

CORRECTION

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Correction to: Long read sequencing reveals novel isoforms and insights into splicing regulation during cell state changes

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Following publication of the original article [1], it was that there was an error in Fig. 4. There was an overlaid panel in the center of the image. The correct Fig. 4 is provided in this Correction and the original article [1] has been updated.

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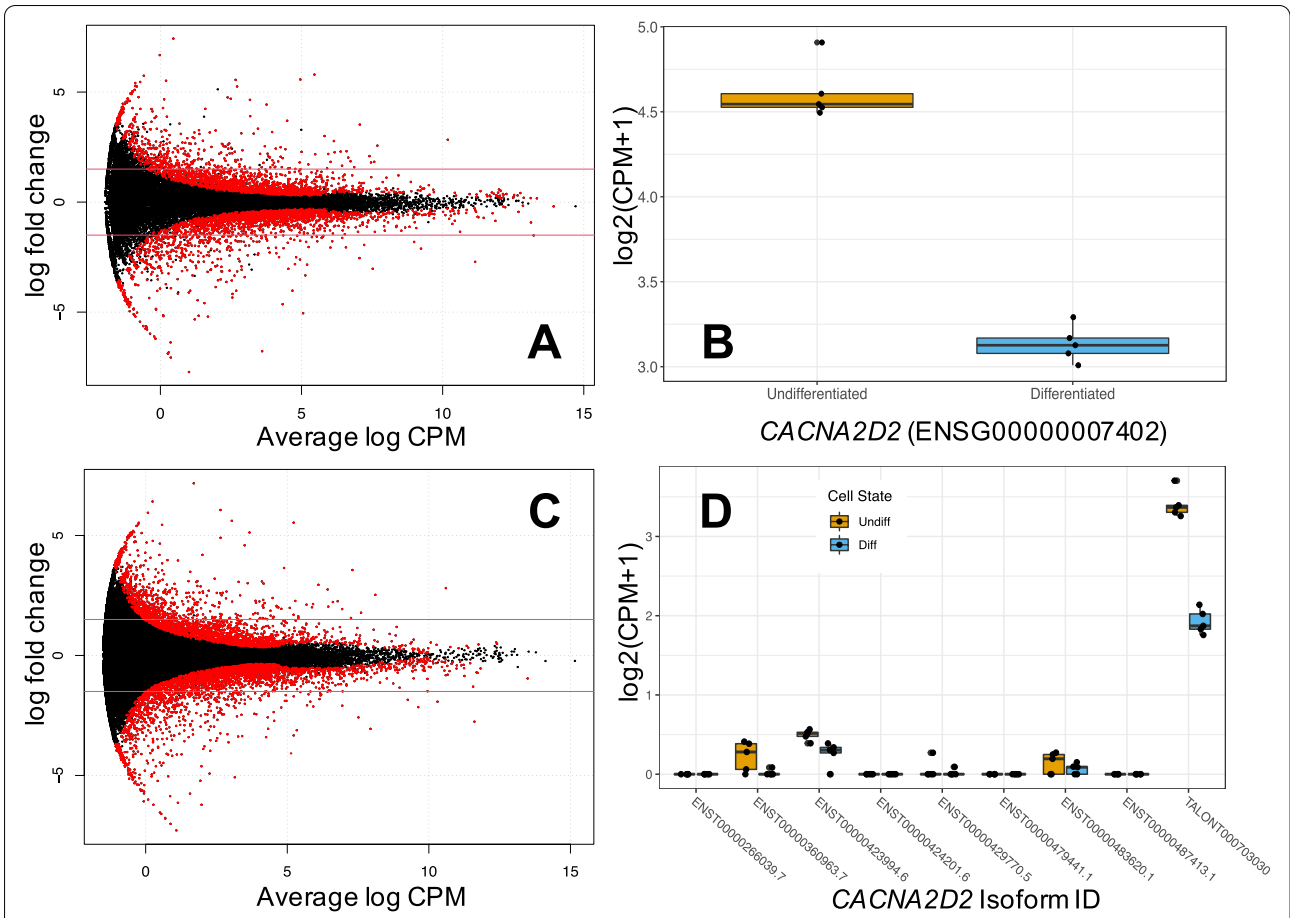


Fig. 4 Panel of **A** gene-level differential expression (DE) smear plot, solid red lines highlighting ± 1.5 $\log_2(\text{FC})$ threshold, **B** gene-level DE of *CACNA2D2* during differentiation, **C** isoform-level DE smear plot with ± 1.5 $\log_2(\text{FC})$ threshold, **D** *CACNA2D2* isoform expression, showing novel TALON isoform with highest read count. Red points on smear plots indicate significant differential expression ($\text{FDR} < 0.05$). Boxplots display median and IQR. Short and long read mapping example provided in Fig. S8