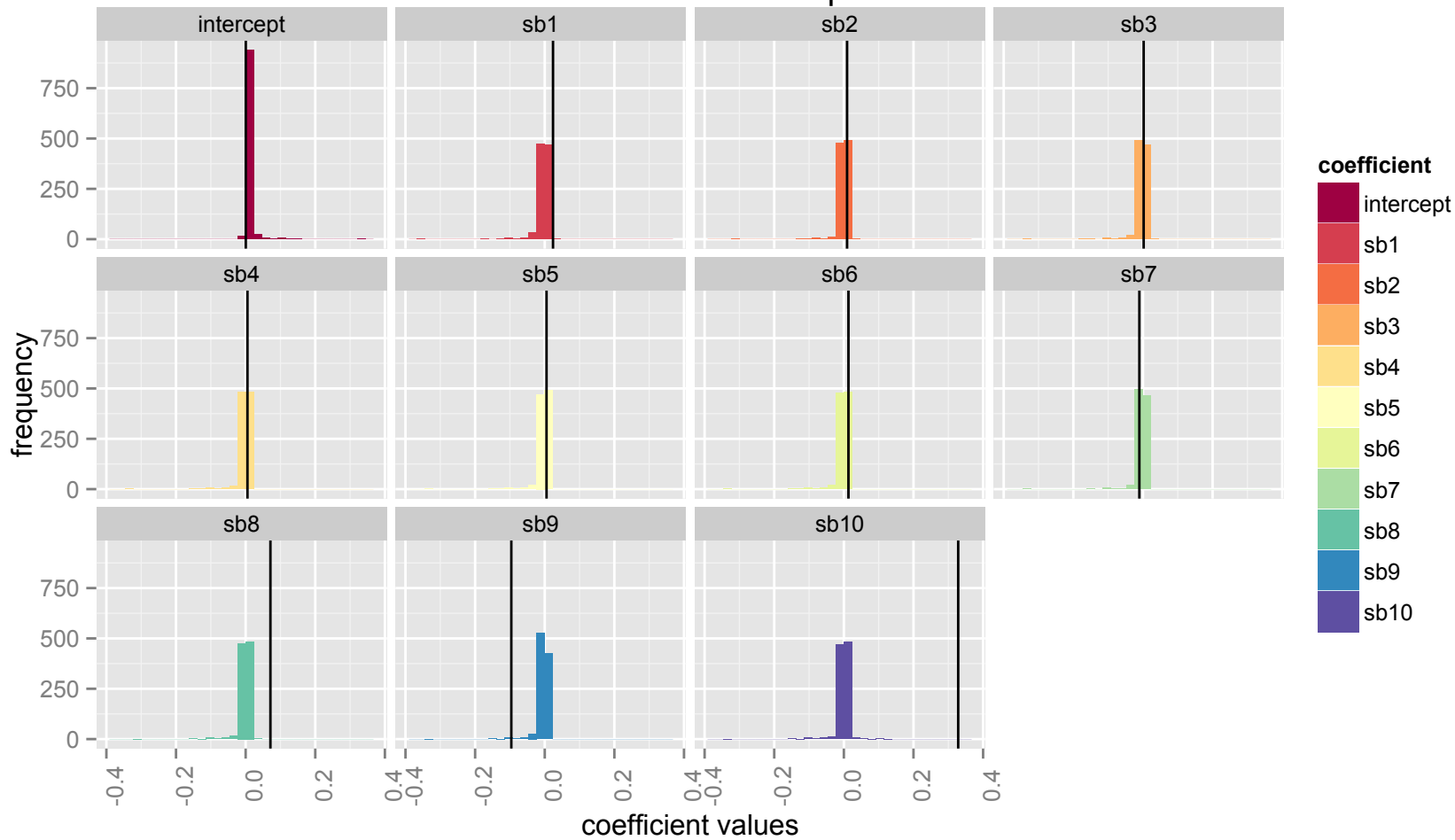


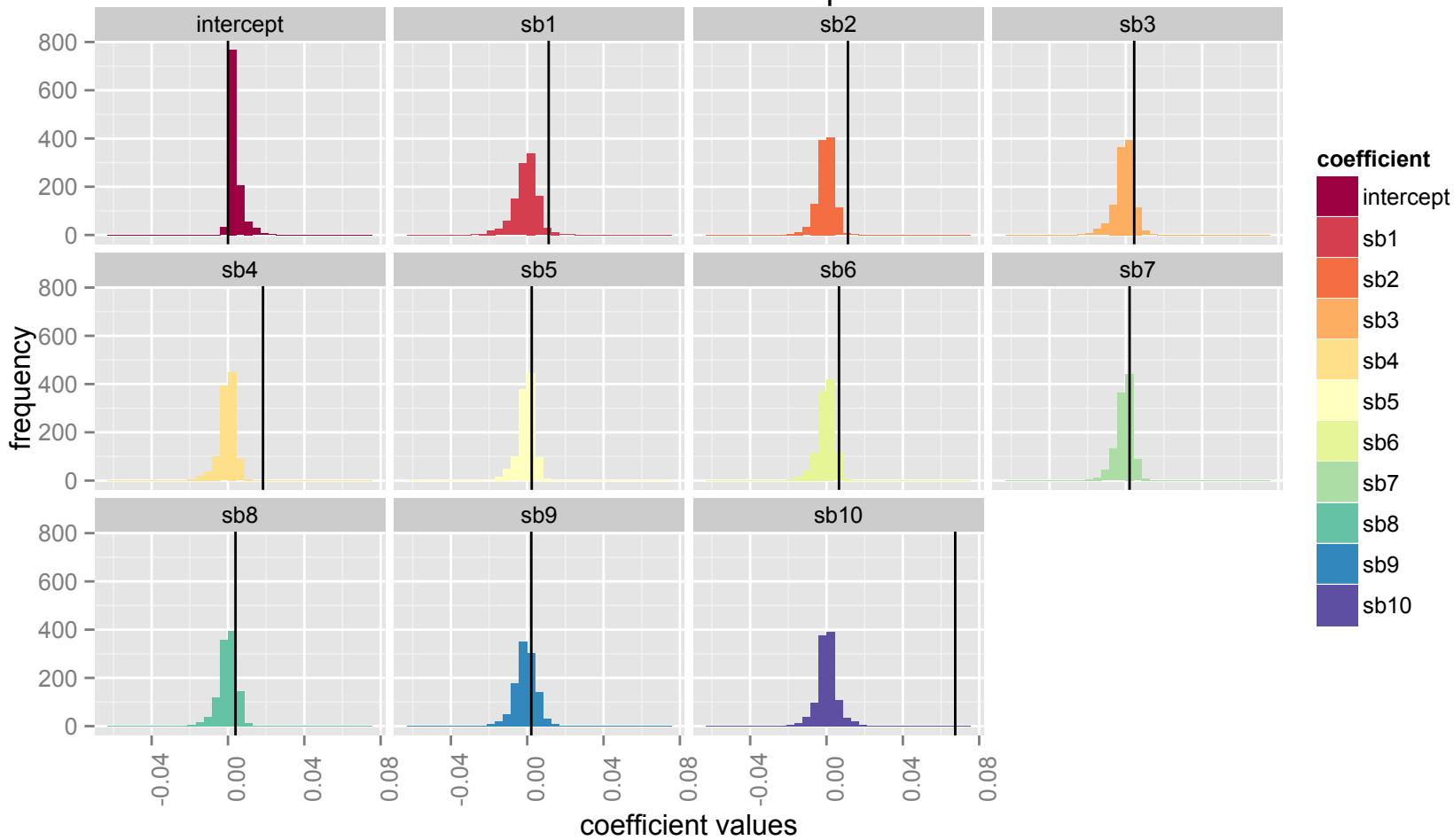
**Supplementary Figure S3. Null distributions for each regression coefficients**

corresponding to each of the 10 knots of the B-spline design, as well as the observed fitted coefficient (black vertical lines) for each one of the 14 chromosomes in *Plasmodium vivax*. We consider significant only those observed fitted values that are above or below 99% of the regression coefficients from the bootstrapped pseudo-random samples.

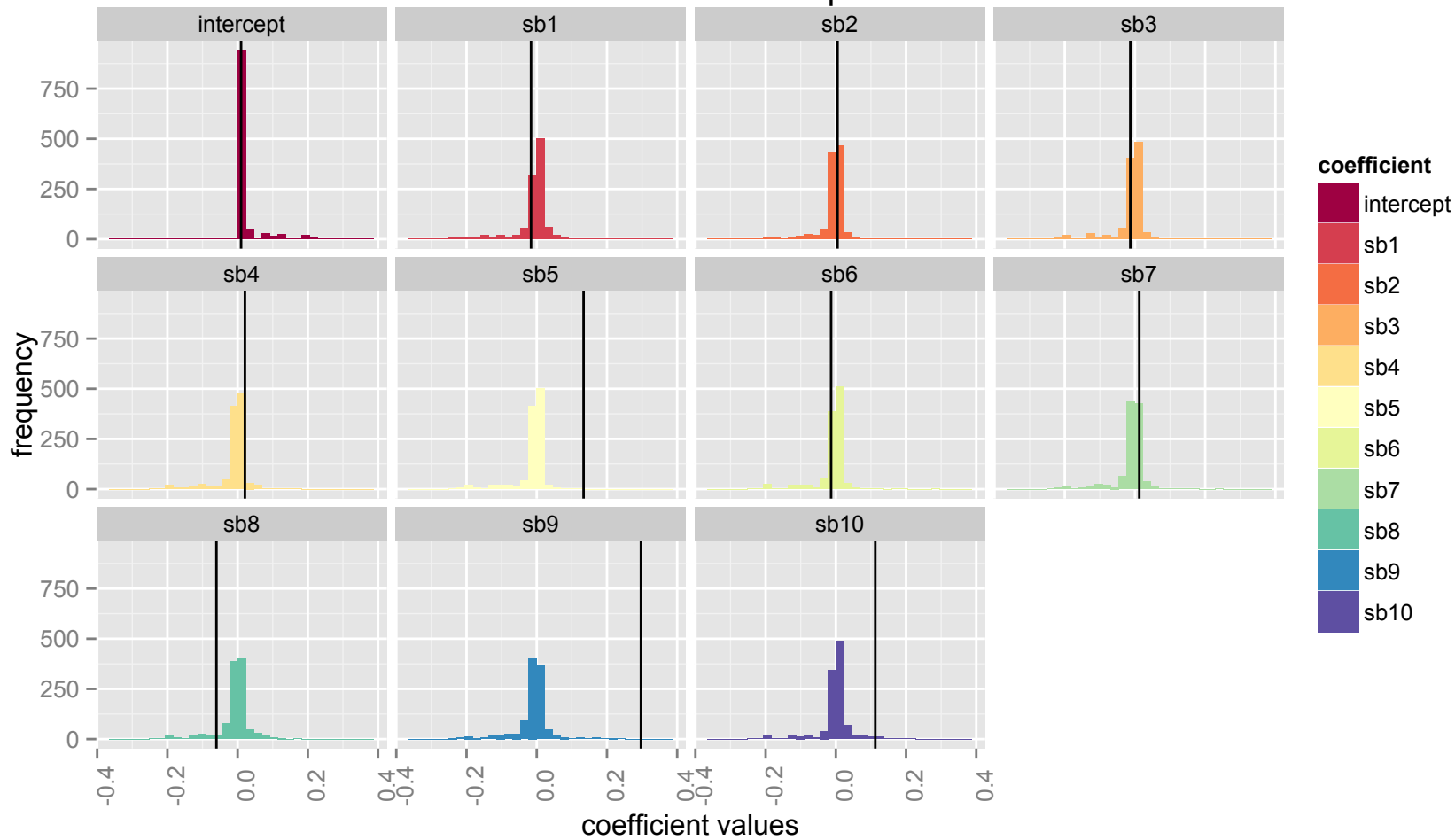
# chromosome 1 bootstrap test



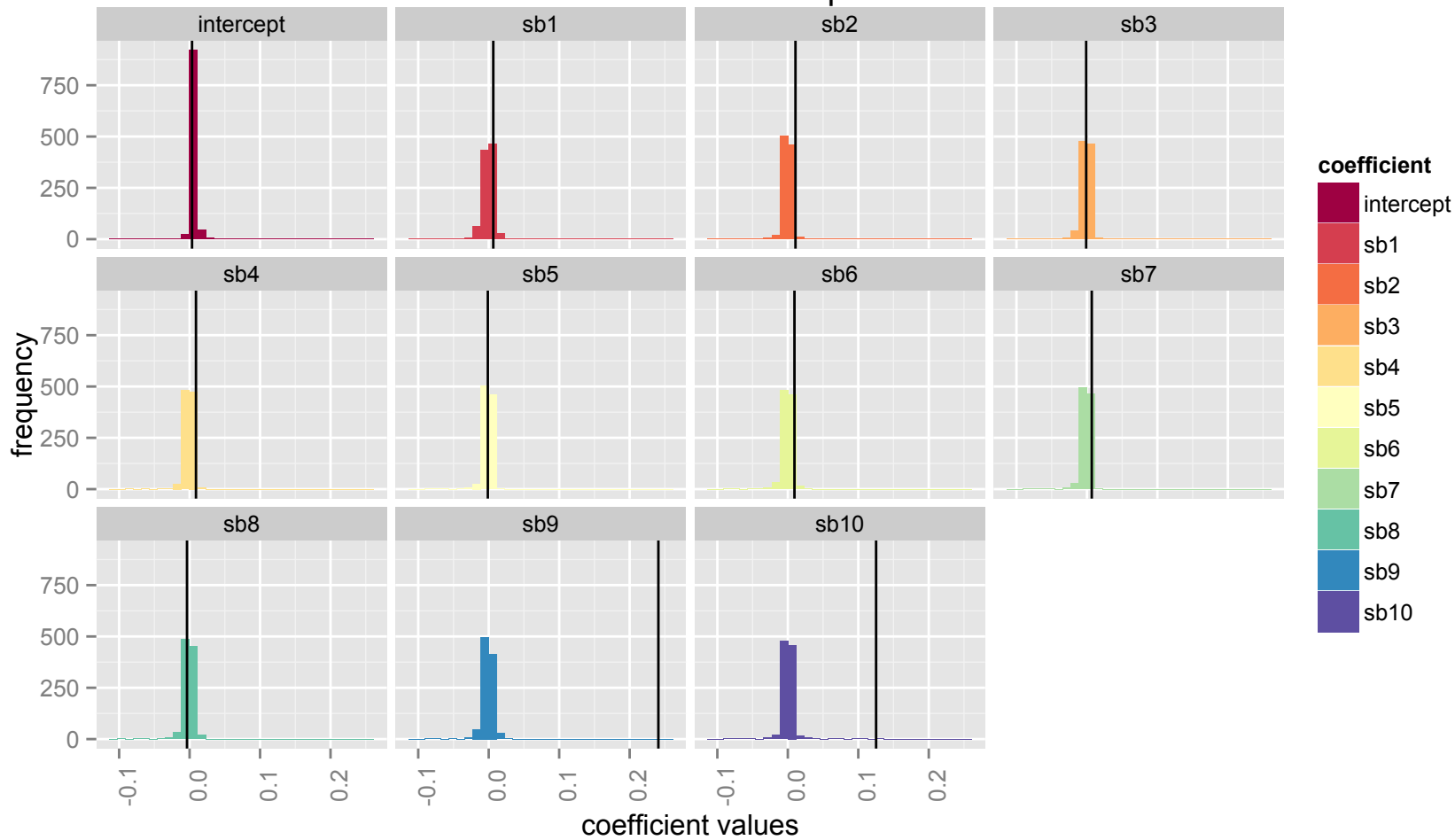
# chromosome 2 bootstrap test



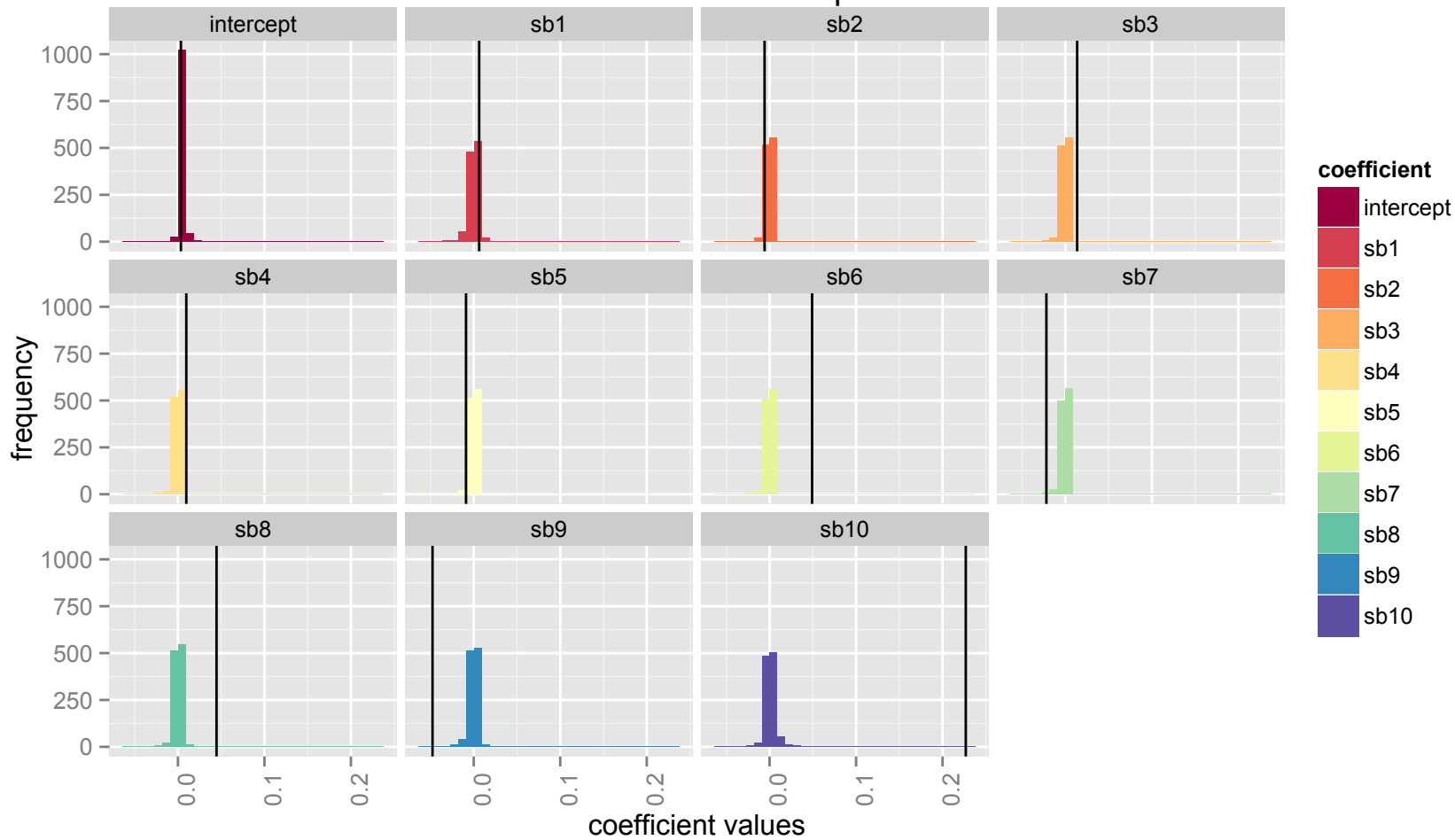
# chromosome 3 bootstrap test



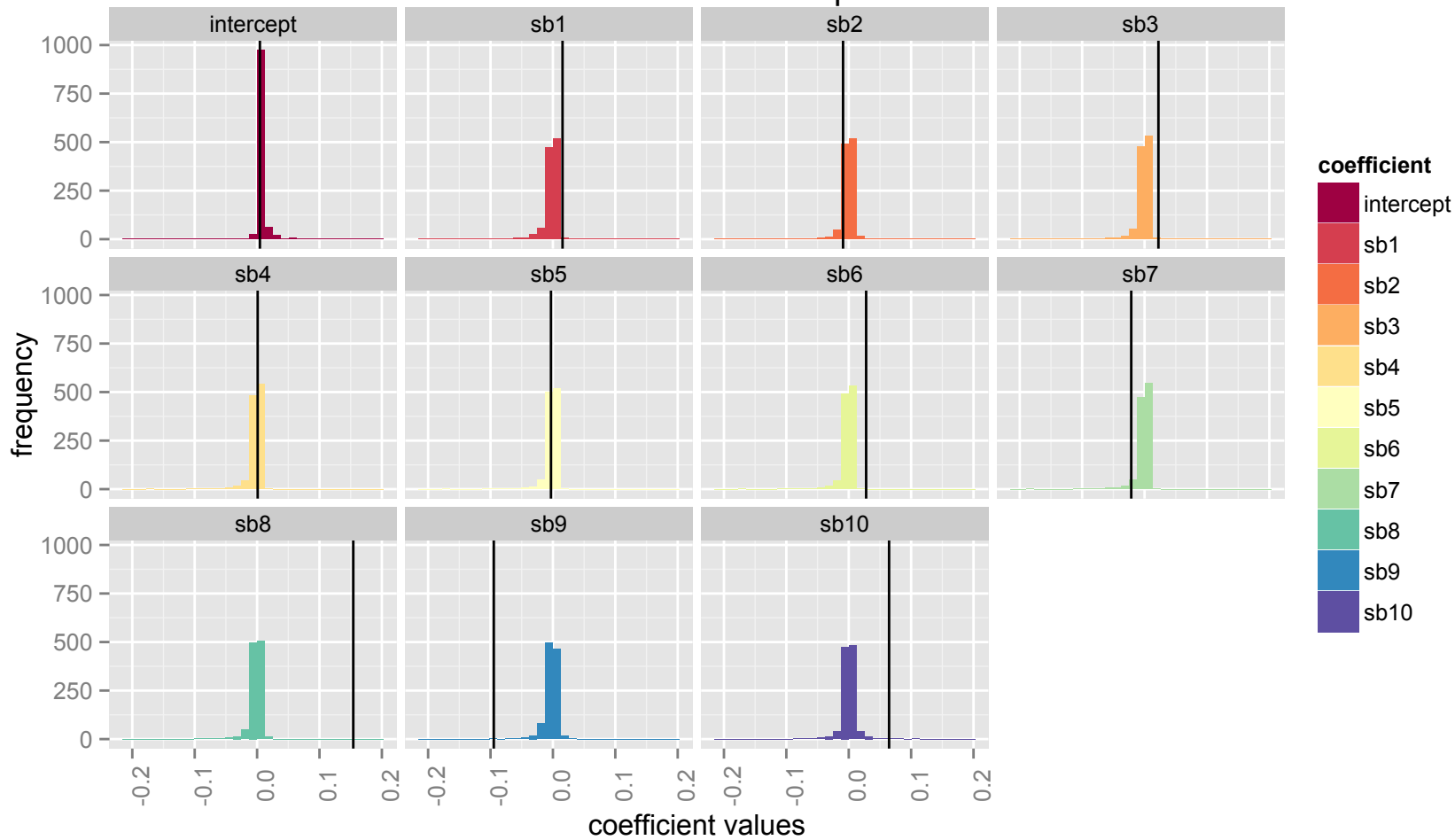
# chromosome 4 bootstrap test



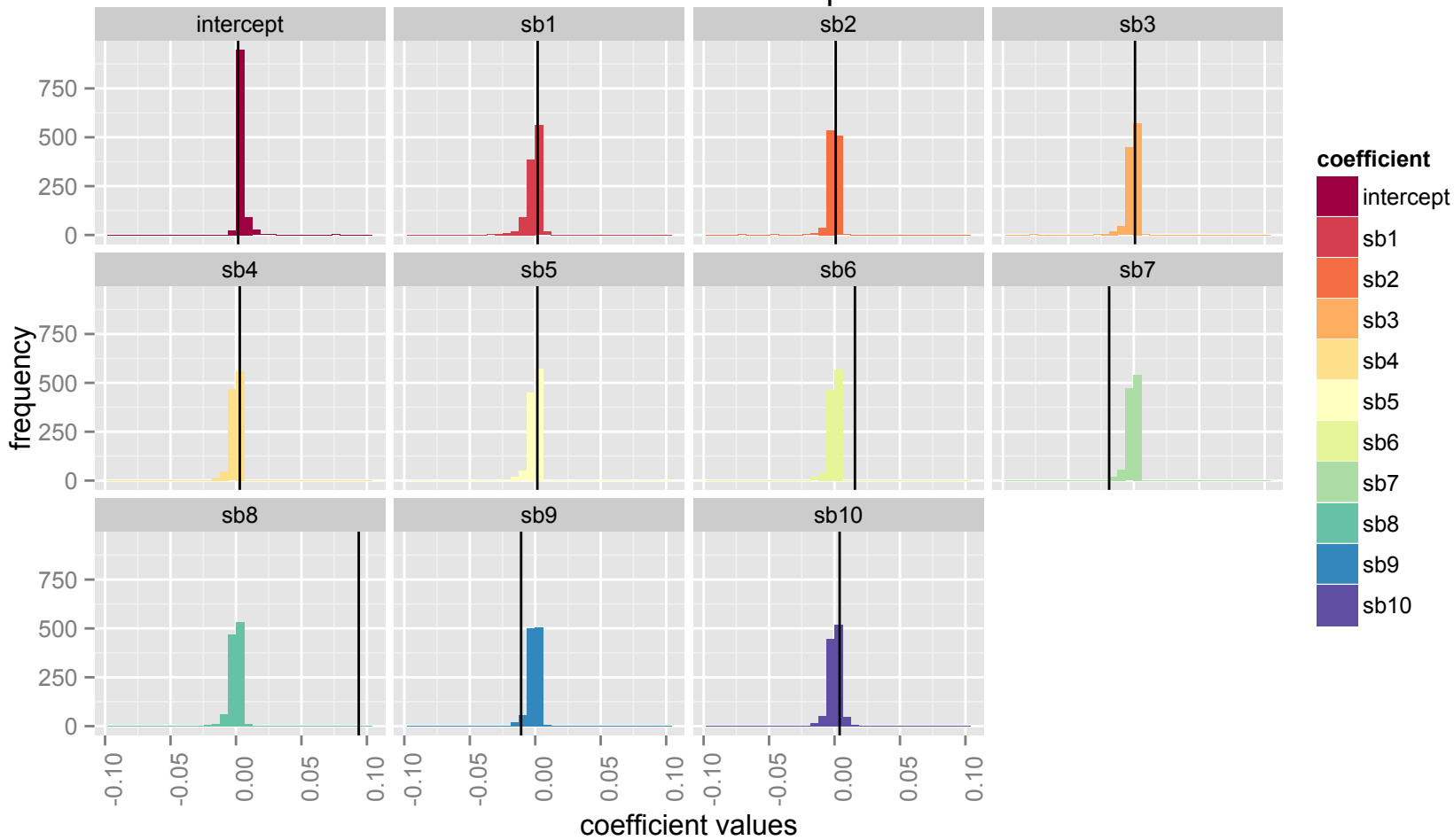
# chromosome 5 bootstrap test



# chromosome 6 bootstrap test

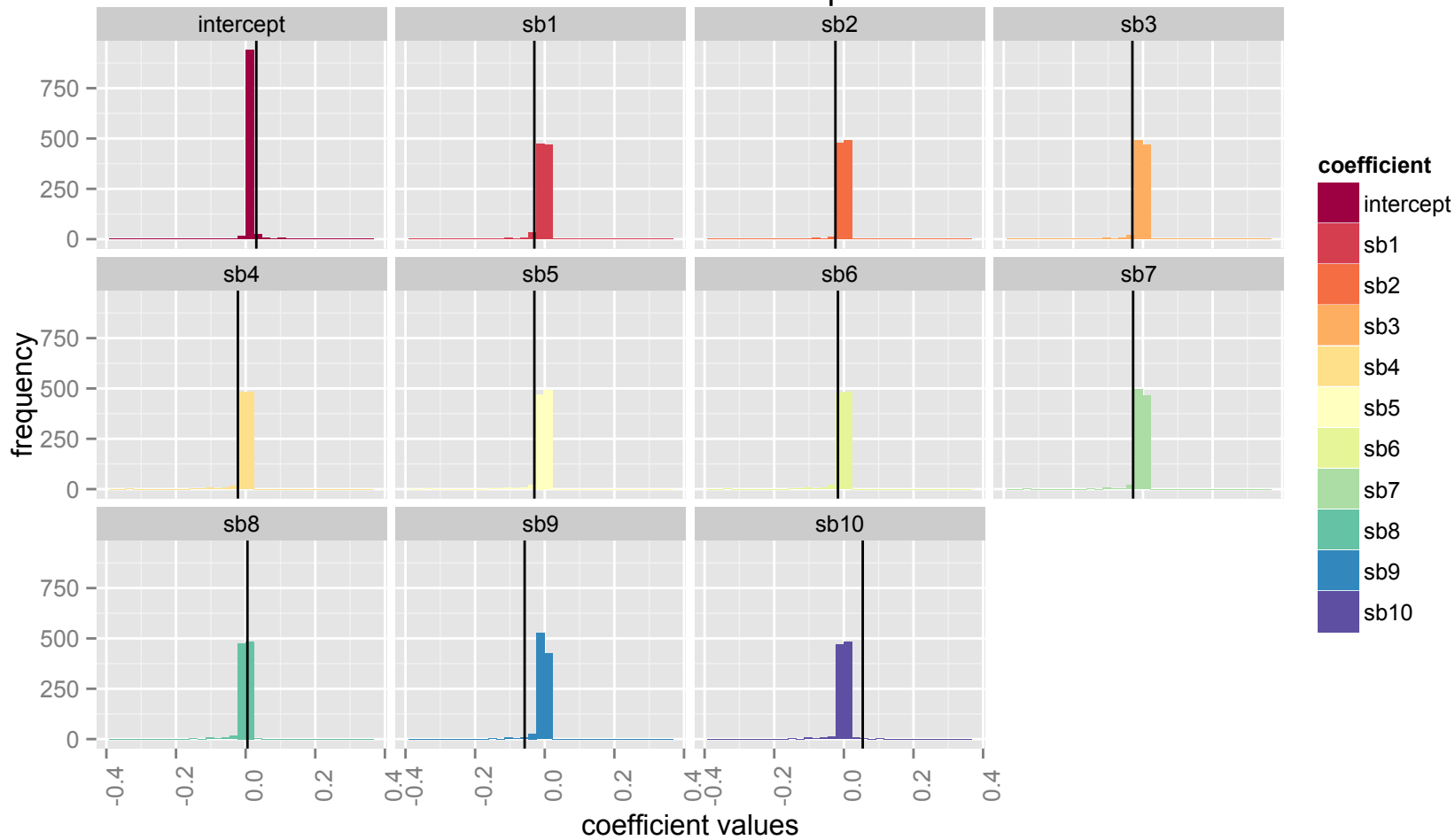


# chromosome 7 bootstrap test

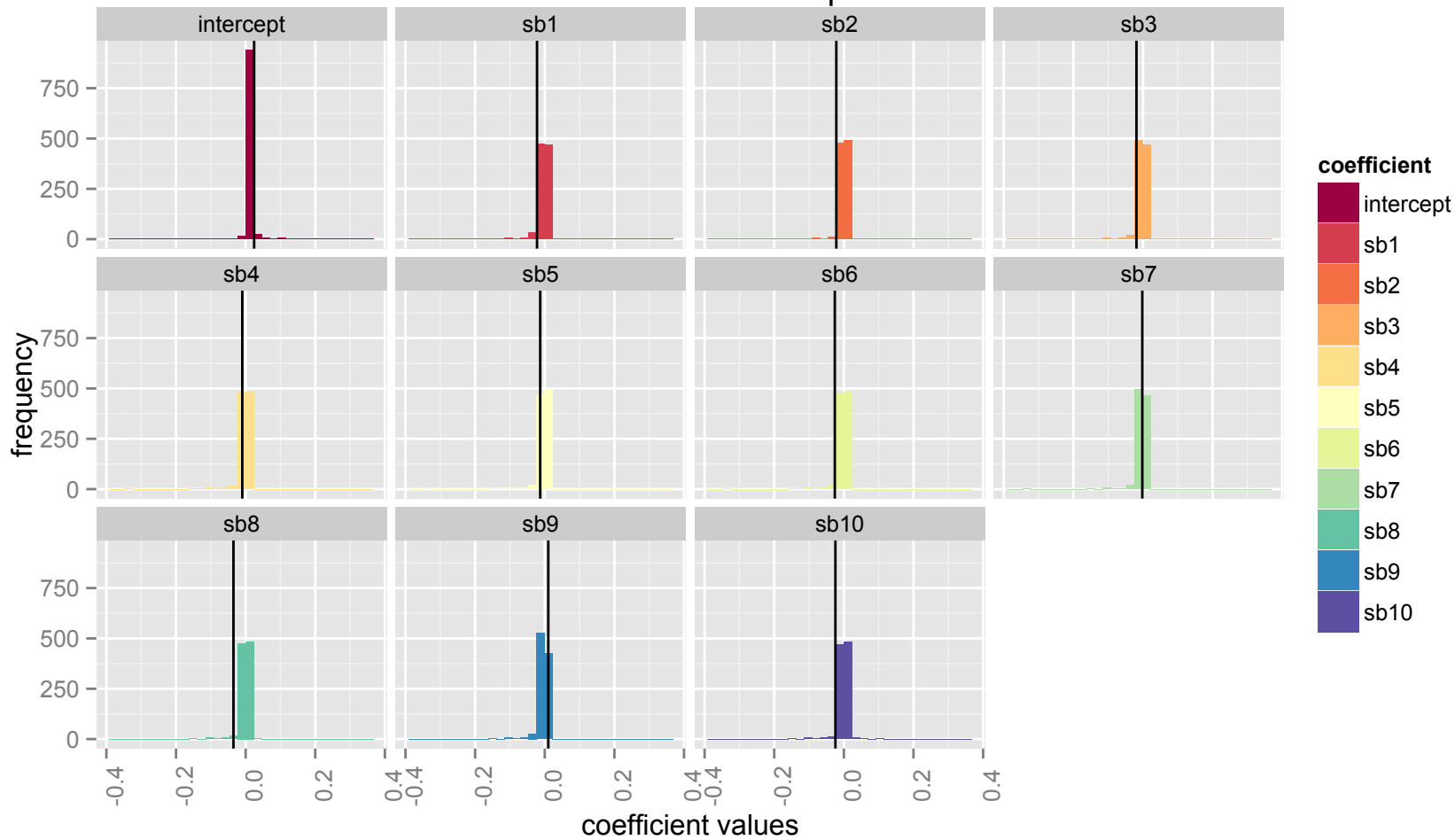




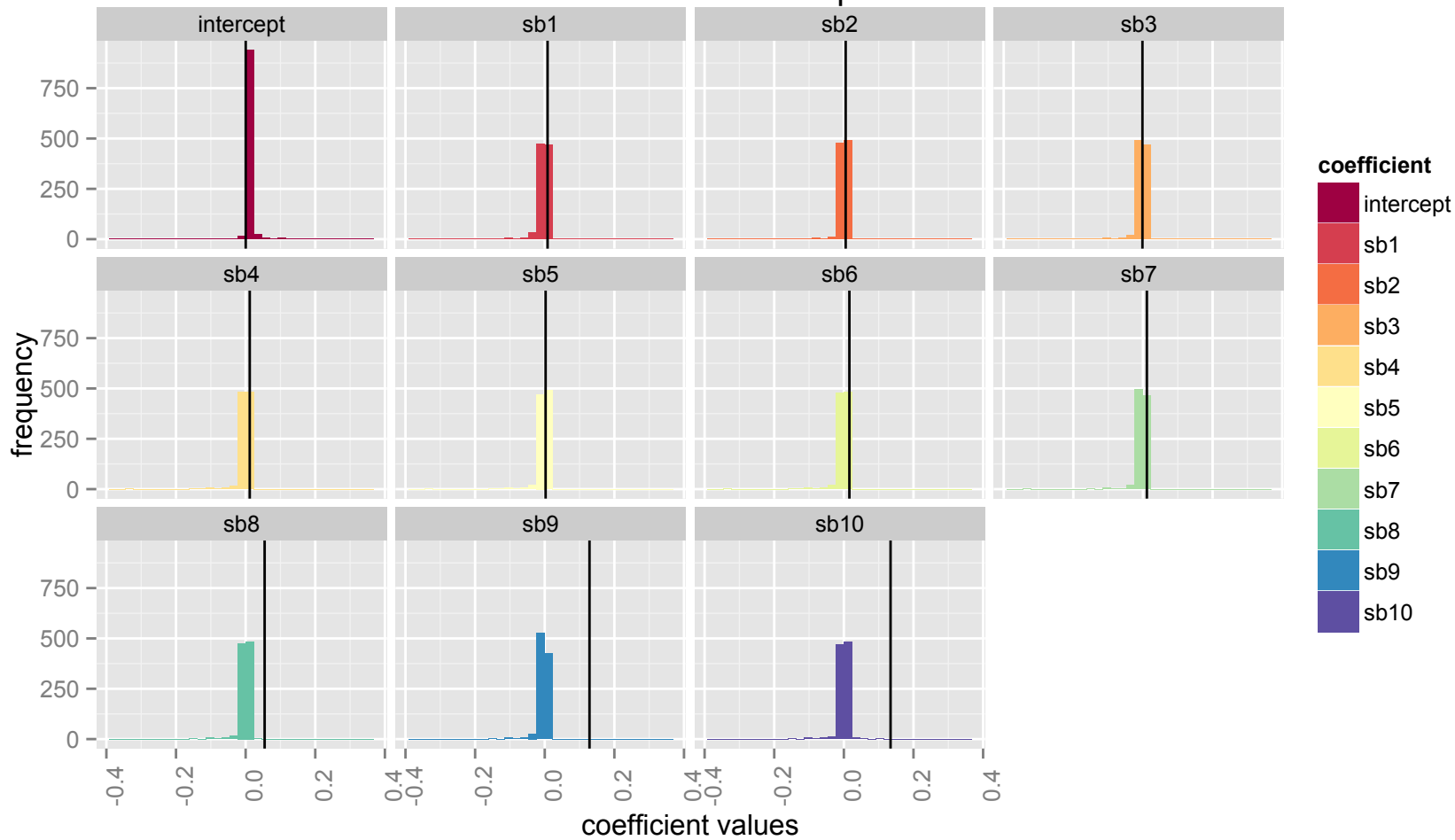
# chromosome 8 bootstrap test



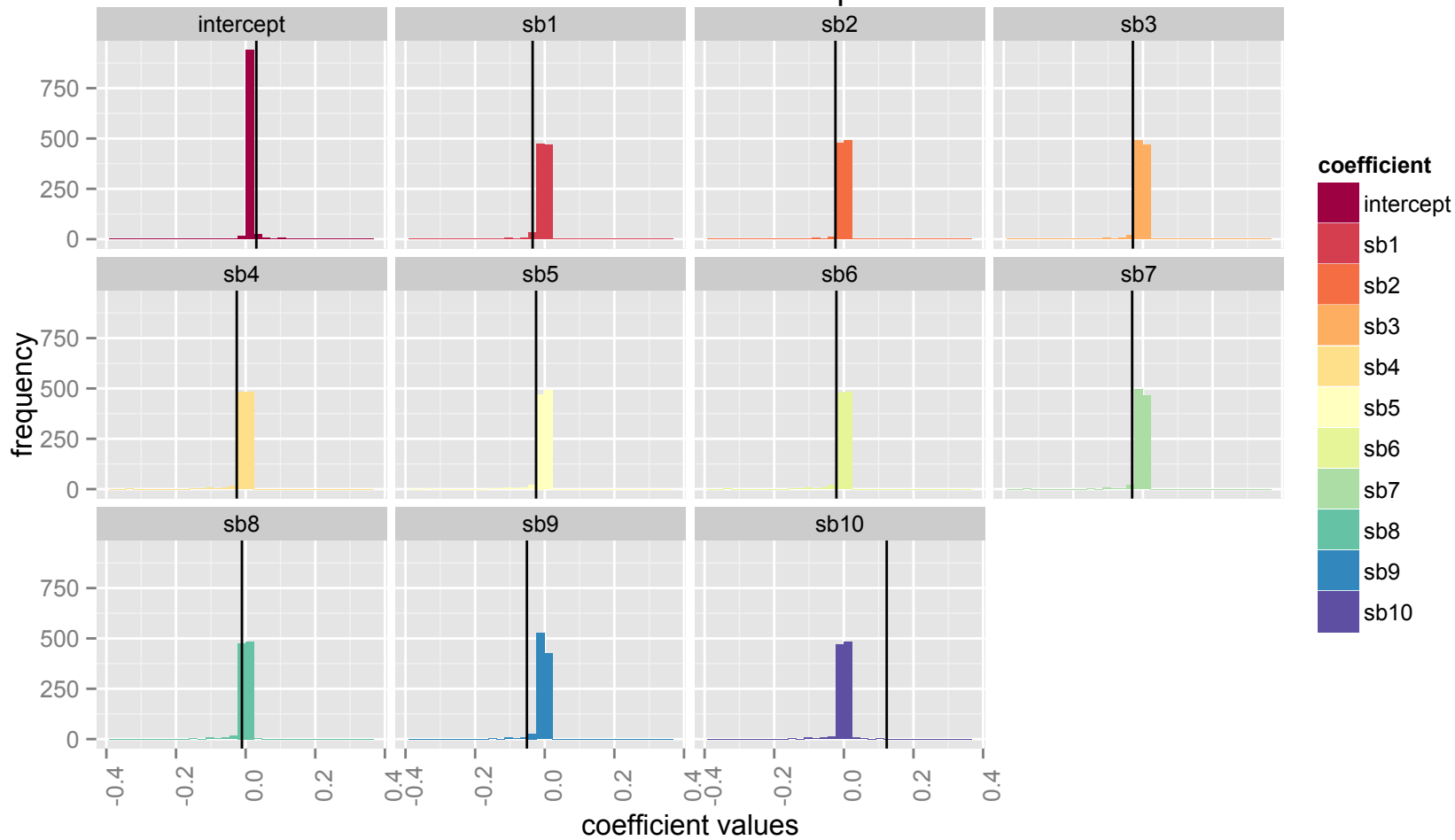
# chromosome 9 bootstrap test



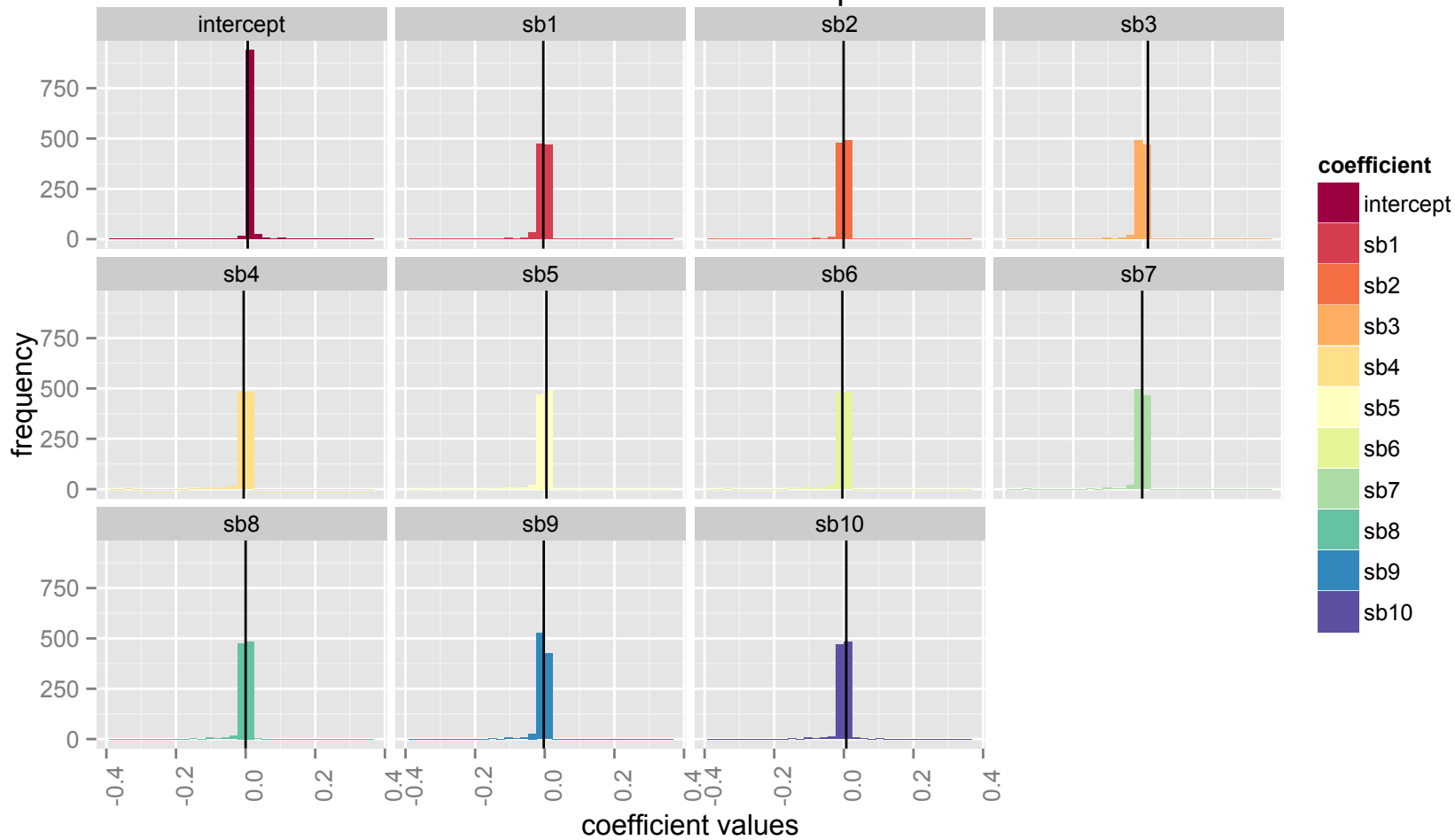
# chromosome 10 bootstrap test



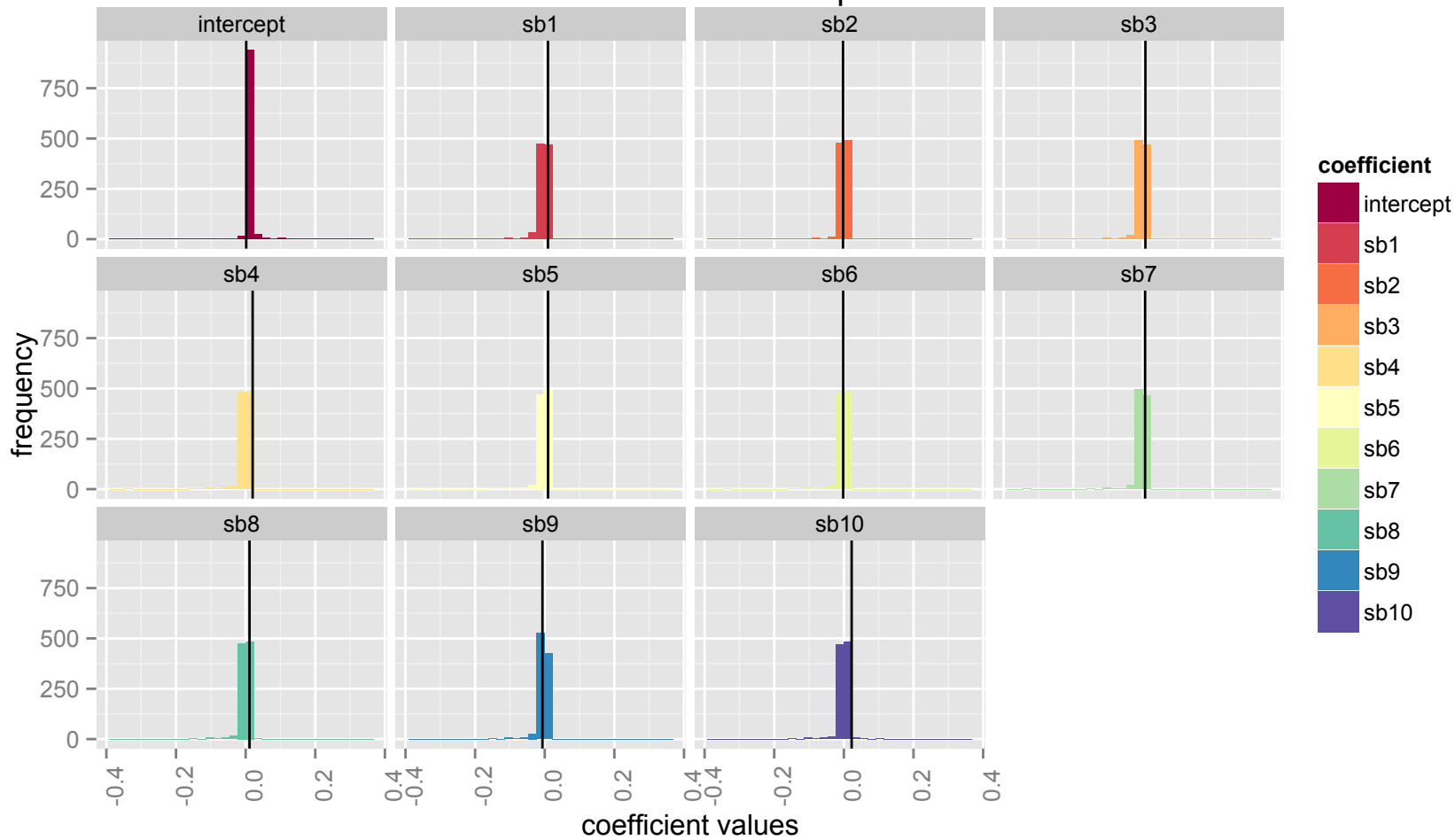
# chromosome 11 bootstrap test



# chromosome 12 bootstrap test



# chromosome 13 bootstrap test



# chromosome 14 bootstrap test

