

Description of Additional Supplementary Files:

Supplementary Data 1: Results of the (a) DEXSeq intra-trajectory analysis, (b) DEXSeq inter-trajectory analysis, (c) Isosceles `isoform switching` analysis.

Supplementary Data 2: RNA-Seq samples quantified in VastDB (sources extracted from Supplemental Table 1 of Tapial et al. 2017; doi: 10.1101/gr.220962.117) that were used to create Fig. 4e and Supplementary Fig. 8