Target of Rapamycin Complex 2 regulates cell growth via Myc in Drosophila

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Supplementary Figure S1. Confirmation of RNA-seq results by Quantitative PCR. (A-C) Quantitative PCR of selected gene expression identified from the comparison of RNA-seq results. Total cDNA and mRNA were extracted and inversed from flies with genotypes of wild-type control (*w*), *Ist8*¹ (*I*), *dm*^{P0} (*d*), *Ist8*¹ *dm*^{P0} (*D*) and *rictor*^{Δ 1} (*r*).



Supplementary Figure S2. Gene ontology classification of the 214 genes affected by the *lst8*¹ and *lst8*¹ dm^{P0} mutations but not by dm^{P0} mutations.



Supplementary Figure S3. Venn diagrams comparing of downregulated (A) and upregulated (B) genes between *Ist8*¹ versus control and *Ist8*¹ dm^{P0} versus dm^{P0}.