

Supplementary Table 5. Transcription factor ChIP-seq peaks located within 100 bp upstream and 20 bp downstream of sRNA 5' ends.

sRNA	New nomenclature	Transcription Factor	5' end - peak distance
B55 ^a	ncRv10609A	Rv3249c	-29 ^e
B11 ^a	ncRv13660Ac	Rv0081 ^d	-56
B11 ^a	ncRv13660Ac	Rv0081 ^d	-45
B11 ^a	ncRv13660Ac	Rv3249c	-25
B11 ^a	ncRv13660Ac	Rv2034	-20.5
Mpr5 ^b	ncBCG11109A/ncRv11051A	Rv0324	-27
Mcr11 ^b	ncBCG11323Ac/ncRv11264A	Rv3574	-52
Mcr8/Mpr4 ^b	ncBCG13719A/ncRv13661A	Rv0821c (PhoY2)	-38 ^e
Bo29 ^c	ncBCG11603Ac	Rv2021c	-19
Bo132 ^c	ncBCG13885A	Rv2887	-44 ^e

^a Arnvig and Young, 2009

^b DiChiara *et al.*, 2010

^c this work

^d replicate ChIP-seq datasets are available for Rv0081 and we required that sRNA-proximal ChIP-seq peaks be present in both replicates for inclusion in this table

^e unambiguously associated with an sRNA 5' end