

**Figure S4
(A)**

Protein	Residues	Sequence	Organism
RsbV	51-63	V S Y M D S T G L G V	<i>B. subtilis</i>
BldG	52-64	V D F L D S T G L G V	<i>S. coelicolor</i>
Slr1859	52-64	L E Y M S S A G L R I	<i>Synechocystis</i> PCC 6803
Slr1856	49-61	V P Y M S S A G L R L	<i>Synechocystis</i> PCC 6803
SpoIIAA	53-65	L S F M D S S G L G V	<i>B. subtilis</i>
RsbR	200-212	V P V V D T M V A H H	<i>B. subtilis</i>
RsbS	54-66	V D M I D S F I A K V	<i>B. subtilis</i>
0516c	76-88	L S H F S G A G I S L	<i>M. tuberculosis</i>

(B)

Family Member Sequence

Rv1904	M G S C A Y A
Rv2638	M G C C A F A
Rv1365c	M G C C A V A
Rv0941c	M P V Y P T E
Rv0516c	F S G A G I S
Rv3687c	L G S V G L K
SpoIIAA	M D S S G L G
RsbV	M D S T G L G