



Figure S1 Polycistronic locus with monocistronic, dicistronic and tricistronic alternative transcripts. All annotated introns are supported by cDNA and RNA-Seq junction data; downstream transcription start sites are supported by RAMPAGE TSS data. A GBrowse view showing (top to bottom) the gene extents and the gene models; unstranded RNA-Seq coverage data corresponding to a developmental series (early embryos, top, to adults, bottom). More information on data presented in GBrowse may be found at http://flybase.org/wiki/FlyBase:GBrowse_Tracks#General.