

## CORRECTION

# Correction: Zebrafish *dazl* regulates cystogenesis and germline stem cell specification during the primordial germ cell to germline stem cell transition

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There were errors in *Development* (2021) **148**, dev187773 (doi:10.1242/dev.187773).

The *dazl*<sup>ae34</sup> allele was incorrect in Fig. 2B.

The corrected and original panels are shown below.

**B**

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WT      GACGCCCAACACACTGTCGTCGGCGGTATTGATATGA
dazlΔ7 GACGCCCAACAC-----GTTGGCGGTATTGATATGA (-7)

WT      TGCAGCTGCTGTCTCCAGGCCTGGCCCCACCATCACCCGTATTCAGTGGAGGAA
dazlae57 TGCAGCTGCTGTCTCCA-----CCTGTCCTGCCTGGCCCCACCATCACCTGTATTTAG (-2, +9, 2s)
dazlae34 TGCAGCTGCTGTCTCCACCATCACCCGTATTC-----AGTGGAGGAA (+15, -12)
  
```

**Fig. 2B (corrected panel). *dazl* mutagenesis and fertility phenotypes.** (B) *dazl* alleles generated by ZFN (*dazl*<sup>Δ7</sup>) and CRISPR (*dazl*<sup>ae57</sup> and *dazl*<sup>ae34</sup>). Partial ZFNs and Cas9 binding sites are highlighted in red. Dashed lines represent deletions and orange denotes substitutions.

**B**

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WT      GACGCCCAACACACTGTCGTCGGCGGTATTGATATGA
dazlΔ7 GACGCCCAACAC-----GTTGGCGGTATTGATATGA (-7)

WT      TGCAGCTGCTGTCTCCAGGCCTGGCCCCACCATCACCCGTATTCAGTGGAGGAA
dazlae57 TGCAGCTGCTGTCTCCA-----CCTGTCCTGCCTGGCCCCACCATCACCTGTATTTAG (-2, +9, 2s)
dazlae34 TGCAGCTGCTGTCTCCA-----CCATCACCCGTATTCAGTGGAGGAA (-12, +15)
  
```

**Fig. 2B (original panel). *dazl* mutagenesis and fertility phenotypes.** (B) *dazl* alleles generated by ZFN (*dazl*<sup>Δ7</sup>) and CRISPR (*dazl*<sup>ae57</sup> and *dazl*<sup>ae34</sup>). Partial ZFNs and Cas9 binding sites are highlighted in red. Dashed lines represent deletions and orange denotes substitutions.

On p. 3, the length of the amino acid deletion in the *dazl*<sup>ae34</sup> was incorrect.

In Fig. S4, the *dazl*<sup>ae34</sup> allele was incorrect.

The authors apologise to readers for these errors, which do not impact the results or conclusions of this article. Both the online full-text and PDF versions of the article have been corrected.