Supplemental Fig. S2. (A) *d*NET-seq/S5P profile over the U6 snRNA gene in the late dataset (replicate 1). **(B)** Frequency of peaks corresponding to splicing intermediates (green) detected by *d*NET-seq/S5Pon exons of genes expressed in late embryos. Genes were grouped into quartiles (Q) based on their *d*NET-seqread density.