

Corrigendum

The Mapping of *Drosophila melanogaster* mutant A.4.4. microPublication Biology.
<https://doi.org/10.17912/MICROPUB.BIOLOGY.000069>

The authors of the above-noted microPublication submit the following corrections.

1. In the description of genetic mapping of mutant A.4.4 it is stated:

In summary mutant A.4.4 mapped to 2R:22,592,996..22,661,827 via deficiency mapping where there are 29 protein coding genes. After additional complementation tests ruled out eight additional genes, A.4.4 likely resides in one of the remaining 21 genes in the region that we were unable to map via complementation.

This should be corrected to read:

In summary mutant A.4.4 mapped to 2R:22,592,996..22,661,827 via deficiency mapping where there are 28 protein coding genes. After additional complementation tests ruled out seven additional genes, A.4.4 likely resides in one of the remaining 21 genes in the region that we were unable to map via complementation.

2. Table 1 lists *CG3927* as the associated allele with BDSC stock 17065, it should be *dnr1* as the affected gene. A corrected version of Table 1 is below:

Table 1: Complementation between mutant A.4.4 and individual candidate alleles

Stock number BDSC	Gene affected	Genotype	Mating with A.4.4
12060	<i>Vps20</i>	<i>P{PZ}Vps20^{G270}, l(2)rG270rG270/CyO</i>	Complement
16199	<i>CG4294</i>	<i>y¹ w¹¹¹⁸; PBac{5HPw⁺}CG4294^{B316}/CyO</i>	Complement
17065	<i>dnr1</i>	<i>w¹¹¹⁸; P{w^{+mC}=EP}EP2515/CyO</i>	Complement
17739	<i>Ugt58Fa</i>	<i>w¹¹¹⁸; PBac{PB}Ugt58Fa^{c05973}/CyO</i>	Complement
23049	<i>CG33143</i>	<i>y¹ w^{67c23}; Mi{ET1}CG33143^{MB01293}/CyO</i>	Complement
29511	<i>RpS24</i>	<i>w*; P{FRT(whs)}G13 P{lacW}RpS24^{SH2053}/CyO</i>	Complement
63874	<i>RpS16</i>	<i>w¹¹¹⁸; PBac{IT.GAL4}RpS16^{0887-G4}/CyO</i>	Complement
67706	<i>Vps20</i>	<i>w*; Vps20^{l3}/CyO</i>	Complement

These corrections do not alter the conclusions or discussion of this microPublication.

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