Figure S4  Clusters of annotations of DE genes for overexpression of TBPH in motor neurons. The terms represented on the vertical axis co-occur in the gene sets represented on the horizontal axis, with enrichment scores reflecting the average log p-value for the group (Huang et al., 2009). Green squares indicate positive association reported between the gene and the annotation, black squares indicate no association yet reported. Red highlighted genes are orthologs of mammalian TDP targets identified by CLIP-seq (Sephton et al., 2011). Red highlighted terms were also enriched among TBPH-regulated Drosophila orthologs of vertebrate targets.