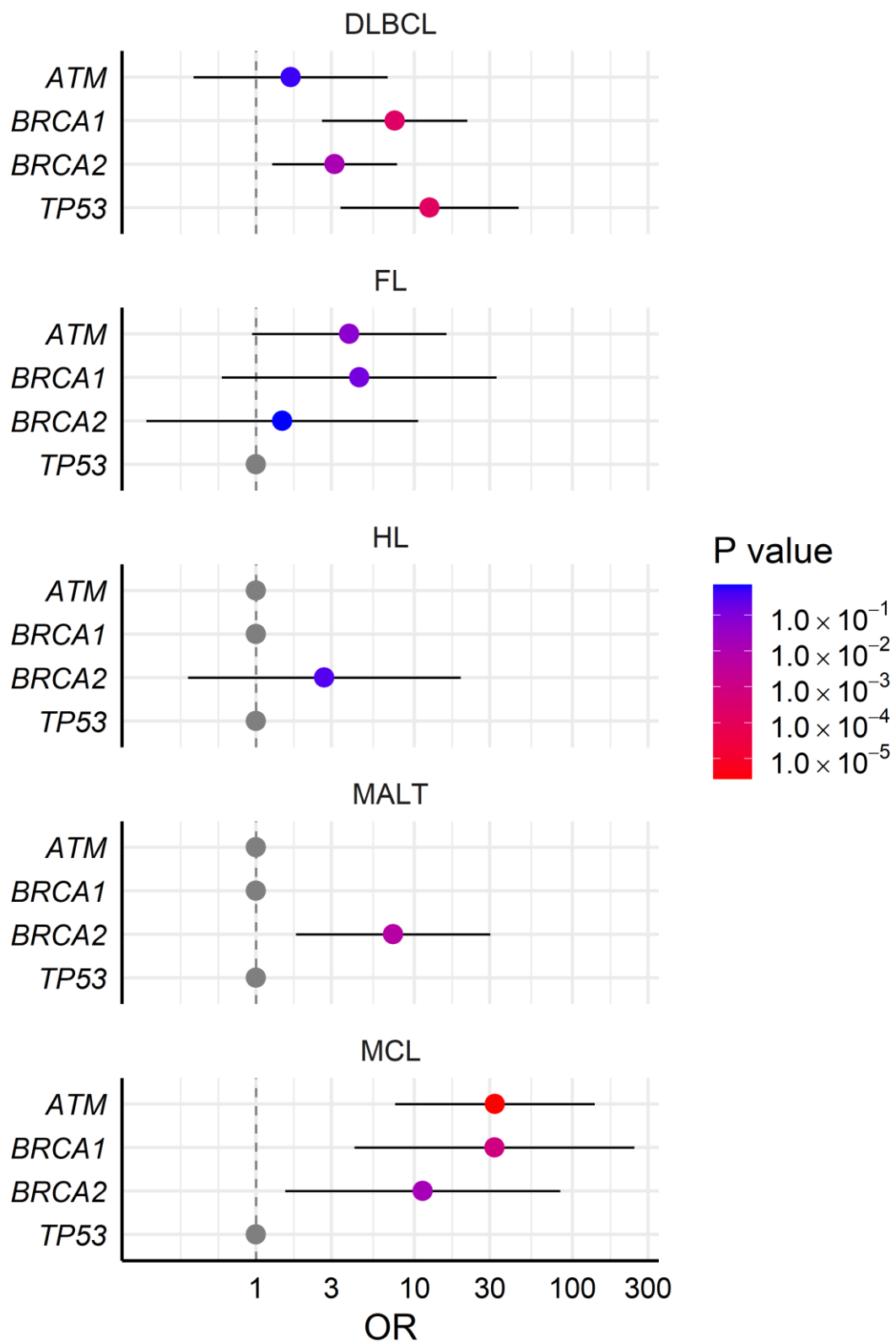


Figure S3. Association of germline pathogenic variants with each lymphoma risk based on different genes



The associations between germline pathogenic variants and lymphoma risk based on different genes were

evaluated using a logistic regression model adjusted for age and sex. Error bars represent 95% CIs.

Abbreviations: CI, confidence interval; MCL, mantle cell lymphoma; OR, odds ratio.