

Accession number	Gene name	Forward primer	Reverse primer
Tb927.8.6660	PFC1	ggggacaactttgtacaaaaaagtggcatgATGACGACGATGCAGAG	ggggacaactttgtacaagaaagtgggTAAGGTGCGAAGGCTG
Tb927.8.4970	PFR2	ggggacaactttgtacaaaaaagtggcatgATGAGCGGAAAGGAAAGTT	ggggacaactttgtacaagaaagtgggTACTGAGTGATCTGCGG
Tb927.8.1550	PFC3	ggggacaactttgtacaaaaaagtggcatgATGTCGGCCACTTATCC	ggggacaactttgtacaagaaagtgggTATGCGCACTCCTCTAC
Tb927.6.4140	PFC4	ggggacaactttgtacaaaaaagtggcatgATGCCAAACCTCGTTGT	ggggacaactttgtacaagaaagtgggGAGCGAGTGAATAAAACTTAATGC
Tb927.3.4290	PFR1	ggggacaactttgtacaaaaaagtggcatgGTGACGATGCCACTG	ggggacaactttgtacaagaaagtgggGGTGCAGTGTGTGTTCC
Tb927.3.3770	PFC6	ggggacaactttgtacaaaaaagtggcatgATGAGTGCAACAGAGGC	ggggacaactttgtacaagaaagtgggTAGTACCTAGAGAACCCTGTGTC
Tb927.3.3750	PFC7	ggggacaactttgtacaaaaaagtggcatgTCGACTCAACTTATTTCCGGT	ggggacaactttgtacaagaaagtgggCGCAGCCACGAAAGAA
Tb927.2.5660	ADKA	ggggacaactttgtacaaaaaagtggcatgGCGGTACTGAGCGAAG	ggggacaactttgtacaagaaagtgggCGAAAGCGGTGCTTCT
Tb11.01.6510	PFC9	ggggacaactttgtacaaaaaagtggcatgATGTCGTGTCGACCCCA	ggggacaactttgtacaagaaagtgggTGAGCCTCCGATTTTC
Tb11.01.5100	PAR1	ggggacaactttgtacaaaaaagtggcatgATGATCGAGGTGCAACTAAC	ggggacaactttgtacaagaaagtgggCGAATCGCCCAACTCGA
Tb10.70.7330	ADKB	ggggacaactttgtacaaaaaagtggcatgGACCTTCCGATAATAAGTTTGC	ggggacaactttgtacaagaaagtgggCGGCTCTCCACTGTTT
Tb10.6k15.0810	PFC14	ggggacaactttgtacaaaaaagtggcatgATGGAGAAACACAACCGT	ggggacaactttgtacaagaaagtgggTACTCTTTCACAAGAGCCC
Tb10.61.1260	PFC15	ggggacaactttgtacaaaaaagtggcatgATGGCAAGTTCTACAGCAA	ggggacaactttgtacaagaaagtgggTACATCTCTGGTCTCTCT
Tb10.26.0680	PFC16	ggggacaactttgtacaaaaaagtggcatgATGTCTCACTACCTTTGGTCA	ggggacaactttgtacaagaaagtgggCAGCTATCCTCAGCAAGAAG
Tb927.2.4330	PFR5	ggggacaactttgtacaaaaaagtggcatgATGACGCAATCAGCTCAA	ggggacaactttgtacaagaaagtgggCAATCATCTTCATCACAGGGT
Tb927.7.6970	PFR6	ggggacaactttgtacaaaaaagtggcatgAAAAAGAGTGGTCATGAGGG	ggggacaactttgtacaagaaagtgggTATGAAGACGTTTATGGCGC
Tb10.6k15.0140	PFC19	ggggacaactttgtacaaaaaagtggcatgGTTTCAGGGCCACTTGT	ggggacaactttgtacaagaaagtgggTACGAAGCTTCACTGAGG
Tb927.2.2160	PFC11	ggggacaactttgtacaaaaaagtggcatgATGACATCGGAGGAAGC	ggggacaactttgtacaagaaagtgggTAAATGTATTGCTCCAAATGCTCA
Tb927.8.3790	PFC2	ggggacaactttgtacaaaaaagtggcatgATGGATGCAGAAGAGAATTTCCT	ggggacaactttgtacaagaaagtgggTACGCACTGTTCGCAG
Tb11.01.3000	PFC17	ggggacaactttgtacaaaaaagtggcatgAAGAACAACCCAGTAGTTATGC	ggggacaactttgtacaagaaagtgggCTTCTGTTTCTGCGCTCT
Tb11.01.4623	Calmodulin	ggggacaactttgtacaaaaaagtggcatgATGGCCGATCAACTCTC	ggggacaactttgtacaagaaagtgggTATTGCTCATCATTTTTTCACGA
Tb927.2.3020		ggggacaactttgtacaaaaaagtggcatgATGTCGTGATGCGTCCG	ggggacaactttgtacaagaaagtgggCACTTACCCTCCTCTCC
Tb927.7.1920	PFC5	ggggacaactttgtacaaaaaagtggcatgATGTTACAAGCGGCTGA	ggggacaactttgtacaagaaagtgggCGTCGATTAACCTTTTTCAGGTTT
Tb10.389.0100	PFC20	ggggacaactttgtacaaaaaagtggcATGGATCAATCATGCAACGAA	ggggacaactttgtacaagaaagtgggCGTGGGATCCGATACG
Tb10.6k15.1510	PFC18	ggggacaactttgtacaaaaaagtggcatgGACTGTGTGATAACTCAGTGG	ggggacaactttgtacaagaaagtgggTACTGGCGAGCTAATGC
Tb11.01.3000	PFC17	ggggacaactttgtacaaaaaagtggcatgAAGAACAACCCAGTAGTTATGC	ggggacaactttgtacaagaaagtgggCTTCTGTTTCTGCGCTCT
Tb11.02.2350	PFC12	ggggacaactttgtacaaaaaagtggcATGGCGAAAAAGGGTAAAAAGA	ggggacaactttgtacaagaaagtgggCAGTGTGAAGGCAACATATC
Tb927.6.4140	domain of PFC4	ggggacaactttgtacaaaaaagtggcTATTTGTGTCACAGCAAAACC	ggggacaactttgtacaagaaagtgggTGGTCACAAAGTGTGACTTAATTC
Tb10.26.0680	domain of PFC16	ggggacaactttgtacaaaaaagtggcAAGTTTGAGTACCTTCCCAAC	ggggacaactttgtacaagaaagtgggTGGTCGCAAACTGTCTGTGTC
Tb09.211.4513	KMP-11	ggggacaactttgtacaaaaaagtggcATGCCACCACATACG	ggggacaactttgtacaagaaagtgggGGGAACGGGCATTTCT

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