**Supplementary Table S2. Plasmids used in this study.**

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Name** | **Antibiotic Resistance** | **Description/Sequence** | **Use** | **Source** |
| pMSG430 | *kanR* | contains *attP, tetO,* encodes integrase | To construct *dciAMsm attB::*tet*dciAMtb* and other strains | Described in [1,2] |
| pDB88 | *hygR*, but results in unmarked deletion | Two step allelic exchange (Barkan et al. 2011) | To delete endogenous *dciAMsm* | [3] |
| pDB19 | *zeoR* | contains *attP, tetO,* encodes integrase | Gene-switching/Marker exchange | Described in [1,2] |
| pTetR | *hygR* | Episomal, expresses TetOn Tet Repressor | To create depletion strains | [4] |
| pGEX-6P | *carbR* | Protein purification vector, encocdes N-terminal GST tag | Recombinant expression of mycobacterial proteins in BL21 | GE Healthcare Life Sciences |
| pET-SUMO | *kanR* | Protein purification vector, encocdes N-terminal His-SUMO tag | Recombinant expression of mycobacterial proteins in BL21 | Invitrogen |

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