

Table 2. Percent identity between Mtb and *M. smegmatis* ortholog promoters<sup>a</sup>

MSMEG gene	H37Rv gene	Percent Identity <sup>b</sup>
0545	Rv1359	57.02
3578	Rv2435	46.3
3780	Rv1647	51.94
4279	Rv2212	78.97
4477	Rv1900	54.42
4924	Rv1318c	54.6
4924	Rv1319c/ Rv1320c	82.08
5018	Rv1264	54.01
6154	Rv2645	69.28

<sup>a</sup> Promoter region defined as the 500 nucleotides upstream of translation start site

<sup>b</sup> Determined with T-Coffee multi-sequence alignment