

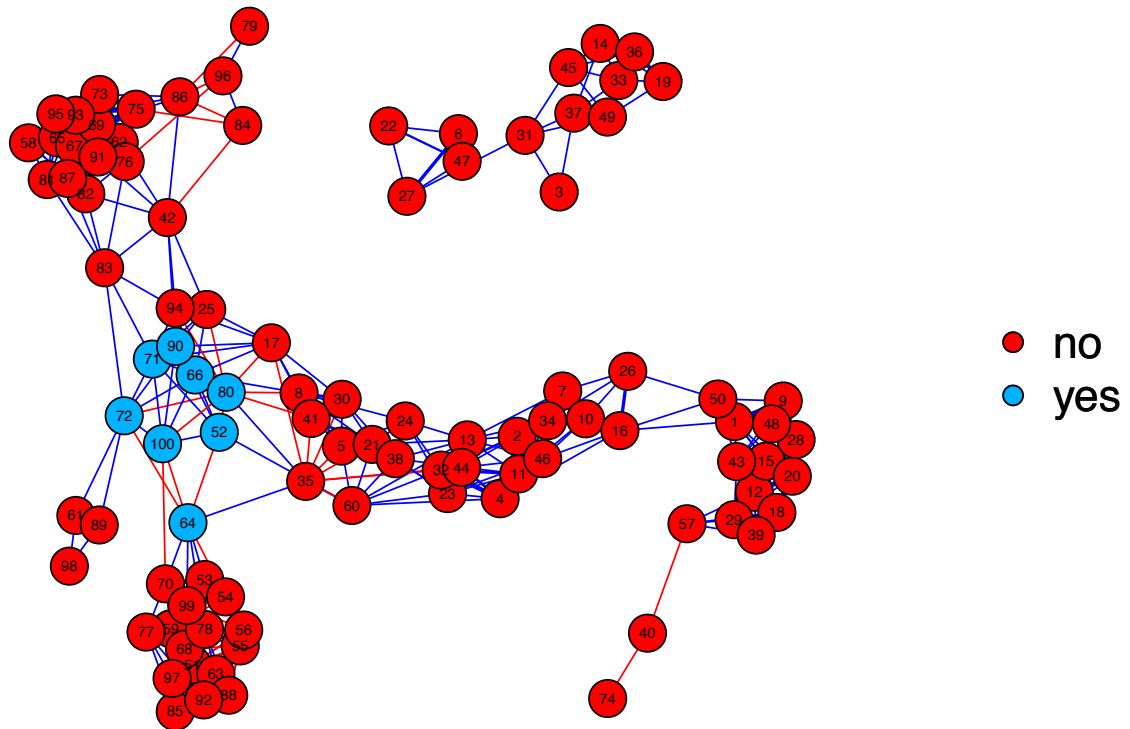


Figure S6. Network analysis of co-expressed genes among LB and SB segregants.

In blue and red are represented DE genes that are part of the network, which respectively present or not significant correlation with component 1 of the PCA analysis, selected as candidate genes for berry weight. Lines in red and blue represent negative and positive correlations, respectively.