Global Patterns of Tissue-Specific Alternative Polyadenylation in *Drosophila*

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http://dx.doi.org/10.1016/j.celrep.2013.03.022

(Cell Reports *1*, 277–289; February 23, 2012)
In the original version of this article, the GEO accession number given in the Experimental Procedures section was written as GSE3390, but the correct number is GSE33905. The corrected paragraph appears below:

One nanogram of the strand-specific RNA-Seq libraries were reamplified by PCR using a primer complementary to the 5′ adaptor and a second primer complementary to the 3′ adaptor with six T residues at the 3′ end. After 10 rounds of amplification, the 3′ primer with the T extension was replaced with a 3′ primer complementary to the adaptor with a 5′ extension containing a 6 nt index sequence and a sequence complementary to the flow cell primer. After an additional 15 rounds of amplication, the libraries were quantitated, 10–12 libraries were pooled together and sequenced on an Illumina HiSeq2000 using paired-end 100 bp and 6 bp index read chemistry. Reads were split into the respective samples using the index sequence and aligned as described above. **All of the raw fastq data and alignments of poly(A)-spanning reads from the poly(A)-enriched libraries were deposited at NCBI Gene Expression Omnibus under Series GSE33905.**

The authors regret this error.

Cell Reports *3*, 968–969, March 28, 2013 ©2013 The Authors **969**