

Terminals	Accession Numbers	Terminals	Accession Numbers
SMG-1 <i>Homo sapiens</i>	NP_055907.3	ATM <i>Harpegnathos saltator</i>	EFN88218.1
SMG-1 <i>Pan troglodytes</i>	XP_001152224.1	ATM <i>Aedes aegypti</i>	XP_001649972.1
SMG-1 <i>Canis familiaris</i>	XP_851552.1	ATM <i>Branchiostoma floridae</i>	ACG68443.1
SMG-1 <i>Bos taurus</i>	XP_603490.3	ATM <i>Stronylocentrotus purpuratus</i>	NP_001107666.1
SMG-1 <i>Mus musculus</i>	NP_001026984.1	ATM <i>Daphnia pulex</i>	EFX88021.1
SMG-1 <i>Gallus gallus</i>	XP_414907.2	ATM <i>Nematostella vectensis</i>	XP_001628467.1
SMG-1 <i>Danio rerio</i>	NP_001073513.2	ATM <i>Trichoplax adhaerens</i>	XP_002118267.1
SMG-1 <i>Ailuropoda melanoleuca</i>	XP_002920515.1	ATM <i>Tribolium castaneum</i>	EFA11787.1
SMG-1 <i>Xenopus tropicalis</i>	XP_002932267.1	ATM <i>Schizosaccharomyces pombe</i>	NP_588126.1
SMG-1 <i>Tribolium castaneum</i>	XP_969532.1	ATM <i>Schistosoma mansoni</i>	XP_002577847
SMG-1 <i>Nematostella vectensis</i>	XP_001640500.1	TRRAP <i>Homo sapiens</i>	NP_003487.1
SMG-1 <i>Branchiostoma floridae</i>	XP_002607176.1	TRRAP <i>Pan troglodytes</i>	XP_001136894.1
SMG-1 <i>Camponotus floridanus</i>	EFN74897.1	TRRAP <i>Canis familiaris</i>	XP_860949.1
SMG-1 <i>Solenopsis invicta</i>	EFZ20104.1	TRRAP <i>Bos taurus</i>	XP_583735.3
SMG-1 <i>Apis mellifera</i>	XP_001122895.1	TRRAP <i>Mus musculus</i>	NP_001074831.1
SMG-1 <i>Acyrtosiphon pisum</i>	XP_001947123.1	TRRAP <i>Rattus norvegicus</i>	XP_213706.4
SMG-1 <i>Drosophila virilis</i>	XP_002057072.1	TRRAP <i>Gallus gallus</i>	XP_414752.2
SMG-1 <i>Trichoplax adhaerens</i>	XP_002116869.1	TRRAP <i>Danio rerio</i>	XP_001919276.2
SMG-1 <i>Daphnia pulex</i>	EFX69271.1	TRRAP <i>Drosophila melanogaster</i>	NP_001097192.1
SMG-1 <i>Schistosoma mansoni</i>	XP_002575156.1	TRRAP <i>Anopheles gambiae</i>	XP_556172.2
SMG-1 <i>Ixodes scapularis</i>	XP_002414133.1	TRRAP <i>Saccharomyces cerevisiae</i>	NP_011967.1
TOR <i>Homo sapiens</i>	NP_004949.1	TRRAP <i>Daphnia pulex</i>	EFX90201.1
TOR <i>Mus musculus</i>	NP_064393.1	TRRAP <i>Tribolium castaneum</i>	EFA05277.1
TOR <i>Danio rerio</i>	NP_001070679.2	DNA-PK <i>Homo sapiens</i>	P78527.3
TOR <i>Drosophila melanogaster</i>	NP_524891.1		
TOR <i>Anopheles gambiae</i>	XP_317619.2		
TOR <i>Branchiostoma floridae</i>	EEA56493.1		
TOR <i>Apis mellifera</i>	XP_625130.1		
TOR <i>Tribolium castaneum</i>	XP_971819.1		
TOR <i>Acyrtosiphon pisum</i>	XP_001948118.1		
TOR <i>Trichoplax adhaerens</i>	XP_002113370.1		
TOR <i>Aedes aegypti</i>	XP_001650255.1		
TOR <i>Nematostella vectensis</i>	XP_001637040.1		
TOR1 <i>Saccharomyces cerevisiae</i>	CAA52849.1		
TOR2 <i>Saccharomyces cerevisiae</i>	NP_012719.2		
TOR <i>Schistosoma mansoni</i>	XP_002571630.1		
ATR <i>Homo sapiens</i>	NP_001175.2		
ATR <i>Danio rerio</i>	XP_696163.4		
ATR <i>Schistosoma mansoni</i>	XP_002578622.1		
ATR <i>Trichoplax adhaerens</i>	XP_002110572.1		
ATR <i>Daphnia pulex</i>	EFX77801.1		
ATR <i>Branchiostoma floridae</i>	XP_002594547.1		
ATR <i>Tribolium castaneum</i>	EFA09227.1		
ATR <i>Drosophila melanogaster</i>	AAC46881.1		
RAD3 <i>Schizosaccharomyces pombe</i>	NP_595357.1		
MEC1P <i>Saccharomyces cerevisiae</i>	NP_009694.1		
ATM <i>Homo sapiens</i>	NP_000042.3		
Tel1p <i>Saccharomyces cerevisiae</i>	AAA69802.1		
ATM <i>Bruqia malayi</i>	XP_001899536.1		