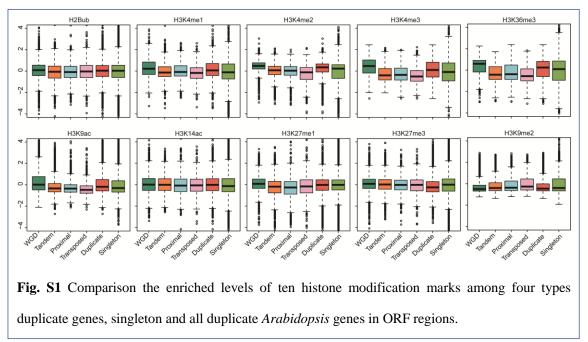
Additional file 1



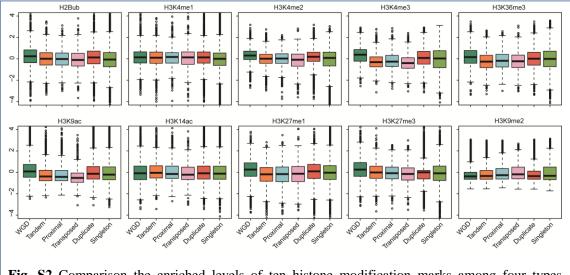


Fig. S2 Comparison the enriched levels of ten histone modification marks among four types duplicate genes, singleton and all duplicate *A. thaliana* genes in promoter regions.

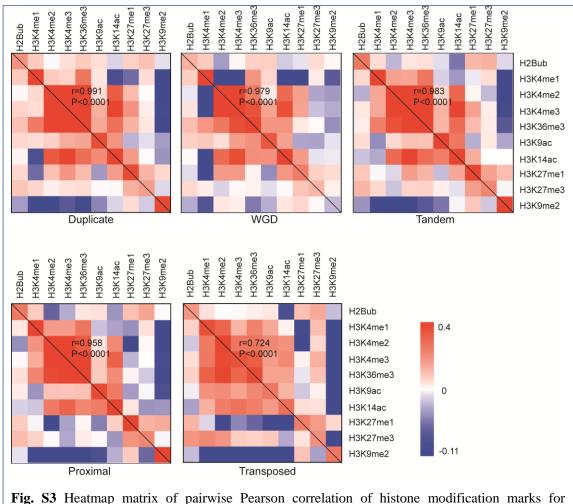


Fig. S3 Heatmap matrix of pairwise Pearson correlation of histone modification marks for different types of paralog pairs in promoter regions.

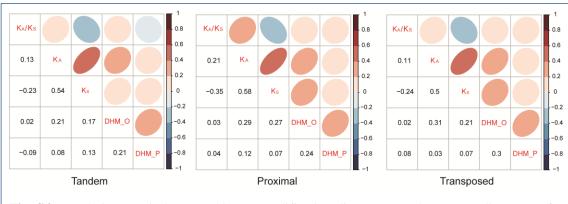


Fig. S4 Correlation matrix between histone modification divergence and sequence divergence for tandem, proximal and transposed duplicate genes.

