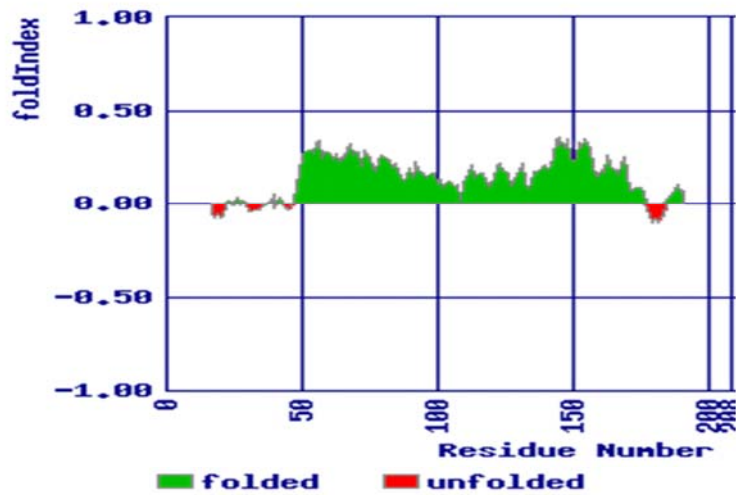


Supplemental Figure Legends

Fig. S1. Prediction of *Mtb* SodC intrinsically folded regions by FoldIndex©.



1 **CSSPQHASTV** **PGTTPSIW**TG**** **SPAPSGLS**GH**** **DEES**PG**AQSL** **TSTLTAPD**GT****
 51 **KVATA**KFE**F**A**** **NGYATV**TI**A**T**** **TGV**GK**LTP**GF**** **HGLH**IQ**V**GK**** **CEPNSV**AP**T**G****
 101 **GAPGN**FL**S**AG**** **GHYH**VP**G**HT**G** **TPAS**GD**L**AS**L** **QVR**GD**S**AML**** **VTTT**DA**F**TMD****
 151 **DLLS**GAK**T**AI**** **IIHAG**AD**N**F**A** **NIP**PER**Y**VQ**V** **NGTP**GP**DE**TT**** **LTT**GD**A**GK**R**V****
 201 **ACGV**IG**S**G****

(Red denotes the predicted disordered segment)

Table S1: rSodC peptide mapping

Trypsin										Precursor Mass (m/z)	Spectrum charge	Actual peptide mass (AMU)	Actual minus calculated peptide mass (AMU)	Peptide start index	Peptide stop index
Peptide	Sequence	Previous amino acid	Next amino acid	Peptide identification probability	SEQUEST XCorr score	SEQUEST DcN score	X! Tandem -log(e) score	Modifications							
Try1	DEESFGAQLSTSLTAPDGTK	H	V	95.00%	5.55	0	0		1053.86	2	2105.71	1.7	23	43	
Try1	DEESFGAQLSTSLTAPDGTK	H	V	95.00%	5.07	0	0		1053.85	2	2105.68	1.7	23	43	
Try1	DEESFGAQLSTSLTAPDGTK	H	V	95.00%	4.86	0	0		1053.42	2	2104.82	0.84	23	43	
Try2	TSLTAPDGTK	L	V	95.00%	2.45	0	0		1092.74	1	1091.73	1.2	33	43	
Try2	TSLTAPDGTK	L	V	95.00%	2.35	0	0		1092.39	1	1091.38	0.83	33	43	
Try2	TSLTAPDGTK	L	V	95.00%	2.13	0	0		1091.69	1	1090.68	0.13	33	43	
Try2	TSLTAPDGTK	L	V	95.00%	2.27	0	0		1091.68	1	1090.67	0.12	33	43	
Try2	TSLTAPDGTK	L	V	95.00%	2.67	0	0		547.23	2	1092.44	1.9	33	43	
Try2	TSLTAPDGTK	L	V	95.00%	2.16	0.915	0		546.76	2	1091.5	0.95	33	43	
Try2	TSLTAPDGTK	L	V	95.00%	2.3	0.904	0		546.74	2	1091.46	0.91	33	43	
Try3	VATAK	K	F	95.00%	2.3	0	0.602		490.66	1	489.65	1.4	44	48	
Try3	VATAK	K	F	95.00%	1.91	0	0.602		490.54	1	489.53	1.2	44	48	
Try3	VATAK	K	F	95.00%	1.98	0	0.678		490.35	1	489.34	1.1	44	48	
Try3	VATAK	K	F	95.00%	1.94	0	0.658		489.96	1	488.95	0.66	44	48	
Try3	VATAK	K	F	95.00%	1.67	0	0.62		489.91	1	488.9	0.61	44	48	
Try3	VATAK	K	F	95.00%	2	0	0.678		489.66	1	488.65	0.36	44	48	
Try3	VATAK	K	F	95.00%	1.87	0	0.678		489.61	1	488.6	0.31	44	48	
Try3	VATAK	K	F	95.00%	1.75	0	0.585		489.6	1	488.59	0.3	44	48	
Try3	VATAK	K	F	95.00%	1.7	0	0.602		489.58	1	488.57	0.28	44	48	
Try3	VATAK	K	F	95.00%	1.98	0	0.638		489.57	1	488.56	0.27	44	48	
Try3	VATAK	K	F	95.00%	1.54	0	0.62		489.56	1	488.55	0.26	44	48	
Try3	VATAK	K	F	95.00%	2.03	0	0.678		489.54	1	488.53	0.24	44	48	
Try3	VATAK	K	F	95.00%	1.86	0	0.638		489.53	1	488.52	0.23	44	48	
Try3	VATAK	K	F	95.00%	1.49	0	0.62		489.51	1	488.5	0.21	44	48	
Try3	VATAK	K	F	95.00%	1.46	0	0.658		489.5	1	488.49	0.2	44	48	
Try3	VATAK	K	F	95.00%	1.37	0	0.602		489.48	1	488.47	0.18	44	48	
Try3	VATAK	K	F	95.00%	1.77	0	0.699		489.47	1	488.46	0.17	44	48	
Try3	VATAK	K	F	95.00%	1.82	0	0.678		489.47	1	488.46	0.17	44	48	
Try3	VATAK	K	F	95.00%	1.93	0	0.678		489.46	1	488.46	0.17	44	48	
Try3	VATAK	K	F	95.00%	1.89	0	0.658		489.46	1	488.45	0.16	44	48	
Try3	VATAK	K	F	95.00%	1.79	0	0.658		489.45	1	488.44	0.15	44	48	
Try3	VATAK	K	F	95.00%	1.74	0	0.678		489.41	1	488.4	0.14	44	48	
Try3	VATAK	K	F	95.00%	1.51	0	0.658		489.17	1	488.16	-0.13	44	48	
Try3	VATAK	K	F	95.00%	1.51	0	0.638		488.87	1	487.86	-0.43	44	48	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	3.44	0	6.85		1950.04	1	1949.03	3.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.97	0	4.04		1949.14	1	1948.13	2.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.94	0	4.27		1949.08	1	1948.07	2.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.81	0	4.11		1949.09	1	1948.08	2.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	3.13	0	5.55		1949.08	1	1948.07	2.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.65	0	4.72		1949.05	1	1948.04	2.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.49	0	3.96		1949.05	1	1948.04	2.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.57	0	3.16		1949.04	1	1948.03	2.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.47	0	5.38		1948.6	1	1947.59	1.6	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.86	0	4.8		1948.23	1	1947.22	1.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	3.01	0	5.11		1948.1	1	1947.09	1.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.06	0	1.39		1948.03	1	1947.02	1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	2.81	0	6.34		1947.99	1	1946.98	1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.36		976.25	2	1950.48	4.5	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.35		976.25	2	1950.48	4.5	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.35		976.22	2	1950.43	4.5	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.92		976.21	2	1950.41	4.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.35		976.2	2	1950.38	4.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.92		976.19	2	1950.37	4.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.92		976.18	2	1950.34	4.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.92		976.18	2	1950.34	4.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.4		976.17	2	1950.32	4.3	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.89		976.1	2	1950.18	4.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	7.14		976.09	2	1950.17	4.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.89		976.08	2	1950.14	4.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	12		976.07	2	1950.12	4.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.19		976.04	2	1950.07	4.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.96		976	2	1949.98	4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	11.2		976	2	1949.99	4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.92		975.99	2	1949.97	4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.17		975.98	2	1949.94	4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	9.68		975.98	2	1949.95	4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	9.62		975.97	2	1949.92	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	7.52		975.96	2	1949.9	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	11.2		975.96	2	1949.9	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	12		975.96	2	1949.91	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	12.7		975.95	2	1949.88	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.96		975.95	2	1949.88	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.19		975.95	2	1949.89	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.96		975.94	2	1949.89	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.89		975.94	2	1949.86	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	9.6		975.92	2	1949.83	3.9	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	10.5		975.9	2	1949.78	3.8	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.57	0	8.96		975.83	2	1949.64	3.7	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.74	0	12		975.73	2	1949.44	3.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.71	0	10.5		975.72	2	1949.43	3.5	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.84	0	9.12		975.7	2	1949.38	3.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.71	0	8.96		975.7	2	1949.39	3.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.63	0	9.64		975.7	2	1949.39	3.4	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.95	0	11.2		975.64	2	1949.26	3.3	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.64	0	10.5		975.63	2	1949.25	3.3	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.48	0	6.92		975.61	2	1949.21	3.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.63	0	10.4		975.59	2	1949.16	3.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.63	0	10.4		975.59	2	1949.17	3.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.56	0	8.96		975.58	2	1949.14	3.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.92	0	12		975.58	2	1949.15	3.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.62	0	10.5		975.58	2	1949.15	3.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.88	0	11.2		975.58	2	1949.15	3.2	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.38	0	10.5		975.57	2	1949.13	3.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.61	0	9		975.54	2	1949.07	3.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	6.02	0	10.5		975.53	2	1949.04	3.1	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.96	0	12.7		975.52	2	1949.02	3	49	67	
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.52	0	11.2		975.52	2	1949.03	3.1	49	67	

Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.75	0	11.2				1949.03	3.1	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.87	0	9.66		975.51	2	1949.01	9.6	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.99	0	10.5		975.49	2	1948.97	3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.65	0	10.4		975.48	2	1948.94	3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.53	0	12		975.47	2	1948.93	3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.99	0	11.2		975.47	2	1948.93	3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.67	0	10.5		975.47	2	1948.93	3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.54	0	10.5		975.46	2	1948.9	3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.62	0	10.5		975.44	2	1948.87	2.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.64	0	11.2		975.44	2	1948.87	2.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.87	0	9.77		975.43	2	1948.84	2.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.6	0	11.9		975.43	2	1948.84	2.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.74	0	11.2		975.42	2	1948.83	2.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.44	0	7.52		975.41	2	1948.8	2.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.95	0	12		975.39	2	1948.77	2.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.54	0	8.96		975.37	2	1948.72	2.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.81	0	10.5		975.36	2	1948.7	2.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.8	0	8.18		975.34	2	1948.67	2.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.86	0	9.64		975.3	2	1948.59	2.6	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.78	0	10.4		975.29	2	1948.57	2.6	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.9	0	12		975.29	2	1948.57	2.6	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.47	0	9.64		975.29	2	1948.57	2.6	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.87	0	11.9		975.24	2	1948.47	2.5	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.66	0	10.5		975.14	2	1948.27	2.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.55	0	11.2		975.12	2	1948.22	2.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.68	0	12		975.1	2	1948.19	2.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.62	0	11.2		975.08	2	1948.15	2.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.6	0	11.2		975.08	2	1948.15	2.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	6.13	0	11.9		975.07	2	1948.13	2.1	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.8	0	11.9		975.05	2	1948.08	2.1	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.58	0	10.6		975.01	2	1948	2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.83	0	10.5		975.07	2	1948.01	2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.66	0	12		974.98	2	1947.95	2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.85	0	11.2		974.98	2	1947.95	2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.89	0	11.2		974.98	2	1947.95	2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.7	0	10.5		974.97	2	1947.93	1.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.59	0	12		974.95	2	1947.89	1.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.72	0	11.2		974.93	2	1947.84	1.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.6	0	10.5		974.92	2	1947.82	1.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.98	0	11.2		974.92	2	1947.82	1.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.72	0	11.2		974.92	2	1947.83	1.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.62	0	12		974.91	2	1947.8	1.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.64	0	11.2		974.9	2	1947.78	1.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.79	0	11.2		974.88	2	1947.75	1.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.84	0	12		974.88	2	1947.7	1.8	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.42	0	12		974.8	2	1947.58	1.6	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.44	0	11.2		974.77	2	1947.53	1.6	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.65	0	11.9		974.73	2	1947.44	1.5	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.49	0	10.5		974.73	2	1947.45	1.5	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.67	0	11.2		974.72	2	1947.42	1.5	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.47	0	12		974.7	2	1947.38	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.66	0	12		974.7	2	1947.39	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.71	0	11.2		974.69	2	1947.36	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.69	0	10.5		974.69	2	1947.37	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.77	0	11.9		974.68	2	1947.34	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.53	0	11.2		974.68	2	1947.35	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.44	0	11.2		974.67	2	1947.32	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.35	0	11.2		974.67	2	1947.32	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.66	0	11.2		974.67	2	1947.33	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.83	0	12.7		974.67	2	1947.33	1.4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.6	0	12.8		974.66	2	1947.3	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.63	0	10.4		974.66	2	1947.3	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.55	0	12		974.66	2	1947.31	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.47	0	11.2		974.66	2	1947.31	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.62	0	11.2		974.65	2	1947.28	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.52	0	11.2		974.65	2	1947.28	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.48	0	11.2		974.65	2	1947.29	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.52	0	12.8		974.64	2	1947.27	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.66	0	11.2		974.63	2	1947.24	1.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.59	0	11.2		974.61	2	1947.2	1.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.71	0	11.2		974.6	2	1947.18	1.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.23	0	11.3		974.6	2	1947.19	1.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.54	0	11.2		974.59	2	1947.17	1.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.42	0	10.5		974.58	2	1947.14	1.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.45	0	11.2		974.53	2	1947.04	1.1	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.39	0	10.5		974.49	2	1946.96	0.98	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.69	0	12		974.39	2	1946.77	0.79	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.33	0	12.8		974.22	2	1946.42	0.44	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	4.86	0	12.8		974.13	2	1946.24	0.26	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	4.79	0	12		974.12	2	1946.23	0.25	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	5.05	0	12		974.11	2	1946.2	0.22	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.33		651.1	3	1950.21	4.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.68		651.1	3	1950.27	4.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	10.3		651.1	3	1950.28	4.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	7.47		651.1	3	1950.28	4.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	10.3		651.09	3	1950.23	4.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	6.85		651.09	3	1950.24	4.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	10.4		651.08	3	1950.21	4.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.57		651.08	3	1950.21	4.2	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	10.5		651.05	3	1950.23	4.3	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	8.33		651.01	3	1950	4	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	10.5		650.98	3	1949.9	3.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	9.1		650.95	3	1949.83	3.9	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	9.89		650.91	3	1949.71	3.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	0	0	9.57		650.9	3	1949.66	3.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	4.52	0	8.27		650.89	3	1949.64	3.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	4.55	0	12.3		650.89	3	1949.65	3.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	4.49	0	11.8		650.89	3	1949.65	3.7	49	67
Try4	FEFANGYATVHIATTGVK	K	L	95.00%	4.54	0	10.4		650.84	3	1949.61	3.6	49	67
Try4														

Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.6	0	9.72	650.84	3	1949.5	3.5	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.72	0	11.6	650.83	3	1949.46	3.5	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.15	0	10.9	650.81	3	1949.39	3.4	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.58	0	10.9	650.81	3	1949.4	3.4	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.69	0	8.6	650.78	3	1949.3	3.3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.71	0	12.4	650.77	3	1949.27	3.3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.45	0	12.2	650.75	3	1949.21	3.2	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.23	0	11.8	650.75	3	1949.21	3.2	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.57	0	10.4	650.75	3	1949.23	3.3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.55	0	10.4	650.72	3	1949.14	3.2	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.72	0	10.6	650.71	3	1949.1	3.1	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.6	0	12.7	650.7	3	1949.07	3.1	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.52	0	11.1	650.68	3	1949	3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.64	0	10.5	650.66	3	1948.95	3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.79	0	11.2	650.66	3	1948.96	3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.6	0	10.7	650.65	3	1948.93	3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.51	0	10.9	650.64	3	1948.89	2.9	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.66	0	12.3	650.64	3	1948.9	2.9	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.26	0	11.6	650.64	3	1948.9	2.9	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.17	0	13	650.59	3	1948.73	2.8	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.33	0	11.8	650.58	3	1948.72	2.7	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.68	0	12.4	650.56	3	1948.66	2.7	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.36	0	10.9	650.55	3	1948.63	2.6	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.41	0	10.9	650.53	3	1948.55	2.6	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.54	0	10.9	650.53	3	1948.58	2.6	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.84	0	12.3	650.52	3	1948.52	2.5	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.49	0	11.2	650.51	3	1948.49	2.5	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.98	0	10.5	650.5	3	1948.48	2.5	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.8	0	11.6	650.47	3	1948.39	2.4	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	5.02	0	9.8	650.44	3	1948.3	2.3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.25	0	12.5	650.42	3	1948.24	2.3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.17	0	12.9	650.42	3	1948.24	2.3	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.55	0	11.6	650.37	3	1948.09	2.1	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.44	0	13.1	650.36	3	1948.04	2.1	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.88	0	12.6	650.36	3	1948.04	2.1	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.3	0	11.6	650.35	3	1948.05	2.1	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.5	0	12.6	650.35	3	1948.02	2	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.45	0	10.4	650.31	3	1947.9	1.9	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.54	0	13.2	650.26	3	1947.74	1.8	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.44	0	14	650.23	3	1947.65	1.7	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.3	0	11.7	650.23	3	1947.65	1.7	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.16	0	11.7	650.23	3	1947.66	1.7	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.67	0	11	650.21	3	1947.6	1.6	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.63	0	10.9	650.21	3	1947.6	1.6	49	67	
Try4	FEFANGYATVTHATTGVK	K	L	95.00%	4.43	0	11.6	650.19	3	1947.55	1.6	49	67	
Try5	LTPGFHGLHHQVVK	K	C	95.00%	0	0	6.35	548.53	3	1642.56	2.7	68	82	
Try5	LTPGFHGLHHQVVK	K	C	95.00%	0	0	6.5	548.23	3	1641.65	1.8	68	82	
Try5	LTPGFHGLHHQVVK	K	C	95.00%	0	0	6.59	547.6	3	1639.77	-0.12	68	82	
Try6	HHQVVK	L	C	95.00%	2.41	0.981	0	410.44	2	818.87	1.4	76	82	
Try6	HHQVVK	L	C	95.00%	2.41	0.88	0	409.91	2	817.81	0.36	76	82	
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.1	M6: Oxidation (+16), M16: Oxidation (+16)	1226.36	2	2450.7	2.6	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.52	M6: Oxidation (+16), M16: Oxidation (+16)	1226.01	2	2450	1.9	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.14	M6: Oxidation (+16), M16: Oxidation (+16)	1225.97	2	2449.93	1.6	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.28	M6: Oxidation (+16), M16: Oxidation (+16)	1225.69	2	2449.36	1.3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.68	M6: Oxidation (+16), M16: Oxidation (+16)	1225.52	2	2449.03	0.93	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.18	M6: Oxidation (+16), M16: Oxidation (+16)	1225.52	2	2449.03	0.93	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.48	M6: Oxidation (+16), M16: Oxidation (+16)	1225.49	2	2448.96	0.86	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	9.64	M6: Oxidation (+16), M16: Oxidation (+16)	1225.48	2	2448.95	0.85	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.54	M16: Oxidation (+16)	1219.06	2	2436.1	4	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.8	M6: Oxidation (+16)	1218.98	2	2435.94	3.8	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.89	M6: Oxidation (+16)	1218.97	2	2435.32	3.2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	9.11	M6: Oxidation (+16)	1218.6	2	2435.18	3.1	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.54	M16: Oxidation (+16)	1218.59	2	2435.17	3.1	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.89	M6: Oxidation (+16)	1218.58	2	2435.15	3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.12	M6: Oxidation (+16)	1218.58	2	2435.15	3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.28	M6: Oxidation (+16)	1218.55	2	2435.08	3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.52	M6: Oxidation (+16)	1218.55	2	2435.09	3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.89	M6: Oxidation (+16)	1218.55	2	2435.09	3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.72	M6: Oxidation (+16)	1218.54	2	2435.06	3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.57	M6: Oxidation (+16)	1218.54	2	2435.07	3	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.8	M6: Oxidation (+16)	1218.53	2	2435.04	2.9	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.82	M16: Oxidation (+16)	1218.5	2	2434.99	2.9	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	9.15	M6: Oxidation (+16)	1218.49	2	2434.97	2.9	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.41	M6: Oxidation (+16)	1218.48	2	2434.94	2.8	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.57	M6: Oxidation (+16)	1218.48	2	2434.94	2.8	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.72	M6: Oxidation (+16)	1218.48	2	2434.95	2.8	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.32	M6: Oxidation (+16)	1218.47	2	2434.92	2.8	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	9.72	M6: Oxidation (+16)	1218.47	2	2434.93	2.8	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.77	M6: Oxidation (+16)	1218.43	2	2434.84	2.7	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.17	M6: Oxidation (+16)	1218.42	2	2434.83	2.7	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.6	M16: Oxidation (+16)	1218.37	2	2434.73	2.6	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.8	M6: Oxidation (+16)	1218.35	2	2434.69	2.6	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.7	M6: Oxidation (+16)	1218.3	2	2434.59	2.5	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.17	M6: Oxidation (+16)	1218.26	2	2434.51	2.4	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.72	M6: Oxidation (+16)	1218.17	2	2434.32	2.2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	9.02	M6: Oxidation (+16)	1218.17	2	2434.32	2.2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.54	M6: Oxidation (+16)	1218.14	2	2434.27	2.2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.4	M6: Oxidation (+16)	1218.12	2	2434.22	2.1	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	9	M6: Oxidation (+16)	1218.09	2	2434.16	2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.4	M6: Oxidation (+16)	1218.09	2	2434.16	2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	10	M6: Oxidation (+16)	1218.07	2	2434.12	2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.96	M6: Oxidation (+16)	1218.07	2	2434.12	2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	9.02	M6: Oxidation (+16)	1218.07	2	2434.12	2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	8.18	M6: Oxidation (+16)	1218.06	2	2434.1	2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	7.25	M6: Oxidation (+16)	1218.06	2	2434.11	2	126	149
Try7	GDGSAMLVTTFDAFTMDDLSSGAK	R	T	95.00%	0	0	6.92	M6: Oxidation (+16)	1218.06	2	2434			

Chy15	ANIPPERY	F	V	95.00%	2.14	0	2.59	480.59	2	959.17	0.69	162	169
Chy15	ANIPPERY	F	V	95.00%	2.13	0	2.59	480.55	2	959.09	0.61	162	169
Chy15	ANIPPERY	F	V	95.00%	2.1	0	2.59	480.49	2	958.97	0.49	162	169
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.37	0	3.62	1429.9	1	1428.89	2.2	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.34	0	3.6	1429.81	1	1428.8	2.1	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.47	0	4.12	1429.76	1	1428.75	2.1	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.08	0	3.5	1428.41	1	1427.4	0.71	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.2	0	3.54	1428.39	1	1427.38	0.69	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2	0	3.54	1428.37	1	1427.36	0.67	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	3.14	0	2.89	715.8	2	1429.59	2.9	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	3.07	0	6.01	715.21	1	1428.4	1.7	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.87	0	6.01	714.95	2	1427.88	1.2	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.53	0	6.66	714.87	2	1427.73	1	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	1.75	0	4.01	714.84	2	1427.67	0.98	170	183
Chy16	VQVNGTPGPDFTLL	Y	T	95.00%	2.25	0	3.77	714.81	2	1427.6	0.91	170	183

Thermolysin

Peptide sequence	Previous amino acid	Next amino acid	Peptide identification probability	SEQUEST_XC score	SEQUEST_FC score	XU-Tandem_log(c) score	Modifications	Peptidom Mass (m/z)	Spectrum charge	Actual peptide mass (AMU)	Actual minus calculated peptide mass (AMU)	Peptide start index	Peptide stop index
TL1	LTAPDGTK	T	V	95.00%	1.21	0	1.89	802.8	1	801.79	0.37	36	43
TL1	LTAPDGTK	T	V	95.00%	1.42	0	1.85	802.67	1	801.66	0.24	36	43
TL1	LTAPDGTK	T	V	95.00%	1.4	0	1.85	802.53	1	801.52	0.099	36	43
TL1	LTAPDGTK	T	V	95.00%	1.4	0	1.85	802.46	1	801.45	0.029	36	43
TL2	IATFGGK	T	L	95.00%	1.56	0	1.13	746.52	1	745.51	0.079	60	67
TL3	VAPTGGAGNPF	S	L	95.00%	2.56	0	3.72	989.53	1	988.52	2	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	1.85	0	3.72	987.73	1	986.72	0.24	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	1.52	0	2.62	987.7	1	986.69	0.21	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	1.79	0	3.72	987.62	1	986.61	0.13	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	1.57	0	3.13	987.6	1	986.59	0.11	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	1.73	0	4.28	987.52	1	986.51	0.03	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	1.88	0	4.39	987.51	1	986.5	0.02	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	1.72	0	1.55	987.5	1	986.49	0.0099	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	2.42	0	6.72	494.98	2	987.95	1.5	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	2.74	0	6.07	494.55	2	987.08	0.6	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	2.73	0	6.72	494.48	2	986.94	0.46	88	98
TL3	VAPTGGAGNPF	S	L	95.00%	2.64	0	6.72	494.46	2	986.91	0.43	88	98
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	3.19	0	7.7	1095.85	2	2189.69	1.7	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	3.47	0	7.03	1095.42	2	2188.82	0.79	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	3.04	0	8.43	1095.34	2	2188.67	0.64	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	2.85	0	7.03	1095.3	2	2188.59	0.56	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	1.33	0	4.04	730.9	3	2189.68	1.7	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	1.75	0	4.85	730.75	3	2189.23	1.2	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	1.88	0	5.75	730.69	3	2189.05	1	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	1.8	0	5.7	730.6	3	2188.78	0.75	99	121
TL4	LSAGGHYVPGHTGTPASGDLS	F	L	95.00%	2.3	0	7.92	730.11	3	2187.3	-0.73	99	121
TL5	LSAGGHYVPGHTGTPASGD	F	L	95.00%	3.61	0	7.96	640.81	3	1919.41	2.5	99	118
TL5	LSAGGHYVPGHTGTPASGD	F	L	95.00%	4.09	0	8.14	640.5	3	1918.48	1.6	99	118
TL6	VRGDGSAML	Q	V	95.00%	1.93	0	2.08	905.7	1	904.69	0.25	124	132
TL6	VRGDGSAML	Q	V	95.00%	1.78	0	2.08	905.63	1	904.62	0.18	124	132
TL7	VQVNGTPGDETT	Y	L	95.00%	1.84	0.931	3.96	1315.56	1	1314.55	0.94	170	182
TL7	VQVNGTPGDETT	Y	L	95.00%	2	0.97	6	658.56	2	1315.1	1.5	170	182
TL8	LITGDAGR	T	V	95.00%	1.74	0	1.85	918.56	1	917.55	0.059	183	191
TL8	LITGDAGR	T	V	95.00%	2.51	0	2.72	460.2	2	918.39	0.9	183	191
TL8	LITGDAGR	T	V	95.00%	2	0	3.96	460.17	2	918.33	0.84	183	191
TL8	LITGDAGR	T	V	95.00%	2.37	0	3.48	460.11	2	918.21	0.72	183	191
TL8	LITGDAGR	T	V	95.00%	2.77	0	3.48	460.03	2	918.04	0.55	183	191
TL8	LITGDAGR	T	V	95.00%	2.27	0	3.02	459.97	2	917.92	0.43	183	191
TL8	LITGDAGR	T	V	95.00%	1.99	0	2.47	459.92	2	917.83	0.34	183	191
TL9	LHHHHHH	K	-	95.00%	2.19	0	1.7	478.19	2	954.36	0.91	202	208
TL9	LHHHHHH	K	-	95.00%	2.43	0	2.21	478.06	2	954.1	0.65	202	208
TL9	LHHHHHH	K	-	95.00%	2.28	0	1.57	478	2	953.98	0.53	202	208
TL9	LHHHHHH	K	-	95.00%	2.69	0	2.28	477.99	2	953.96	0.51	202	208
TL9	LHHHHHH	K	-	95.00%	2.28	0	2.38	319.15	3	954.42	0.97	202	208

Table SII. Plasmids and primers used in this study

	Description/Sequence ^a	Reference or Source
Plasmid		
pCR4Blunt-TOPO	cloning vector, Amp ^r , Kan ^r	Invitrogen
pET23b	<i>E. coli</i> expression vector, Amp ^r	Novagen
pVV16	<i>Mycobacteria</i> expression vector, Kan ^r , Hyg ^r	Schulbach <i>et al.</i> (2001)
pMRLB60	<i>rv0432</i> gene fragment lacking signal peptide sequence in pet23b	This study
pMRLB61	Full-length <i>rv0432</i> gene in pVV16	This study
pMRLB61.1	Derivative of pMRLB61 encoding Rv0432 Thr ₄₁ to Ala ₄₁ substitution	This study
pMRLB61.2	Derivative of pMRLB61 encoding Rv0432 Thr ₄₅ to Ala ₄₅ substitution	This study
pMRLB61.3	Derivative of pMRLB61 encoding Rv0432 Thr ₄₆ to Ala ₄₆ substitution	This study
pMRLB61.4	Derivative of pMRLB61 encoding Rv0432 Thr ₄₅ Thr ₄₆ to Ala ₄₅ Ala ₄₆ substitution	This study
pMRLB61.5	Derivative of pMRLB61 encoding Rv0432 Thr ₅₁ to Ala ₅₁ substitution	This study
pMRLB61.6	Derivative of pMRLB61 encoding Rv0432 Thr ₁₃₁ to Ala ₁₃₁ substitution	This study
pMRLB62	<i>rv0432</i> gene fragment lacking signal peptide sequence in pVV16	This study
Primer		
SodC(-SP)F	<u>5'-CATATGTGCTCGTCGCCGAG</u>	This study
SodCR1	<u>5'-CTCGAGGCCGGAACCAATGAC</u>	This study
SodC(+SP)F	<u>5'-CATATGCCAAAGCCCGCGAT</u>	This study
SodCR2	<u>5'-AAGCTTGCCGGAACCAATGAC</u>	This study
SodCT41AF	5'-GCCGCAGCACGCGT TCG CCGTTCCGGGTACCCACGCC	This study
SodCT41AR	5'-GGCGTGTACCCGGA AGGG CAGACGCGTGCTGCGGC	This study
SodCT45AF	5'-CTACAGTTCGGGT GGC ACGCCGTCGATTTGGACC	This study
SodCT45AR	5'-GGTCAAATCGACGGCG TGG CACCCGGA ACT GTAG	This study
SodCT46AF	5'-CTACAGTTCGGGTACCG CGC CGTCGATTTGGACC	This study
SodCT46AR	5'-GGTCAAATCGAC GGC CGGTACCCGGA ACT GTAG	This study
SodCTTAAF	5'-CTACAGTTCGGGT GGC CGCGCCGTCGATTTGGACC	This study
SodCTTAAR	5'-GGTCAAATCGAC GGC CGCGCCGGA ACT GTAG	This study
SodCT51AF	5'-GCCGTCGATTTGG CCG GATCGCCCGCGCC	This study
SodCT51AR	5'-GGCGGGCGGAT CCGG CCCAAATCGACGGC	This study
SodCT131AF	5'-CAACTCGGTTGCC CCG CCGGCGGTGCGCC	This study
SodCT131AR	5'-GGCGCACCG CGG CGGGGGCAACCGAGTTG	This study

a. Restriction endonuclease sites are underlined, site-directed mutations are shown in bold