

Organism	Domain	KsgA/Dim1	mtTFB	mtTFB1	mtTFB2	Pfc1
<i>Arabidopsis thaliana</i> (at)	Eukarya	NP_182264				NP_171690
<i>Dictyostelium discoideum</i> (dd)	Eukarya	XP_638864	XP_637775			
<i>Leishmania braziliensis</i> (lb)	Eukarya	XP_001566655				
<i>Giardia lamblia</i> (gl)	Eukarya	XP_001704233				
<i>Plasmodium vivax</i> (pv)	Eukarya	XP_001616795				
<i>Homo sapiens</i> (hs)	Eukarya	NP_055288		NP_057104	NP_071761	
<i>Saccharomyces cerevisiae</i> (sc)	Eukarya	NP_015057	P14908			
<i>Drosophila melanogaster</i> (dm)	Eukarya			NP_996062	NP_649971	
<i>Caenorhabditis elegans</i> (ce)	Eukarya			NP_491242		
<i>Methanopyrus kandleri</i> (mk)	Archaea	P_614217				
<i>Methanosaeta thermophila</i> (mth)	Archaea	YP_843421				
<i>Haloquadratum walsbyi</i> (hw)	Archaea	YP_658586				
<i>Methanoculleus marisnigri</i> (mma)	Archaea	YP_001047866				
<i>Methanocaldococcus jannaschii</i> (mj)	Archaea	NP_248023				
<i>Pyrococcus horikoshii</i> (ph)	Archaea	NP_143656				
<i>Methanospaera stadtmanae</i> (ms)	Archaea	YP_448554				
<i>Picrophilus torridus</i> (pt)	Archaea	YP_023164				
<i>Archaeoglobus fulgidus</i> (af)	Archaea	NP_070611				
<i>Aeropyrum pernix</i> (ap)	Archaea	NP_147322				
<i>Sulfolobus solfataricus</i> (ss)	Archaea	NP_342254				
<i>Pyrobaculum aerophilum</i> (pa)	Archaea	NP_560585				
<i>Cenarchaeum symbiosum</i> (cs)	Archaea	YP_875407				
<i>Synechococcus elongatus</i> (se)	Bacteria	YP_171912				
<i>Bacillus subtilis</i> (bs)	Bacteria	NP_387923				
<i>Mycobacterium tuberculosis</i> (mtu)	Bacteria	NP_215526				
<i>Thermus thermophilus</i> (tt)	Bacteria	YP_005890				
<i>Bacteroides fragilis</i> (bf)	Bacteria	YP_098063				
<i>Chlamydia trachomatis</i> (ct)	Bacteria	NP_219862				
<i>Borrelia burgdorferi</i> (bb)	Bacteria	NP_212724				

<i>Escherichia coli</i> (ec)	Bacteria	NP_414593				
<i>Anopheles gambiae</i> (ag)	Eukarya			XP_320403		
<i>Apis mellifera</i> (am)	Eukarya			XP_001121333		
<i>Xenopus laevis</i> (xl)	Eukarya			NP_001079850		
<i>Takifugu rubripes</i> (tr)	Eukarya			ENSTRUP0000024475*	ENSTRUP0000026212*	
<i>Ciona intestinalis</i> (ci)	Eukarya			ENSCINP0000008063*		
<i>Rattus norvegicus</i> (rn)	Eukarya				P_001008294	
<i>Pan troglodytes</i> (pt)	Eukarya				XP_514318	
<i>Mus musculus</i> (mmu)	Eukarya				NP_032275	
<i>Bos taurus</i> (bt)	Eukarya				NP_001033216	
<i>Tetraodon nigroviridis</i> (tn)	Eukarya				CAG11026	
<i>Schizosaccharomyces pombe</i> (sp)	Eukarya		NP_593495			
<i>Kluyveromyces lactis</i> (kl)	Eukarya		XP_451899			
<i>Eremothecium gossypii</i> (eg)	Eukarya		NP_986111			
<i>Candida albicans</i> (ca)	Eukarya		XP_722572			
<i>Trypanosoma brucei</i> (tb)	Eukarya		AAM97535			
<i>Leishmania major</i> (lm)	Eukarya		XP_843186			
Erm enzymes						
Organism	Protein	Accession				
<i>Staphylococcus aureus</i>	ErmA	CAA26964				
<i>Enterococcus faecalis</i>	ErmB	CAA68299				
<i>Bacillus subtilis</i>	ErmC	P13956				
<i>Bacillus licheniformis</i>	ErmD	AAA22599				
<i>Saccharopolyspora erythraea</i>	ErmE	AAA26492				
<i>Bacteroides fragilis</i>	ErmF	AAA98217				
<i>Lysinibacillus sphaericus</i>	ErmG	AAA22419				
<i>Streptomyces thermotolerans</i>	ErmH	AAC32026				

<i>Streptomyces fradiae</i>	ErmN	CAA66307				
<i>Streptomyces coelicolor</i>	ErmO	NP_630197				
<i>Clostridium perfringens</i>	ErmQ	AAC36915				
<i>Aeromicrobium erythreum</i>	ErmR	P09891				
<i>Streptomyces fradiae</i>	ErmS	AAA26742				
<i>Lactobacillus reuteri</i>	ErmT	Q00014				
<i>Streptomyces lincolnensis</i>	ErmU	AAB23456				
<i>Streptomyces viridochromogenes</i>	ErmV	AAB51440				
<i>Micromonospora griseorubida</i>	ErmW	BAA03402				
<i>Corynebacterium jeikeium</i>	ErmX	AAK28910				
<i>Staphylococcus aureus</i>	ErmY	BAB20748				
<i>Streptomyces ambofaciens</i>	ErmZ	CAM96571				
<i>Streptomyces venezuelae</i>	Erm30	AAC69328				
<i>Streptomyces venezuelae</i>	Erm31	AAC69327				
<i>Staphylococcus sciuri</i>	Erm33	CAC86410				
<i>Bacillus clausii</i>	Erm34	AAP74657				
<i>Bacteroides coprosuis</i>	Erm35	AAK07612				
<i>Micrococcus luteus</i>	Erm36	AAL68827				
<i>Mycobacterium tuberculosis</i>	Erm37	CAA98396				
<i>Mycobacterium smegmatis</i>	Erm38	AAN86837				
<i>Mycobacterium fortuitum</i>	Erm39	AAR92235				
<i>Mycobacterium mageritense</i>	Erm40	AAS76623				
<i>Mycobacterium abscessus</i>	Erm41	ABW06859				

* Indicates Ensembl peptide ID numbers; all others are NCBI accession numbers.