Figure S1. Top twenty-five genes selected by Lasso in the penalized logistic regression model

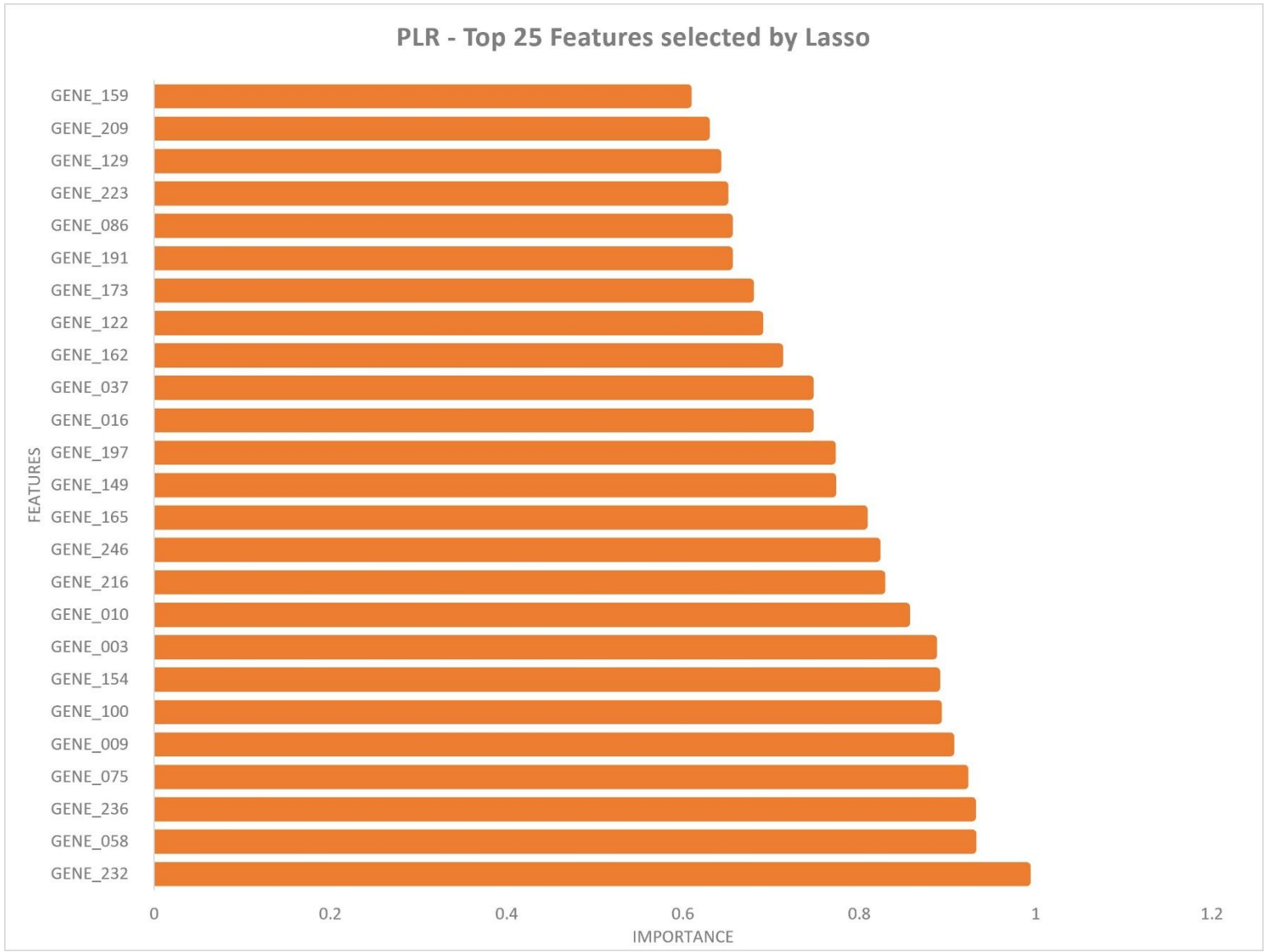


Figure S2. Top twenty-five genes selected by permutation importance in the random forest model

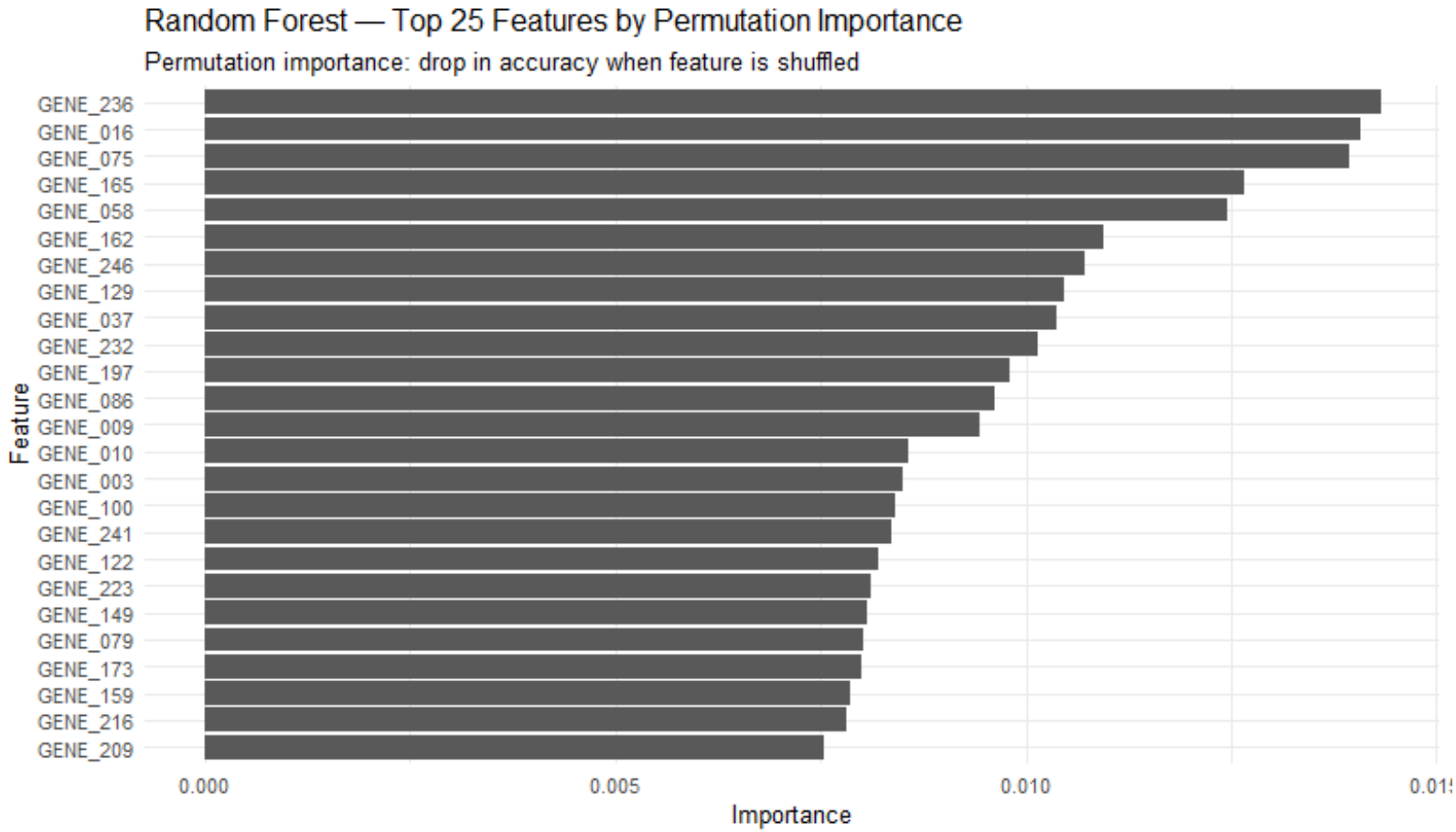


Figure S3. Top twenty-five genes selected by average gain in the extreme gradient boosting model

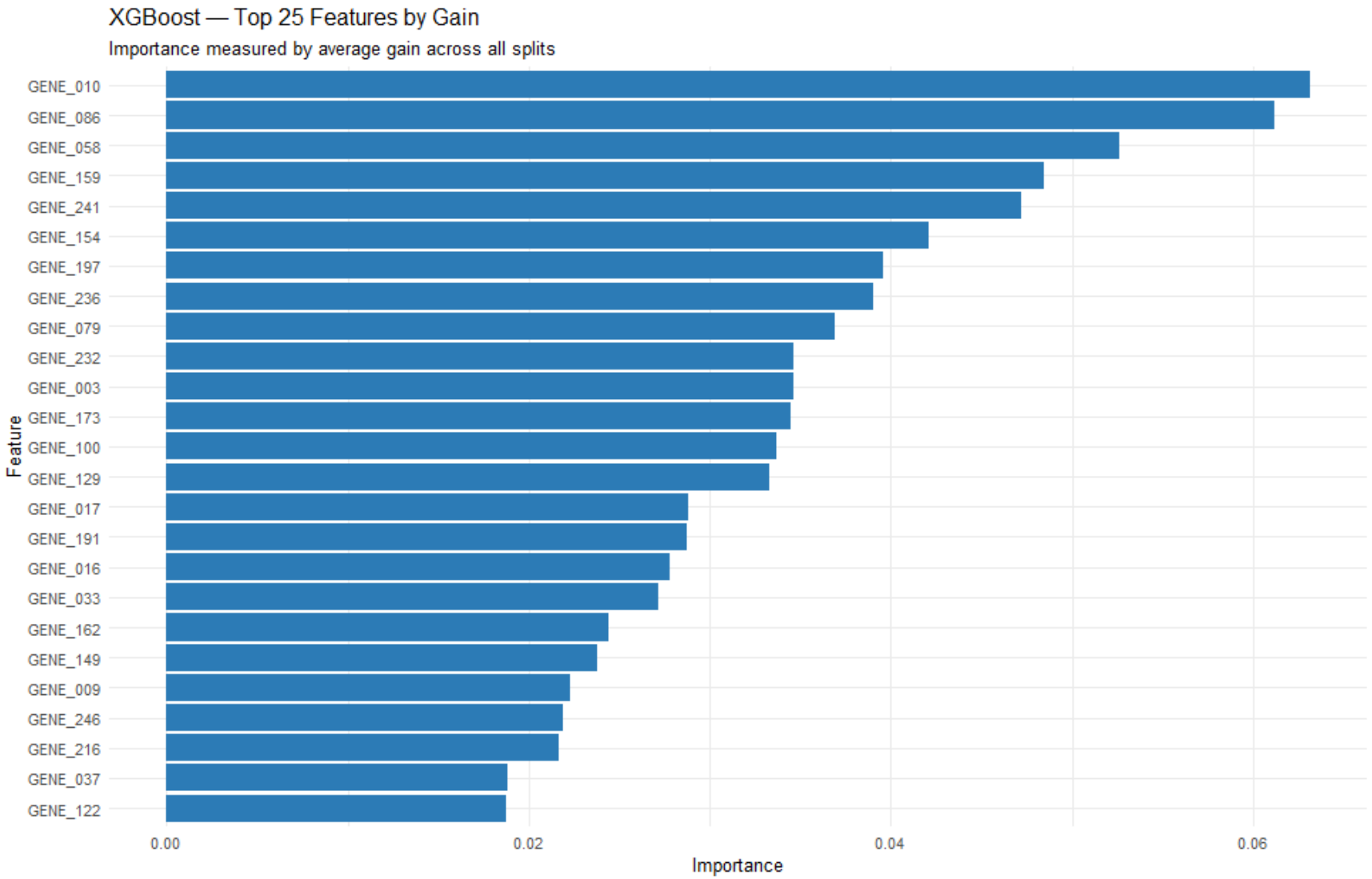


Figure S4. Top twenty-five features selected by permutation importance in the support vector machine model

