**Additional file 8: Table S2.** Log fold changes (LogFC) of genes encoding predicted membrane transporters which are DE upon antibiotic exposure. Non-DE = not differentially expressed in this condition. APC = Amino Acid-Polyamine-Organocation, ABC = ATP-binding cassette, DMT = Drug/Metabolite Transporter, MFS = Major Facilitator Superfamily, MmpL = Mycobacterial membrane protein Large, NRAMP = Metal Ion (Mn2+-iron) Transporter, GAP = Peptidoglycolipid Addressing Protein, Rhtp = Resistance to Homoserine/Threonine

|  |  |  |  |
| --- | --- | --- | --- |
| **Gene** | **LogFC** | | **Transporter type** |
| **Kanamycin** | **Erythromycin** |
| MAB\_3829c | Non-DE | -1.02 | APC |
| MAB\_4323 | Non-DE | 1.32 | APC |
| MAB\_1007c | Non-DE | -1.22 | ABC |
| MAB\_1008c | Non-DE | -1.08 | ABC |
| MAB\_1009c | Non-DE | -1.13 | ABC |
| MAB\_1846 | 1.37 | 2.04 | ABC |
| MAB\_2177 | 1.11 | 1.96 | ABC |
| MAB\_2355c | Non-DE | 2.95 | ABC |
| MAB\_2622c | Non-DE | -1.05 | ABC |
| MAB\_4155c | -1.05 | Non-DE | ABC |
| MAB\_4564c | Non-DE | -1.14 | ABC |
| MAB\_4566c | Non-DE | -1.18 | ABC |
| MAB\_0677c | 1.44 | 2.65 | DMT |
| MAB\_0766 | 1.33 | 4.16 | DMT |
| MAB\_2640c | Non-DE | 1.04 | DMT |
| MAB\_3180 | Non-DE | 1.53 | DMT |
| MAB\_3762 | Non-DE | 1.05 | DMT |
| MAB\_0880 | Non-DE | 2.36 | MFS |
| MAB\_1395 | Non-DE | 2.47 | MFS |
| MAB\_1396 | Non-DE | 1.89 | MFS |
| MAB\_1409c | Non-DE | 3.11 | MFS |
| MAB\_2273 | 1.39 | 3.70 | MFS |
| MAB\_2780c | Non-DE | 2.13 | MFS |
| MAB\_2958 | -2.07 | -1.26 | MFS |
| MAB\_4716c | -1.31 | Non-DE | NRAMP |
| MAB\_2275 | Non-DE | 1.20 | MmpL |
| MAB\_4698 | Non-DE | -1.27 | GAP |
| MAB\_3913 | Non-DE | 2.53 | RhtB |