

FlyPrimerBank: An Online Database for *Drosophila melanogaster* Gene Expression Analysis and Knockdown Evaluation of RNAi Reagents

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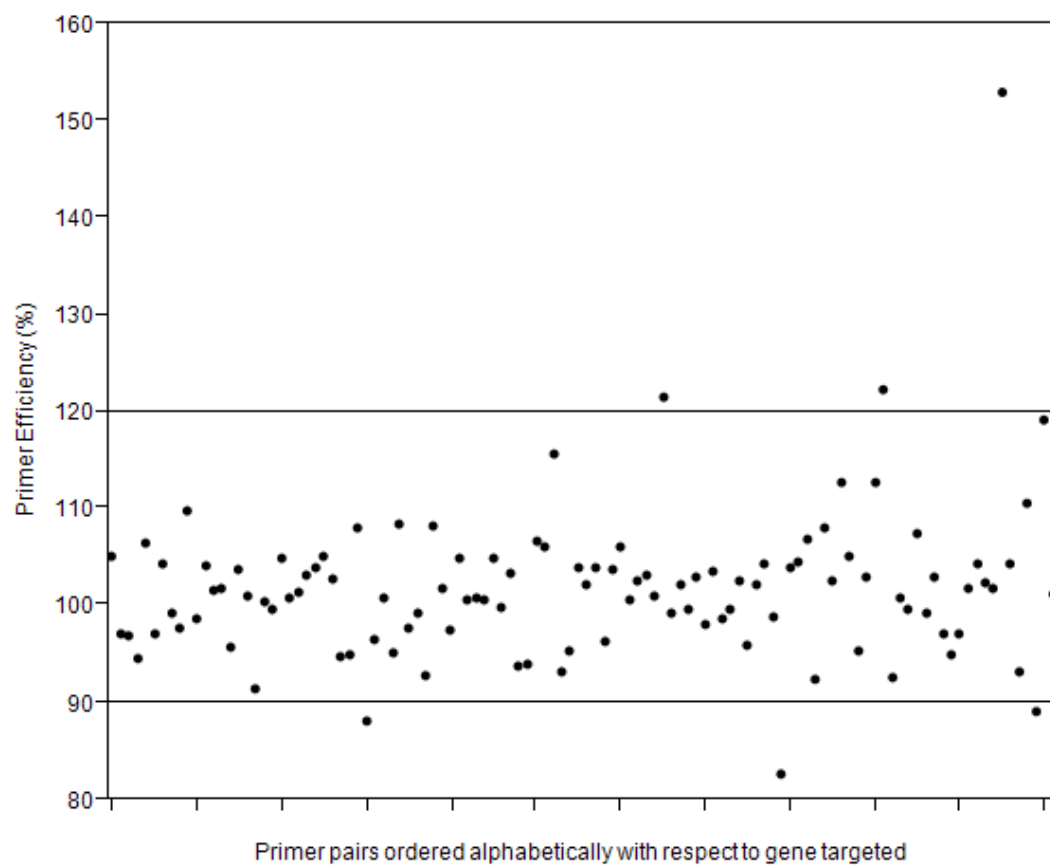
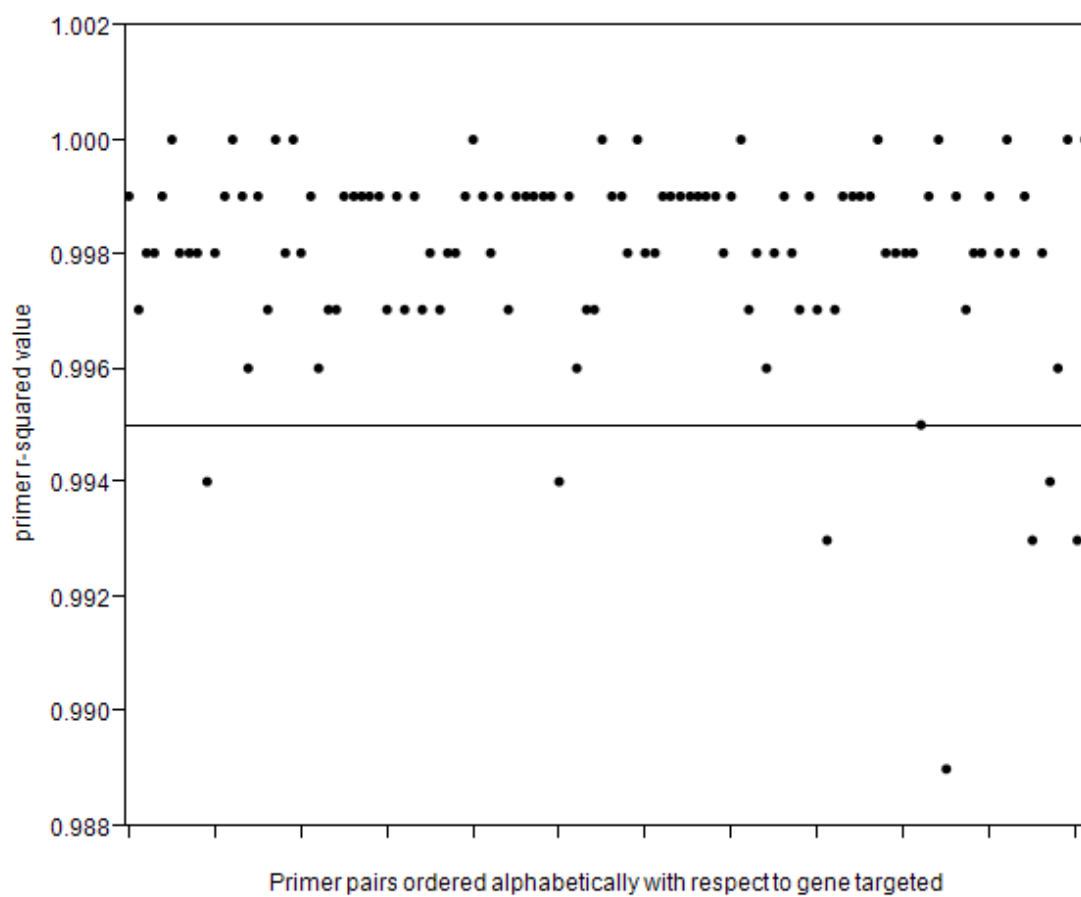
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Figure S1 Primer suitability evaluation. (a) The upper and lower bounds for acceptable primer efficiency were 120% and 90%, respectively. Higher percentages suggest non-specific products, whereas lower percentages may reflect reaction inhibition. (b) Primers with an R-squared value (a measure of reproducibility) below 0.995 were considered unacceptable.

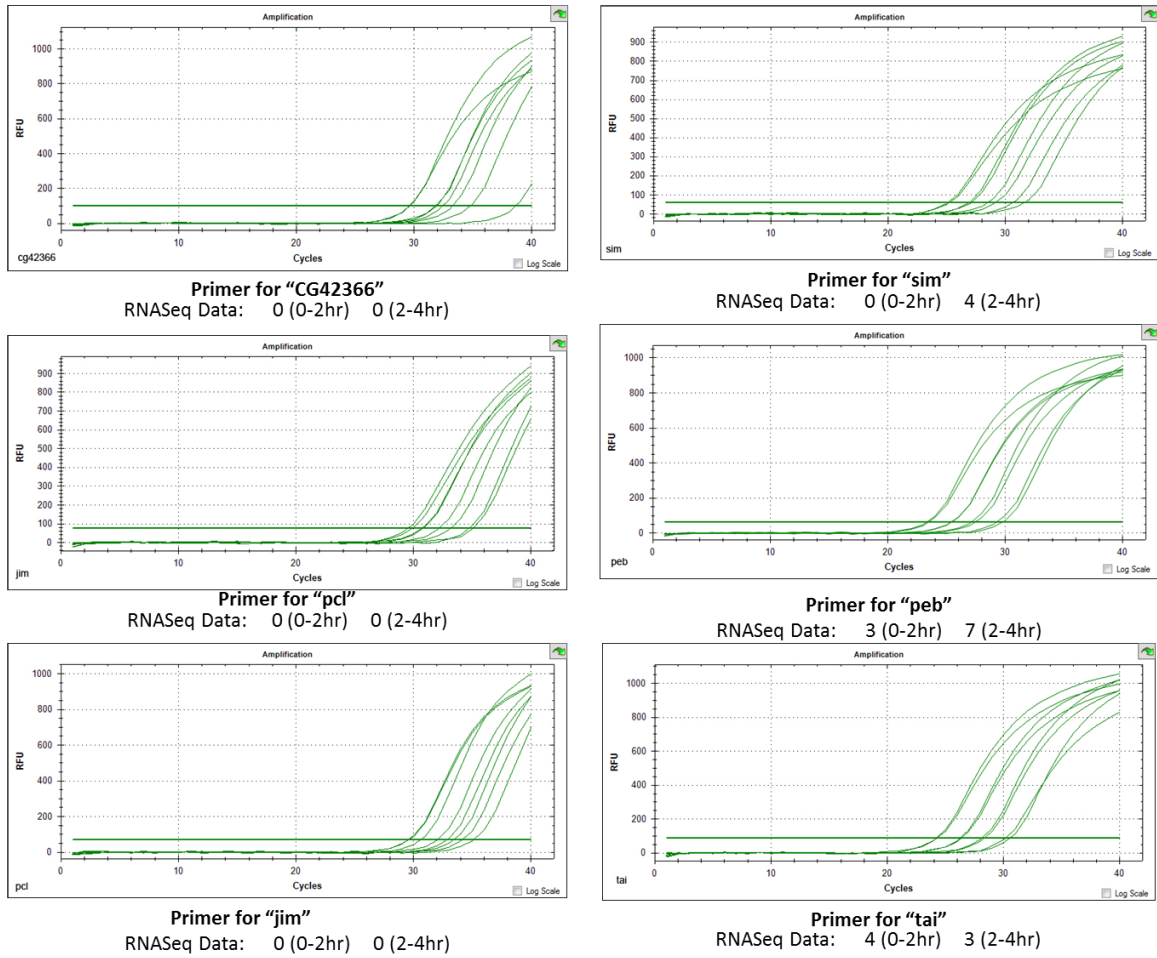


Figure S2 Determining the expression cutoff for primer evaluation. Genes represented in the left panels (*CG42866*, *pcl* and *jim*) do not express and are not suitable for primer evaluation in *Drosophila* early embryos while the genes in the right panels (*sim*, *peb* and *tai*) express and are suitable for primer evaluation.

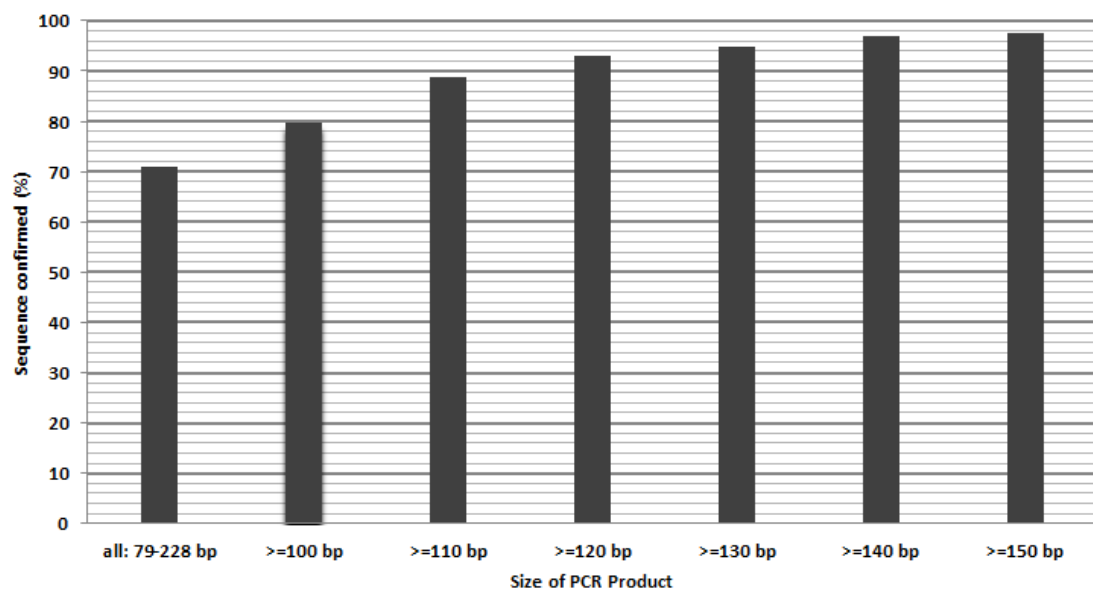


Figure S3 Sequence validation success is related to PCR product size.

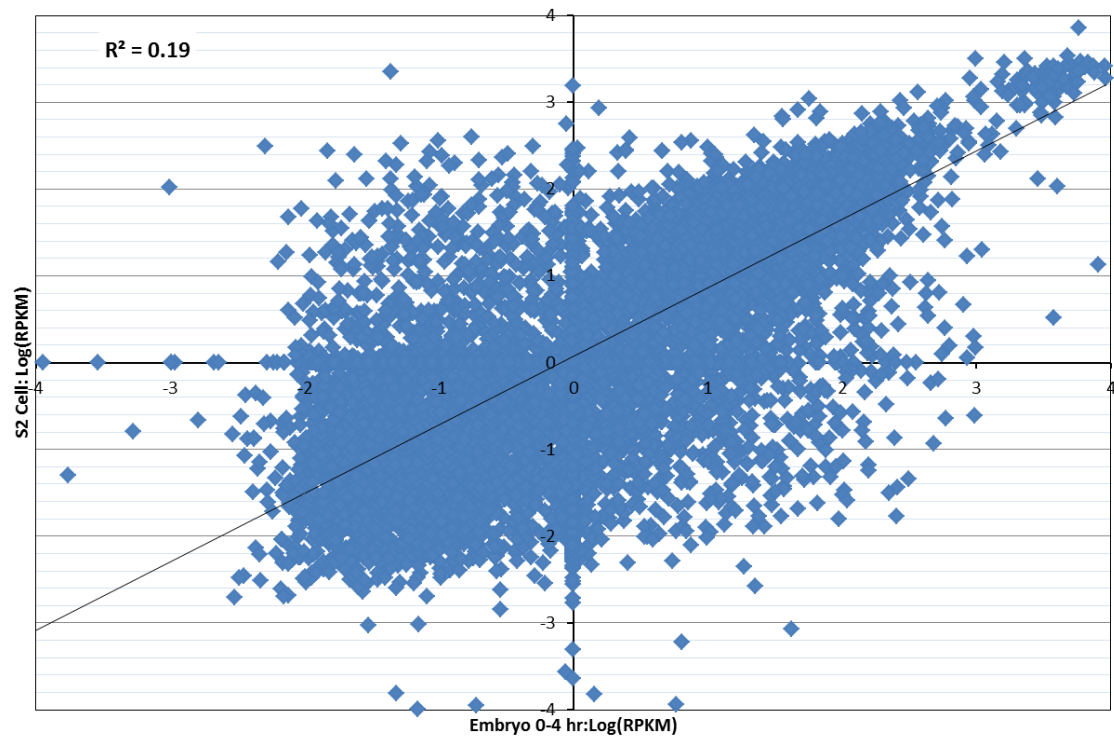


Figure S4 Early embryos and S2 cells have different transcriptomes. This graph is based on RNA-Seq data from the modEncode consortium.

Tables S1 and S2

Table S1: Results of primer pair testing.

Table S2: Protein kinase and phosphatase list

Available for download at <http://www.g3journal.org/lookup/suppl/doi:10.1534/g3.113.007021/-/DC1>.