

Retro-transposon	Accession numbers	RT-domain sequences	Key reference
<i>Pao</i>	S33901 AB042118 AB042119	LVAKGYAEPAPKTKTENRTWYLPHFVAVNPPKPEKLRVVHDAAARTRG VALNDMLLKGPNNLLQSLPGVIMRFRQHNITATADIKEMFIQVKLRPEDKD ALRYLWRKDQRDNKPPEENRMTSLIFGASSSPSTAIYVKNLNAQKHEAT HPEAAATIQNRHYVDDYLDSEFKTLKDAVRITTDVRRRIHEKAHFEKQWK SNSLSLLETLAGENENGNGVELYKPEEKTERVLGLIW	27
<i>Boudicca</i>	AY308018 AY308019 AY308021 AY308022	MDLGIIRPSNSPWASPLHMVPPKDSNDWRPTGDYRRLNAKTIPDRYPL PHIHDLTATLKGTTVFSKIDLVKAYNQIPMAPDDIPKTAITPFGLYEFLRM PFGLRNAAQTFQRFIDDVFRGLNFVHAYVDDCLIASSDRESHLKHLDIVF DRLQRHGITVNIQKCCQIGTNSLDFLGHTI	20, 22
<i>Sinbad</i>	AY506538	HLSKGYIIEASKEGFDRDAVCWYIPHPVINPXKAWKTQNCFFDCAAVY QGFSLNQFLRGPNTVNSLFGVLLRFRGLNIALAADIEEMFLQVRIPRQ DRGAFRLWWEDGDMKRTAKEYCLTVHPFGAVSSPFCANFALKKTVDI FGKEFNRIQEVVDNSFYVDDYLASIDNVQDAIELAKTLGLLLKGGFRL TKWISSCLQVLESIHPEERAEAVGEIDFERLPTERTLGLFWN	This study
<i>D. Rerio</i> <i>Sinbad</i> -like element	BX537152	IIEKGYAVRVPTEHLNRNDNKVWYIPHHGVYHPKKNKIRVVFDCASFO GMSLNSQLLQGPNTLNTLIGVLTRFREEPIAVMADVESMFYQVKVPEED TDLLRFLWWPDGNLNAPIEEYRMTVHLFGATSSPSCASYALRKTAEDR KHVASQKAVDTVLNNFYVDDCLKSVSNEQEAIVLVEIQDLCLEGGFRL TKWVSNNRKVLLSIPEDQRASGVKDLDLDQDSLPIERALGMQWC	This study
<i>A. gambiae</i> <i>BEL</i> -like element	XM_308737	YITLGHMTLVPREREDCSGAFYLPHPVLKDSSTTKVRVVFDDGSAKTS TGKSLNDALLVGPVVQEELLTLIRFRKYEIALIADIEKMYRQVTMNPTDR HLQRILWRFDNSQPIRTYELGTVTYGLAPSAFLATRTLILQLANDEGDKYL KARSVIKENVYVDDLLAGANNIPETIELRNQLNALLQKGGFLLRKWCSN APAVLADLPPELVATHSSVNFDPDESIKTLGICW	This study
<i>C. briggsae</i> <i>Tas</i> -like element	AC084491	SLEFIEEVPDESISDGPVVSYPHPVIKETSSTKVRIVFDGSAKTNKVS HSLNDHLHTGERLLPDIAAIMLRIRQHNILISGDIEKAFLQLVLHQTDRDA TRFLWKDPTDGHLCYRYRRVPFGLKPSPYLLNKTVRTHLESYDHPWA QAMINSFYVDNVFMGLDTAEEMDFYVFAKKVFAEAQMNLCQFCSNSS TVNTFFVKHEKDTTIDRNQKILGISW	This study
<i>D. rerio</i> <i>Suzu</i> -like element	BX005079	MLDRGAAVKLPESSIANWAGPVWYVSHLIAPNPHSVTTTPVRLVWNSSQ RFRGVSMNDLLIKGPDVLNQIRAVLLRFRSGVYAALGDIKKMYNSVWLE DQEVHLHRFLWRDTENEELGEYAITRVNIGDKPAGCIAQLAMRETANLP SFAHLKEERRVIQHDSYVDDILTSHNDLDQLQSIVANTELILKAGGFHLK PWVFGQSGREKSDDKSCIKIKEKVMVLPNQMHDDDNKALGLGY	This study