

TPR

1qqe_5a	ESNFLKSLIDAVNEGDSEQLSEHCKEFDNFMRLD
1kt0_3a	EKGLYRRGEAQLLMNEFESAKGDFEKLVEVNAAR
1e96b_3b	CEVLYNIAFMYAKKEEWKKAEEQLALATSMKSEP
1fch_1a	HPQPFEEGLRRLQEGDLNPAVLLFEAAVQQDPKH
1hxi_1a	EEAWRSLGLTQAENEKDGLAIIALNHARLDPKDX
1elr_3a	AKAYARIGNSYFKEEKYKDAIHFNKSLAEH RTP
1a17_3	IKGYRRAASNMA LGKFRAALRDYETVVKVKPHD
1elw_3a	GKGYSRKA AALEFLNRFEEAKRTYEEGLKHEANN
1iyg_1a	RDYVFYLA VGNYRLKEYEKALKYVRGLLQTEPQN
1fch_2a	MEAWQYLGTTQAENEQELLAISALRRCLELKP DN
1fch_5a	YLLWNKLGATLANGNQSEEAVAAYRRALELQPGY
1fch_4a	PDVQCGLGVLFNLSGEYDKAVDCFTAALSVRPND
1kt1_3a	EKGLYRRGEAQLLMNEFESAKGDFEKLVEVNPQN
1ihg_2a	LSCVLNIGACKLKMSDWQGAVDSCLEALEIDPSN
1kt0_2a	LAAFLNLAMCYLKLREYTKAVECCDKALGLDSAN
1kt1_2a	LAAFLNLAMCYLKLREYTKAVECCDKALGLDSAN
1a17_1	AEELKTQANDYFKAKDYENAIKFYSQAIELNPSN
1elr_1a	ALKEKELGNDAYKKKDFDTALKHYDKAKELDPTN
1a17_2	AIYYGNRSLAYLRTECYGYALGDATRAIELDKKY
1e96b_1b	SRICFNIGCMYTILKNMTEAEKAFTRSINRDKHL
1elw_2a	HVLYSNRSAAYAKKGDYQKAYEDGCKTVDLKPDW
1elw_1a	VNELKEKGNKALSVGNIDDALQCYSEAIKLDPHN
1ihg_3a	TKALYRRAQGWQGLKEYDQALADLKKAQEIAPED
1kt0_1a	AAIVKEKGT VYFKGGKYMQAVIQYGKIVSWLEME
1kt1_1a	AAIVKEKGT VYFKGGKYVQAVIQYGKIVSWLEME
1qqe_4a	NKCFIKCADLKALDGQYIEASDIYSKLIKSSMGN
1fch_6a	IRSRYNLGISCINLGAHREAVEHFLEALNMQRKS
1ihg_1a	SEDLKNIGNTFFKSQNWEMA IKKYTKVLRVVEGS
1qqe_2a	GNTYVEAYKCFKSGGNSVNAVDSLENAIQIFTHR
1elr_2a	MTYITNQAAVYFEKGDYNKCRELCEKAIEVGREN
1qqe_1a	ADLCVQAAT IYRLRKELNLAGDSFLKAADYQKKA
1e96b_2b	AVAYFQRGMLYYQTEKYDLAIKDLKEALIQLRGN
1fch_3a	QTALMALAVSFTNESLQRQACEILRDWLRYTPAY

SEL1-like repeats

1klx_1a	NGCRFLGDFYENGKYVKKDLRKAAQYYSKACGLNDQ
1klx_2a	DGCLILGYKQYAGKG VVKNEKQAVKTFEKACRLGSE
hcpc_2a	SGCFNLGVLYYQGQGV EKNLKKAA SFYAKACDLNYS
hcpc_4a	DGCTILGSLYDAGR GTPKDLKKALASYDKACDLKDS
hcpc_3a	EGCASLGGIYHDGKV VTRDFKKAVEYFTKACDLNDG
hcpc_5a	PGCFNAGNMYHHGEGATKNFKEALARYSKACELENG
hcpc_6a	GGCFNLGAMQYNGEGVTRNEKQAIENFKKGCKLGAK

PPR

pfam PPR alignment was used.