992 | NZ_CP039576 | 4350426 | 4771 |
| Salmonella enterica Typhimurium 22495 | NZ_CP017617 | 4265308 | 4724 |
| Salmonella enterica Typhimurium 33676 | NZ_CP012681 | 4235788 | 4581 |
| Salmonella enterica Typhimurium ATCC 13311 | NZ_CP009102 | 4219591 | 4551 |

| Name | Accession Num. | Sequence Length | CDS number |
|---|------------------|-----------------|------------|
| Salmonella enterica Typhimurium E2840153550 | NZ_CP040321 | 4338443 | 4746 |
| Salmonella enterica Typhimurium FORC015 | $NZ_CP011365$ | 4186788 | 4547 |
| Salmonella enterica Typhimurium NC983 | $NZ_CP015157$ | 4268681 | 4625 |
| Salmonella enterica Typhimurium RM10961 | $NZ_{-}CP013702$ | 4281474 | 4651 |
| Salmonella enterica Typhimurium SO2 | $NZ_{-}CP014356$ | 4337925 | 4716 |
| Salmonella enterica Typhimurium SO3 | $NZ_CP014536$ | 4333931 | 4708 |
| Salmonella enterica Typhimurium UGA14 | $NZ_CP021462$ | 4352481 | 4915 |
| Salmonella enterica Typhimurium VNP20009 | NZ_CP007804 | 4194027 | 4547 |
| Salmonella enterica Typhimurium YU15 | $NZ_{-}CP014358$ | 4347319 | 4755 |
| Shigella flexneri 1c Y394 | $NZ_CP020753$ | 4007861 | 4545 |
| Shigella flexneri 2002017 | NC_017328 | 4071936 | 4622 |
| Shigella flexneri 2003036 | $NZ_{-}CP004056$ | 4017841 | 4575 |
| Shigella flexneri 2a 981 | $NZ_{-}CP012137$ | 4074506 | 4635 |
| Shigella flexneri 4c 1602 | $NZ_{-}CP020336$ | 4075350 | 4643 |
| Shigella flexneri 5 8401 | NC_008258 | 4006104 | 4521 |
| Shigella flexneri G1663 | $NZ_{-}CP007037$ | 4017117 | 4552 |
| Shigella flexneri Shi06HN006 | NZ_CP004057 | 4038738 | 4610 |
| Shigella flexneri Y PE577 | NZ_CP042980 | 4015236 | 4547 |
| Shigella flexneri SFL1520 | $NZ_CP032513$ | 4008317 | 4640 |
| Shigella sonnei 53G main | NC_016822 | 4445319 | 5389 |
| Shigella sonnei Ss046 | $NC_{-}007384$ | 4280935 | 5174 |
| Shigella sonnei FORC011 | NZ_CP010829 | 4280962 | 5173 |

Table S3: The second dataset

| Name | Accession Num. | Sequence Length | CDS number |
|---------------------------------------|------------------|-----------------|------------|
| Acinetobacter baumannii 09A16CRGN0014 | NZ_CP034242 | 3943275 | 3738 |
| Acinetobacter baumannii 09A16CRGN003B | NZ_CP034243 | 3943105 | 3737 |
| Acinetobacter baumannii1A1213CRGN008 | NZ_CP035186 | 3944921 | 3719 |
| Acinetobacter baumannii1A1213CRGN055 | $NZ_{-}CP035185$ | 3934099 | 3706 |
| Acinetobacter baumannii1A1213CRGN064 | NZ_CP043419 | 3935331 | 3709 |
| Acinetobacter baumannii1A1314CRGN088 | NZ_CP035184 | 3937511 | 3715 |
| Acinetobacter baumannii1A1314CRGN089 | NZ_CP043418 | 3935083 | 3708 |
| Acinetobacter baumannii1A14CRGN003 | NZ_CP035183 | 3940649 | 3722 |
| Acinetobacter baumannii KAB04 | NZ_CP017648 | 4006639 | 3815 |
| Acinetobacter baumannii TG22182 | NZ_CP039993 | 3975485 | 3753 |
| Acinetobacter baumannii TG22627 | $NZ_{-}CP039520$ | 3915428 | 3674 |
| Acinetobacter baumannii TG22653 | NZ_CP039518 | 3891074 | 3641 |
| Acinetobacter baumannii TG29392 | NZ_CP039930 | 3963137 | 3735 |
| Acinetobacter baumannii TG31302 | NZ_CP039343 | 4044095 | 3814 |
| Acinetobacter baumannii TG31986 | NZ_CP039341 | 4049647 | 3823 |
| Acinetobacter baumannii WCHAB005133 | $NZ_{-}CP026750$ | 3893621 | 3642 |
| Acinetobacter baumannii XH386 | NZ_CP021326 | 4108882 | 3935 |
| Acinetobacter baumannii XH386 | NZ_CP010779 | 4087343 | 3916 |
| Acinetobacter baumannii XH731 | NZ_CP021321 | 3914578 | 3675 |
| Acinetobacter baumannii XH731 | $NZ_{-}CP019217$ | 3909773 | 3667 |
| Acinetobacter baumannii XH856 | NZ_CP014541 | 3908399 | 3651 |
| Acinetobacter baumannii XH860 | NZ_CP014538 | 3846621 | 3598 |
| Acinetobacter baumannii XH906 | NZ_CP023140 | 3891370 | 3655 |
| Bacillus cereus ATCC4579 | $NC_{-}004722$ | 5411809 | 5210 |
| Bacillus cereus ATCC4579 | NZ_CP034551 | 5416249 | 5458 |

continued table S3

| continued table S3 | | | |
|--------------------------------------|----------------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Bacillus cereus B4264 | $NC_{-}011725$ | 5419036 | 5535 |
| Bacillus cereus C1L | $NZ_{-}CP022445$ | 5312355 | 5517 |
| Bacillus cereus000305 | $NZ_{-}CP028516$ | 5330209 | 5290 |
| Bacillus cereus CMCC P0011 | NZ_CP011153 | 5506876 | 5645 |
| Bacillus cereus CMCC P0021 | NZ_CP011151 | 5521782 | 5648 |
| Bacillus cereus D12-2 | $NZ_{-}CP016315$ | 5369104 | 5469 |
| Bacillus cereus FORC087 | $NZ_{-}CP029454$ | 5271204 | 5345 |
| Bacillus cereus FORC60 | $NZ_{-}CP020383$ | 5361178 | 5542 |
| Bacillus cereus FORC-005 | $NZ_{-}CP009686$ | 5349617 | 5347 |
| Bacillus cereus FORC-024 | $NZ_{-}CP012691$ | 5430118 | 5518 |
| Bacillus cereus FORC-047 | NZ_CP017060 | 5365433 | 5515 |
| Bacillus cereus FORC-048 | $NZ_{-}CP017234$ | 5358595 | 5510 |
| Bacillus cereus FORC-086 | $NZ_{-}CP029468$ | 5340498 | 5281 |
| Bacillus cereus K8 | $NZ_{-}CP016595$ | 5329684 | 5497 |
| Bacillus cereus M13 | NZ_CP016360 | 5498820 | 5670 |
| Bacillus cereus SB1 | $NZ_{-}CP045606$ | 5461601 | 5366 |
| Bacillus cereus ZB201708 | NZ_CP030982 | 5466652 | 5484 |
| Clostridioides difficile 09-00072 | NZ_CP026599 | 4119929 | 3714 |
| Clostridioides difficile0-00071 | NZ_CP026596 | 4119720 | 3715 |
| Clostridioides difficile0-00078 | NZ_CP026597 | 4119749 | 3717 |
| Clostridioides difficile0-00253 | NZ_CP026598 | 4120065 | 3715 |
| Clostridioides difficile2-00008 | NZ_CP026593 | 4119604 | 3714 |
| Clostridioides difficile CD-10-00484 | NZ_CP026592 | 4119484 | 3715 |
| Clostridioides difficile CD-17-01474 | NZ_CP026591 | 4201330 | 3812 |
| Clostridioides difficile CD10010 | NZ_CP033213 | 4049580 | 3657 |
| Clostridioides difficile CD161 | NZ_CP029154 | 4295210 | 3925 |
| Clostridioides difficile CD21062 | NZ_CP033216 | 3996135 | 3610 |
| Clostridioides difficile CDT4 | NZ_CP029152 | 4240954 | 3836 |
| Clostridioides difficile DSM02859 | NZ_CP020378 | 4247015 | 3922 |
| Clostridioides difficile DSM02860 | NZ_CP020379 | 4253785 | 3943 |
| Clostridioides difficile DSM02978 | NZ_CP020380 | 4253783 | 3943 |
| Clostridioides difficile DSM 27638 | NZ_CP011846 | 4229698 | 3838 |
| Clostridioides difficile DSM 27639 | NZ_CP011847 | 4263997 | 3899 |
| Clostridioides difficile DSM 27640 | NZ_CP011848 | 4229629 | 3836 |
| Clostridioides difficile DSM 28196 | NZ_CP012320 | 4205365 | 3820 |
| Clostridioides difficile DSM 28666 | NZ_CP012321 | 4122585 | 3701 |
| Clostridioides difficile DSM 28668 | NZ_CP012321 NZ_CP012309 | 4214812 | 3772 |
| Clostridioides difficile DSM 28669 | NZ_CP012323 | 4137693 | 3780 |
| Clostridioides difficile DSM 28670 | NZ_CP012323 NZ_CP012312 | 4191653 | 3833 |
| Clostridioides difficile DSM 29020 | NZ_CP012312 NZ_CP012325 | 4139418 | 3776 |
| Clostridioides difficile DSM 29627 | | | |
| | NZ_CP016102 | 4209547 | 3788 |
| Clostridioides difficile DSM 29629 | NZ_CP016104 | 4111235 | 3674 |
| Clostridioides difficile DSM 29632 | NZ_CP019860 | 4175826 | 3763 |
| Clostridioides difficile DSM 29637 | NZ_CP016106 | 4114585 | 3716 |
| Clostridioides difficile DSM 29688 | NZ_CP019858 | 4228964 | 3913 |
| Clostridioides difficile DSM 29745 | NZ_CP019857 | 4240646 | 3847 |
| Clostridioides difficile DSM 29747 | NZ_CP019864 | 4071596 | 3681 |
| Clostridioides difficile Mta-79 | NZ_CP042267 | 4123685 | 3719 |
| Clostridioides difficile R1 | NZ_CP026613 | 4093143 | 3707 |
| Clostridioides difficile R2 | NZ_CP026614 | 4093145 | 3707 |
| Enterococcus faecium 2014-VREF-63 | NZ_CP019988 | 2907074 | 2864 |
| Enterococcus faecium 2014-VREF-41 | NZ_CP019208 | 3009007 | 2985 |

| continued table S3 | | | |
|---|----------------------------|-----------------|--------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Enterococcus faecium AUSMDU00004024 | $NZ_{-}CP027517$ | 2863087 | 2844 |
| Enterococcus faecium AUSMDU00004028 | $NZ_CP027512$ | 2855729 | 2833 |
| Enterococcus faecium AUSMDU00004055 | $NZ_{-}CP027506$ | 2731844 | 2678 |
| Enterococcus faecium AUSMDU00004142 | $NZ_{-}CP027501$ | 2912017 | 2892 |
| Enterococcus faecium AUSMDU00004167 | $NZ_{-}CP027497$ | 2883877 | 2850 |
| Enterococcus faecium Efaecium-ER039333A | NZ_CP023780 | 2863296 | 2861 |
| Enterococcus faecium Efaecium-ER041203A | NZ_CP023784 | 2869275 | 2866 |
| Enterococcus faecium Efaecium-ER044623A | NZ_CP023789 | 2867770 | 2866 |
| Enterococcus faecium Efaecium-ER044843A | NZ_CP023794 | 2869064 | 2867 |
| Enterococcus faecium Efaecium-ER045263A | NZ_CP023808 | 2870954 | 2869 |
| Enterococcus faecium Efaecium-ER045265A | NZ_CP023799 | 2867667 | 2865 |
| Enterococcus faecium Efaecium-ER046193A | NZ_CP023804 | 2870397 | 2869 |
| Enterococcus faecium F17E0263 | NZ_CP040849 | 2659111 | 2661 |
| Enterococcus faecium ISMMS-VRE-1 | NZ_CP012430 | 3130373 | 3074 |
| Enterococcus faecium ISMMS-VRE-7 | NZ_CP012460 | 2850245 | 2756 |
| Enterococcus faecium ISMMS-VRE-9 | NZ_CP018830 | 2802604 | 2694 |
| Enterococcus faecium K60-39 | NZ_CP023423 | 2739582 | 2704 |
| Enterococcus faecium LAC72 | NZ_CP045012 | 2625745 | 2521 |
| Enterococcus faecium RBWH1 | NZ_CP033206 | 2876110 | 2749 |
| Enterococcus faecium SC4 | NZ_CP035200 NZ_CP025425 | 2772425 | 2749 2751 |
| Enterococcus faecium UAMSEF-01 | NZ_CP025425 NZ_CP035648 | 2824869 | 2803 |
| | | | |
| Enterococcus faecium UAMSEF-08 | NZ_CP035654 | 2827987 | 2806 |
| Enterococcus faecium UAMSEF-09 | NZ_CP035660 | 2912202 | 2920 |
| Enterococcus faecium UAMSEF-20 | NZ_CP035666 | 2912170 | 2921 |
| Enterococcus faecium V2937 | NZ_CP044274 | 2896393 | 2794 |
| Enterococcus faecium VB3240 | NZ_CP040368 | 2805984 | 2759 |
| Enterococcus faecium VRE001 | NZ_CP018071 | 2933966 | 2896 |
| Enterococcus faecium VRE1 | NZ_CP040740 | 2718395 | 2659 |
| Enterococcus faecium VVEswe-R | NZ_CP041261 | 2851560 | 2834 |
| Enterococcus faecium VVEswe-S | NZ_CP041270 | 2851769 | 2836 |
| Escherichia coli O103:H2 FWSEC0007 | NZ_CP031908 | 5397605 | 5368 |
| Escherichia coli O104:H4 FWSEC0009 | NZ_CP031902 | 5277234 | 5151 |
| Escherichia coli O113:H21 FWSEC0010 | $NZ_{-}CP031898$ | 4955402 | 4751 |
| Escherichia coli O121:H19 FWSEC0006 | NZ_CP031910 | 5398870 | 5345 |
| Escherichia coli O145:NM FWSEC0002 | NZ_CP031919 | 5695528 | 5679 |
| Escherichia coli O157:H7 FWSEC0004 | NZ_CP031913 | 5406250 | 5341 |
| Escherichia coli O16:H48 PG20180052 | $NZ_{-}CP043209$ | 4615351 | 4385 |
| Escherichia coli O16:H48 PG20180053 | $NZ_{-}CP043207$ | 4615322 | 4384 |
| Escherichia coli O16:H48 PG20180054 | $NZ_{-}CP043205$ | 4615343 | 4385 |
| Escherichia coli O16:H48 PG20180058 | $NZ_{-}CP043203$ | 4656753 | 4418 |
| Escherichia coli O16:H48 PG20180059 | $NZ_{-}CP043201$ | 4656146 | 4416 |
| Escherichia coli O16:H48 PG20180063 | NZ_CP043193 | 4656756 | 4419 |
| Escherichia coli O16:H48 PG20180170 | NZ_CP043187 | 4656750 | 4417 |
| Escherichia coli O16:H48 PG20180172 | NZ_CP043183 | 4656755 | 4417 |
| Escherichia coli O16:H48 PG20180173 | NZ_CP043191 | 4656753 | 4418 |
| Escherichia coli O16:H48 PG20180175 | NZ_CP043189 | 4656553 | 4420 |
| Escherichia coli O26:H11 FWSEC0001 | NZ_CP031922 | 5697154 | 5784 |
| Escherichia coli O91:H21 FWSEC0008 | NZ_CP031906 | 4972544 | 4781 |
| Escherichia coli AR2022 | NZ_CP043946 | 4847893 | 4563 |
| Escherichia coli AR2162b | NZ_CP043942 | 4849144 | 4565 |
| Escherichia coli AR242b | NZ_CP035944 | 4666029 | 4416 |
| | 112201 0000 IT | 1000020 | 1110 |

| Name | Accession Num. | Sequence Length | CDS numbe |
|---|------------------|-----------------|-----------|
| Escherichia coli FORC 064 | NZ_CP022664 | 4873055 | 498 |
| Escherichia coli FORC-029 | NZ_CP013185 | 5189944 | 530 |
| Escherichia coli FORC-031 | NZ_CP013190 | 4830073 | 462 |
| Escherichia coli FORC-041 | NZ_CP016628 | 4867152 | 461 |
| Escherichia coli FORC-042 | $NZ_{-}CP025318$ | 5052887 | 520 |
| Escherichia coli FORC-043 | NZ_CP016828 | 5244443 | 538 |
| Escherichia coli FORC-044 | $NZ_{-}CP016755$ | 5483726 | 572 |
| Escherichia coli FORC-069 | NZ_CP023061 | 5189917 | 530 |
| Escherichia coli FORC-082 | NZ_CP026641 | 4728307 | 456 |
| Mycobacterium tuberculosis CCDC5079 | NC_017523 | 4398812 | 424 |
| Mycobacterium tuberculosis CCDC5079 | $NC_{-}021251$ | 4414325 | 424 |
| Mycobacterium tuberculosis CCDC5180 | $NC_{-}017522$ | 4405981 | 425 |
| Mycobacterium tuberculosis CCDC5180 | $NZ_{-}CP002885$ | 4414346 | 424 |
| Mycobacterium tuberculosis CDC1551 | NC_002755 | 4403837 | 424 |
| Mycobacterium tuberculosis Beijing-like | NZ_CP010873 | 4411216 | 420 |
| Mycobacterium tuberculosis Beijing-like/1104 | $NZ_{-}CP017598$ | 4380156 | 425 |
| Mycobacterium tuberculosis Beijing-like/35049 | $NZ_{-}CP017593$ | 4427062 | 42 |
| Mycobacterium tuberculosis Beijing-like/36918 | NZ_CP017594 | 4441591 | 420 |
| Mycobacterium tuberculosis Beijing-like/38774 | $NZ_{-}CP017595$ | 4431885 | 42 |
| Mycobacterium tuberculosis Beijing-like/50148 | $NZ_{-}CP017597$ | 4444417 | 420 |
| Mycobacterium tuberculosis Beijing/391 | $NZ_{-}CP017596$ | 4406925 | 42 |
| Mycobacterium tuberculosis Beijing2014PNGD | $NZ_{-}CP022704$ | 4404064 | 420 |
| Mycobacterium tuberculosis CSV10399 | $NZ_{-}CP023576$ | 4411180 | 42 |
| Mycobacterium tuberculosis CSV11678 | $NZ_{-}CP023577$ | 4411382 | 423 |
| Mycobacterium tuberculosis CSV3611 | $NZ_{-}CP023600$ | 4411439 | 423 |
| Mycobacterium tuberculosis CSV383 | $NZ_{-}CP023599$ | 4411115 | 423 |
| Mycobacterium tuberculosis CSV4519 | NZ_CP023573 | 4411288 | 423 |
| Mycobacterium tuberculosis CSV4644 | $NZ_{-}CP023574$ | 4411271 | 423 |
| Mycobacterium tuberculosis CSV5769 | $NZ_{-}CP023575$ | 4411312 | 423 |
| Mycobacterium tuberculosis CSV9577 | NZ_CP023601 | 4411230 | 423 |
| Mycobacterium tuberculosis GG-109-10 | $NZ_{-}CP025602$ | 4411463 | 42 |
| Mycobacterium tuberculosis GG-111-10 | $NZ_{-}CP025593$ | 4411563 | 42 |
| Mycobacterium tuberculosis GG-121-10 | $NZ_{-}CP025603$ | 4411510 | 42 |
| Mycobacterium tuberculosis GG-129-11 | $NZ_{-}CP025604$ | 4411413 | 42 |
| Mycobacterium tuberculosis GG-134-11 | $NZ_{-}CP025605$ | 4411399 | 42 |
| Mycobacterium tuberculosis GG-137-10 | $NZ_{-}CP025606$ | 4411446 | 42 |
| Mycobacterium tuberculosis GG-186-10 | $NZ_{-}CP025607$ | 4411478 | 42 |
| Mycobacterium tuberculosis GG-20-11 | $NZ_{-}CP025595$ | 4411504 | 42 |
| Mycobacterium tuberculosis GG-229-10 | NZ_CP025608 | 4411519 | 42 |
| Mycobacterium tuberculosis GG-27-11 | $NZ_{-}CP025596$ | 4411443 | 42 |
| Mycobacterium tuberculosis GG-36-11 | $NZ_{-}CP025597$ | 4411469 | 42 |
| Mycobacterium tuberculosis GG-37-11 | $NZ_{-}CP025598$ | 4411526 | 42 |
| Mycobacterium tuberculosis GG-45-11 | $NZ_{-}CP025599$ | 4411469 | 42 |
| Mycobacterium tuberculosis GG-5-10 | $NZ_{-}CP025594$ | 4411442 | 42^{4} |
| Mycobacterium tuberculosis GG-77-11 | $NZ_{-}CP025600$ | 4411508 | 42 |
| Mycobacterium tuberculosis GG-90-10 | $NZ_{-}CP025601$ | 4411602 | 42 |
| Mycobacterium tuberculosis MDRDM1098 | $NZ_{-}CP023585$ | 4411148 | 423 |
| Mycobacterium tuberculosis MDRDM260 | $NZ_{-}CP023583$ | 4411280 | 423 |
| Mycobacterium tuberculosis MDRDM627 | $NZ_{-}CP023584$ | 4411215 | 423 |
| Mycobacterium tuberculosis MDRDM827 | $NZ_{-}CP023622$ | 4411315 | 423 |
| Mycobacterium tuberculosis MDRMA1565 | NZ_CP023626 | 4411159 | 423 |
| Mycobacterium tuberculosis MDRMA2019 | $NZ_{-}CP023627$ | 4411229 | 42 |

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| Name | Accession Num. | Sequence Length | CDS number |
| Mycobacterium tuberculosis MDRMA203 | NZ_CP023623 | 4411290 | 4234 |
| Mycobacterium tuberculosis MDRMA2082 | NZ_CP023628 | 4411479 | 4234 |
| Mycobacterium tuberculosis MDRMA2260 | NZ_CP023629 | 4411352 | 4237 |
| Mycobacterium tuberculosis MDRMA2441 | NZ_CP023630 | 4411454 | 4238 |
| Mycobacterium tuberculosis MDRMA2491 | $NZ_{-}CP023586$ | 4411121 | 4229 |
| Mycobacterium tuberculosis MDRMA701 | $NZ_{-}CP023624$ | 4411338 | 4235 |
| Mycobacterium tuberculosis MDRMA863 | $NZ_{-}CP023625$ | 4411432 | 4236 |
| Mycobacterium tuberculosis SLM036 | $NZ_{-}CP023592$ | 4411342 | 4236 |
| Mycobacterium tuberculosis SLM040 | $NZ_{-}CP023593$ | 4411408 | 4236 |
| Mycobacterium tuberculosis SLM056 | $NZ_{-}CP023594$ | 4411306 | 4232 |
| Mycobacterium tuberculosis SLM060 | $NZ_{-}CP023595$ | 4411134 | 4231 |
| Mycobacterium tuberculosis SLM063 | $NZ_{-}CP023596$ | 4411337 | 4234 |
| Mycobacterium tuberculosis SLM088 | $NZ_{-}CP023597$ | 4411385 | 4236 |
| Mycobacterium tuberculosis SLM100 | $NZ_{-}CP023598$ | 4411394 | 4234 |
| Mycobacterium tuberculosis TBDM1506 | $NZ_{-}CP023631$ | 4411157 | 4230 |
| Mycobacterium tuberculosis TBDM2189 | $NZ_{-}CP023632$ | 4411316 | 4234 |
| Mycobacterium tuberculosis TBDM2444 | $NZ_{-}CP023633$ | 4411284 | 4237 |
| Mycobacterium tuberculosis TBDM2487 | $NZ_{-}CP023634$ | 4411314 | 4237 |
| Mycobacterium tuberculosis TBDM2489 | $NZ_{-}CP023635$ | 4411369 | 4238 |
| Mycobacterium tuberculosis TBDM2699 | $NZ_{-}CP023636$ | 4411457 | 4235 |
| Mycobacterium tuberculosis TBDM2717 | $NZ_{-}CP023637$ | 4411442 | 4236 |
| Mycobacterium tuberculosis TBDM425 | $NZ_{-}CP023588$ | 4411143 | 4235 |
| Mycobacterium tuberculosis TBV4766 | $NZ_{-}CP023638$ | 4411310 | 4234 |
| Mycobacterium tuberculosis TBV4768 | $NZ_{-}CP023639$ | 4411173 | 4229 |
| Mycobacterium tuberculosis TBV4952 | $NZ_{-}CP023640$ | 4411414 | 4234 |
| Mycobacterium tuberculosis TBV5000 | $NZ_{-}CP023589$ | 4411318 | 4234 |
| Mycobacterium tuberculosis TBV5362 | $NZ_{-}CP023590$ | 4411331 | 4235 |
| Mycobacterium tuberculosis TBV5365 | $NZ_{-}CP023591$ | 4411398 | 4234 |
| Mycobacteroides abscessus FLAC003 | $NZ_{-}CP014950$ | 4826045 | 4724 |
| Mycobacteroides abscessus FLAC005 | $NZ_{-}CP014952$ | 4869298 | 4772 |
| Mycobacteroides abscessus FLAC006 | $NZ_{-}CP016188$ | 4891993 | 4810 |
| Mycobacteroides abscessus FLAC013 | $NZ_{-}CP014955$ | 5074222 | 5013 |
| Mycobacteroides abscessus FLAC028 | NZ_CP016190 | 5188101 | 5151 |
| Mycobacteroides abscessus FLAC029 | $NZ_{-}CP014956$ | 5188507 | 5151 |
| Mycobacteroides abscessus FLAC031 | $NZ_{-}CP014957$ | 5146255 | 5061 |
| Mycobacteroides abscessus FLAC046 | $NZ_{-}CP016192$ | 5214168 | 5180 |
| Mycobacteroides abscessus FLAC049 | $NZ_{-}CP014960$ | 4799801 | 4704 |
| Salmonella entericaTyphi 311189-201186 | $NZ_{-}CP029958$ | 4783542 | 4778 |
| Salmonella entericaTyphi 311189-202186 | $NZ_{-}CP029956$ | 4799946 | 4809 |
| Salmonella entericaTyphi 311189-203186 | $NZ_{-}CP029942$ | 4786232 | 4783 |
| Salmonella entericaTyphi 311189-204186 | $NZ_{-}CP029954$ | 4784767 | 4784 |
| Salmonella entericaTyphi 311189-205186 | $NZ_{-}CP029950$ | 4821321 | 4846 |
| Salmonella entericaTyphi 311189-206186 | $NZ_{-}CP029940$ | 4786487 | 4782 |
| Salmonella entericaTyphi 311189-207186 | $NZ_{-}CP029938$ | 4784615 | 4778 |
| Salmonella entericaTyphi 311189-208103 | $NZ_{-}CP029932$ | 4783092 | 4776 |
| Salmonella entericaTyphi 311189-208186 | $NZ_{-}CP029949$ | 4783337 | 4775 |
| Salmonella entericaTyphi 311189-209186 | $NZ_{-}CP029952$ | 4782252 | 4775 |
| Salmonella entericaTyphi 311189-210186 | $NZ_{-}CP029946$ | 4785705 | 4784 |
| Salmonella entericaTyphi 311189-211186 | $NZ_{-}CP029945$ | 4786194 | 4782 |
| Salmonella entericaTyphi 311189-212186 | NZ_CP029944 | 4784815 | 4777 |
| Salmonella entericaTyphi 311189-213186 | $NZ_{-}CP029936$ | 4783243 | 4777 |
| Salmonella entericaTyphi 311189-214186 | $NZ_{-}CP029933$ | 4782854 | 4777 |

continued table S3

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|--|----------------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Salmonella entericaTyphi 311189-215186 | NZ_CP029930 | 4782608 | 4774 |
| Salmonella entericaTyphi 311189-216103 | NZ_CP029928 | 4799426 | 4803 |
| Salmonella entericaTyphi 311189-217103 | $NZ_{-}CP029927$ | 4778473 | 4771 |
| Salmonella entericaTyphi 311189-217186 | NZ_CP029646 | 4782348 | 4781 |
| Salmonella entericaTyphi 311189-218186 | $NZ_{-}CP029925$ | 4786711 | 4783 |
| Salmonella entericaTyphi 311189-219186 | NZ_CP029911 | 4784370 | 4779 |
| Salmonella entericaTyphi 311189-220186 | $NZ_{-}CP029886$ | 4783528 | 4775 |
| Salmonella entericaTyphi 311189-221186 | $NZ_{-}CP029923$ | 4782756 | 4779 |
| Salmonella entericaTyphi 311189-222186 | $NZ_{CP029909}$ | 4842715 | 4867 |
| Salmonella enterica Typhi 311189-223186 | $NZ_{-}CP029880$ | 4783762 | 4778 |
| Salmonella entericaTyphi 311189-224186 | $NZ_{-}CP029878$ | 4782665 | 4776 |
| Salmonella entericaTyphi 311189-231186 | $NZ_{-}CP029904$ | 4807430 | 4815 |
| Salmonella entericaTyphi 311189-232103 | $NZ_{-}CP029918$ | 4778368 | 4765 |
| Salmonella entericaTyphi 311189-239103 | $NZ_{-}CP029908$ | 4778864 | 4768 |
| Salmonella entericaTyphi 311189-252186 | $NZ_{-}CP029896$ | 4781801 | 4773 |
| Salmonella entericaTyphi 311189-255186 | $NZ_{-}CP029885$ | 4808675 | 4824 |
| Salmonella entericaTyphi 311189-256186 | $NZ_{-}CP029917$ | 4800376 | 4800 |
| Salmonella entericaTyphi 311189-268103 | $NZ_{-}CP029902$ | 4804086 | 4815 |
| Salmonella enterica Typhi 311189-268186 | NZ_CP029883 | 4807681 | 4815 |
| Salmonella enterica Typhi 311189-269186 | $NZ_{-}CP029922$ | 4782083 | 4773 |
| Salmonella entericaTyphi 311189-282186 | $NZ_{-}CP029920$ | 4799956 | 4800 |
| Salmonella enterica Typhi 311189-291186 | $NZ_{-}CP029894$ | 4806999 | 4815 |
| Salmonella enterica Typhi 343076-202113 | $NZ_{-}CP029915$ | 4799269 | 4801 |
| Salmonella enterica Typhi 343076-217103 | $NZ_{-}CP029914$ | 4775784 | 4776 |
| Salmonella enterica Typhi 343076-227128 | $NZ_{-}CP029875$ | 4800199 | 4800 |
| Salmonella entericaTyphi 343076-232188 | $NZ_{-}CP029900$ | 4800621 | 4803 |
| Salmonella enterica Typhi 343076-241106 | $NZ_{-}CP029899$ | 4775702 | 4775 |
| Salmonella enterica Typhi 343076-248190 | $NZ_{-}CP029882$ | 4773823 | 4777 |
| Salmonella enterica Typhi 343076-249107 | $NZ_{-}CP029913$ | 4783891 | 4776 |
| Salmonella entericaTyphi 343076-252143 | $NZ_{-}CP029892$ | 4808285 | 4817 |
| Salmonella entericaTyphi 343076-253155 | NZ_CP029890 | 4791587 | 4800 |
| Salmonella entericaTyphi 343076-269157 | NZ_CP029881 | 4791108 | 4798 |
| Salmonella entericaTyphi 343076-294172 | NZ_CP029888 | 4809171 | 4820 |
| Salmonella entericaTyphi 343077-211171 | NZ_CP029873 | 4782825 | 4777 |
| Salmonella enterica Typhi 343077-212138 | NZ_CP029919 | 4774558 | 4773 |
| Salmonella entericaTyphi 343077-212159 | NZ_CP029870 | 4795465 | 4804 |
| Salmonella entericaTyphi 343077-213147 | NZ_CP029897 | 4897593 | 4940 |
| Salmonella entericaTyphi 343077-214135 | NZ_CP029872 | 4783204 | 4774 |
| Salmonella entericaTyphi 343077-214162 | NZ_CP029862 | 4814028 | 4816 |
| Salmonella entericaTyphi 343077-215174 | NZ_CP029868 | 4785665 | 4779 |
| Salmonella entericaTyphi 343077-228140 | NZ_CP029866 | 4790593 | 4799 |
| Salmonella entericaTyphi 343077-228157 | NZ_CP029864 | 4807408 | 4818 |
| Salmonella entericaTyphi 343077-255118 | NZ_CP029907 | 4774500 | 4776 |
| Salmonella enterica Typhi 343077-260153 | NZ_CP029861 | 4783691 | 4778 |
| Salmonella entericaTyphi 343077-267164 | NZ_CP029906 | 4775386 | 4775 |
| Salmonella enterica Typhi 343077-278127 | NZ_CP029863 | 4807245 | 4816 |
| Salmonella entericaTyphi 343077-281186 | NZ_CP029853 | 4829489 | 4852 |
| Salmonella enterica Typhi 343077-285138 | NZ_CP029858 | 4800928 | 4805 |
| Salmonella enterica Typhi 343077-286126 | NZ_CP029856 | 4797628 | 4804 |
| Salmonella entericaTyphi 343077-292148 | NZ_CF029855 NZ_CP029855 | 4800586 | 4802 |
| Salmonella enterica Typhi 343077-292148 Salmonella enterica Typhi 343078-201101 | NZ_CP029853 NZ_CP029852 | 4782235 | 4774 |
| Salmonella enterica Typhi 343078-201101 Salmonella enterica Typhi 343078-203125 | | | |
| Dannonena enterica i ypiil 545076-205125 | $NZ_{-}CP029850$ | 4801819 | 4803 |

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|---|----------------------------|--------------------|---------------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Salmonella entericaTyphi 343078-211126 | NZ_CP029848 | 4799748 | 4803 |
| Salmonella entericaTyphi 343078-223175 | $NZ_{-}CP029964$ | 4782707 | 4780 |
| Salmonella entericaTyphi 343078-228140 | $NZ_{-}CP029962$ | 4783156 | 4775 |
| Salmonella entericaTyphi 343078-251131 | $NZ_{-}CP029960$ | 4799747 | 4804 |
| Salmonella entericaTyphi 343078-256191 | $NZ_{-}CP029959$ | 4795890 | 4803 |
| Salmonella entericaTyphi 343078-273110 | NZ_CP029846 | 4801146 | 4804 |
| Staphylococcus aureus AR461 | NZ_CP029087 | 2909586 | 2984 |
| Staphylococcus aureus AR466 | NZ_CP029080 | 2786146 | 2777 |
| Staphylococcus aureus AR-0215 | NZ_CP029680 | 2910130 | 2986 |
| Staphylococcus aureus AR-0216 | NZ_CP029678 | 2921067 | 3009 |
| Staphylococcus aureus AR-0219 | NZ_CP029675 | 2878431 | 2935 |
| Staphylococcus aureus AR-0220 | NZ_CP029673 | 2828139 | 2848 |
| Staphylococcus aureus AR-0222 | NZ_CP029671 | 2692211 | 2670 |
| Staphylococcus aureus AR-0225 | NZ_CP029667 | 2897877 | 2981 |
| Staphylococcus aureus AR-0228 | NZ_CP029663 | 2867357 | 2926 |
| Staphylococcus aureus AR-0467 | NZ_CP029658 | 2910633 | 2989 |
| Staphylococcus aureus AR-0468 | NZ_CP029657 | 2911284 | 2983 |
| Staphylococcus aureus AR-0470 | NZ_CP029653 | 2963657 | 2964 |
| Staphylococcus aureus AR-474 | NZ_CP030326 | 2906506 | 2977 |
| Staphylococcus aureus CFSAN007835 | NZ_CP017685 | 2676924 | 2634 |
| Staphylococcus aureus CFSAN007847 | NZ_CP017684 | 2857060 | 2871 |
| Staphylococcus aureus CFSAN007850 | NZ_CP017682 | 2786129 | 2798 |
| Staphylococcus aureus CFSAN007850 Staphylococcus aureus CFSAN007851 | NZ_CP017680 | 2748822 | $\frac{2798}{2751}$ |
| Staphylococcus aureus CFSAN007883 | NZ_CP017680 NZ_CP017679 | 2758960 | $\frac{2731}{2729}$ |
| Staphylococcus aureus CFSAN007894 | NZ_CP017679 NZ_CP017677 | | $\frac{2729}{2723}$ |
| | NZ_CP017077 NZ_CP020467 | 2730395 2855675 | 2860 |
| Staphylococcus aureus CFSAN007896 | NZ_CP020467 NZ_CP028165 | 2855075 2790673 | |
| Staphylococcus aureus CFSAN064037 | | | $2738 \\ 2823$ |
| Staphylococcus aureus FDAARGOS-48 | NZ_CP026953 | 2834575 | |
| Staphylococcus aureus FDAARGOS-6 | NZ_CP026962 | 2697644 | 2634 |
| Staphylococcus aureus USA300-SUR1 | NZ_CP014362 | 2904958 | 2929 |
| Staphylococcus aureus USA300-SUR10 | NZ_CP014397 | 2918010 | 2940 |
| Staphylococcus aureus USA300-SUR11 | NZ_CP014402 | 2918010 | 2940 |
| Staphylococcus aureus USA300-SUR12 | NZ_CP014407 | 2876325 | 2878 |
| Staphylococcus aureus USA300-SUR13 | NZ_CP014409 | 2919720 | 2945 |
| Staphylococcus aureus USA300-SUR14 | NZ_CP014412 | 2919827 | 2944 |
| Staphylococcus aureus USA300-SUR15 | NZ_CP014415 | 2919910 | 2943 |
| Staphylococcus aureus USA300-SUR16 | NZ_CP014420 | 2919827 | 2943 |
| Staphylococcus aureus USA300-SUR17 | NZ_CP014423 | 2919826 | 2944 |
| Staphylococcus aureus USA300-SUR18 | NZ_CP014426 | 2919827 | 2943 |
| Staphylococcus aureus USA300-SUR19 | $NZ_{-}CP014429$ | 2919828 | 2944 |
| Staphylococcus aureus USA300-SUR2 | $NZ_{-}CP014365$ | 2904958 | 2929 |
| Staphylococcus aureus USA300-SUR20 | $NZ_{-}CP014432$ | 2915434 | 2937 |
| Staphylococcus aureus USA300-SUR21 | $NZ_{-}CP014435$ | 2919720 | 2946 |
| Staphylococcus aureus USA300-SUR22 | $NZ_{-}CP014438$ | 2915433 | 2936 |
| Staphylococcus aureus USA300-SUR23 | $NZ_{-}CP014441$ | 2915433 | 2936 |
| Staphylococcus aureus USA300-SUR24 | NZ_CP014444 | 2919910 | 2943 |
| Staphylococcus aureus USA300-SUR3 | $NZ_{-}CP014368$ | 2919706 | 2945 |
| Staphylococcus aureus USA300-SUR4 | NZ_CP014371 | 2918010 | 2940 |
| Staphylococcus aureus USA300-SUR5 | $NZ_{-}CP014376$ | 2918010 | 2940 |
| Staphylococcus aureus USA300-SUR6 | $NZ_{-}CP014381$ | 2919719 | 2944 |
| Staphylococcus aureus USA300-SUR7 | NZ_CP014384 | 2919670 | 2943 |
| Staphylococcus aureus USA300-SUR9 | $NZ_CP014392$ | 2918010 | 2940 |

| Name | Accession Num. | Sequence Length | CDS number |
|--|------------------|-----------------|------------|
| Staphylococcus aureus USA400-0051 | NZ_CP019574 | 2811886 | 2844 |
| Vibrio cholerae 2740-80 | NZ_CP016324 | 2987030 | 2676 |
| Vibrio cholerae 2012Env-9 | NZ_CP012997 | 1124711 | 1178 |
| Vibrio cholerae A1552 | NZ_CP025936 | 3015092 | 2713 |
| Vibrio cholerae A1552 I | NZ_CP024867 | 3015093 | 2702 |
| Vibrio cholerae C5 | NZ_CP013301 | 2996195 | 2682 |
| Vibrio cholerae CRC711 | NZ_CP013303 | 2993255 | 2687 |
| Vibrio cholerae E4 | NZ_CP033515 | 3075249 | 2766 |
| Vibrio cholerae E506 | NZ_CP013307 | 2969901 | 2659 |
| Vibrio cholerae E7946 | NZ_CP024162 | 2992571 | 2701 |
| Vibrio cholerae E9120 | NZ_CP013313 | 2961701 | 2664 |
| Vibrio cholerae M2140 | NZ_CP013315 | 2939488 | 2615 |
| Vibrio cholerae NCTC5395 | $NZ_{CP013317}$ | 2981414 | 2670 |
| Vibrio cholerae NCTC9420 | NZ_CP013319 | 3008486 | 2695 |
| Yersinia pestis412 | NZ_CP006783 | 4524943 | 4140 |
| Yersinia pestis CO92 | $NC_{-}003143$ | 4653728 | 3798 |
| Yersinia pestis CO92 | $NZ_{-}CP009973$ | 4556337 | 4195 |
| Yersinia pestis D106004 | $NC_{-}017154$ | 4640720 | 4237 |
| Yersinia pestis D182038 | NC_017160 | 4626944 | 4225 |
| Yersinia pestis Nepal516 | NC_008149 | 4534590 | 4145 |
| Yersinia pestis Pestoides G | $NZ_{-}CP010247$ | 4529728 | 4140 |
| Yersinia pestis Z176003 | NC_014029 | 4553586 | 4184 |
| Yersinia pestis biovar Medievalis Harbin 35 | $NC_{-}017265$ | 4532063 | 4171 |
| Yersinia pestis Pestoides B | $NZ_{-}CP010023$ | 4607660 | 4180 |
| Yersinia pestis FDAARGOS-601 | $NZ_{-}CP033699$ | 4708493 | 4318 |
| Yersinia pestis SCPM-O-B-5935 (I-1996) | NZ_CP045154 | 4629837 | 4207 |
| Yersinia pestis SCPM-O-B-6291 (C-25) | NZ_CP045163 | 4503487 | 4109 |
| Yersinia pestis SCPM-O-DNA-18 (I-3113) | NZ_CP045149 | 4546217 | 4153 |
| Yersinia pestis subsp pestis by Medievalis SCPM-O-B-6530 | NZ_CP045158 | 4596548 | 4197 |

Table S4: The third dataset

| Name | Accession Num. | Sequence Length | CDS number |
|------------------------------------|------------------|-----------------|------------|
| Pseudomonas aeruginosa DHS01 | NZ_CP013993 | 7055752 | 6828 |
| Pseudomonas aeruginosa DK2 | NC_018080 | 6402658 | 6036 |
| Pseudomonas aeruginosa DSM 50071 | NZ_CP012001 | 6317050 | 5900 |
| Pseudomonas aeruginosa LES431 | $NC_{-}023066$ | 6550070 | 6144 |
| Pseudomonas aeruginosa LESB58 | $NC_{-}011770$ | 6601757 | 6228 |
| Pseudomonas aeruginosa M18 | $NC_{-}017548$ | 6327754 | 5935 |
| Pseudomonas aeruginosa MTB-1 | NC_023019 | 6580038 | 6213 |
| Pseudomonas aeruginosa NCGM2S1 DNA | $NC_{-}017549$ | 6764661 | 6420 |
| Pseudomonas aeruginosa PA1 | $NC_{-}022808$ | 6498072 | 6121 |
| Pseudomonas aeruginosa PA7 | $NC_{-}009656$ | 6588339 | 6233 |
| Pseudomonas aeruginosa SCV20265 | $NC_{-}023149$ | 6725183 | 6371 |
| Pseudomonas aeruginosa UCBPP-PA14 | NC_008463 | 6537648 | 6112 |
| Pseudomonas aeruginosa 12939 | $NZ_{-}CP024477$ | 6621378 | 6270 |
| Pseudomonas aeruginosa 1334/14 | $NZ_{-}CP035739$ | 6902135 | 6556 |
| Pseudomonas aeruginosa 24Pae112 | $NZ_{-}CP029605$ | 7097241 | 6774 |
| Pseudomonas aeruginosa 60503 | NZ_CP041774 | 6809062 | 6469 |
| Pseudomonas aeruginosa 97 | $NZ_{-}CP031449$ | 6925889 | 6610 |
| Pseudomonas aeruginosa A681 | NZ_CP041771 | 6616247 | 6272 |

continued table S4

| Name Accession Num. Sequence Length CDS number Pseudomonas aeruginosa AES1R NZ_CP037926 6373893 6015 Pseudomonas aeruginosa AR441 NZ_CP029093 6584825 6207 Pseudomonas aeruginosa AR445 NZ_CP029088 7125975 6796 Pseudomonas aeruginosa AR-0110 NZ_CP029745 6799785 6416 Pseudomonas aeruginosa ATCC 15692 NZ_CP017149 6276434 5846 Pseudomonas aeruginosa ATCC 27853 NZ_CP015117 6827737 6473 Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CR1 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa F23197 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F5677 |
|--|
| Pseudomonas aeruginosa AR441 NZ_CP029093 6584825 6207 Pseudomonas aeruginosa AR445 NZ_CP029088 7125975 6796 Pseudomonas aeruginosa AR-0110 NZ_CP029745 6799785 6416 Pseudomonas aeruginosa ATCC 15692 NZ_CP017149 6276434 5846 Pseudomonas aeruginosa ATCC 27853 NZ_CP015117 6827737 6473 Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CR1 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa AR445 NZ_CP029088 7125975 6796 Pseudomonas aeruginosa AR-0110 NZ_CP029745 6799785 6416 Pseudomonas aeruginosa ATCC 15692 NZ_CP017149 6276434 5846 Pseudomonas aeruginosa ATCC 27853 NZ_CP015117 6827737 6473 Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CR1 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa AR-0110 NZ_CP029745 6799785 6416 Pseudomonas aeruginosa ATCC 15692 NZ_CP017149 6276434 5846 Pseudomonas aeruginosa ATCC 27853 NZ_CP015117 6827737 6473 Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa F23197 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa ATCC 15692 NZ_CP017149 6276434 5846 Pseudomonas aeruginosa ATCC 27853 NZ_CP015117 6827737 6473 Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa F90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa ATCC 27853 NZ_CP015117 6827737 6473 Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa ATCC 27853 NZ_CP011857 6833187 6463 Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa AZPAE15042 NZ_CP041354 6527298 6169 Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa B14130 NZ_CP034435 6759594 6340 Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa CCUG 51971 NZ_CP043328 7012798 6690 Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa CCUG 70744 NZ_CP023255 6859232 6533 Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa CR1 NZ_CP020560 6118054 5712 Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa E80 NZ_CP031677 6819140 6462 Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa E90 NZ_CP044006 6813135 6463 Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| Pseudomonas aeruginosa F23197 NZ_CP008856 6517340 6051 |
| |
| 1 Seddonionas del agniosa 1 5077 172201 020000 00 19227 00001 |
| Pseudomonas aeruginosa F63912 NZ_CP008858 6618768 6299 |
| Pseudomonas aeruginosa F9670 NZ_CP008873 6794354 6388 |
| Pseudomonas aeruginosa GIMC5015:PAKB6 NZ_CP034429 6258491 5824 |
| Pseudomonas aeruginosa H25883 NZ_CP033686 6706800 6337 |
| Pseudomonas aeruginosa H26023 NZ_CP033685 6729216 6347 |
| - |
| |
| |
| Pseudomonas aeruginosa H47921 NZ_CP008861 6836415 6468 |
| Pseudomonas aeruginosa H5708 NZ_CP008859 6334378 5906 |
| Pseudomonas aeruginosa HOU1 NZ_CP042269 6123373 5701 |
| Pseudomonas aeruginosa HS9 NZ_CP030861 6876988 6532 |
| Pseudomonas aeruginosa IMP66 NZ_CP028959 6486336 6093 |
| Pseudomonas aeruginosa IMP67 NZ_CP028848 6501414 6113 |
| Pseudomonas aeruginosa IMP68 NZ_CP028849 6481697 6086 |
| Pseudomonas aeruginosa L10 NZ_CP019338 6661962 6217 |
| Pseudomonas aeruginosa PA11803 NZ_CP015003 7006578 6738 |
| Pseudomonas aeruginosa PA298 NZ_CP040127 6539894 6141 |
| Pseudomonas aeruginosa PA34 NZ_CP032552 6810079 6462 |
| Pseudomonas aeruginosa PA7790 NZ_CP014999 7018690 6681 |
| Pseudomonas aeruginosa PA8281 NZ_CP015002 6928736 6562 |
| Pseudomonas aeruginosa PA83 NZ_CP017293 6816227 6385 |
| Pseudomonas aeruginosa PABL017 NZ_CP031660 6503460 6091 |
| Pseudomonas aeruginosa PABL048 NZ_CP039293 6879622 6522 |
| Pseudomonas aeruginosa PAER4-119 NZ_CP013113 6504659 6056 |
| Pseudomonas aeruginosa PAO1161 NZ_CP032126 6383803 5962 |
| Pseudomonas aeruginosa PA-150577 NZ_CP017306 6334472 5928 |
| Pseudomonas aeruginosa PA-D16 NZ_CP012581 6681975 6247 |
| Pseudomonas aeruginosa PA-D2 NZ_CP012578 6642996 6204 |
| Pseudomonas aeruginosa PA-D21 NZ_CP012582 6639108 6202 |
| Pseudomonas aeruginosa PA-D22 NZ_CP012583 6681981 6248 |
| Pseudomonas aeruginosa PA-D25 NZ_CP012584 6683204 6250 |
| Pseudomonas aeruginosa PA-D5 NZ_CP012579 6681992 6248 |
| Pseudomonas aeruginosa PB350 NZ_CP025055 6752870 6340 |
| Pseudomonas aeruginosa PB354 NZ_CP025053 6434352 6066 |
| Pseudomonas aeruginosa S86968 NZ_CP008865 6934277 6577 |
| Pseudomonas aeruginosa SCVFeb NZ_CP013477 6213026 5777 |

| Name | Accession Num. | Sequence Length | CDS number |
|--|------------------|-----------------|------------|
| Pseudomonas aeruginosa T38079 | NZ_CP008866 | 6795741 | 6391 |
| Pseudomonas aeruginosa W16407 | NZ_CP008869 | 6808844 | 6421 |
| Pseudomonas aeruginosa Y89 | NZ_CP030913 | 6868832 | 6493 |
| Pseudomonas fluorescens L321 | NZ_CP015637 | 6641144 | 5984 |
| Pseudomonas fluorescens Pf275 | NZ_CP031648 | 6610362 | 6025 |
| Pseudomonas fluorescens Pt14 | $NZ_{-}CP017296$ | 5841722 | 5274 |
| Pseudomonas fluorescens SIK-W1 | NZ_CP031450 | 6791087 | 6112 |
| Pseudomonas putida GB-1 | $NC_{-}010322$ | 6078430 | 5515 |
| Pseudomonas putida H8234 | $NC_{-}021491$ | 6870827 | 6498 |
| Pseudomonas putida JB | $NZ_{-}CP016212$ | 5846746 | 5380 |
| Pseudomonas putida KT2440 | $NC_{-}002947$ | 6181873 | 5564 |
| Pseudomonas putida NBRC 14164 DNA | $NC_{-}021505$ | 6156701 | 5622 |
| Pseudomonas putida ND6 | $NC_{-}017986$ | 6085449 | 5625 |
| Pseudomonas putida SJTE-1 | $NZ_{-}CP015876$ | 6077201 | 5606 |
| Pseudomonas putida B4 | $NZ_CP022561$ | 6011680 | 5525 |
| Pseudomonas putida E41 | $NZ_{-}CP024085$ | 6093023 | 5430 |
| Pseudomonas putida E46 | $NZ_{CP024086}$ | 6180967 | 5403 |
| Pseudomonas putida PP112420 | $NZ_CP017073$ | 6031212 | 5415 |
| Pseudomonas stutzeri A1501 | $NC_{-}009434$ | 4567418 | 4206 |
| Pseudomonas stutzeri CCUG 29243 | $NC_{-}018028$ | 4709064 | 4356 |
| Pseudomonas stutzeri DSM 10701 | $NC_{-}018177$ | 4174118 | 3861 |
| Pseudomonas stutzeri DSM 4166 | NC_017532 | 4689946 | 4366 |
| Pseudomonas stutzeri RCH2 | NC_019936 | 4575057 | 4241 |
| Pseudomonas stutzeri SGAir0442 | $NZ_{-}CP025149$ | 4524655 | 4241 |
| Pseudomonas syringae pv actinidiae ICMP 18884 | $NZ_{CP011972}$ | 6555569 | 6272 |
| Pseudomonas syringae pv actinidiae ICMP 9853 | NZ_CP018202 | 6439609 | 6047 |
| Pseudomonas syringae pv actinidiae CRAFRU 1229 | NZ_CP019730 | 6545549 | 6145 |
| Pseudomonas syringae pv actinidiae CRAFRU 1408 | $NZ_{-}CP019732$ | 6546028 | 6148 |
| Pseudomonas syringae pv actinidiae MAFF212063 | $NZ_{CP024712}$ | 6556999 | 6191 |
| Pseudomonas syringae pv actinidiae NZ-45 | $NZ_{-}CP017007$ | 6665031 | 6285 |
| Pseudomonas syringae pv actinidiae NZ-47 | NZ_CP017009 | 6545910 | 6157 |

Table S5: The fourth dataset

| Name | Accession Num. | Sequence Length | CDS number |
|------------------------------------|------------------|-----------------|------------|
| Streptococcus agalactiae NGBS061 | NZ_CP007631 | 2221207 | 2220 |
| Streptococcus agalactiae NGBS572 | $NZ_{-}CP007632$ | 2061426 | 2023 |
| Streptococcus agalactiae GBS6 | $NZ_{-}CP007572$ | 2231475 | 2210 |
| Streptococcus agalactiae GBS1-NY | $NZ_{-}CP007570$ | 2243708 | 2210 |
| Streptococcus agalactiae GBS2-NM | NZ_CP007571 | 2214297 | 2186 |
| Streptococcus agalactiae SS1 | NZ_CP010867 | 2092071 | 2090 |
| Streptococcus agalactiae H002 | $NZ_{-}CP011329$ | 2147416 | 2138 |
| Streptococcus agalactiae HN016 | $NZ_{-}CP011325$ | 2064722 | 2037 |
| Streptococcus agalactiae GBS85147 | NZ_CP010319 | 1996151 | 1972 |
| Streptococcus agalactiae GBS ST-1 | $NZ_{-}CP013202$ | 2165968 | 2178 |
| Streptococcus agalactiae SG-M1 | NZ_CP012419 | 2116811 | 2143 |
| Streptococcus agalactiae NGBS128 | $NZ_{-}CP012480$ | 2074179 | 2036 |
| Streptococcus agalactiae CU-GBS-98 | $NZ_{-}CP010875$ | 2029669 | 2041 |
| Streptococcus agalactiae CU-GBS-08 | NZ_CP010874 | 2084511 | 2113 |
| Streptococcus agalactiae FWL1402 | NZ_CP016391 | 2090292 | 2082 |
| Streptococcus agalactiae NGBS357 | $NZ_{-}CP012503$ | 2172875 | 2144 |

continued table S5

| continued table S5 | | G . | GDG ' |
|--|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Streptococcus agalactiae Sag158 | NZ_CP019979 | 2096882 | 2086 |
| Streptococcus agalactiae GBS-M002 | NZ_CP013908 | 2092570 | 2065 |
| Streptococcus agalactiae Sag37 | NZ_CP019978 | 2198785 | 2188 |
| Streptococcus agalactiae SG-M8 | NZ_CP021868 | 2167120 | 2163 |
| Streptococcus agalactiae CUGBS591 | NZ_CP021862 | 2227680 | 2258 |
| Streptococcus agalactiae SG-M29 | $NZ_{-}CP021866$ | 2116773 | 2142 |
| Streptococcus agalactiae SG-M163 | $NZ_{-}CP021863$ | 2116810 | 2141 |
| Streptococcus agalactiae SG-M50 | $NZ_{-}CP021865$ | 2116810 | 2143 |
| Streptococcus agalactiae SG-M25 | $NZ_{-}CP021867$ | 2208337 | 2226 |
| Streptococcus agalactiae SG-M6 | $NZ_{-}CP021869$ | 2106018 | 2112 |
| Streptococcus agalactiae SG-M158 | NZ_CP021864 | 2116811 | 2143 |
| Streptococcus agalactiae SG-M4 | $NZ_{-}CP021870$ | 2072007 | 2049 |
| Streptococcus agalactiae C001 | $NZ_{-}CP008813$ | 2121372 | 2098 |
| Streptococcus agalactiae 874391 | $NZ_{-}CP022537$ | 2153937 | 2119 |
| Streptococcus agalactiae SGEHI2015-25 | $NZ_{-}CP025029$ | 2054713 | 2069 |
| Streptococcus agalactiae SGEHI2015-113 | $NZ_{-}CP025026$ | 2043851 | 2052 |
| Streptococcus agalactiae SGEHI2015-107 | $NZ_{-}CP025027$ | 2016805 | 2015 |
| Streptococcus agalactiae SGEHI2015-95 | $NZ_{-}CP025028$ | 2116769 | 2143 |
| Streptococcus agalactiae SA79 | NZ_CP019809 | 1841946 | 1857 |
| Streptococcus agalactiae B509 | $NZ_{-}CP021769$ | 2060637 | 2054 |
| Streptococcus agalactiae S73 | $NZ_{-}CP030845$ | 2059915 | 2085 |
| Streptococcus agalactiae GX026 | $NZ_{-}CP011328$ | 1840649 | 1843 |
| Streptococcus agalactiae S25 | $NZ_{-}CP015976$ | 1838989 | 1847 |
| Streptococcus agalactiae SA20 | NC_019048 | 1841952 | 1855 |
| Streptococcus agalactiae SA16 | $NZ_{-}CP019807$ | 1841859 | 1856 |
| Streptococcus agalactiae SA209 | $NZ_{-}CP019822$ | 1841857 | 1857 |
| Streptococcus agalactiae SA289 | $NZ_{-}CP019828$ | 1848987 | 1862 |
| Streptococcus agalactiae SA75 | $NZ_{-}CP019808$ | 1849016 | 1860 |
| Streptococcus agalactiae SA201 | $NZ_{-}CP019821$ | 1841835 | 1853 |
| Streptococcus agalactiae SA220 | $NZ_{-}CP019825$ | 1841963 | 1854 |
| Streptococcus agalactiae SA245 | $NZ_{-}CP019826$ | 1848956 | 1859 |
| Streptococcus agalactiae S13 | $NZ_{-}CP018623$ | 1835156 | 1854 |
| Streptococcus agalactiae SA256 | $NZ_{-}CP019827$ | 1848972 | 1862 |
| Streptococcus agalactiae SA132 | $NZ_{-}CP019815$ | 1852032 | 1864 |
| Streptococcus agalactiae SA333 | $NZ_{-}CP019830$ | 1842037 | 1854 |
| Streptococcus agalactiae SA346 | $NZ_{-}CP019833$ | 1841984 | 1857 |
| Streptococcus agalactiae SA1 | $NZ_{-}CP019804$ | 1841943 | 1855 |
| Streptococcus agalactiae SA81 | NZ_CP019810 | 1840363 | 1854 |
| Streptococcus agalactiae SA218 | $NZ_{-}CP019824$ | 1849985 | 1862 |
| Streptococcus agalactiae SA343 | $NZ_{-}CP019832$ | 1841977 | 1855 |
| Streptococcus agalactiae SA627 | $NZ_{-}CP019837$ | 1842066 | 1855 |
| Streptococcus agalactiae SA212 | $NZ_{-}CP019823$ | 1841962 | 1855 |
| Streptococcus agalactiae SA623 | NZ_CP019836 | 1842115 | 1857 |
| Streptococcus agalactiae SA53 | $NZ_{-}CP019802$ | 1848971 | 1863 |
| Streptococcus agalactiae SA5 | $NZ_{-}CP019805$ | 1841945 | 1854 |
| Streptococcus agalactiae SA95 | $NZ_{-}CP019812$ | 1856590 | 1866 |
| Streptococcus agalactiae SA9 | $NZ_{-}CP019806$ | 1841929 | 1855 |
| Streptococcus agalactiae 32790-3A | $NZ_{-}CP029561$ | 2148904 | 2131 |
| Streptococcus agalactiae PLGBS13 | $NZ_{-}CP029749$ | 2095031 | 2098 |
| Streptococcus agalactiae SA30 | $NZ_{-}CP019800$ | 1841729 | 1856 |
| Streptococcus agalactiae SA33 | $NZ_{-}CP019801$ | 1841628 | 1857 |
| Streptococcus agalactiae SA191 | NZ_CP019819 | 1848676 | 1864 |

continued table S5

| Name | continued table S5 | | | |
|--|------------------------------------|-----------------|-----------------|------------|
| Streptococcus agalactiae SA374 NZ. CP019831 1842153 1856 Streptococcus agalactiae SA45 NZ. CP019811 184989 1859 Streptococcus agalactiae SA155 NZ. CP019803 1841715 1858 Streptococcus agalactiae SA136 NZ. CP019816 1841715 1858 Streptococcus agalactiae SA136 NZ. CP019816 1849103 1846 Streptococcus agalactiae SA370 NZ. CP019813 1856410 1848 Streptococcus agalactiae SA373 NZ. CP019829 1842081 1856 Streptococcus agalactiae SA373 NZ. CP019829 1842081 1856 Streptococcus agalactiae CA375 NZ. CP019829 1842081 1856 Streptococcus agalactiae COCTC 10/84 NZ. CP009835 1812219 1856 Streptococcus agalactiae CNCTC 10/84 NZ. CP006910 2013842 2008 Streptococcus agalactiae CNCTC 10/84 NZ. CP006910 2013842 2008 Streptococcus agalactiae AC-2713 NC. 016646 2063112 2036 Streptococcus dysgalactiae AC-2713 NC. 01666 206312 2179445 2058 <td></td> <td>Accession Num.</td> <td>Sequence Length</td> <td>CDS number</td> | | Accession Num. | Sequence Length | CDS number |
| Streptococcus agalactiae SA341 NZ. CP019811 1841213 1861 Streptococcus agalactiae SA155 NZ. CP019820 1841715 1858 Streptococcus agalactiae SA136 NZ. CP019820 1841715 1858 Streptococcus agalactiae SA36 NZ. CP019816 1848839 1836 Streptococcus agalactiae SA376 NZ. CP019813 1856410 1848 Streptococcus agalactiae SA375 NZ. CP019829 1842081 1856 Streptococcus agalactiae SA375 NZ. CP019835 1842219 1859 Streptococcus agalactiae A090 NC. 004116 2160267 2127 Streptococcus agalactiae CNCTC 10/84 NZ. CP006910 2013842 2008 Streptococcus agalactiae CD201008-001 NC. 018646 2063112 2036 Streptococcus dysgalactiae AC-2713 NC. 019042 2179445 2058 Streptococcus dysgalactiae AC-2713 NC. 019042 2179445 2058 Streptococcus dysgalactiae ATCC 12944 NC. 017567 2159491 2056 Streptococcus dysgalactiae RE378 DNA NC. 018712 2151145 2027 | Streptococcus agalactiae SA184 | NZ_CP019818 | 1841893 | 1858 |
| Streptococcus agalactiae SA85 NZ. CP019810 1841715 1858 Streptococcus agalactiae SA136 NZ. CP019803 1841715 1858 Streptococcus agalactiae SA136 NZ. CP019816 18419103 1848 Streptococcus agalactiae SA379 NZ. CP019813 1856410 1848 Streptococus agalactiae SA330 NZ. CP019829 1842081 1856 Streptococus agalactiae SA375 NZ. CP019835 1842219 1856 Streptococcus agalactiae 2603V/R NC.004116 2160267 2127 Streptococcus agalactiae CNCTC 10/84 NZ. CP006910 2013842 2108 Streptococcus agalactiae CNCTC 10/84 NZ. CP006910 2013842 2008 Streptococcus agalactiae NEM316 NC.004368 2211485 2190 Streptococcus dysgalactiae AC-2713 NC.019042 2179445 2058 Streptococcus dysgalactiae AC-2713 NC.017567 2159491 2056 Streptococcus dysgalactiae AC-2713 NC.017567 2259491 2056 Streptococcus gylactiae AC-2713 NC.017567 2362241 2277 <th< td=""><td>Streptococcus agalactiae SA374</td><td>NZ_CP019834</td><td>1842255</td><td>1856</td></th<> | Streptococcus agalactiae SA374 | NZ_CP019834 | 1842255 | 1856 |
| Streptococcus agalactiae SA195 NZ. CP019820 1841715 1858 Streptococcus agalactiae SA373 NZ. CP019816 184839 1836 Streptococcus agalactiae SA376 NZ. CP019816 1849103 1840 Streptococcus agalactiae SA370 NZ. CP019829 1842081 1856 Streptococcus agalactiae SA375 NZ. CP019835 1842219 1859 Streptococcus agalactiae A376 NC. 004116 2160267 212 Streptococcus agalactiae A375 NZ. CP019835 1842219 1859 Streptococcus agalactiae CNCTC 10/84 NZ. CP006910 2013842 2008 Streptococcus agalactiae CNCTC 10/84 NZ. CP006910 2013842 2008 Streptococcus agalactiae ACNIA NC. 018646 2063112 2036 Streptococcus agalactiae ACC-2713 NC. 019042 2179445 205 Streptococcus dysgalactiae ATCC 12394 NC.017567 2159491 2056 Streptococcus dysgalactiae RE378 DNA NC.018712 2151145 2027 Streptococcus gallolyticus ATCC BAA-2069 NC.01576 2362241 2277 | Streptococcus agalactiae SA341 | NZ_CP019831 | 1842113 | 1861 |
| Streptococcus agalactiae SA73 NZ. CP019803 1848839 1836 Streptococcus agalactiae SA136 NZ. CP019813 1856410 1848 Streptococcus agalactiae SA37 NZ. CP019813 1856410 1848 Streptococcus agalactiae SA375 NZ. CP019829 1842081 1856 Streptococcus agalactiae SA375 NZ. CP019829 1842081 1859 Streptococcus agalactiae SA375 NZ. CP019823 184221 1859 Streptococcus agalactiae SA375 NZ. CP019823 184221 1859 Streptococcus agalactiae 2603V/R NC. 004116 2160267 2127 Streptococcus agalactiae CNCTC 10/84 NZ. CP000910 2013842 2008 Streptococcus agalactiae CNCTC 10/84 NZ. CP000910 2013842 2008 Streptococcus agalactiae CNCTC 10/84 NZ. CP000910 2013842 2008 Streptococcus agalactiae CD201008-001 NC. 018646 2063112 2036 Streptococcus agalactiae AC-2713 NC. 019042 2179445 2056 Streptococcus dysgalactiae AC-2713 NC. 019042 2179445 2058 Streptococcus dysgalactiae RE378 DNA NC. 018767 2159491 2056 Streptococcus dysgalactiae RE378 DNA NC. 018767 2159491 2056 Streptococcus gallolyticus ATCC 43143 DNA NC. 018776 2362241 2277 Streptococcus gallolyticus ATCC BAA-2069 NC. 015215 235644 2250 Streptococcus gallolyticus DSM 16831 NZ. CP018822 2492900 2406 Streptococcus gallolyticus DSM 16831 NZ. CP018822 2492900 2406 Streptococcus pneumoniae NT-110-58 NZ. CP007593 225071 2251 Streptococcus pneumoniae SP49 NZ. CP018138 2073113 2143 Streptococcus pneumoniae SP49 NZ. CP018138 2073113 2143 Streptococcus pneumoniae SP49 NZ. CP018138 2073113 2143 Streptococcus pneumoniae Aga35 NZ. CP026576 2221315 2301 Streptococcus pneumoniae Aga35 NZ. CP026576 2221315 2301 Streptococcus pneumoniae Aga35 NZ. CP026576 2221315 2301 Streptococcus pneumoniae Aga365 NZ. CP026588 2168552 2249 Streptococcus pneumoniae BA2666 NZ. CP031248 2169584 2291 Streptococcus pneumoniae AP200 NZ. CP026588 2168552 2249 Streptococcus pneumoniae AP200 NZ | Streptococcus agalactiae SA85 | NZ_CP019811 | 1849989 | 1859 |
| Streptococcus agalactiae SA136 NZ.CP019813 1840 Streptococcus agalactiae SA97 NZ.CP019829 1842081 1856 Streptococcus agalactiae SA330 NZ.CP019829 1842081 1856 Streptococcus agalactiae SA375 NZ.CP019835 1842219 1859 Streptococcus agalactiae A009 NC.007432 2127839 2134 Streptococcus agalactiae AD09 NC.007432 2127839 2134 Streptococcus agalactiae GD201008-001 NC.01846 2063112 2036 Streptococcus agalactiae ACC2713 NC.019042 217845 2190 Streptococcus dysgalactiae ATCC 12394 NC.017567 2159491 2056 Streptococcus dysgalactiae ATCC 12394 NC.017567 2159491 2056 Streptococcus gallolyticus ATCC 43143 DNA NC.018712 2151145 2027 Streptococcus gallolyticus ATCC 43143 DNA NC.017576 2356444 2250 Streptococcus gallolyticus DSM 16831 NZ.CP018822 2492900 2406 Streptococcus gallolyticus UCN34 NC.013798 2350911 2551 Streptococcus pn | Streptococcus agalactiae SA195 | $NZ_{CP019820}$ | 1841715 | 1858 |
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| Streptococcus agalactiae SA330 NZ.CP019825 1842219 1855 Streptococcus agalactiae 2603V/R NZ.CP019835 1842219 1859 Streptococcus agalactiae A909 NC.007432 2127839 2134 Streptococcus agalactiae CNCTC 10/84 NZ.CP006910 2013842 2008 Streptococcus agalactiae GNCTC 10/84 NZ.CP006910 2013842 2008 Streptococcus agalactiae GD201008-001 NC.018646 2063112 2036 Streptococcus dysaglactiae NEM316 NC.004368 2211485 2190 Streptococcus dysaglactiae AC-2713 NC.019042 2179445 2056 Streptococcus dysaglactiae ATCC 12394 NC.017567 2159491 2056 Streptococcus gallolyticus ATCC 43143 DNA NC.018712 2151145 2027 Streptococcus gallolyticus ATCC BAA-2069 NC.015215 2356444 2250 Streptococcus gallolyticus UCN34 NC.013798 2350911 2251 Streptococcus gallolyticus UCN34 NC.013798 2350911 2251 Streptococcus gallolyticus UCN34 NC.2013798 2350911 2251 | Streptococcus agalactiae SA136 | NZ_CP019816 | 1849103 | 1840 |
| Streptococcus agalactiae SA375 NZ_CP019835 1842219 1859 Streptococcus agalactiae 2603V/R NC_001116 2160267 2127 | Streptococcus agalactiae SA97 | NZ_CP019813 | 1856410 | 1848 |
| Streptococcus agalactiae 2603V / R | Streptococcus agalactiae SA330 | NZ_CP019829 | 1842081 | 1856 |
| Streptococcus agalactiae 2603V / R | Streptococcus agalactiae SA375 | NZ_CP019835 | 1842219 | 1859 |
| Streptococcus agalactiae A909 NC.007432 2127839 2134 Streptococcus agalactiae CNCTC 10/84 NZ.CP006910 2013842 2008 Streptococcus agalactiae GD201008-001 NC.018646 2003112 2036 Streptococcus dysgalactiae ACPCI13 NC.019042 2178445 2058 Streptococcus dysgalactiae AC-2713 NC.019042 2179445 2058 Streptococcus dysgalactiae ACC-2713 NC.019767 2159491 2056 Streptococcus dysgalactiae RE378 DNA NC.018712 2151145 2027 Streptococcus gallolyticus ATCC 43143 DNA NC.017576 2362241 2277 Streptococcus gallolyticus DSM 16831 NZ.CP018822 249200 2406 Streptococcus pallolyticus UCN34 NC.013798 2350911 2251 Streptococcus pneumoniae NT-110-58 NZ.CP018836 2206644 2314 Streptococcus pneumoniae SP49 NZ.CP018136 2206644 2314 Streptococcus pneumoniae SP64 NZ.CP018138 2073113 2143 Streptococcus pneumoniae M26365 NZ.CP025266 2168552 2249 <t< td=""><td></td><td>NC_004116</td><td>2160267</td><td>2127</td></t<> | | NC_004116 | 2160267 | 2127 |
| Streptococcus agalactiae CNCTC 10/84 NZ_CP006910 2013842 2008 Streptococcus agalactiae GD201008-001 NC_018646 2063112 2036 Streptococcus agalactiae NEM316 NC_004368 2211485 2190 Streptococcus dysgalactiae AC-2713 NC_019042 2179445 2058 Streptococcus dysgalactiae AC-2713 NC_017567 2159491 2056 Streptococcus dysgalactiae ATCC 12394 NC_018712 2151145 2027 Streptococcus gallolyticus ATCC 43143 DNA NC_018712 2151145 2207 Streptococcus gallolyticus ATCC BAA-2069 NC_015215 2356444 2250 Streptococcus gallolyticus UCN34 NC_0713798 2350911 2251 Streptococcus gallolyticus UCN34 NC_07013798 2350911 2251 Streptococcus pneumoniae SP49 NZ_CP018136 2206644 2314 Streptococcus pneumoniae SP64 NZ_CP018138 2073113 2143 Streptococcus pneumoniae 335 NZ_CP02556 2168552 2249 Streptococcus pneumoniae M26365 NZ_CP027540 2046572 2118 <tr< td=""><td></td><td>NC_007432</td><td>2127839</td><td>2134</td></tr<> | | NC_007432 | 2127839 | 2134 |
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| Streptococcus pneumoniae SPN XDR SMC1710-32 NZ_CP025838 2057144 2172 Streptococcus pneumoniae HKU1-14 NZ_CP019299 2038121 2125 Streptococcus pneumoniae EF3030 NZ_CP035897 2101621 2152 Streptococcus pneumoniae R6CIB17 NZ_CP038808 2038617 2111 Streptococcus pneumoniae 670-6B NC_014498 2240045 2366 Streptococcus pneumoniae 70585 NC_012468 2184682 2299 Streptococcus pneumoniae AP200 NC_014494 2130580 2222 Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_010853 2046116 2118 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_017591 2142122 2247 Streptococcus pneumoniae INV104 genome NC_017593 2093317 2153 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 <td< td=""><td></td><td></td><td></td><td></td></td<> | | | | |
| Streptococcus pneumoniae HKU1-14 NZ_CP019299 2038121 2125 Streptococcus pneumoniae EF3030 NZ_CP035897 2101621 2152 Streptococcus pneumoniae R6CIB17 NZ_CP038808 2038617 2111 Streptococcus pneumoniae 670-6B NC_014498 2240045 2366 Streptococcus pneumoniae 70585 NC_012468 2184682 2299 Streptococcus pneumoniae AP200 NC_014494 2130580 2222 Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae G54 NC_010582 2209198 2278 Streptococcus pneumoniae gamPNI0373 NC_018630 2064116 2118 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome NC_017593 2093317 2153 Streptococcus pneumoniae OXC141 NC_012466 2120234 2211 Str | | | | |
| Streptococcus pneumoniae EF3030 NZ_CP035897 2101621 2152 Streptococcus pneumoniae R6CIB17 NZ_CP038808 2038617 2111 Streptococcus pneumoniae 670-6B NC_014498 2240045 2366 Streptococcus pneumoniae 70585 NC_012468 2184682 2299 Streptococcus pneumoniae AP200 NC_014494 2130580 2222 Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae Hungary19A-6 NC_018630 2064154 2180 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae JJA NC_017593 2093317 2153 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumonia | | | | |
| Streptococcus pneumoniae R6CIB17 NZ_CP038808 2038617 2111 Streptococcus pneumoniae 670-6B NC_014498 2240045 2366 Streptococcus pneumoniae 70585 NC_012468 2184682 2299 Streptococcus pneumoniae AP200 NC_014494 2130580 2222 Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae JJA NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae 670-6B NC_014498 2240045 2366 Streptococcus pneumoniae 70585 NC_012468 2184682 2299 Streptococcus pneumoniae AP200 NC_014494 2130580 2222 Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae JJA NC_017593 2093317 2153 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae 70585 NC_012468 2184682 2299 Streptococcus pneumoniae AP200 NC_014494 2130580 2222 Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae OXC141 NC_012466 2120234 2211 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae AP200 NC_014494 2130580 2222 Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae ATCC 700669 NC_011900 2221315 2305 Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae CGSP14 NC_010582 2209198 2278 Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae D39 NC_008533 2046116 2118 Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae G54 NC_011072 2078953 2157 Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae gamPNI0373 NC_018630 2064154 2180 Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae Hungary19A-6 NC_010380 2245615 2335 Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae INV104 genome. NC_017591 2142122 2247 Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae INV200 genome. NC_017593 2093317 2153 Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae JJA NC_012466 2120234 2211 Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae OXC141 NC_017592 2036867 2151 Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae P1031 NC_012467 2111882 2238 Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae R6 NC_003098 2038615 1814 Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| Streptococcus pneumoniae SPN034183 NC_021028 2037254 2151 | | | | |
| | | | | |
| 5treptococcus phenimoniae 5trin994058 NG UZTUZ6 ZUZ6Z39 Z141 | Streptococcus pneumoniae SPN994038 | NC_021026 | 2026239 | 2141 |
| Streptococcus pneumoniae SPN994039 NC_021005 2026505 2141 | | | | |

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|---|----------------------------|-----------------|--------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Streptococcus pneumoniae ST556 | NC_017769 | 2150813 | 2242 |
| Streptococcus pneumoniae Taiwan19F-14 | NC_012469 | 2112148 | 2108 |
| Streptococcus pneumoniae TIGR4 | $NC_{-}003028$ | 2160842 | 2271 |
| Streptococcus pyogenes HKU360 | $NZ_{-}CP009612$ | 1944537 | 1897 |
| Streptococcus pyogenes D471 | $NZ_{-}CP011415$ | 1811968 | 1763 |
| Streptococcus pyogenes NGAS743 | NZ_CP007560 | 1915554 | 1895 |
| Streptococcus pyogenes NGAS596 | NZ_CP007561 | 1791306 | 1711 |
| Streptococcus pyogenes NGAS327 | $NZ_{-}CP007562$ | 1702054 | 1611 |
| Streptococcus pyogenes STAB10015 | NZ_CP011068 | 1950454 | 1902 |
| Streptococcus pyogenes HKU488 | NZ_CP012045 | 1943415 | 1924 |
| Streptococcus pyogenes NGAS638 | NZ_CP010450 | 1791401 | 1740 |
| Streptococcus pyogenes NGAS322 | NZ_CP010449 | 1950469 | 1898 |
| Streptococcus pyogenes MEW123 | NZ_CP014139 | 1878699 | 1794 |
| Streptococcus pyogenes AP53 | NZ_CP013672 | 1860554 | 1848 |
| Streptococcus pyogenes MGAS23530 | NZ_CP013839 | 1709394 | 1623 |
| Streptococcus pyogenes MGAS27061 | NZ_CP013840 | 1741348 | 1660 |
| Streptococcus pyogenes MGAS11027 | NZ_CP013838 | 1786874 | 1718 |
| Streptococcus pyogenes STAB09014 | NZ_CP011069 | 1862487 | 1801 |
| Streptococcus pyogenes M28PF1 | NZ_CP011535 | 1896976 | 1852 |
| Streptococcus pyogenes NS53 | NZ_CP015238 | 1765123 | 1696 |
| Streptococcus pyogenes STAB14018 | NZ_CP014542 | 1890465 | 1877 |
| Streptococcus pyogenes GURSA1 | NZ_CP022206 | 1893171 | 1883 |
| Streptococcus pyogenes GUR Streptococcus pyogenes GUR | NZ_CP022354 | 1890204 | 1870 |
| Streptococcus pyogenes STAB090229 | NZ_CP020027 | 1846347 | 1811 |
| Streptococcus pyogenes STAB120304 | NZ_CP020082 | 1890354 | 1875 |
| Streptococcus pyogenes STAB120304 Streptococcus pyogenes HarveyGAS | NZ_CP023769 | 1852980 | 1788 |
| Streptococcus pyogenes IfalveyGAS Streptococcus pyogenes JS12 | NZ_CP021640 | 1810283 | 1722 |
| Streptococcus pyogenes TJ11-001 | NZ_CP021040 NZ_CP028148 | 1939844 | 1899 |
| | NZ_CP028140 | 1790938 | 1725 |
| Streptococcus pyogenes NGAS979 Streptococcus pyogenes DMG1800716 | NZ_CP027771 | 1869673 | 1723 1794 |
| Streptococcus pyogenes SASM4-Duke | NZ_CP031770 | 1903997 | 1875 |
| | | | |
| Streptococcus pyogenes ABC020055975 | NZ_CP029694 | 1866513 | 1859 |
| Streptococcus pyogenes FDAARGOS-534 | NZ_CP033767 | 1837737 | 1786 |
| Streptococcus pyogenes Duke-Large | NZ_CP033907 | 1903852 | 1874 |
| Streptococcus pyogenes RLGH | NZ_CP033908 | 1903972 | 1875 |
| Streptococcus pyogenes MGAS28085 | NZ_CP032666 | 1851471 | 1783 |
| Streptococcus pyogenes MGAS27961 | NZ_CP032665 | 1853912 | 1790 |
| Streptococcus pyogenes CCUG 4207 | NZ_CP028841 | 1914862 | 1920 |
| Streptococcus pyogenes TSPY453 | NZ_CP033337 | 1976011 | 1930 |
| Streptococcus pyogenes TSPY165 | NZ_CP033336 | 1812036 | 1761 |
| Streptococcus pyogenes TSPY556 | NZ_CP032700 | 1828132 | 1755 |
| Streptococcus pyogenes TSPY208 | NZ_CP033335 | 1801696 | 1732 |
| Streptococcus pyogenes M75 | NZ_CP033621 | 1852894 | 1807 |
| Streptococcus pyogenes MGAS29409 | NZ_CP031617 | 1894040 | 1840 |
| Streptococcus pyogenes MGAS28669 | NZ_CP031623 | 1905059 | 1857 |
| Streptococcus pyogenes MGAS28278 | $NZ_{-}CP031629$ | 1893446 | 1832 |
| Streptococcus pyogenes MGAS29064 | NZ_CP031620 | 1839848 | 1761 |
| Streptococcus pyogenes SP1336 | NZ_CP031738 | 1878827 | 1842 |
| Streptococcus pyogenes MGAS8347 | $NZ_{-}CP031638$ | 1833822 | 1742 |
| Streptococcus pyogenes MGAS29326 | $NZ_{-}CP031618$ | 1940684 | 1884 |
| Streptococcus pyogenes MGAS28650 | $NZ_{-}CP031624$ | 1851876 | 1784 |
| Streptococcus pyogenes MGAS28386 | NZ_CP031626 | 1837594 | 1759 |

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|--|----------------|-----------------|--------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Streptococcus pyogenes MGAS10786 | NZ_CP031637 | 1937069 | 1895 |
| Streptococcus pyogenes MGAS29284 | NZ_CP031619 | 1946265 | 1919 |
| Streptococcus pyogenes MGAS28533 | $NZ_CP031625$ | 1893263 | 1851 |
| Streptococcus pyogenes MGAS28191 | NZ_CP031631 | 1892473 | 1832 |
| Streptococcus pyogenes MGAS11115 | NZ_CP031633 | 1851403 | 1788 |
| Streptococcus pyogenes MGAS28078 | NZ_CP031632 | 1880780 | 1812 |
| Streptococcus pyogenes MGAS11052 | NZ_CP031635 | 1907750 | 1833 |
| Streptococcus pyogenes MGAS7914 | NZ_CP031639 | 1885126 | 1832 |
| Streptococcus pyogenes MGAS7888 | NZ_CP031640 | 1962429 | 1898 |
| Streptococcus pyogenes MGAS28746 | NZ_CP031621 | 1853620 | 1785 |
| Streptococcus pyogenes MGAS28686 | NZ_CP031622 | 1894087 | 1851 |
| Streptococcus pyogenes MGAS28360 | NZ_CP031627 | 1893041 | 1833 |
| Streptococcus pyogenes MGAS28330 | NZ_CP031628 | 1851619 | 1776 |
| Streptococcus pyogenes MGAS11108 | NZ_CP031634 | 1853812 | 1783 |
| Streptococcus pyogenes MGAS10826 | NZ_CP031636 | 1852263 | 1783 |
| Streptococcus pyogenes emm78.3 | NZ_CP035437 | 1806521 | 1724 |
| Streptococcus pyogenes emm97.1 | NZ_CP035447 | 1812090 | 1744 |
| Streptococcus pyogenes emm22.8 | NZ_CP035438 | 1950616 | 1920 |
| Streptococcus pyogenes emm123 | NZ_CP035452 | 1768816 | 1662 |
| Streptococcus pyogenes emm75.1 | NZ_CP035434 | 1887700 | 1815 |
| Streptococcus pyogenes emm90.5 | NZ_CP035444 | 1764969 | 1667 |
| Streptococcus pyogenes emm58 | NZ_CP035443 | 1888806 | 1846 |
| Streptococcus pyogenes emm11 | NZ_CP035432 | 1809631 | 1738 |
| Streptococcus pyogenes emm?5 | NZ_CP035442 | 1835714 | 1786 |
| Streptococcus pyogenes emm197 | NZ_CP035455 | 1797172 | 1764 |
| Streptococcus pyogenes emm157 Streptococcus pyogenes emm55 | NZ_CP035430 | 1899479 | 1849 |
| Streptococcus pyogenes emm77 | NZ_CP035439 | 1806314 | 1740 |
| Streptococcus pyogenes TSPY155 | NZ_CP032699 | 1882327 | 1817 |
| Streptococcus pyogenes 131 1133 Streptococcus pyogenes emm230 | NZ_CP035451 | 1826832 | 1775 |
| Streptococcus pyogenes emm250 Streptococcus pyogenes emm1 | NZ_CP035441 | 1834762 | 1801 |
| Streptococcus pyogenes emmi Streptococcus pyogenes MGAS28271 | NZ_CP031630 | 1842830 | 1771 |
| | | | |
| Streptococcus pyogenes emm93.4 | NZ_CP035450 | 1701466 | 1613 1746 |
| Streptococcus pyogenes emm89.14 | NZ_CP035436 | 1806344 | |
| Streptococcus pyogenes emm100 | NZ_CP035453 | 1839531 | 1800 |
| Streptococcus pyogenes emm92 | NZ_CP035445 | 1758778 | 1683 |
| Streptococcus pyogenes emm54 | NZ_CP035454 | 1742565 | 1664 |
| Streptococcus pyogenes emm64.3 | NZ_CP035435 | 1779766 | 1743 |
| Streptococcus pyogenes emmSTG866.1 | NZ_CP035428 | 1816007 | 1736 |
| Streptococcus pyogenes emm105 | NZ_CP035431 | 1800116 | 1716 |
| Streptococcus pyogenes emm124 | NZ_CP035440 | 1790668 | 1706 |
| Streptococcus pyogenes emm74 | NZ_CP035427 | 1861037 | 1827 |
| Streptococcus pyogenes emm57 | NZ_CP035426 | 1756622 | 1676 |
| Streptococcus pyogenes emm65 | NZ_CP035433 | 1789855 | 1709 |
| Streptococcus pyogenes emm56 | NZ_CP035449 | 1825427 | 1785 |
| Streptococcus pyogenes emmNA | NZ_CP035429 | 1788166 | 1749 |
| Streptococcus pyogenes emm68.2 | NZ_CP035446 | 1747924 | 1659 |
| Streptococcus pyogenes JRS4 | NZ_CP011414 | 1811968 | 1763 |
| Streptococcus pyogenes MGAS10270 | $NC_{-}008022$ | 1928252 | 1899 |
| Streptococcus pyogenes MGAS10394 | $NC_{-}006086$ | 1899877 | 1859 |
| Streptococcus pyogenes MGAS10750 | $NC_{-}008024$ | 1937111 | 1901 |
| Streptococcus pyogenes MGAS2096 | $NC_{-}008023$ | 1860355 | 1787 |
| Streptococcus pyogenes MGAS5005 | NC_007297 | 1838562 | 1806 |

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|--------------------------------------|------------------|-----------------|------------|
| Name | Accession Num. | Sequence Length | CDS number |
| Streptococcus pyogenes MGAS6180 | $NC_{-}007296$ | 1897578 | 1852 |
| Streptococcus pyogenes MGAS9429 | $NC_{-}008021$ | 1836467 | 1785 |
| Streptococcus sobrinus SL1 | $NZ_{-}CP029490$ | 2198571 | 2060 |
| Streptococcus sobrinus 10919 | $NZ_{-}CP029491$ | 2145290 | 2005 |
| Streptococcus sobrinus NIDR 6715-7 | $NZ_{-}CP029560$ | 2145017 | 1997 |
| Streptococcus sobrinus NIDR 6715-15 | $NZ_{-}CP029559$ | 2144746 | 1996 |
| Streptococcus suis ZY05719 | $NZ_{-}CP007497$ | 2094898 | 2014 |
| Streptococcus suis NSUI002 | NZ_CP011419 | 2255345 | 2257 |
| Streptococcus suis DN13 | $NZ_{-}CP015557$ | 2142184 | 2075 |
| Streptococcus suis GZ0565 | $NZ_CP017142$ | 2131882 | 2073 |
| Streptococcus suis SS2-1 | NZ_CP018908 | 2094030 | 2043 |
| Streptococcus suis SC19 | $NZ_{-}CP020863$ | 2095966 | 2035 |
| Streptococcus suis LSM102 | $NZ_{-}CP016175$ | 2067932 | 2011 |
| Streptococcus suis CS100322 | $NZ_{-}CP024050$ | 2137649 | 2082 |
| Streptococcus suis HA0609 | NZ_CP024126 | 2247350 | 2224 |
| Streptococcus suis CZ130302 | $NZ_{-}CP024974$ | 2538894 | 2508 |
| Streptococcus suis AH681 | $NZ_{-}CP025043$ | 2404013 | 2310 |
| Streptococcus suis HN105 | $NZ_{-}CP029398$ | 2196724 | 2092 |
| Streptococcus suis 0061 | NZ_CP017666 | 2138420 | 1990 |
| Streptococcus suis 1081 | $NZ_{-}CP017667$ | 2228089 | 2097 |
| Streptococcus suis SH0104 | $NZ_{-}CP025419$ | 2243598 | 2233 |
| Streptococcus suis HA1003 | $NZ_{-}CP030125$ | 2229183 | 2133 |
| Streptococcus suis SH1510 | NZ_CP030124 | 2214693 | 2156 |
| Streptococcus suis WUSS351 | $NZ_{-}CP039462$ | 2283286 | 2157 |
| Streptococcus suis SRD478 | NZ_CP017088 | 2062916 | 1986 |
| Streptococcus suis 6407 | $NZ_{-}CP008921$ | 2292360 | 2245 |
| Streptococcus suis A7 | $NC_{-}017622$ | 2038409 | 1977 |
| Streptococcus suis BM407 | $NC_{-}012926$ | 2146229 | 1932 |
| Streptococcus suis D12 | $NC_{-}017621$ | 2183059 | 2132 |
| Streptococcus suis D9 | $NC_{-}017620$ | 2177656 | 2121 |
| Streptococcus suis GZ1 | $NC_{-}017617$ | 2038034 | 1981 |
| Streptococcus suis JS14 | $NC_{-}017618$ | 2137435 | 2083 |
| Streptococcus suis P1/7 | $NC_{-}012925$ | 2007491 | 1951 |
| Streptococcus suis S735 | NC_018526 | 1980887 | 1908 |
| Streptococcus suis SC070731 | $NC_{-}020526$ | 2138568 | 2083 |
| Streptococcus suis SC84 | $NC_{-}012924$ | 2095898 | 2041 |
| Streptococcus suis SS12 | $NC_{-}017619$ | 2096866 | 2068 |
| Streptococcus suis ST1 | NC_017950 | 2034321 | 2049 |
| Streptococcus suis ST3 | NC_015433 | 2028815 | 1983 |
| Streptococcus suis T15 | $NC_{-}022665$ | 2240234 | 2189 |
| Streptococcus suis TL13 | $NC_{-}021213$ | 2038146 | 1984 |
| Streptococcus suis YB51 | $NC_{-}022516$ | 2043655 | 1995 |
| Streptococcus thermophilus SMQ-301 | $NZ_{-}CP011217$ | 1861792 | 1895 |
| Streptococcus thermophilus MN-BM-A02 | NZ_CP010999 | 1850434 | 1891 |
| Streptococcus thermophilus ST3 | NZ_CP017064 | 1865056 | 1891 |
| Streptococcus thermophilus EPS | $NZ_{-}CP025400$ | 1812305 | 1848 |
| Streptococcus thermophilus GABA | NZ_CP025399 | 1857468 | 1862 |
| Streptococcus thermophilus DGCC 7710 | $NZ_{-}CP025216$ | 1851207 | 1887 |
| Streptococcus thermophilus ST109 | $NZ_{-}CP031545$ | 1788866 | 1817 |
| Streptococcus thermophilus ST106 | NZ_CP031881 | 1856083 | 1917 |
| Streptococcus thermophilus CNRZ1066 | NC_006449 | 1796226 | 1837 |
| Streptococcus thermophilus JIM 8232 | NC_017581 | 1929905 | 1944 |

| Name | Accession Num. | Sequence Length | CDS number |
|---------------------------------------|----------------|-----------------|------------|
| Streptococcus thermophilus LMD-9 | NC_008532 | 1856368 | 1892 |
| Streptococcus thermophilus LMG 18311 | $NC_{-}006448$ | 1796846 | 1819 |
| Streptococcus thermophilus MN-ZLW-002 | $NC_{-}017927$ | 1848520 | 1884 |
| Streptococcus thermophilus ND03 | $NC_{-}017563$ | 1831949 | 1875 |

Table S6: 101 bacteria in third dataset and bacteria closest to them

| Bacterium i | Bacteria j closest to Bacteria i | The minimum Manhattan distance |
|----------------------------|----------------------------------|--------------------------------|
| P.aeruginosa DHS01 | P.aeruginosa 97 | 0.0317 |
| P.aeruginosa DK2 | P.aeruginosa MTB-1 | 0.0248 |
| P.aeruginosa DSM 50071 | P.aeruginosa ATCC 15692 | 0.0189 |
| P.aeruginosa LES431 | P.aeruginosa LESB58 | 0.0149 |
| P.aeruginosa LESB58 | P.aeruginosa LES431 | 0.0149 |
| P.aeruginosa M18 | P.aeruginosa PA150577 | 0.0183 |
| P.aeruginosa MTB-1 | P.aeruginosa PA7790 | 0.0239 |
| P.aeruginosa NCGM2S1 | P.aeruginosa B14130 | 0.0446 |
| P.aeruginosa PA1 | P.aeruginosa AES1R | 0.0197 |
| P.aeruginosa PA7 | P.aeruginosa CR1 | 0.0258 |
| P.aeruginosa SCV20265 | P.aeruginosa ATCC 15692 | 0.0203 |
| P.aeruginosa UCBPP-PA14 | P.aeruginosa SCVFeb | 0.0219 |
| P.aeruginosa 12939 | P.aeruginosa Y89 | 0.0229 |
| P.aeruginosa 1334/14 | P.fluorescens L321 | 0.0521 |
| P.aeruginosa 24Pae112 | P.aeruginosa CCUG 51971 | 0.0278 |
| P.aeruginosa 60503 | P.aeruginosa MTB-1 | 0.0276 |
| P.aeruginosa 97 | P.aeruginosa PA11803 | 0.0262 |
| P.aeruginosa A681 | P.aeruginosa LES431 | 0.0226 |
| P.aeruginosa AES1R | P.aeruginosa LESB58 | 0.0188 |
| P.aeruginosa AR441 | P.aeruginosa IMP66 | 0.0274 |
| P.aeruginosa AR445 | P.aeruginosa H27930 | 0.0303 |
| P.aeruginosa AR0110 | P.aeruginosa PA83 | 0.0318 |
| P.aeruginosa ATCC 15692 | P.aeruginosa GIMC5015PAKB6 | 0.0070 |
| P.aeruginosa ATCC 27853 | P.aeruginosa F9670 | 0.0436 |
| P.aeruginosa ATCC 27853 | P.aeruginosa S86968 | 0.0322 |
| P.aeruginosa AZPAE15042 | P.aeruginosa IMP67 | 0.0215 |
| P.aeruginosa B14130 | P.aeruginosa NCGM2S1 | 0.0446 |
| P.aeruginosa CCUG 51971 | P.aeruginosa S86968 | 0.0246 |
| P.aeruginosa CCUG 70744 | P.aeruginosa HS9 | 0.0564 |
| P.aeruginosa CR1 | P.aeruginosa PA7 | 0.0258 |
| P.aeruginosa E80 | P.aeruginosa S86968 | 0.0312 |
| P.aeruginosa E90 | P.aeruginosa SCV20265 | 0.0224 |
| P.aeruginosa F23197 | P.aeruginosa IMP66 | 0.0179 |
| P.aeruginosa F5677 | P.aeruginosa HOU1 | 0.0295 |
| P.aeruginosa F63912 | P.aeruginosa PA298 | 0.0197 |
| P.aeruginosa F9670 | P.aeruginosa ATCC 27853 | 0.0436 |
| P.aeruginosa GIMC5015PAKB6 | P.aeruginosa ATCC 15692 | 0.0070 |
| P.aeruginosa H25883 | P.aeruginosa PABL048 | 0.0309 |
| P.aeruginosa H26023 | P.aeruginosa LESB58 | 0.0219 |
| P.aeruginosa H26027 | P.aeruginosa F23197 | 0.0258 |
| P.aeruginosa H27930 | P.aeruginosa PABL017 | 0.0169 |
| P.aeruginosa H47921 | P.fluorescens L321 | 0.0619 |
| P.aeruginosa H5708 | P.aeruginosa PB354 | 0.0206 |
| P.aeruginosa HOU1 | P.aeruginosa IMP68 | 0.0272 |

continued table S6

| Bacterium i | Bacterium j closest to bacterium i The m | ninimum Manhattan distance |
|-----------------------|--|----------------------------|
| P.aeruginosa HS9 | P.aeruginosa ATCC 27853 | 0.0529 |
| P.aeruginosa IMP66 | P.aeruginosa IMP68 | 0.0075 |
| P.aeruginosa IMP67 | P.aeruginosa IMP68 | 0.0062 |
| P.aeruginosa IMP68 | P.aeruginosa IMP67 | 0.0062 |
| P.aeruginosa L10 | P.aeruginosa UCBPP-PA14 | 0.0022 |
| P.aeruginosa PA11803 | P.aeruginosa 97 | 0.0262 |
| P.aeruginosa PA298 | P.aeruginosa ATCC 15692 | 0.0202 |
| P.aeruginosa PA34 | P.aeruginosa PAD21 | 0.0336 |
| P.aeruginosa PA7790 | P.aeruginosa PA8281 | 0.0194 |
| P.aeruginosa PA8281 | P.aeruginosa PA7790 | 0.0194 |
| P.aeruginosa PA83 | P.aeruginosa SCV20265 | 0.0245 |
| P.aeruginosa PABL017 | P.aeruginosa H27930 | 0.0169 |
| P.aeruginosa PABL048 | P.aeruginosa F23197 | 0.0245 |
| P.aeruginosa PAER4119 | P.aeruginosa PA298 | 0.0254 |
| P.aeruginosa PAO1161 | P.aeruginosa GIMC5015PAKB6 | 0.0234 |
| P.aeruginosa PA150577 | P.aeruginosa M18 | 0.0173 |
| P.aeruginosa PAD16 | P.aeruginosa PAD5 | 0.0183 |
| | · · | 0.0013 |
| P.aeruginosa PAD2 | P.aeruginosa PAD21 | 0.0021 |
| P.aeruginosa PAD21 | P.aeruginosa PAD2 | |
| P.aeruginosa PAD22 | P.aeruginosa PAD5 | 0.0023 |
| P.aeruginosa PAD25 | P.aeruginosa PAD22 | 0.0025 |
| P.aeruginosa PAD5 | P.aeruginosa PAD16 | 0.0013 |
| P.aeruginosa PB350 | P.aeruginosa H5708 | 0.0352 |
| P.aeruginosa PB354 | P.aeruginosa H5708 | 0.0206 |
| P.aeruginosa S86968 | P.aeruginosa CCUG 51971 | 0.0246 |
| P.aeruginosa SCVFeb | P.aeruginosa M18 | 0.0216 |
| P.aeruginosa T38079 | P.aeruginosa S86968 | 0.0314 |
| P.aeruginosa W16407 | P.aeruginosa AZPAE15042 | 0.0238 |
| P.aeruginosa Y89 | P.aeruginosa 12939 | 0.0229 |
| P.fluorescens L321 | P.aeruginosa 1334/14 | 0.0521 |
| P.fluorescens Pf275 | P.syringae actinidiae ICMP 985 | 0.0699 |
| P.fluorescens Pt14 | P.aeruginosa 1334/14 | 0.0715 |
| P.fluorescens SIKW1 | P.fluorescens L321 | 0.0694 |
| P.putida GB-1 | P.putida PP112420 | 0.0542 |
| P.putida H8234 | P.aeruginosa ATCC 27853 | 0.0842 |
| P.putida JB | P.putida E46 | 0.0515 |
| P.putida KT2440 | P.aeruginosa LES431 | 0.0553 |
| P.putida NBRC 14164 | P.putida E46 | 0.0577 |
| P.putida ND6 | P.aeruginosa CCUG 70744 | 0.0731 |
| P.putida SJTE-1 | P.aeruginosa PA7 | 0.0420 |
| P.putida B4 | P.putida SJTE-1 | 0.0554 |
| P.putida E41 | P.putida PP112420 | 0.0451 |

| Bacterium i | Bacterium j closest to Bacteria i | The minimum Manhattan distance |
|---|-----------------------------------|--------------------------------|
| P.putida E46 | P.aeruginosa PA1 | 0.0476 |
| P.putida PP112420 | P.putida E41 | 0.0451 |
| P.stutzeri A1501 | P.stutzeri SGAir0442 | 0.0494 |
| P.stutzeri CCUG 29243 | P.aeruginosa ATCC 27853 | 0.0546 |
| P.stutzeri DSM 10701 | P.stutzeri A1501 | 0.0588 |
| P.stutzeri DSM 4166 | P.stutzeri SGAir0442 | 0.0408 |
| P.stutzeri RCH2 | P.fluorescens Pt14 | 0.0805 |
| P.stutzeri SGAir0442 | P.stutzeri DSM 4166 | 0.0408 |
| P.syringae actinidiae ICMP 18884P.syringae actinidiae NZ-47 | | 0.0336 |
| P.syringae actinidiae ICMP 9853 P.aeruginosa S86968 | | 0.0376 |
| P.syringae actinidiae CRAFRU 122 P.syringae actinidiae CRAFRU 140 | | 0.0247 |
| P.syringae actinidiae CRAFRU 140 P.syringae actinidiae CRAFRU 122 | | 0.0247 |
| P.syringae actinidiae MAFF212063 P.aeruginosa T38079 | | 0.0587 |
| P.syringae actinidiae NZ-45 | P.syringae actinidiae NZ-47 | 0.0134 |
| P.syringae actinidiae NZ-47 | P.syringae actinidiae NZ-45 | 0.0134 |