

Specification of dopaminergic subsets involves interplay of En1 and Pitx3

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There was an error published in *Development* **140**, 3373-3384.

On p. 3374, a section of the microarray analysis text should read as follows:

Analysis was performed as described (Roepman et al., 2005). Data were analyzed using ANOVA (R version 2.2.1/MAANOVA version 0.98-7) (Wu et al., 2003). Genes with $P < 0.05$ after false discovery rate correction were considered to be significantly changed. Mouse Whole Genome Gene Expression Microarrays V1 (Agilent Technologies, Belgium) containing mouse 60-mer probes were used and data have been deposited in ArrayExpress under accession number E-TABM-1104.

The authors apologise to readers for this mistake.