

transposon	representative targets	sequence (+1 ~ +30)	position	sequence identity			type of repeat	times targeted
				100%	99.9%-97.0%	total		
<i>piggyBac</i>	B89-4	TTAAGACCCGTCTCTTAAAAAAAAAAAA	chr16(30140496)	4	0	4	NF	1
	B87-4	TTAAGAATGTTGAATATTGGCCCCACTCT	chr3(55059477)	1	510	511	NF	1
	B75-4	TTAAGAAACAAGTTAACACCGAAGCCAGA	chr17(60391785)	1	1	2	NF	2
	B85-4	TTAAAAAGGCATTATTTTCGAGCTATCT	chr6(125167100)	0	2	2	NF	1
	B42-3	TTAATTTACTTAAGATAATGGCCTCCACAC	chr22(15414966)	5	3	8	NF	1
	B100-1	TTAAGAAAGGAGTTGAATTAAGCTCAGGTT	chr1(120900969)	0	2	2	NF	1
	B90-1	TTAATATCCACCTTTGCACAGTAGACAAT	chr3(137392502)	2	0	2	NF	1
	B92-2	TTAAACACACACTTAGAGGGAAATAATTCA	chr18(11877795)	0	2	2	NF	1
	B82-3	TTAAGAATATAAGGCCAAGCACAGTGGCT	chr11(27402368)	1	1	2	NF	1
	B89-3	TTAAATAAGAGATAATAACTAACCATGGC	chr3(57972969)	1	1	2	NF	1
	B92-1	TTAAGAAACAAGTTAACACCGAAGCCAGA	chr17(60391785)	2	2	4	NF	1
	B79-1	TTAATGAGCAGTAGCATTTTGACGTGATGA	chr1(231429889)	0	2	2	NF	1
	B77-4	TTAAGACCCGTCTCTTAAAAAAAAAAAA	chr16(29401156)	0	4	4	NF	1
	B78-1	TTAATGGAACTTAAATCTAGTTGATTAAA	chr22(19948556)	1	6	7	NF	1
	B79-2	TTAAGGGGGGAAACAGTTCAGGGCCAACA	chr14(55711916)	1	1	2	NF	1
	B84-1	TTAATGTAAATTACAAACACTGTTTTATC	chr18(29434150)	0	2	2	NF	1
	B85-1	TTAAGCACAGTATCAGTGATAAAAAATAGCT	chr1(201276407)	1	1	2	NF	1
	B82-1	TTAAGCTGAATCTGTTTTTCCAGTGCCCC	chr2(231461932)	0	2	2	NF	1
<i>To12</i>	T111-3	TTTAAGAATGTTGAGTATTGGCCCCCACTC	chr8(9712455)	1	8	9	LINE	1
	T147-1	ATCCTGAGCAGCCGAATCTGCAATCATCTT	chr7(154665283)	1	1	2	LTR	1
	T3-2	TCCACCACAGCATGAGTTAAACCAAAGTCT	chr7(76097237)	2	0	2	LINE	1
	T88-3	GTCTGTACTGCTGCAAAGCTTCACAGACAG	chr10(98485313)	1	2	3	NF	1
	T115-4	AATTTATCCATTTCTTCTAGATTTTCTAGT	chr20(1472025)	0	3	3	LINE	5
	T103-2	GGCGCCCGCCACTACGCCTGGCTAATTTTT	chr21(13847935)	1	3	4	SINE	1
	T113-2	AATTTATCCATTTCTTCTAGATTTTCTAGT	chr20(1472025)	0	3	3	LINE	1
	T111-3	TTTAAGAATGTTGAGTATTGGCCCCCACTC	chr8(9712455)	1	8	9	LINE	1
	T162-3	CCAGAGACCTTTGTTCACTTGTTTATCTGC	chr20(33206406)	202	ND*	202	> Low_complexity	1
	T157-1	CTCGTACGTAAGTTTTAGTGTGAACATATA	chr4(48956277)	4	3	7	LINE	1
	T157-2	GTTAACAGTGACCTATTTGGGAGAAGGGGA	chr7(66290199)	1	1	2	LINE	1
	T107-2	AATATATGAGTAGCTAAACAACCTCTATAAG	chr2(97613844)	1	1	2	LINE	1
	T104-1	GAACACATGGACACAGGAAGGGGAACATCA	chr3(134171282)	1	52	53	LINE	1
	T25-3	ACCCCATCTCTACTAAAAATACAAAAAATT	chr6(166733349)	1	1	2	SINE	1

NF designated as "not found"