

Table S3

Differentially expressed predicted antisense RNA plus large and small RNA products.

| Start ^a | Stop ^b | Length | Strand | Product | FC ^c | q ^d | Comments |
|--------------------|-------------------|--------|--------|-------------------------------------|-----------------|----------------|-----------------|
| | | | | | ≥1.5 ≥-1.5 | ≤0.05 | |
| 64255 | 63730 | 525 | - | antisense: PA14_00630 | 2.4 | 2.8E-25 | |
| 328092 | 327954 | 138 | - | antisense: PA14_03630 | -2.3 | 5.78E-08 | |
| 369305 | 369090 | 215 | - | antisense: PA14_04100 | 3.0 | 2.01E-30 | |
| 653645 | 653892 | 247 | + | antisense: PA14_07580 | 2.1 | 4.16E-19 | |
| 734616 | 733096 | 1520 | - | antisense: PA14_08570 ^e | 4.0 | 2.17E-138 | 16S rRNA |
| 737953 | 735100 | 2853 | - | antisense: PA14_08600 | 6.6 | 1.04E-291 | 23S rRNA |
| 765237 | 764974 | 263 | - | antisense: <i>rpIE</i> | 2.9 | 5.59E-26 | |
| 775136 | 774767 | 369 | - | antisense: <i>bfrA</i> ^f | 2.2 | 1.12E-17 | nc ^g |
| 775296 | 775611 | 315 | + | antisense: <i>uvrA</i> | 2.3 | 4.93E-20 | |
| 805283 | 805500 | 217 | + | antisense: phzS | -2.4 | 5.82E-12 | 1.9 |
| 1016946 | 1016918 | 28 | - | antisense: PA14_11730 | -2.9 | 2.64E-07 | |
| 1053400 | 1053276 | 124 | - | antisense: PA14_12170 | 3.0 | 1.03E-15 | |
| 1222942 | 1223625 | 683 | + | antisense: PA14_14340 | -54.0 | 0.00E+00 | nc |
| 1274815 | 1275275 | 460 | + | antisense: <i>leuA</i> | 3.6 | 5.35E-55 | -2.1 |
| 1522104 | 1521977 | 127 | - | antisense: <i>rpmE2</i> | 3.8 | 9.95E-30 | |
| 1836723 | 1837064 | 341 | + | antisense: PA14_21220 | 6.0 | 3.89E-86 | |
| 1849687 | 1849567 | 120 | - | antisense: <i>fadD2</i> | 2.5 | 3.15E-09 | |
| 1863998 | 1863886 | 112 | - | antisense: PA14_21490 | 2.0 | 6.42E-07 | |
| 1870523 | 1870313 | 210 | - | antisense: PA14_21570 | 6.0 | 1.57E-144 | 3.0 |
| | | | | PA14_21580 | | | 2.8 |
| 1902333 | 1902404 | 71 | + | antisense: <i>tsp</i> | 6.0 | 4.75E-46 | |
| 1905417 | 1905055 | 362 | - | antisense: PA14_21890 | 3.2 | 8.70E-48 | nc |
| 1962801 | 1962597 | 204 | - | antisense: PA14_22550 | 2.2 | 6.91E-12 | |
| 1962801 | 1962597 | 204 | - | antisense: PA14_22550 | 2.2 | 6.91E-12 | |
| 2120706 | 2120697 | 9 | - | antisense: PA14_24360 | 2.4 | 6.18E-06 | |
| 2153404 | 2153293 | 111 | - | antisense: PA14_24630 | 2.0 | 1.00E-06 | |
| 2190534 | 2190794 | 260 | + | antisense: PA14_25040 | 2.0 | 6.52E-11 | |
| 2194052 | 2193777 | 275 | - | antisense: <i>fadB</i> | 2.3 | 8.09E-14 | |
| 2204351 | 2204396 | 45 | + | antisense: PA14_25220 | 2.0 | 1.45E-03 | |
| 2320151 | 2320248 | 97 | + | antisense: PA14_26640 | 3.0 | 7.74E-16 | |
| 2356351 | 2356217 | 134 | - | antisense: PA14_27120 | 2.0 | 4.34E-09 | |
| 2360088 | 2360104 | 16 | + | antisense: PA14_27150 | -2.3 | 8.91E-04 | |
| 2802689 | 2803426 | 737 | + | antisense: PA14_32250 | 4.3 | 2.45E-204 | 5.2 |
| 3009991 | 3010452 | 461 | + | antisense: PA14_33870 | 2.3 | 3.55E-36 | 1.8 |
| 3084412 | 3086372 | 1960 | + | antisense: PA14_34720 | 7.1 | 0.00E+00 | |
| | | | | PA14_34730 PA14_34740 | | | |
| 3239166 | 3239282 | 116 | + | antisense: PA14_36370 | 2.0 | 5.16E-06 | |
| 3418139 | 3417966 | 173 | - | antisense: PA14_38340 | 3.3 | 3.02E-27 | |
| 3484578 | 3484761 | 183 | + | antisense: PA14_39100 | -2.1 | 4.01E-06 | |
| 3518240 | 3518513 | 273 | + | antisense: PA14_39520 | -22.0 | 0.00E+00 | nc |
| 3612543 | 3612599 | 56 | + | antisense: PA14_40430 | 2.9 | 7.30E-09 | |
| 3656240 | 3656520 | 280 | + | antisense: PA14_41010 | 2.1 | 1.34E-11 | |
| 3686273 | 3686214 | 59 | - | antisense: PA14_41290 | 2.2 | 2.17E-05 | |
| 3739392 | 3739278 | 114 | - | antisense: PA14_41860 | 2.0 | 3.96E-07 | |
| 3854106 | 3853766 | 340 | - | antisense: PA14_43250 | 2.4 | 6.44E-19 | |
| 3924407 | 3924741 | 334 | + | antisense: PA14_44080 | 2.1 | 1.36E-15 | |

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|---------|---------|------|---|--|------------|-----------|---|
| 3924510 | 3924046 | 464 | - | antisense: <i>gltA</i> | 2.0 | 1.27E-12 | |
| 4049209 | 4048834 | 375 | - | antisense: PA14_45470 | 2.0 | 1.46E-11 | |
| 4492049 | 4491818 | 231 | - | antisense: <i>braG</i> | -2.3 | 6.21E-08 | |
| 4563787 | 4562873 | 914 | - | antisense: <i>mvfR</i> | 3.0 | 6.19E-58 | nc |
| 4625023 | 4624794 | 229 | - | antisense: <i>lpxO2</i> | -2.2 | 1.10E-08 | |
| 4695835 | 4695098 | 737 | - | antisense: PA14_52940 | 2.0 | 3.53E-16 | |
| 4808699 | 4809117 | 418 | + | antisense: <i>pruR</i> | 2.6 | 6.63E-37 | |
| 4952318 | 4957474 | 5156 | + | antisense: PA14_55631 PA14_55634 PA14_55635 PA14_55637 | 6.0 | 3.25E-275 | 23S rRNA; tRNA-Ala; tRNA-Ile; 16S rRNA |
| 5060140 | 5060202 | 62 | + | antisense: PA14_56800 | -2.3 | 1.14E-04 | |
| 5266416 | 5266240 | 176 | - | antisense: PA14_59140 PA14_59150 | 3.3 | 8.44E-27 | |
| 5350100 | 5349555 | 545 | - | antisense: PA14_60070 | 3.4 | 5.17E-74 | |
| 5384678 | 5385115 | 437 | + | antisense: PA14_60390 | 2.3 | 8.14E-24 | |
| 5399019 | 5398890 | 129 | - | antisense: PA14_60570 | 5.5 | 6.14E-47 | |
| 5474606 | 5474583 | 23 | - | antisense: PA14_61300 | -2.2 | 7.49E-04 | |
| 5536325 | 5539178 | 2853 | + | antisense: PA14_62060 | 6.6 | 4.75E-292 | 23S rRNA |
| 5539662 | 5541182 | 1520 | + | antisense: PA14_62090 | 4.0 | 2.17E-138 | 16S rRNA |
| 5608393 | 5608523 | 130 | + | antisense: <i>ftsH</i> | 4.0 | 5.86E-32 | |
| 5714456 | 5714335 | 121 | - | antisense: <i>accC</i> | 2.1 | 3.04E-07 | |
| 6140565 | 6140680 | 115 | + | antisense: <i>gcvP1</i> | 2.2 | 7.33E-10 | |
| 6154122 | 6154310 | 188 | + | antisense: <i>pepP</i> | -18.0 | 0.00E+00 | nc |
| 6312796 | 6315649 | 2853 | + | antisense: PA14_70880 | 6.6 | 1.04E-291 | 23S rRNA |
| 6316133 | 6317653 | 1520 | + | antisense: PA14_70910 | 4.0 | 2.17E-138 | 16S rRNA |
| 6356929 | 6356474 | 455 | - | antisense: PA14_71350 PA14_71360 | -2.9 | 3.27E-20 | |
| 6382634 | 6382509 | 125 | - | antisense: <i>aspA</i> | -2.8 | 3.22E-08 | |
| 2786835 | 2787657 | 822 | + | sense: PA14_32015 ^h | 6.6 | 0.00E+00 | CzcA amino terminus |
| 2787806 | 2789915 | 2109 | + | sense: PA14_32025 | 6.2 | 0.00E+00 | CzcA carboxy terminus |
| 57035 | 57139 | 104 | + | - | 2.3 | 1.94E-06 | |
| 61368 | 61149 | 219 | - | - | -3.0 | 1.51E-26 | |
| 200534 | 200500 | 34 | - | - | -2.4 | 5.66E-04 | |
| 226488 | 226476 | 12 | - | - | -2.3 | 1.83E-04 | |
| 244004 | 243946 | 58 | - | - | 3.0 | 6.27E-08 | |
| 249913 | 249934 | 21 | + | - | -4.6 | 1.70E-28 | |
| 269220 | 269202 | 18 | - | - | 3.3 | 1.79E-18 | |
| 279717 | 279566 | 151 | - | - | 2.0 | 1.78E-07 | |
| 309605 | 309748 | 143 | + | - | 4.8 | 7.12E-32 | |
| 329965 | 329956 | 9 | - | - | 2.3 | 1.02E-05 | |
| 330918 | 330785 | 133 | - | - | 2.0 | 2.76E-17 | |
| 332114 | 331962 | 152 | - | - | 2.5 | 2.68E-48 | |
| 374503 | 374521 | 18 | + | - | 2.7 | 8.99E-09 | |
| 377253 | 377028 | 225 | - | - | -3.6 | 1.89E-19 | |

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|---------|---------|-----|---|-------|------|-----------|
| 384642 | 384575 | 67 | - | - | -2.6 | 2.50E-05 |
| 403447 | 403506 | 59 | + | - | 2.1 | 1.72E-07 |
| 458075 | 457976 | 99 | - | - | 3.0 | 3.46E-14 |
| 481583 | 481492 | 91 | - | - | -2.3 | 2.65E-05 |
| 514654 | 514466 | 188 | - | - | 2.3 | 1.20E-21 |
| 564678 | 564698 | 20 | + | - | -2.8 | 1.51E-07 |
| 565458 | 565533 | 75 | + | - | -5.7 | 4.86E-41 |
| 565783 | 565792 | 9 | + | - | -2.6 | 3.94E-08 |
| 610279 | 610210 | 69 | - | - | 2.4 | 2.41E-09 |
| 634868 | 634779 | 89 | - | - | 2.3 | 3.78E-08 |
| 673111 | 673135 | 24 | + | - | 3.0 | 7.68E-11 |
| 784463 | 784623 | 160 | + | - | -2.0 | 3.05E-06 |
| 799927 | 800011 | 84 | + | - | -2.8 | 3.88E-54 |
| 802613 | 802792 | 179 | + | - | -2.5 | 2.84E-09 |
| 805206 | 805216 | 10 | + | - | -2.1 | 3.49E-04 |
| 806855 | 806528 | 327 | - | - | 5.0 | 7.65E-100 |
| 807525 | 807504 | 21 | - | - | 6.0 | 1.15E-63 |
| 808375 | 808363 | 12 | - | - | 2.0 | 6.25E-05 |
| 813422 | 813136 | 286 | - | - | 2.3 | 6.65E-23 |
| 840541 | 840449 | 92 | - | - | 3.8 | 2.89E-21 |
| 906820 | 906724 | 96 | - | - | 4.8 | 1.12E-29 |
| 908551 | 908647 | 96 | + | - | 3.1 | 2.23E-18 |
| 933885 | 933825 | 60 | - | - | 3.8 | 1.30E-50 |
| 1042855 | 1042885 | 30 | + | - | 2.4 | 4.88E-05 |
| 1107375 | 1107457 | 82 | + | - | 2.3 | 4.16E-08 |
| 1117707 | 1117870 | 163 | + | - | 2.0 | 1.01E-14 |
| 1210068 | 1210133 | 65 | + | - | 2.0 | 7.89E-05 |
| 1310747 | 1310736 | 11 | - | - | 2.1 | 2.76E-04 |
| 1334440 | 1334545 | 105 | + | - | 2.8 | 3.02E-11 |
| 1349852 | 1349837 | 15 | - | - | -2.0 | 4.94E-03 |
| 1384119 | 1384136 | 17 | + | - | -2.1 | 2.22E-05 |
| 1437572 | 1437554 | 18 | - | - | 2.2 | 5.02E-04 |
| 1455960 | 1455934 | 26 | - | - | -3.8 | 1.01E-16 |
| 1477571 | 1477600 | 29 | + | - | 2.8 | 2.65E-07 |
| 1492951 | 1493022 | 71 | + | - | -2.2 | 4.88E-03 |
| 1505272 | 1505293 | 21 | + | - | -2.0 | 4.32E-03 |
| 1545244 | 1545205 | 39 | - | - | -2.1 | 2.23E-03 |
| 1561325 | 1561359 | 34 | + | - | 3.6 | 3.31E-18 |
| 1597096 | 1596991 | 105 | - | - | 2.0 | 2.51E-06 |
| 1599848 | 1599983 | 135 | + | RealL | 2.0 | 4.03E-09 |
| 1635661 | 1635234 | 427 | - | - | -2.1 | 1.04E-22 |
| 1648377 | 1648572 | 195 | + | - | 2.1 | 1.11E-20 |
| 1676713 | 1676674 | 39 | - | - | 2.1 | 3.94E-04 |
| 1685556 | 1685652 | 96 | + | - | 2.8 | 2.66E-46 |
| 1689131 | 1689248 | 117 | + | - | 3.2 | 6.16E-38 |
| 1699602 | 1699715 | 113 | + | - | 2.0 | 7.38E-05 |
| 1725316 | 1725523 | 207 | + | - | -3.2 | 2.00E-28 |
| 1736707 | 1736732 | 25 | + | - | -2.6 | 3.18E-05 |
| 1749126 | 1749177 | 51 | + | - | 7.4 | 7.42E-94 |
| 1750183 | 1750249 | 66 | + | - | 14.5 | 0.00E+00 |
| 1870104 | 1870121 | 17 | + | - | 2.7 | 3.52E-15 |
| 1872668 | 1872899 | 231 | + | - | 3.2 | 1.40E-64 |

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|---------|---------|-----|---|---|------|-----------|
| 1887235 | 1887202 | 33 | - | - | 2.2 | 1.06E-04 |
| 1934976 | 1934915 | 61 | - | - | 3.6 | 4.10E-17 |
| 1978119 | 1978012 | 107 | - | - | 2.0 | 3.37E-07 |
| 1983463 | 1983606 | 143 | + | - | 2.1 | 1.46E-11 |
| 1993611 | 1993668 | 57 | + | - | -2.0 | 1.59E-03 |
| 2025890 | 2025947 | 57 | + | - | 2.7 | 6.33E-13 |
| 2045697 | 2045716 | 19 | + | - | -3.3 | 6.61E-13 |
| 2253608 | 2253666 | 58 | + | - | 2.1 | 1.80E-05 |
| 2275154 | 2275079 | 75 | - | - | -2.1 | 4.22E-05 |
| 2303834 | 2303550 | 284 | - | - | 2.5 | 4.38E-29 |
| 2362094 | 2362489 | 395 | + | - | 2.7 | 6.32E-32 |
| 2388350 | 2388427 | 77 | + | - | 2.1 | 8.61E-22 |
| 2411667 | 2411678 | 11 | + | - | 5.3 | 1.59E-44 |
| 2411892 | 2411937 | 45 | + | - | 7.4 | 1.31E-85 |
| 2417927 | 2417943 | 16 | + | - | 2.1 | 1.37E-04 |
| 2423522 | 2423639 | 117 | + | - | 3.6 | 5.52E-17 |
| 2436434 | 2436421 | 13 | - | - | -2.1 | 7.44E-05 |
| 2455743 | 2455780 | 37 | + | - | -4.0 | 1.68E-22 |
| 2492542 | 2492412 | 130 | - | - | 2.7 | 4.14E-14 |
| 2538176 | 2538038 | 138 | - | - | -2.3 | 3.94E-11 |
| 2644422 | 2644482 | 60 | + | - | 3.6 | 2.90E-15 |
| 2667798 | 2667876 | 78 | + | - | 2.3 | 4.39E-16 |
| 2670884 | 2670869 | 15 | - | - | 2.2 | 2.17E-04 |
| 2730130 | 2730149 | 19 | + | - | -2.6 | 3.91E-05 |
| 2745826 | 2745686 | 140 | - | - | -2.2 | 5.98E-06 |
| 2757997 | 2758085 | 88 | + | - | -2.2 | 3.31E-14 |
| 2783778 | 2783529 | 249 | - | - | 4.7 | 1.62E-81 |
| 2785328 | 2785379 | 51 | + | - | 9.5 | 1.26E-115 |
| 2802845 | 2802731 | 114 | - | - | 7.8 | 1.27E-138 |
| 2888995 | 2889136 | 141 | + | - | -2.9 | 8.00E-18 |
| 2908551 | 2908568 | 17 | + | - | 2.3 | 2.05E-05 |
| 2912026 | 2912098 | 72 | + | - | -2.0 | 1.84E-11 |
| 2925194 | 2925332 | 138 | + | - | 2.0 | 2.31E-08 |
| 2948952 | 2948991 | 39 | + | - | -2.6 | 4.03E-20 |
| 3014294 | 3014188 | 106 | - | - | 2.4 | 2.95E-09 |
| 3038665 | 3038697 | 32 | + | - | 2.6 | 8.13E-07 |
| 3058178 | 3058157 | 21 | - | - | 6.8 | 5.38E-85 |
| 3059330 | 3059292 | 38 | - | - | 10.3 | 1.01E-245 |
| 3060174 | 3060165 | 9 | - | - | 16.0 | 0.00E+00 |
| 3062547 | 3062494 | 53 | - | - | 7.6 | 2.21E-117 |
| 3075621 | 3075597 | 24 | - | - | 2.6 | 6.85E-06 |
| 3216785 | 3216766 | 19 | - | - | -4.5 | 2.93E-21 |
| 3224444 | 3224524 | 80 | + | - | -2.2 | 5.54E-04 |
| 3230291 | 3230204 | 87 | - | - | 2.2 | 4.62E-11 |
| 3232595 | 3232552 | 43 | - | - | 2.6 | 3.95E-10 |
| 3236469 | 3236397 | 72 | - | - | 2.6 | 8.67E-16 |
| 3246608 | 3246619 | 11 | + | - | 4.5 | 1.87E-39 |
| 3249928 | 3249796 | 132 | - | - | 2.5 | 4.57E-14 |
| 3318288 | 3318311 | 23 | + | - | -2.4 | 5.12E-04 |
| 3319542 | 3319571 | 29 | + | - | -2.7 | 4.38E-05 |
| 3347302 | 3347402 | 100 | + | - | -3.8 | 1.79E-13 |
| 3426030 | 3426062 | 32 | + | - | 3.3 | 2.54E-12 |

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|---------|---------|-----|---|-------|------|-----------|
| 3441642 | 3441742 | 100 | + | - | 2.1 | 9.54E-09 |
| 3512853 | 3513092 | 239 | + | - | -2.1 | 2.17E-07 |
| 3546312 | 3546402 | 90 | + | - | 2.4 | 2.15E-10 |
| 3556128 | 3556107 | 21 | - | - | 6.0 | 1.15E-63 |
| 3556978 | 3556966 | 12 | - | - | 2.0 | 6.25E-05 |
| 3630810 | 3630859 | 49 | + | - | -2.1 | 8.32E-03 |
| 3738075 | 3738049 | 26 | - | - | 2.0 | 2.38E-07 |
| 3897277 | 3897483 | 206 | + | - | -2.0 | 5.09E-12 |
| 3959448 | 3959561 | 113 | + | - | 2.7 | 1.07E-14 |
| 3990202 | 3990270 | 68 | + | - | -2.9 | 1.13E-08 |
| 3992674 | 3992782 | 108 | + | - | -2.2 | 5.52E-10 |
| 3997582 | 3997635 | 53 | + | - | -3.5 | 8.01E-19 |
| 4014792 | 4014865 | 73 | + | - | -3.5 | 1.35E-17 |
| 4027326 | 4027728 | 402 | + | - | -2.1 | 1.90E-10 |
| 4146755 | 4146825 | 70 | + | - | 2.1 | 5.19E-05 |
| 4178838 | 4179010 | 172 | + | - | -2.4 | 2.66E-24 |
| 4182929 | 4182999 | 70 | + | - | -2.1 | 1.11E-18 |
| 4198208 | 4197971 | 237 | - | - | 2.5 | 1.12E-47 |
| 4279322 | 4279182 | 140 | - | - | 2.0 | 4.72E-10 |
| 4285964 | 4286194 | 230 | + | - | -5.6 | 7.20E-172 |
| 4401009 | 4401134 | 125 | + | - | 2.0 | 1.87E-07 |
| 4417716 | 4417839 | 123 | + | - | -2.0 | 3.29E-04 |
| 4428274 | 4428241 | 33 | - | - | 2.1 | 5.57E-05 |
| 4485828 | 4485653 | 175 | - | - | 2.0 | 7.69E-12 |
| 4491974 | 4492001 | 27 | + | - | -3.3 | 9.43E-10 |
| 4568030 | 4568008 | 22 | - | - | 2.0 | 1.23E-15 |
| 4571549 | 4571522 | 27 | - | - | 2.1 | 5.20E-04 |
| 4736602 | 4736563 | 39 | - | - | 2.0 | 1.82E-09 |
| 4743227 | 4743250 | 23 | + | - | 3.2 | 1.61E-12 |
| 4774596 | 4774449 | 147 | - | - | -3.7 | 1.49E-36 |
| 4787163 | 4787256 | 93 | + | - | -2.3 | 2.20E-03 |
| 4801008 | 4801020 | 12 | + | - | -3.1 | 4.66E-12 |
| 4806518 | 4806629 | 111 | + | - | 2.4 | 1.03E-15 |
| 4835040 | 4835068 | 28 | + | - | 2.1 | 1.45E-04 |
| 4861607 | 4861446 | 161 | - | - | 2.6 | 4.34E-23 |
| 4888811 | 4888903 | 92 | + | - | -2.6 | 2.41E-31 |
| 4900204 | 4900284 | 80 | + | - | -3.1 | 8.02E-11 |
| 5007181 | 5007215 | 34 | + | - | -2.3 | 2.50E-06 |
| 5029936 | 5029844 | 92 | - | - | -3.3 | 8.21E-11 |
| 5047121 | 5047156 | 35 | + | - | -2.2 | 6.60E-03 |
| 5119903 | 5120244 | 341 | + | - | 2.0 | 1.11E-20 |
| 5174418 | 5174330 | 88 | - | - | 2.6 | 1.58E-14 |
| 5177144 | 5177056 | 88 | - | - | -2.6 | 1.03E-05 |
| 5264754 | 5264995 | 241 | + | - | 2.0 | 5.85E-11 |
| 5292645 | 5292764 | 119 | + | - | -2.1 | 4.92E-09 |
| 5347847 | 5347968 | 121 | + | - | -2.0 | 2.01E-05 |
| 5370203 | 5370018 | 185 | - | - | 3.2 | 8.81E-24 |
| 5388487 | 5388379 | 108 | - | - | 2.1 | 5.86E-10 |
| 5398440 | 5398591 | 151 | + | - | 2.1 | 3.34E-10 |
| 5521532 | 5521511 | 21 | - | - | -2.4 | 4.52E-04 |
| 5556009 | 5555936 | 73 | - | PrrF1 | 9.3 | 1.48E-113 |
| 5563443 | 5563544 | 101 | + | - | 3.6 | 3.07E-23 |

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|---------|---------|-----|---|---|------|----------|
| 5654193 | 5654308 | 115 | + | - | 2.2 | 4.30E-12 |
| 5673607 | 5673586 | 21 | - | - | -4.7 | 5.72E-35 |
| 5732048 | 5732159 | 111 | + | - | -3.0 | 3.50E-07 |
| 5739000 | 5738964 | 36 | - | - | 3.6 | 1.71E-22 |
| 5749614 | 5749550 | 64 | - | - | 6.6 | 1.21E-58 |
| 5789169 | 5789188 | 19 | + | - | -3.1 | 5.85E-12 |
| 6010689 | 6010573 | 116 | - | - | -2.5 | 4.80E-05 |
| 6012155 | 6012094 | 61 | - | - | -2.8 | 2.76E-07 |
| 6015331 | 6015225 | 106 | - | - | -3.0 | 6.09E-29 |
| 6017106 | 6017012 | 94 | - | - | -2.1 | 4.84E-05 |
| 6036887 | 6036877 | 10 | - | - | 2.3 | 2.35E-05 |
| 6043146 | 6043189 | 43 | + | - | -2.6 | 1.77E-05 |
| 6059019 | 6058871 | 148 | - | - | -2.1 | 3.03E-10 |
| 6224090 | 6224069 | 21 | - | - | 3.0 | 4.54E-12 |
| 6258847 | 6258866 | 19 | + | - | 2.2 | 1.11E-04 |
| 6298198 | 6298223 | 25 | + | - | 3.6 | 1.04E-19 |
| 6348125 | 6348178 | 53 | + | - | -3.5 | 6.27E-11 |
| 6348782 | 6348796 | 14 | + | - | -2.5 | 2.11E-05 |
| 6372966 | 6373058 | 92 | + | - | -2.6 | 1.87E-12 |
| 6374602 | 6374807 | 205 | + | - | -2.2 | 4.66E-08 |
| 6382150 | 6382164 | 14 | + | - | -2.6 | 5.22E-07 |
| 6460839 | 6461078 | 239 | + | - | 2.8 | 1.94E-26 |

^aTranscriptional start site

^bTranscriptional stop site

^cEach fold change value represents the average expression of each RNA in PA14 grown in the blood of the TPs/HvS; upregulated RNAs, red; downregulated RNAs, green.

^d*q* value of HVs versus TPs in scientific notation as all values are 0.00 in numeric form.

^eYellow-highlighted asRNAs are discussed in the text.

^fBolded transcripts were also found in the 2018 study by Eckweiler and Häussler.

^gChange in gene expression; red, upregulated; green, downregulated; nc, no significant change.

^hTranscripts encoding the potential amino and carboxy termini of CzcA are highlighted in blue.