

Table S2. *Mycobacterium bovis* BCG strains used in this study.

Strain:	Description:	Source/reference:
wt-BCG	<i>M. bovis</i> BCG str. Pasteur 1743P2	Laboratory collection
Δ <i>gltBD</i>	wt-BCG pJV53 background in which the <i>gltBD</i> operon was substituted with a hygromycin cassette, <i>hyg^R</i> , <i>kan^R</i>	[1]
Δ <i>gltBD</i> complement	Genetically complemented strain of Δ <i>gltBD</i> , <i>attB</i> ::pGC <i>gltBD</i> , carries pJV53, <i>hyg^R</i> , <i>kan^R</i> , <i>gent^R</i>	[1]
Δ <i>gdh</i>	wt-BCG background in which the <i>Nrul</i> fragment spanning the GDH domain within <i>gdh</i> was deleted.	[1]
Δ <i>gdh</i> complement	Genetically complemented strain of Δ <i>gdh</i> , <i>attB</i> ::pGC <i>gdh</i> , <i>gent^R</i>	[1]

1. Viljoen AJ, Kirsten CJ, Baker B, van Helden PD, Wiid IJF. The Role of Glutamine Oxoglutarate Aminotransferase and Glutamate Dehydrogenase in Nitrogen Metabolism in *Mycobacterium bovis* BCG. PLoS ONE. 2013;8. doi:10.1371/journal.pone.0084452