

Lines with CG15630 affected

Breakpoints coordinates are indicated according to FlyBase released September 3rd, 2015 (FB2015_04)

| Description | Genotype | genome region | ID | Source |
|--|---|-----------------------|---------|----------|
| Perfect <i>PdL</i> excision in mutant 3404a | w[-] | - | Se361 | internal |
| Perfect <i>PdL</i> excision in mutant 3404a | w[+] | - | Se361R | internal |
| Perfect <i>PdL</i> excision in mutant 3404a | w[-] | - | Se363 | internal |
| Perfect <i>PdL</i> excision in mutant 3404a | w[+] | - | Se363R | internal |
| Perfect <i>PdL</i> excision in mutant 3404a | w[-] | - | Se373 | internal |
| Imperfect <i>PdL</i> excision in mutant 3404a | w[-] | 2L: 4793237.. 4793955 | Se362 | internal |
| Imperfect <i>PdL</i> excision in mutant 3404a | w[+] | 2L: 4793237.. 4793955 | Se362R | internal |
| Imperfect <i>PdL</i> excision in mutant 3404a | w[-] | 2L: 4792816.. 4793955 | Se624 | internal |
| Insertion in CG15630 | y1 w*; Mi{MIC}CG15630MI11721 | 2L | 56367 | BL |
| Insertion in CG15630 | w1118; Mi{ET1}CG15630MB07747 | 2L | 25568 | BL |
| Deletion in CG15630 - FBst0009702 | w1118; Df(2L)BSC225/CyO | 2L 25A3--25A7 | 9702 | BL |
| Deletion in CG15630 - FBst0008482 | Df(2L)sc19-10/In(2L)CyLtR In(2R)Cy, Cy1 amosRoi-1 cn2 sp2 | 2L 25A4-25C1 | 8482 | BL |
| Insertion in CG15630 (rev) | W[-]; P{PdL}CG15630[3404z] | 2L: 4793956-for | 3404a | internal |
| Insertion in CG15630 (rev) on background of Cantonized w[1118] | W[1118]; P{PdL}CG15630[3404z] | 2L: 4793956-for | 3404aCs | internal |

BL: Bloomington Drosophila Stock Center, Indiana University, USA