

RNAcompete Hyb_ID	HTR-SELEX Expt_ID	Gene_Name	Systematic_ID	RBP_Type	RNAcompete Consensus Kmer	RNAcompete Logo	HTR-SELEX Logo	HTR-SELEX Consensus Kmer	Plasmid_ID	AA_Seq	AA_Length	Construct Type	Domains	RBD
1988	IFIT2_AA40NCCAGT	IFIT2	ENSG00000119922	ucRBP	RRAUUAUA			CCUUUAUA	pTH11287	MSENKKN	472	Full-length	TPR x 9	Unknown RBD
1982	-	LSM6	ENSG00000164167	ucRBP	DWUUUUA		-	-	pTH11669	MSLRKQTP	81	Full-length	Sm-Like	Unknown RBD
1940	-	NUDT16L1	ENSG00000168101	ucRBP	GGRAYCC		-	-	pTH13040	MAGARRLI	195	Full-length	Nudix Hydrolase-like	Unknown RBD
2005	-	NUPL2	ENSG00000136243	No (cRBP)	GAAAGAG		-	-	pTH11768	MAICQFLI	424	Full-length	zf-CCCH x 1	Unknown RBD
1010	-	PEG10	ENSG00000242265	ucRBP	UYUUGKG		-	-	pTH10760	MTERRRDE	325	Full-length	zf-CCHC x 1	Unknown RBD
2000	-	PRPF31	ENSG00000105618	ucRBP	CGCRGC		-	-	pTH11088	MSLADELL	499	Full-length	Nop, Prp C-term	Unknown RBD
1939	-	PURA	ENSG00000185129	ucRBP	GAAGGAA		-	-	pTH11867	MADRDSG	322	Full-length	pur-alpha DNA-binding	Unknown RBD
1902	-	RPL11	ENSG00000142676	ucRBP	RAGAUUR		-	-	pTH11596	MAQDQGE	179	Full-length	L5	Unknown RBD
1903	RPL22_AA40NCCAG	RPL22	ENSG00000116251	ucRBP	UCCSMGB			??	pTH11667	MAPVKKLV	129	Full-length	L22e	Unknown RBD
1883	RPL30_AC40NGATG	RPL30	ENSG00000156482	ucRBP	WAAARGG			AGUUACA	pTH11634	MVAARKT	116	Full-length	L30e	Unknown RBD
1905	-	RPP25	ENSG00000178718	ucRBP	AWAUGGG		-	-	pTH11809	MENFRKVF	200	Full-length	Alba-like DNA/RNA binding	Unknown RBD
1969	RPS12_TG40NCATA	RPS12	ENSG00000112306	ucRBP	CGAUGAH			GAUGA	pTH11673	MAEEGIAA	133	Full-length	L30e	Unknown RBD
2017	-	RPS15A	ENSG00000134419	ucRBP	SGUAYCB		-	-	pTH11611	MVMNVVL	131	Full-length	S8	Unknown RBD
1889	-	SERP1	ENSG00000142864	No (Control)	DCGCGCG		-	-	pTH11303	MPGHLQEI	408	Full-length	Hyaluronan/mRNA binding	Unknown RBD
2009	-	SSBP1	ENSG00000106028	ucRBP	RAUGUWA		-	-	pTH10483	MFRRPLVC	148	Full-length	ssDNA binding	Unknown RBD
2011	-	PRR3	ENSG00000177707	No (cRBP)	WAAUARC		-	-	pTH11093	MPKRKQI	188	Full-length	zf-CCCH x 1	Unknown RBD
1892	-	ETF1	ENSG00000120705	ucRBP	SCCSGGS		-	-	pTH11824	MADDPSAV	438	Full-length	eRF1 x1, L30e-like	Unknown RBD
1873	-	GAR1	ENSG00000109534	ucRBP	GCGCGB		-	-	pTH11860	MSFRGGGI	217	Full-length	Cbf5 binding domain	Unknown RBD
1506	-	SF3A3	ENSG00000183431	ucRBP	GUWUACH		-	-	pTH11092	METILEQQ	501	Full-length	SF3A3, DUF3449, Telomere_Sde2_2	Unknown RBD
2459	-	HARS2	ENSG00000112855	ucRBP	WGUGDDR		-	-	pTH12151	MPLLGLPI	506	Full-length	tRNA_synth_HIS, HGTP_anticodon	Unknown RBD
2345	-	RPS2	ENSG00000140988	ucRBP	YGCGB		-	-	pTH12237	MADDAGA	293	Full-length	Ribosomal_S5_C	Unknown RBD
2456	-	GTPBP10	ENSG00000105793	ucRBP	SCCUGSB		-	-	pTH12148	MVHCSCVI	387	Full-length	GTP1_OBG, MMR_HSR1	Unknown RBD
1933	-	CNBP	ENSG00000169714	No (Control)	GGAGGDR		-	-	pTH11285	MSSNECFK	179	Full-length	zf-CCHC x 7	Unknown RBD
1948	MBNL2_TT40NTAGC	MBNL2	ENSG00000152601	No (Control)	AWGCUUD			GCUUGC	pTH10757	MALNVAP	373	Full-length	zf-CCCH, zf-CCCH, zf-CCCH, zf-CCCH	Unknown RBD
1907	NHP2L1_AC40NGT	NHP2L1	ENSG00000100138	ucRBP	CRAUGAH			CNAUAGC	pTH11577	MLLVQTEA	133	Full-length	L7e	Unknown RBD
1928	NUDT21_AC40NGT	NUDT21	ENSG00000167005	No (Control)	DWUGUAA			UGUAA	pTH10485	MSVVPNPF	227	Full-length	Nudix Hydrolase	Unknown RBD
1963	ZRANB2_TG40NCC	ZRANB2	ENSG00000132485	No (Control)	AGGUUAD			AGGUAA	pTH10754	MSTKNFRV	330	Full-length	zf-RANBP2 x 2	zf-RANBP2, zf-RANBP2
1825	-	RPL5	ENSG00000122406	ucRBP	GUWCGU		-	-	pTH11782	MGFVKVVI	298	Full-length	Ribosomal_L5e, Ribosomal_L18_C	Unknown RBD
1709	-	RPP25L	ENSG00000164967	ucRBP	AURGRVM		-	-	pTH11620	MEHYRKAC	164	Full-length	Alba	Unknown RBD
-	RPL12_TT40NTAGC	RPL12	ENSG00000197958	ucRBP	-			CGAAG	pTH11680	MPPKFDPA	166	Full-length	Ribosomal_L11_N, Ribosomal_L11	Unknown RBD
-	SUB1_CT40NGCGCT	SUB1	ENSG00000113387	ucRBP	-			GUCUUCU	pTH11732	MPKSKELV	127	Full-length	PC4	Unknown RBD
-	TRIT1_FL_TG40NGA	TRIT1	ENSG00000043514	ucRBP	-			CUNNAAA	pTH13139	MASVAAA	467	Full-length	IPPT, zf-C2H2_jaz	Unknown RBD
-	MRT04_CG40NTCT	MRT04	ENSG00000053372	ucRBP	-			GACCAAG	pTH11728	MPKSKRDK	240	Full-length	Ribosomal_L10, RL10P_insert	Unknown RBD
-	MAPRE1_GG40NTA	MAPRE1	ENSG00000101367	ucRBP	-			ACGUACA	pTH11644	MAVNVYSI	269	Full-length	CH, E81	Unknown RBD
-	RPS27_TG40NCCAT	RPS27	ENSG00000179554	ucRBP	-			ACACUC	pTH11678	MPLAKDLL	85	Full-length	Ribosomal_S27e	Unknown RBD
-	SRCAP_CT40NTCTA	SRCAP	ENSG00000080603	AT-Hook	-			CACCGG	pTH12029	GPCEAAPS	258	RBD	AT-Hook x 3	AT-Hook, AT-Hook, AT-Hook
-	ZNF326_CT40NCCG	ZNF326	ENSG00000162664	zf-C2H2	-			ACGGAC	pTH14046	MEKISLSKS	265	RBD	zf-C2H2_AKAP95 x 2	zf-C2H2_AKAP95, zf-C2H2_AKAP95