

# COVID correlates analysis report CoR

2021-02-03

## 1 Univariate CoR: Cox Proportional Hazards Modeling of Relative and Absolute Risk

All analyses for D57 markers are run on the population with EventTimePrimaryD57 $\geq$ 7. The main regression model is the Cox proportional hazards model. All plots are made with Cox models fit unless specified otherwise. The trichotomized variables in Table 1 are defined with respect to inverse probability weighted quantiles computed using the Hmisc::wtd.quantile function.

Table 1: Inference for Day 57 antibody marker covariate-adjusted correlates of risk of COVID in the vaccine group: Hazard ratios per 10-fold increment in the marker\*

| Mock<br>Immunologic Marker | No. cases /<br>No. at-risk** | HR per 10-fold incr. |             | P-value<br>(2-sided) | q-value | FWER   |
|----------------------------|------------------------------|----------------------|-------------|----------------------|---------|--------|
|                            |                              | Pt. Est.             | 95% CI      |                      |         |        |
| Spike IgG (IU/ml)          | 72/13,254                    | 0.08                 | (0.05-0.12) | <0.001               | <0.001  | <0.001 |
| RBD IgG (IU/ml)            | 72/13,254                    | 0.17                 | (0.12-0.25) | <0.001               | <0.001  | <0.001 |
| PsV-nAb ID50               | 72/13,254                    | 0.26                 | (0.20-0.34) | <0.001               | <0.001  | <0.001 |
| PsV-nAb ID80               | 72/13,254                    | 0.39                 | (0.29-0.52) | <0.001               | <0.001  | <0.001 |

\*Baseline covariates adjusted for: age in years, at risk or not, community of color or not, baseline risk score. Average follow-up time 175 days, maximum follow-up time 185 days.

\*\*No. at-risk = number of per-protocol baseline negative vaccine recipients at-risk for COVID at Day 57; no. cases = number of this cohort with an observed COVID endpoints.

Table 2: Inference for Day 57 antibody marker covariate-adjusted correlates of risk of COVID in the vaccine group: Hazard ratios for Middle vs. Upper tertile vs. Lower tertile\*

| Mock Immunologic Marker | Tertile | No. cases / No. at-risk** | Attack rate | Haz. Ratio Pt. Est. | 95% CI      | P-value (2-sided) | Overall P-value*** | Overall q-value | Overall FWER |
|-------------------------|---------|---------------------------|-------------|---------------------|-------------|-------------------|--------------------|-----------------|--------------|
| Spike IgG (IU/ml)       | Lower   | 67/4,373                  | 0.0153      | 1                   | N/A         | N/A               | <0.001             | <0.001          | <0.001       |
|                         | Middle  | 4/4,449                   | 0.0009      | 0.04                | (0.01-0.11) | <0.001            |                    |                 |              |
|                         | Upper   | 1/4,422                   | 0.0002      | 0.00                | (0.00-0.03) | <0.001            |                    |                 |              |
| RBD IgG (IU/ml)         | Lower   | 45/4,395                  | 0.0102      | 1                   | N/A         | N/A               | <0.001             | <0.001          | <0.001       |
|                         | Middle  | 19/4,433                  | 0.0043      | 0.24                | (0.13-0.43) | <0.001            |                    |                 |              |
|                         | Upper   | 8/4,416                   | 0.0018      | 0.05                | (0.02-0.12) | <0.001            |                    |                 |              |
| PsV-nAb ID50            | Lower   | 56/4,440                  | 0.0126      | 1                   | N/A         | N/A               | <0.001             | <0.001          | <0.001       |
|                         | Middle  | 9/4,416                   | 0.0020      | 0.10                | (0.05-0.22) | <0.001            |                    |                 |              |
|                         | Upper   | 6/4,388                   | 0.0014      | 0.05                | (0.02-0.11) | <0.001            |                    |                 |              |
| PsV-nAb ID80            | Lower   | 40/4,392                  | 0.0091      | 1                   | N/A         | N/A               | <0.001             | <0.001          | <0.001       |
|                         | Middle  | 21/4,436                  | 0.0047      | 0.43                | (0.24-0.78) | 0.005             |                    |                 |              |
|                         | Upper   | 11/4,417                  | 0.0025      | 0.16                | (0.08-0.34) | <0.001            |                    |                 |              |
| Placebo                 |         | 713/13,299                | 0.0536      |                     |             |                   |                    |                 |              |

\*Baseline covariates adjusted for: age in years, at risk or not, community of color or not, baseline risk score. Average follow-up time 175 days, maximum follow-up time 185 days. Cutpoints: Spike IgG (IU/ml) [6.09, 6.7), RBD IgG (IU/ml) [5.68, 6.38), PsV-nAb ID50 [2.8, 3.66), PsV-nAb ID80 [3.08, 3.82).

\*\*No. at-risk = number of per-protocol baseline negative vaccine recipients at-risk for COVID at Day 57; no. cases = number of this cohort with an observed COVID endpoints.

\*\*\*Generalized Wald-test p-value of the null hypothesis that the hazard rate is constant across the Lower, Middle, and Upper tertile groups.

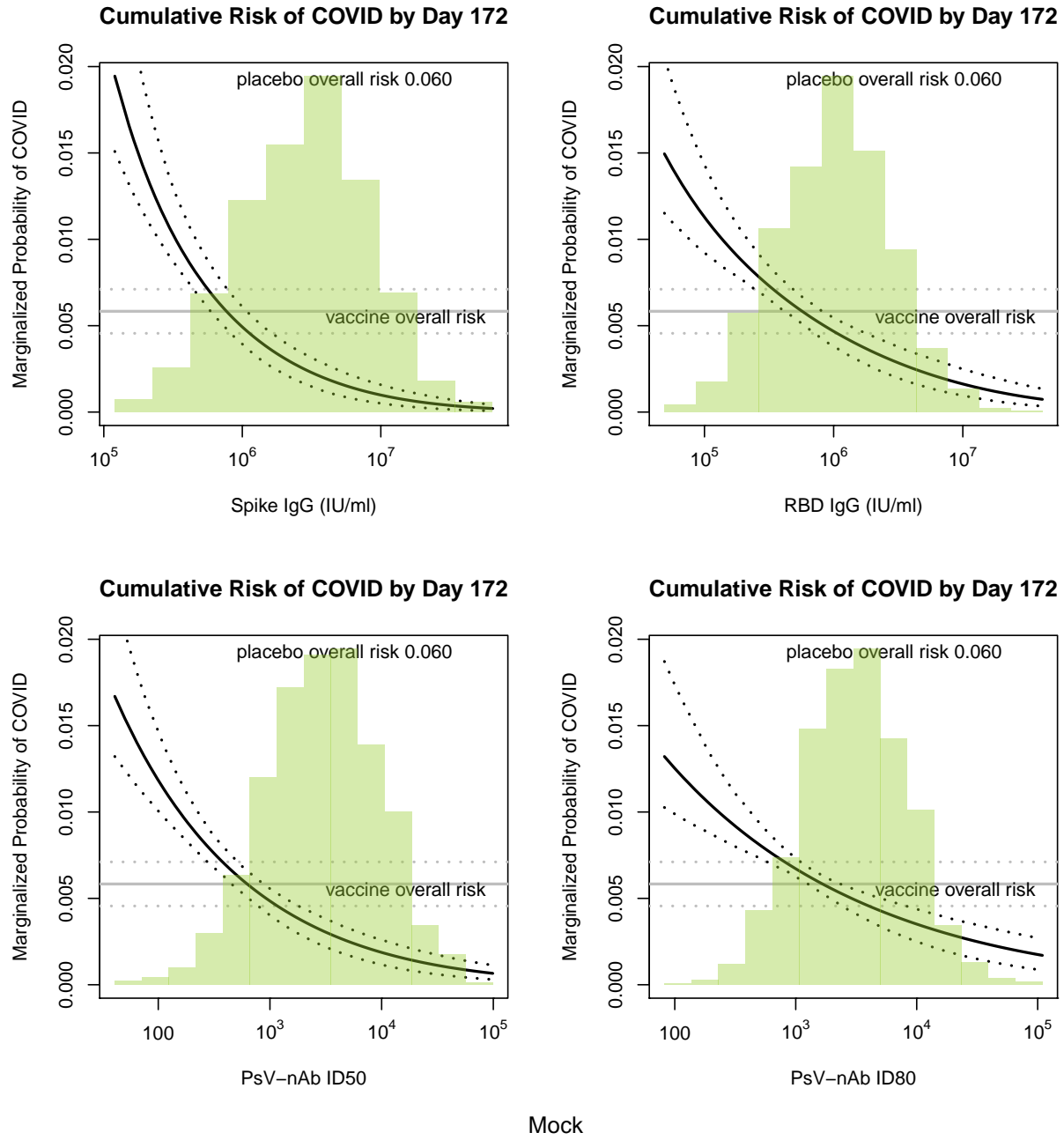


Figure 1: Marginalized cumulative risk by Day 172 as functions of Day 57 markers among baseline seronegative vaccine recipients with 95% bootstrap point-wise confidence bands. The horizontal lines indicate the overall cumulative risk of the vaccine arm by Day 172 and its 95% point-wise confidence interval. Histograms of the immunological markers in the vaccine arm are overlaid.

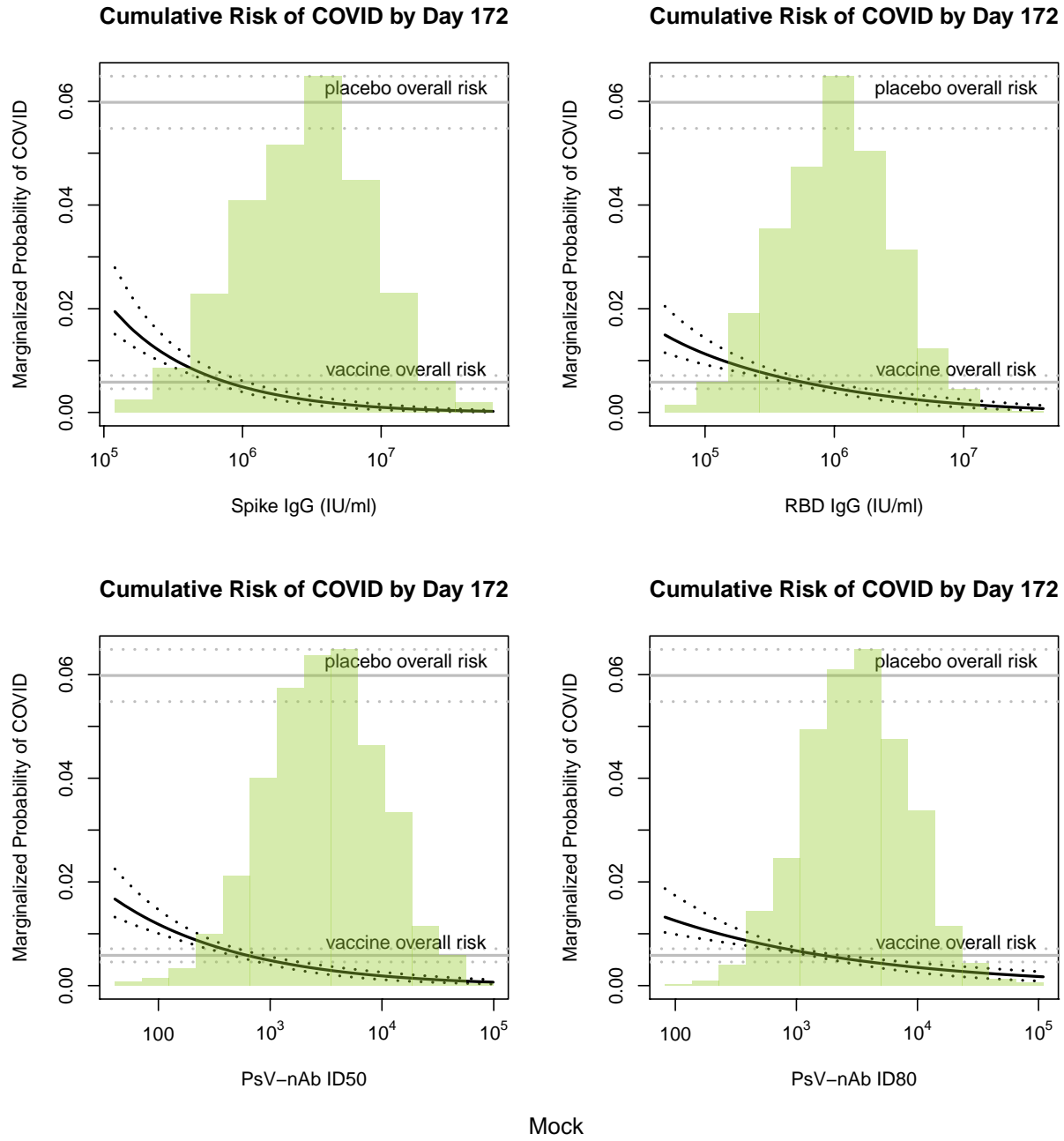


Figure 2: Marginalized cumulative risk by Day 172 as functions of Day 57 markers among baseline seronegative vaccine recipients with 95% bootstrap point-wise confidence bands. The horizontal lines indicate the overall cumulative risk of the placebo and vaccine arms by Day 172 and its 95% point-wise confidence interval. Histograms of the immunological markers in the vaccine arm are overlaid.

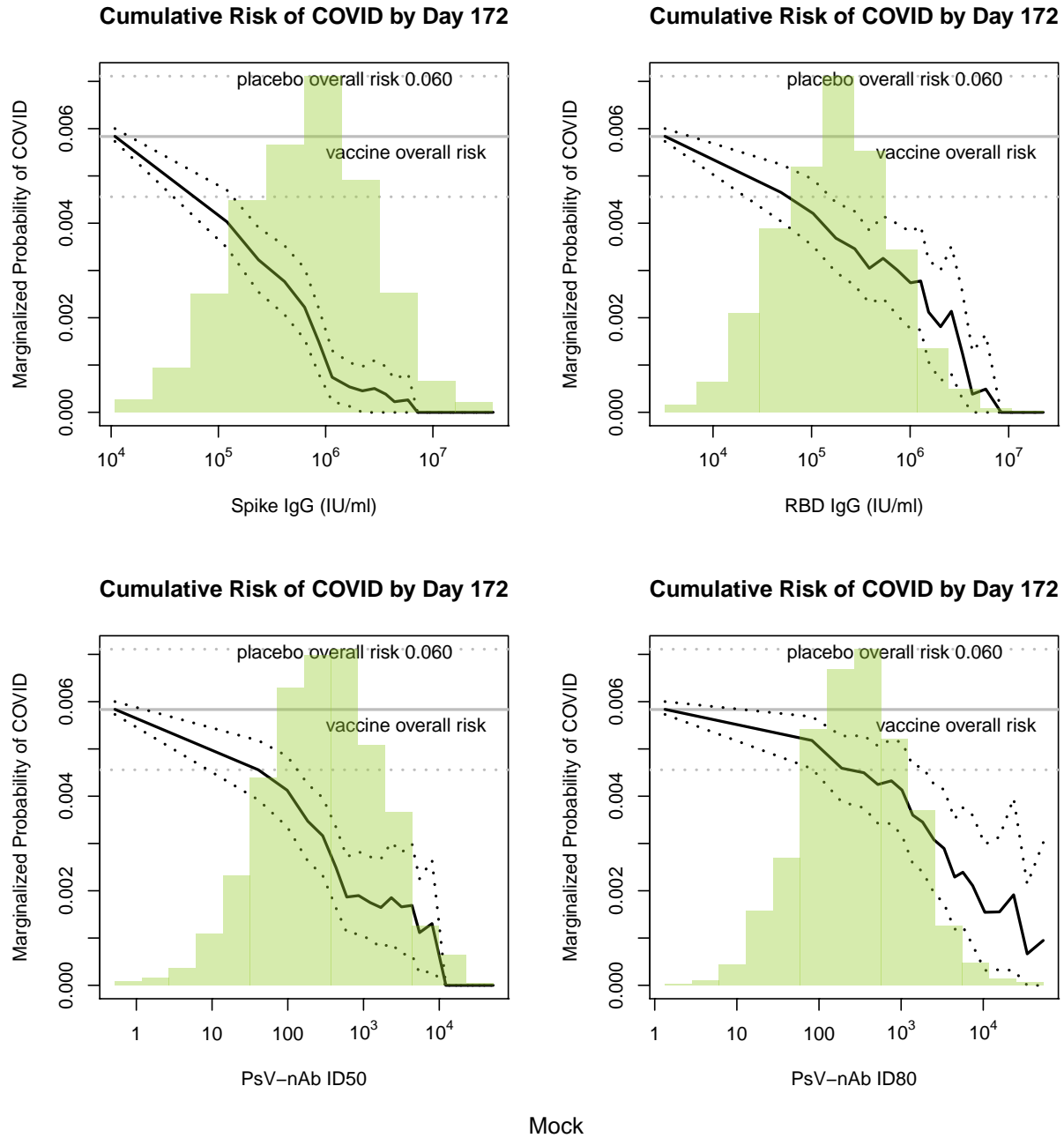


Figure 3: Marginalized cumulative risk by Day 172 as functions of Day 57 markers above a threshold among baseline seronegative vaccine recipients with 95% bootstrap point-wise confidence bands. The horizontal lines indicate the overall cumulative risk of the vaccine arm by Day 172 and its 95% point-wise confidence interval. Histograms of the immunological markers in the vaccine arm are overlaid.

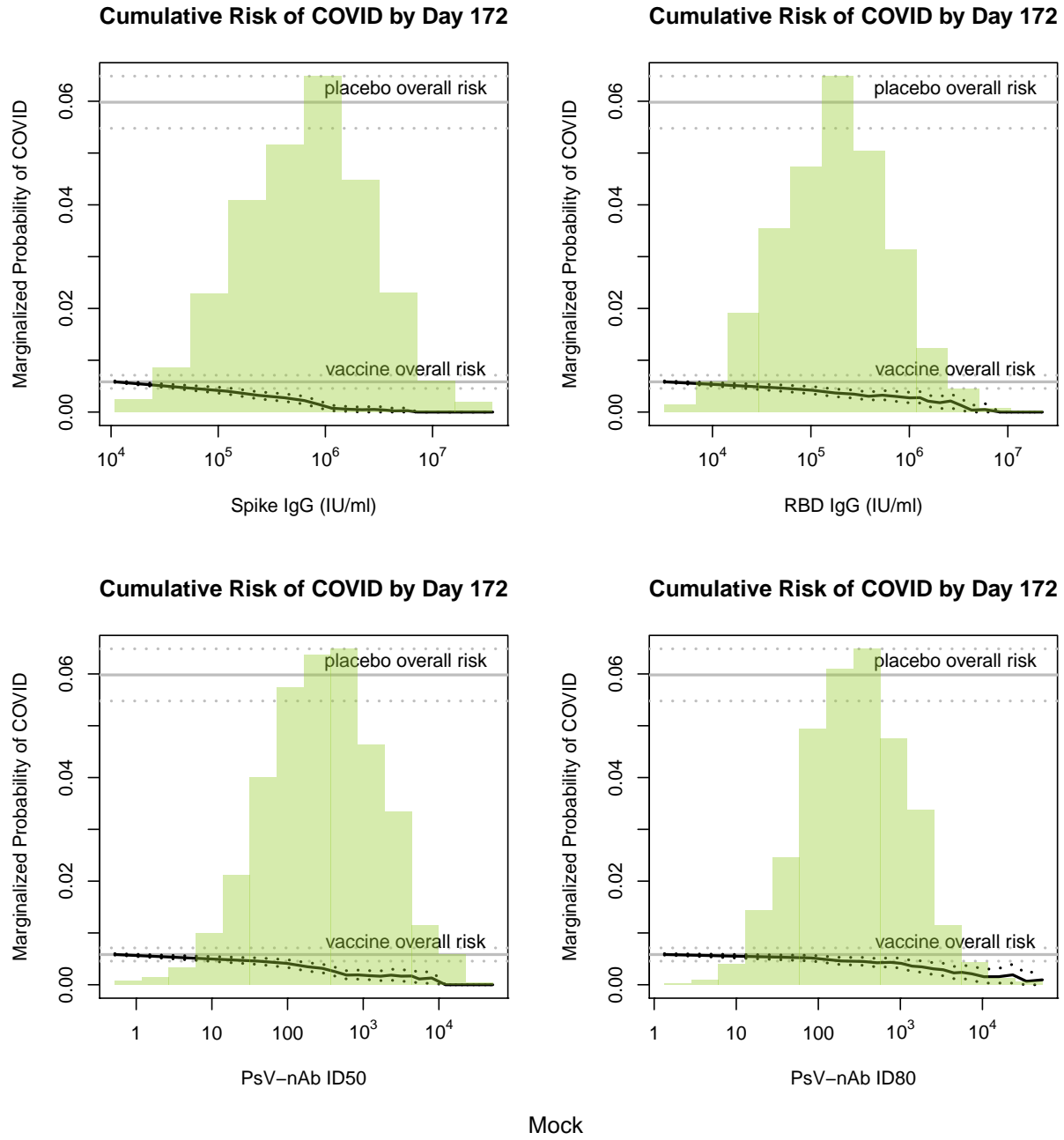


Figure 4: Marginalized cumulative risk by Day 172 as functions of Day 57 markers above a threshold among baseline seronegative vaccine recipients with 95% bootstrap point-wise confidence bands. The horizontal lines indicate the overall cumulative risk of the placebo and vaccine arms by Day 172 and its 95% point-wise confidence interval. Histograms of the immunological markers in the vaccine arm are overlaid.

### Controlled Vaccine Efficacy against COVID by Antibody Titer

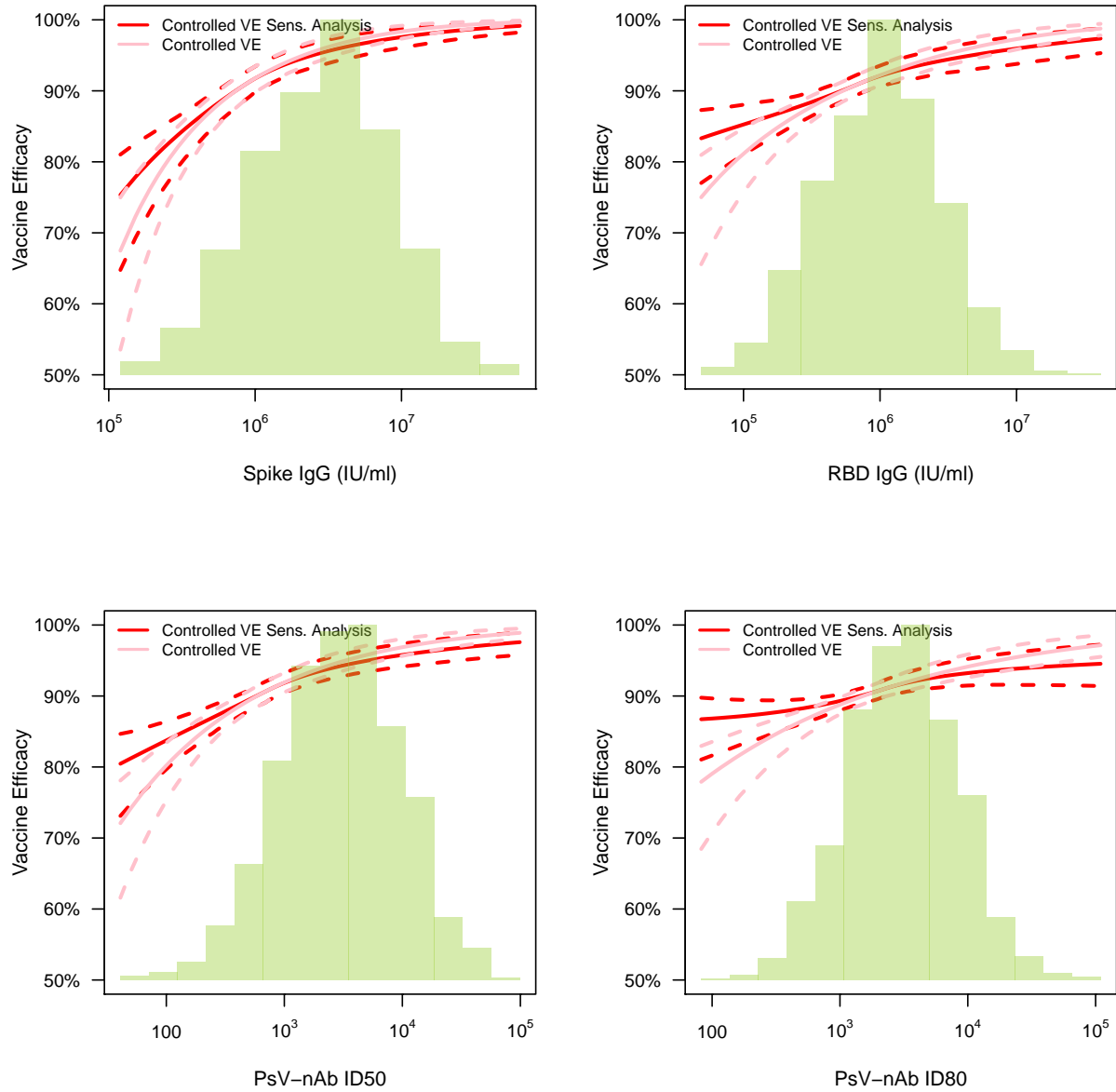


Figure 5: Controlled VE with sensitivity analysis as functions of Day 57 markers among baseline seronegative vaccine recipients with 95% bootstrap point-wise confidence bands. Histograms of the immunological markers in the vaccine arm are overlaid.

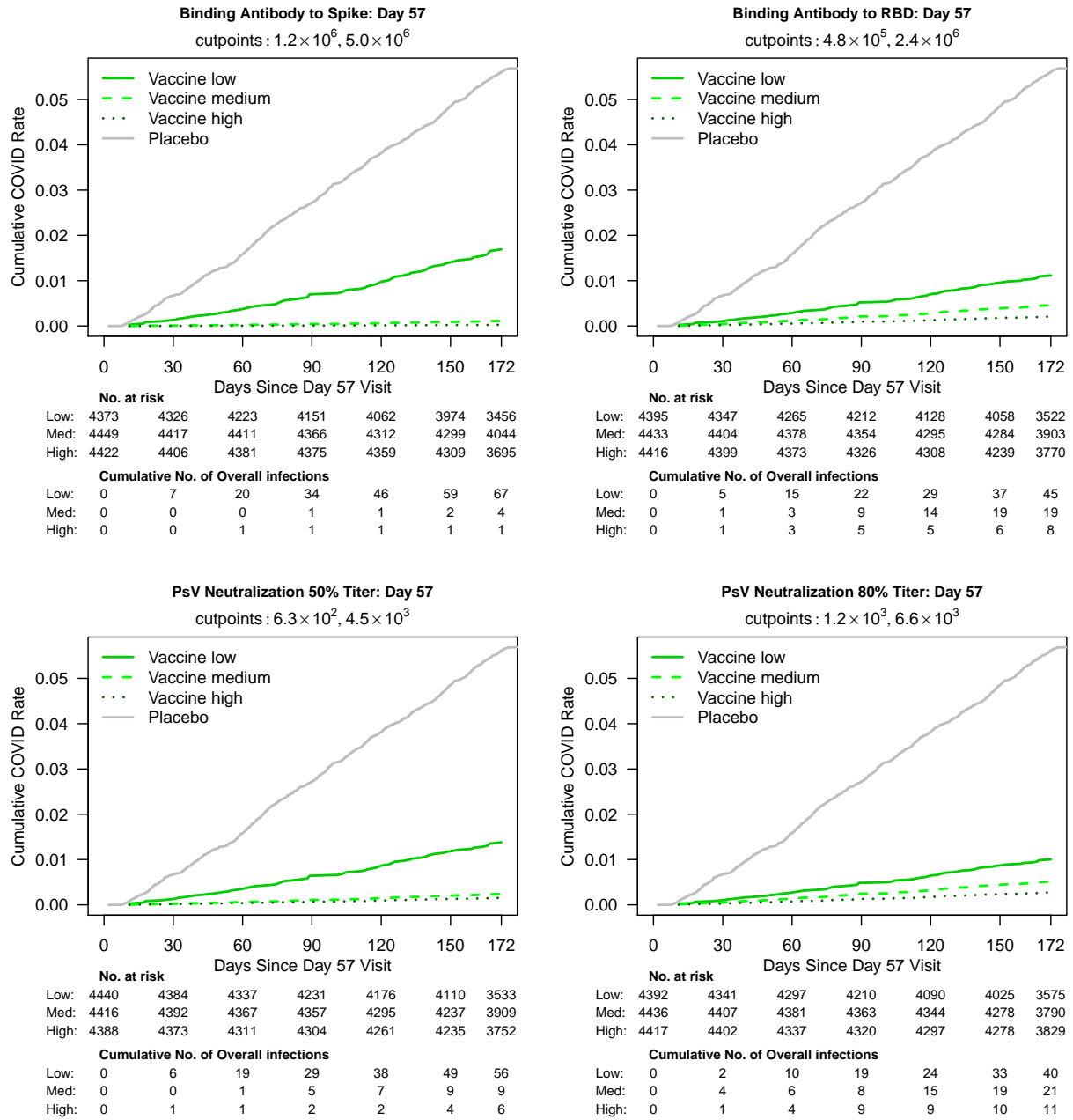


Figure 6: Marginalized cumulative incidence rate curves for trichotomized Day 57 markers among baseline seronegative vaccine recipients. The gray line is the overall cumulative incidence rate curve in the placebo arm.



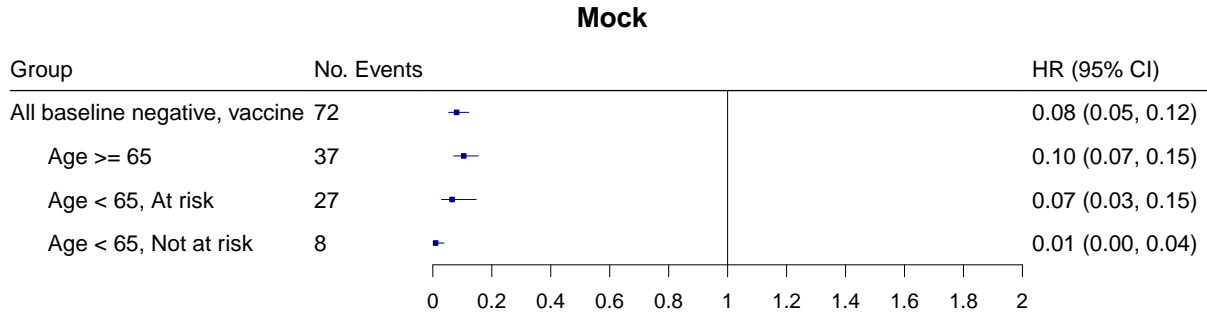


Figure 7: Forest plots of hazard ratios of Day 57 binding Ab to spike markers among baseline seronegative vaccine recipients (top row) and each of the 3 randomization strata with 95% point-wise confidence intervals.

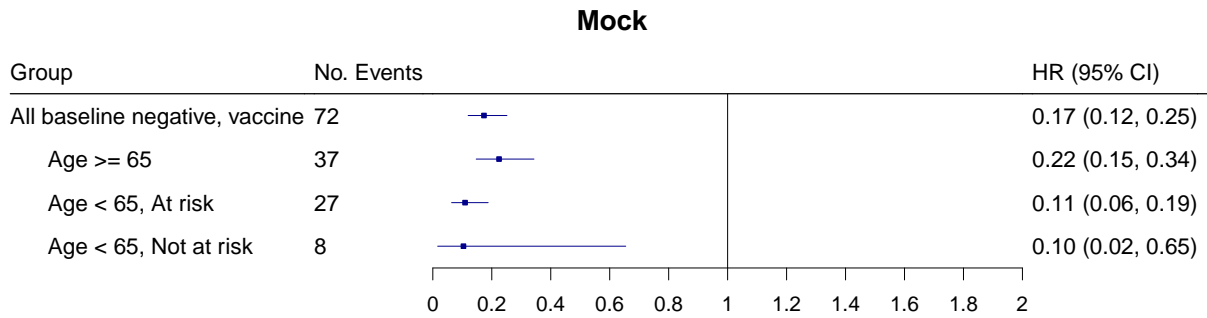


Figure 8: Forest plots of hazard ratios of Day 57 binding Ab to RBD markers among baseline seronegative vaccine recipients (top row) and each of the 3 randomization strata with 95% point-wise confidence intervals.

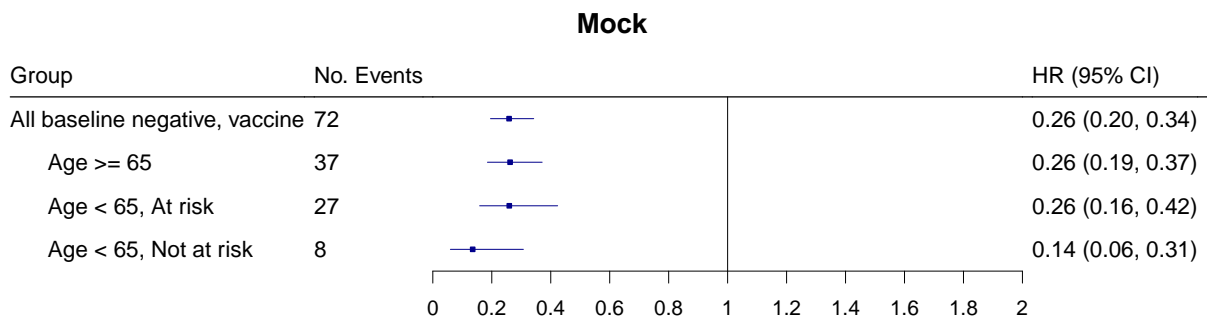


Figure 9: Forest plots of hazard ratios of Day 57 pseudo neut ID50 markers among baseline seronegative vaccine recipients (top row) and each of the 3 randomization strata with 95% point-wise confidence intervals.

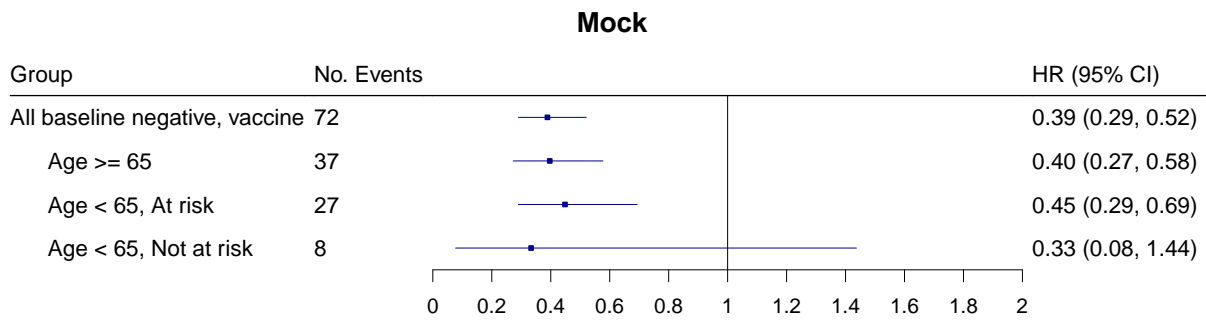


Figure 10: Forest plots of hazard ratios of Day 57 pseudo neut ID80 markers among baseline seronegative vaccine recipients (top row) and each of the 3 randomization strata with 95% point-wise confidence intervals.

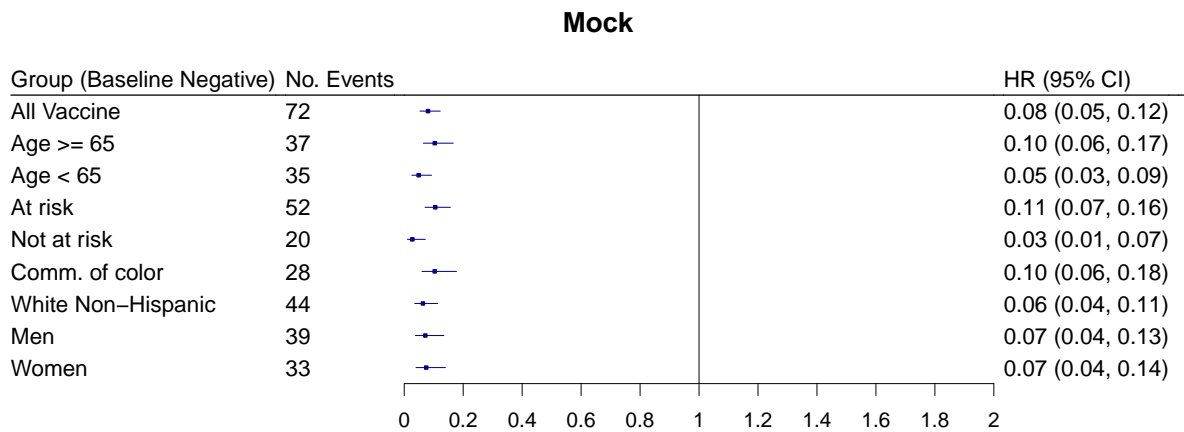


Figure 11: Forest plots of hazard ratios of Day 57 binding Ab to spike markers among baseline seronegative vaccine recipients (top row) and eight subpopulations (row 2-9) with 95% point-wise confidence intervals.

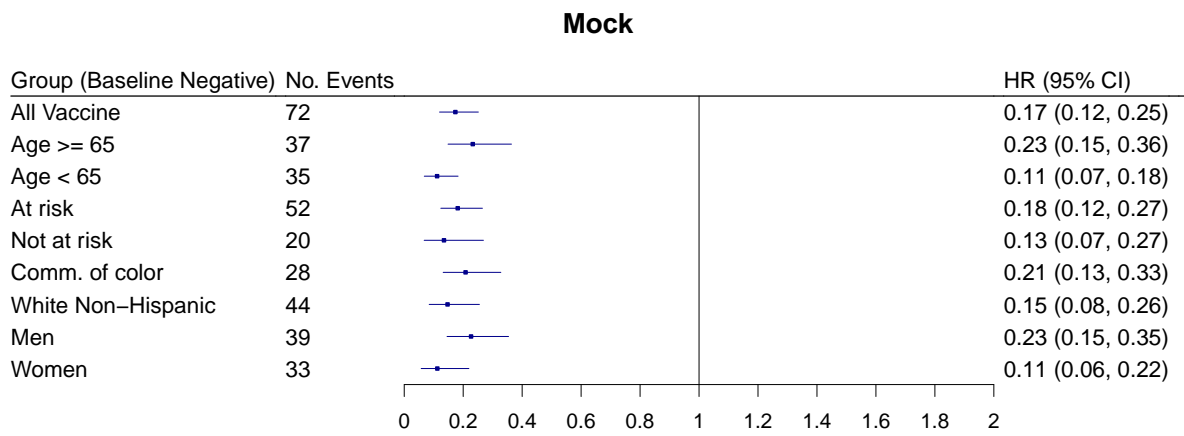


Figure 12: Forest plots of hazard ratios of Day 57 binding Ab to RBD markers among baseline seronegative vaccine recipients (top row) and eight subpopulations (row 2-9) with 95% point-wise confidence intervals.

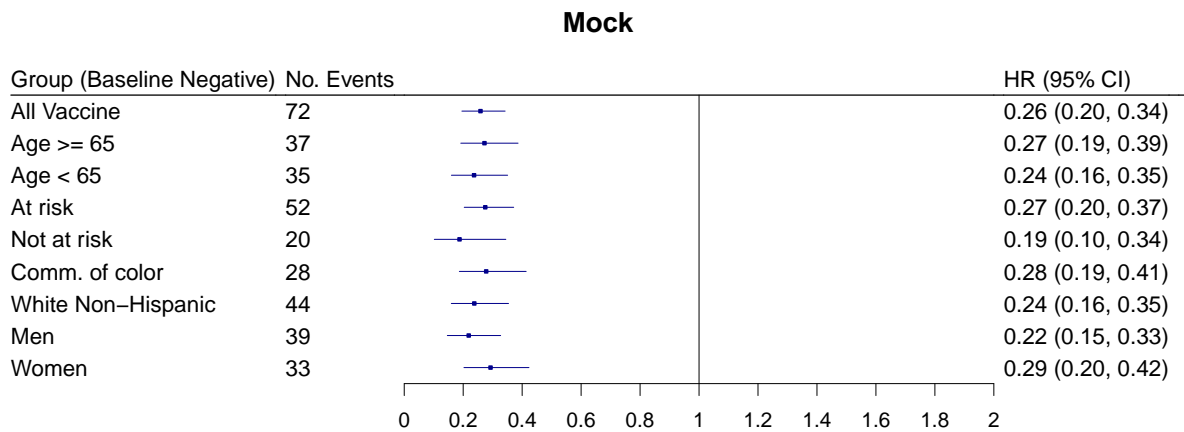


Figure 13: Forest plots of hazard ratios of Day 57 pseudo neut ID50 markers among baseline seronegative vaccine recipients (top row) and eight subpopulations (row 2-9) with 95% point-wise confidence intervals.

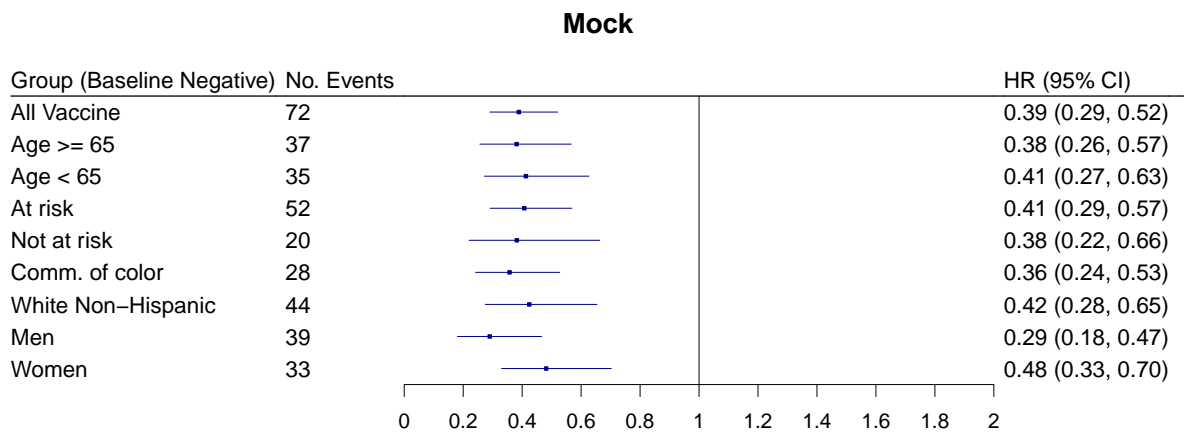


Figure 14: Forest plots of hazard ratios of Day 57 pseudo neut ID80 markers among baseline seronegative vaccine recipients (top row) and eight subpopulations (row 2-9) with 95% point-wise confidence intervals.