Enrichment of reactome-pathway related genes within glial protrusion-localised transcripts Voclano plot: Fisher's exact test against full fly genome L13a-mediated translational silencing of Ceruloplasmin expression Cap-dependent Translation Initiation Eukaryotic Translation Initiation 200 Formation of a pool of free 40S subunits GTP hydrolysis and joining of the 60S ribosomal subuni Nonsense Mediated Decay (NMD) independent of the Exon Junction Complex (EJC) SRP-dependent cotranslational protein targeting to membrane Nonsense Mediated Decay (NMD) enhanced by the Exon Junction Complex (EJ®)onsense—Mediated Decay (NMD) 150 Bonferroni-adjusted p-value (-log10) Localised genes 300 Ribosomal scanning and start codon recognition Activation of the mRNA upon binding of the dap-binding complex and eIFs, and subsequent binding to 435 Translation initiation complex formation 200 Formation of the ternary complex, and subsequently, the 43S complex 100 WG ligand not bound to FZ receptors Ubiquitination and degration of phosphorylated ARM of phosphorylated CI Degradation of PER 50 Insertion of tail-anchored proteins into the endoplasmic reticulum membrane Negative regulation of p = 0.01Gap junction trafficking and regulation Regulated Necrosis -2 log2FoldChange