

COVID correlates analysis report Nonlinear CoR

2021-02-07

1 Nonlinear association

To explore nonlinear association and threshold modeling, we fit smoothing spline models using the `mgcv` R package and segmented models using the `chngpt` R package.

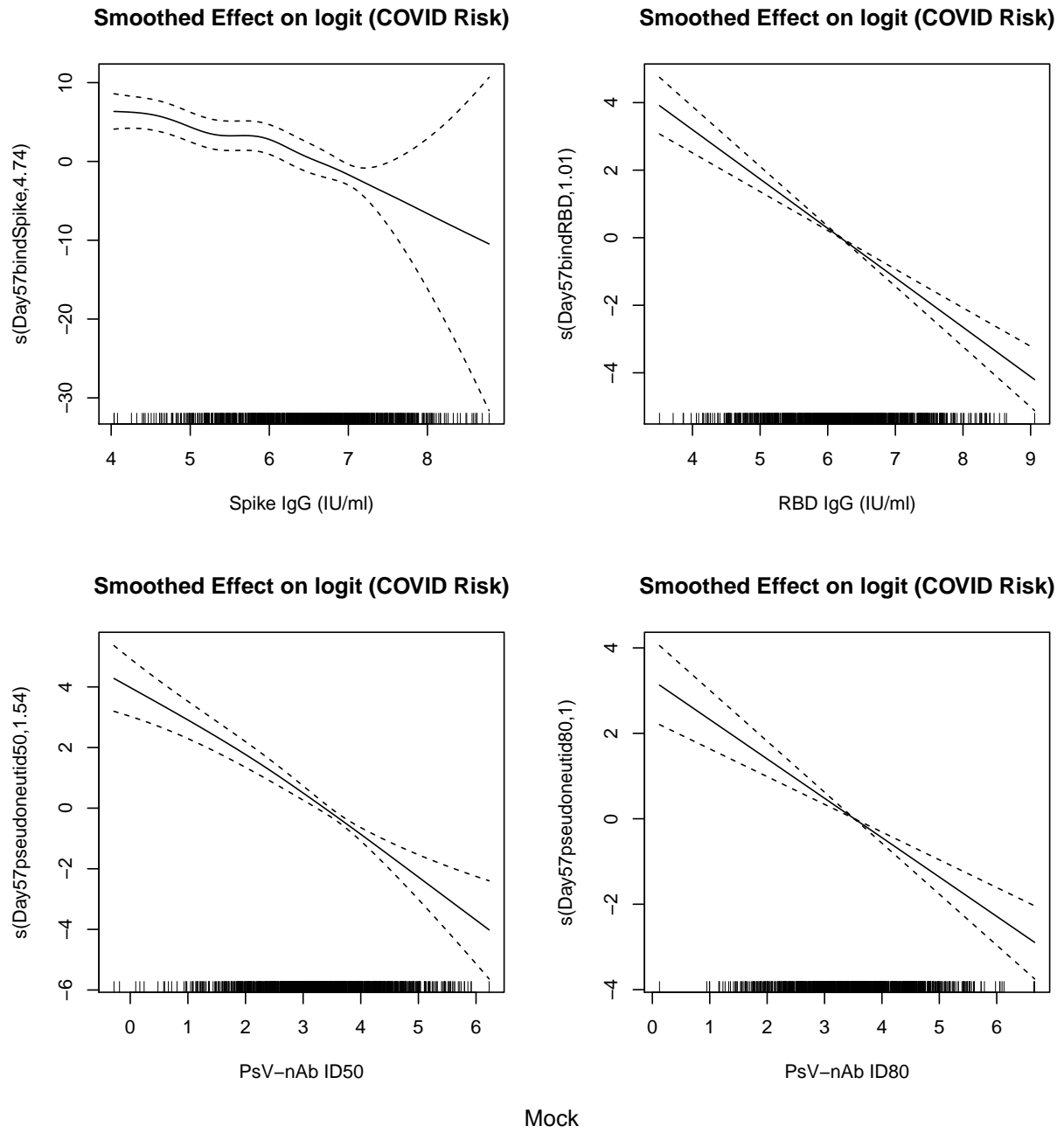


Figure 1: GAM fit of risk of infection versus markers. Clinical covariates are adjusted.

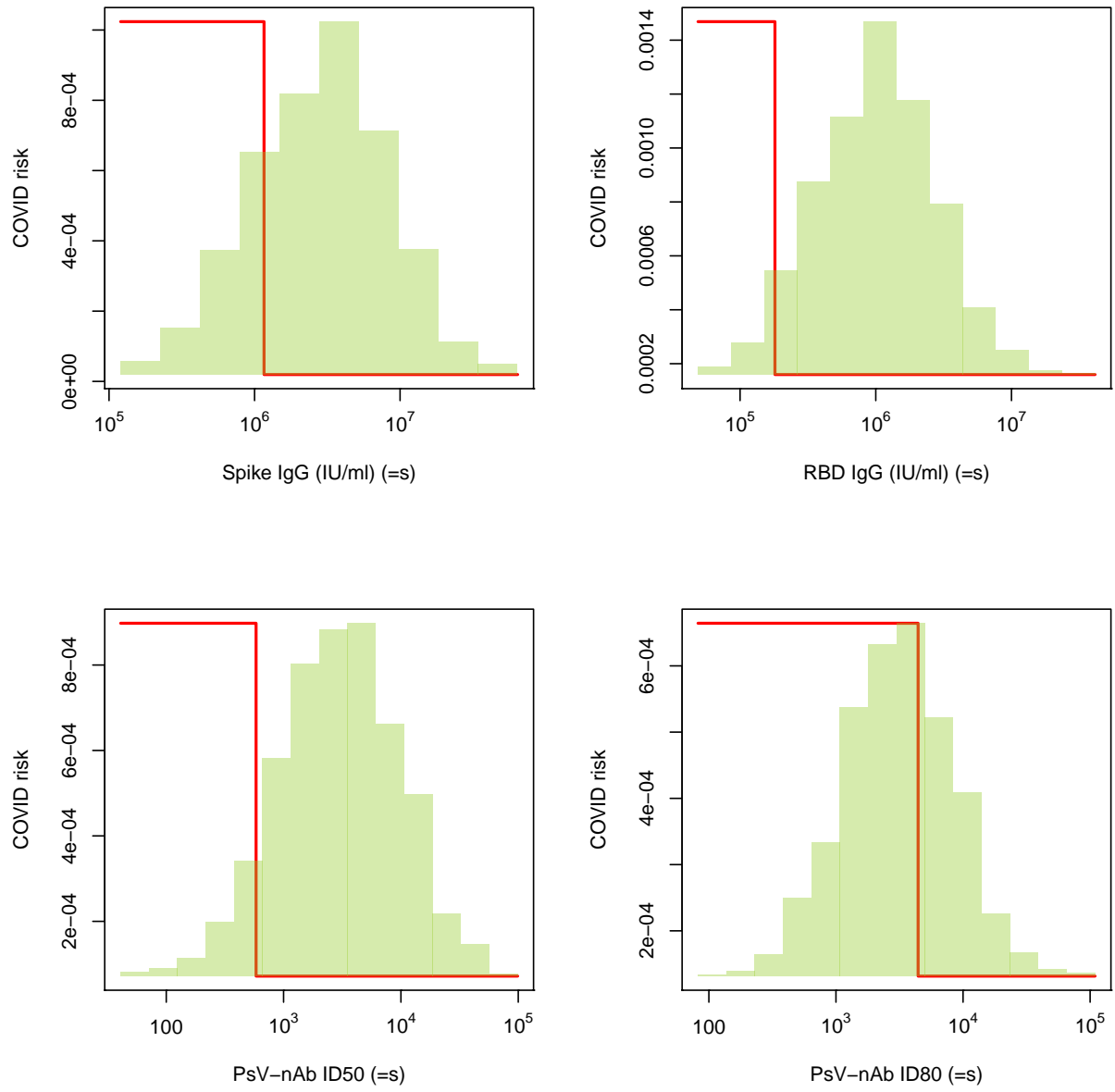


Figure 2: Step model fits. Clinical covariates are adjusted.