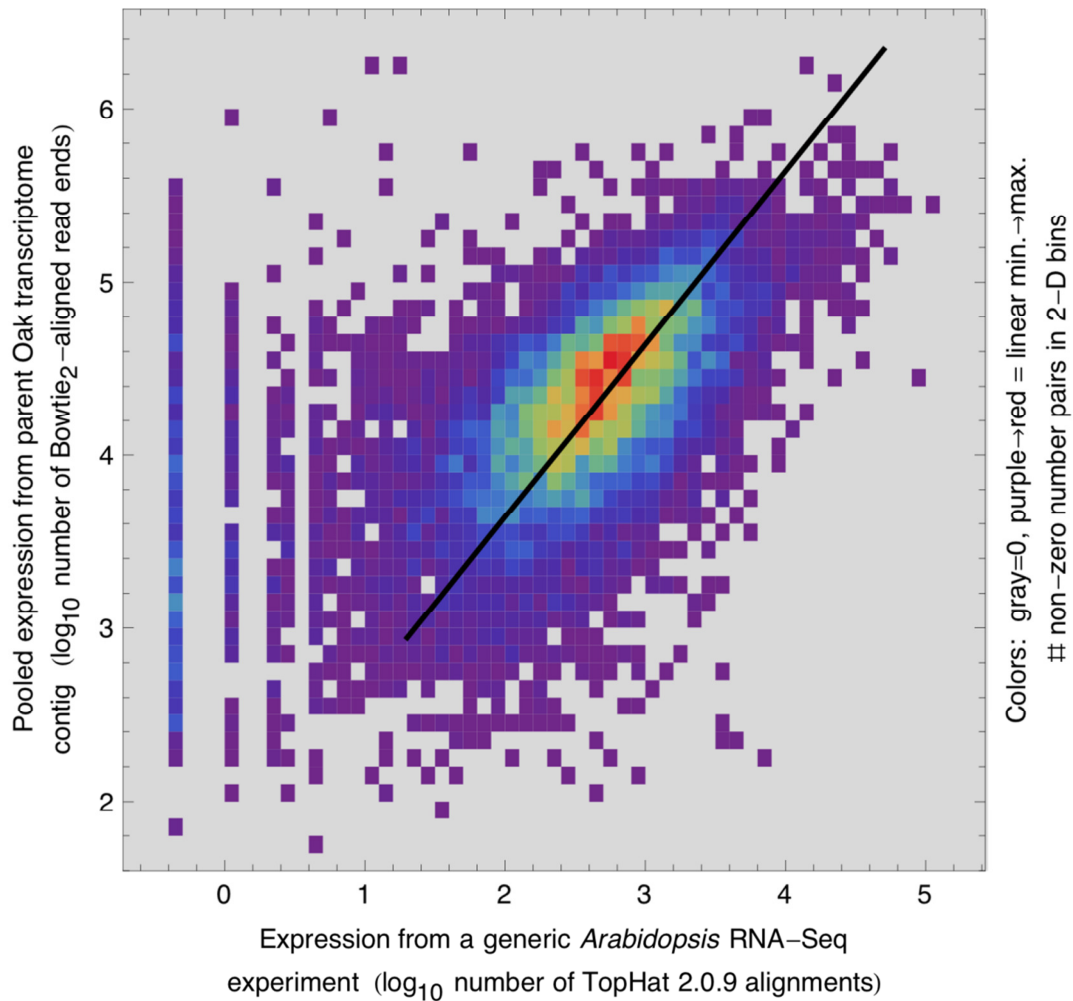


Comparison of expression between members of orthologous pairs,
line = least squares best fit slope 1 on pairs with *Arabidopsis* ≥ 20 :



Additional file 9: Expression level in oak versus *Arabidopsis* orthologs.

Two-dimensional histogram of pooled oak expression from oak transcriptome contigs versus expression of the orthologous *Arabidopsis* gene in a generic *Arabidopsis* RNA-Seq experiment (NCBI SRX145413 [43]).