

Table S1. SALP proteins that were subjected to phylogenetic analysis. Nomenclature is based on highest sequence homology to the *S. coelicolor* SALP homologues. Database tags refer to the locus name given by the respective genome sequencing projects. Contigs refer to short sequence files extracted from the *S. scabies* database (www.sanger.ac.uk/Projects/S_scabies).

Protein name	Organism of origin	Database tag /contig number
SsgA[Scoel]	<i>S. coelicolor</i>	SCO3926
SsgB[Scoel]	<i>S. coelicolor</i>	SCO1541
SsgC[Scoel]	<i>S. coelicolor</i>	SCO7289
SsgD[Scoel]	<i>S. coelicolor</i>	SCO6722
SsgE[Scoel]	<i>S. coelicolor</i>	SCO3158
SsgF[Scoel]	<i>S. coelicolor</i>	SCO7175
SsgG[Scoel]	<i>S. coelicolor</i>	SCO2924
SsgA[Saver]	<i>S. avermitilis</i>	SAV4267
SsgB[Saver]	<i>S. avermitilis</i>	SAV6810
SsgD[Saver]	<i>S. avermitilis</i>	SAV1687
SsgE[Saver]	<i>S. avermitilis</i>	SAV3605
SsgY[Saver]	<i>S. avermitilis</i>	SAV570
SsgZ[Saver]	<i>S. avermitilis</i>	SAV580
SsgA[Sscab]	<i>S. scabies</i>	contig: scab0274d04.q1k
SsgB[Sscab]	<i>S. scabies</i>	contig: scab0975f02.p1k
SsgD[Sscab]	<i>S. scabies</i>	contig: scab0372d01.q1k
SsgE[Sscab]	<i>S. scabies</i>	contig: scab0136f09.q1k
SsgG[Sscab]	<i>S. scabies</i>	contig: sab0313c04.q1k
SsgV[Sscab]	<i>S. scabies</i>	contig: scab0162c08.q1k
SsgW[Sscab]	<i>S. scabies</i>	contig: scab0039b09.q1k
SsgA[Sgris]	<i>S. griseus</i>	SGR3655
SsgB[Sgris]	<i>S. griseus</i>	SGR5997
SsgD[Sgris]	<i>S. griseus</i>	SGR1004
SsgE[Sgris]	<i>S. griseus</i>	SGR4320
SsgG[Sgris]	<i>S. griseus</i>	SGR4615
SGR128[Sgris]	<i>S. griseus</i>	SGR128
SGR41t[Sgris]	<i>S. griseus</i>	SGR41t
SGR7098t[Sgris]	<i>S. griseus</i>	SGR7098t
SsgA[Salbu]	<i>S. albus</i>	AF195771
SsgA[Sgold]	<i>S. goldeniensis</i>	AF195773
SsgA[Snetr]	<i>S. netropsis</i>	AF195772
SsgB[Tfusc]	<i>Thermobifido fusca</i>	Tfu_2111
SsgB[Kradi]	<i>Kineococcus radiotolerans</i>	Krad_3069
SsgB[Nocar]	<i>Nocardioides</i>	Noca_2368
SsgB[Acell]	<i>Acidothermus cellulolyticus</i>	Acel_1369
SsgB[SalTr]	<i>Salinispora tropica</i>	Strop_1600
SsgB[SalAr]	<i>Salinispora arenicola</i>	Sare_1560
SsgB[Falni]	<i>Frankia alni</i> ACN14a	FRAAL2127
FrS1[Falni]	<i>Frankia alni</i> ACN14a	FRAAL5533
FrS2[Falni]	<i>Frankia alni</i> ACN14a	FRAAL4594
FrS3[Falni]	<i>Frankia alni</i> ACN14a	FRAAL5373
FrS4[Falni]	<i>Frankia alni</i> ACN14a	FRAAL6494
SsgB[Fcci3]	<i>Frankia</i> sp. Ccl3	Francci3_1359
FrS1[Fcci3]	<i>Frankia</i> sp. Ccl3	Francci3_3418
FrS2[Fcci3]	<i>Frankia</i> sp. Ccl3	Francci3_1632
SsgB[Fean1]	<i>Frankia</i> sp. EAN1pec	Franean1_5158
FrS2[Fean1]	<i>Frankia</i> sp. EAN1pec	Franean1_2058
FrS5[Fean1]	<i>Frankia</i> sp. EAN1pec	Franean1_4060
SsgB[Sacch]	<i>Saccharopolyspora erythraea</i>	SACE_1961
SacS1[Sacch]	<i>Saccharopolyspora erythraea</i>	SACE_5535