

Table 7. Complete and nearly complete 1360 transposable elements found in different *Drosophila* species

Species	Contig	Coordinates	Orientation	Transposase	TIRa/ TIRb	TSD	Total length	Insertions
<i>D. melanogaster</i>	Chr. 4	809215-812470	Inverted	812470-809907	-/31	GCCATAC/-	3615	—
<i>D. simulans</i>	Chr. U	8185141-8188796	Inverted	8188307-8185783	-/31	CCCGAAC/-	3656	—
<i>D. sechellia</i>	5259	961-5382	Direct	2194-4726	31/31	GTTGAC/ ATTCCAC	4422	—
<i>D. erecta</i>	7407	137172-141324	Inverted	140279-137852	31/31	GTTTGAC	4153	—
<i>D. yakuba</i>	0.40	344624-349004	Direct	345812-348373	31/31	GTCAAAG	4381	—
<i>D. pseudoobscura</i>	784	22941-26100	Direct	23339-25777	31/32	ATCATAT	3160	—
<i>D. persimilis</i>	9857	64151-67135	Inverted	66929-64479	32/31	ATATGAT	2985	—
<i>D. virilis</i>	17537	32378-37583	Inverted	33160-36198	32/31	GTTTGAC	4702	LINE (33361-33864)

TIRa is the TIR at the 5' side of the transposase while TIRb designates the TIR at its 3' end. TSDs are given in the same orientation they have in the contig.