

Table S5. Primers used

Primer	Sequence	Location
Plac1 ^a	CACCCAAGGCTCTGCTCCCACAAT	<i>P</i> element 5' end, outwards orientation
Pry4 ^a	CAATCATATCGCTGTCTCACTCA	<i>P</i> element 3' end, outwards orientation
w11678U ^b	TCATCGCAGATCAGAAGCGG	3' exon of the reporter gene <i>mini-white</i>
CB-0236-3up	ACACGCCGCAAGATGAATAC	2R:11260327..11260346 (+)
CB-0236-3down	GATGACCAATTTCCGTTTGG	2R:11261115..11261096 (-)
5-HA-1995up	AAATTTTTCTGCCTCGCAAA	2R:15613845..15613864 (+)
5-HA-1995down	TCGAGTGGAATGAGTTGTGG	2R:15614470..15614451 (-)

^a <http://www.fruitfly.org/about/methods/inverse.pcr.html>. ^b [1].

Supporting References

1. Golic KG, Golic MM (1996) Engineering the *Drosophila* genome: chromosome rearrangements by design. *Genetics* 144: 1693-1711.