

Figure 1. Alpha diversity metrics (Observed richness, Chao1 estimator and Shannon diversity index) between study groups.

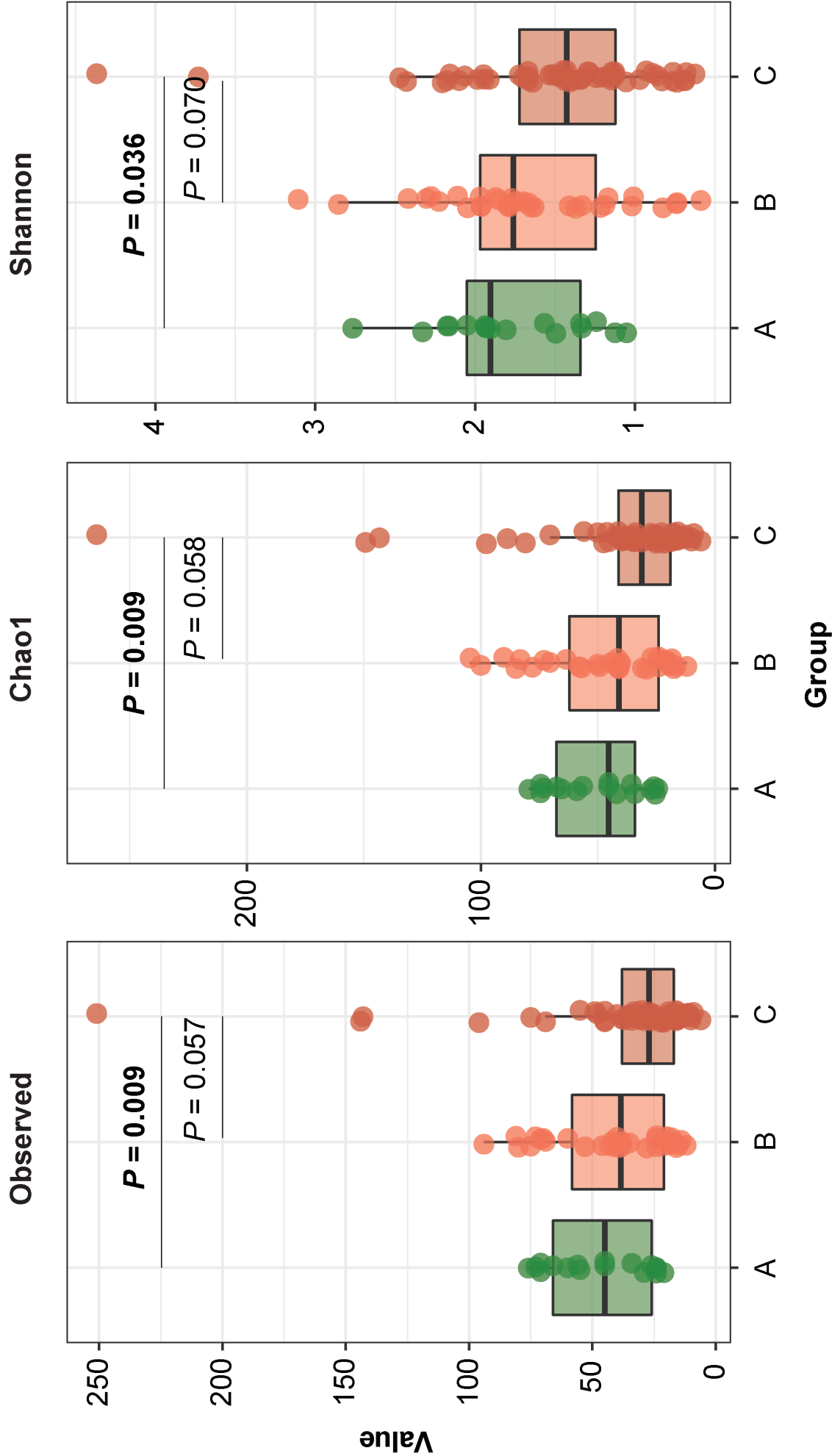


Figure 2. Overall nasopharyngeal bacterial genera composition between study groups.

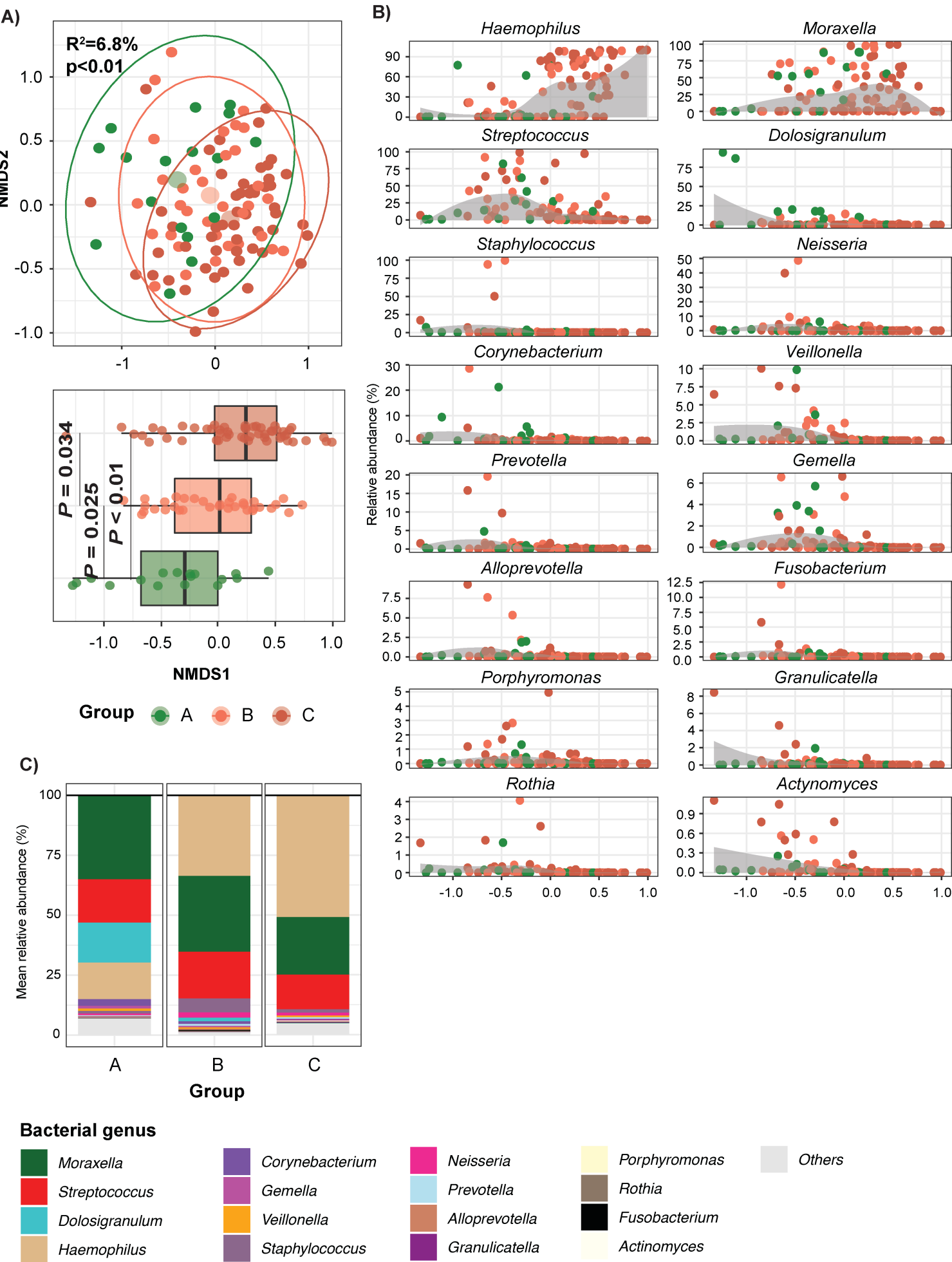
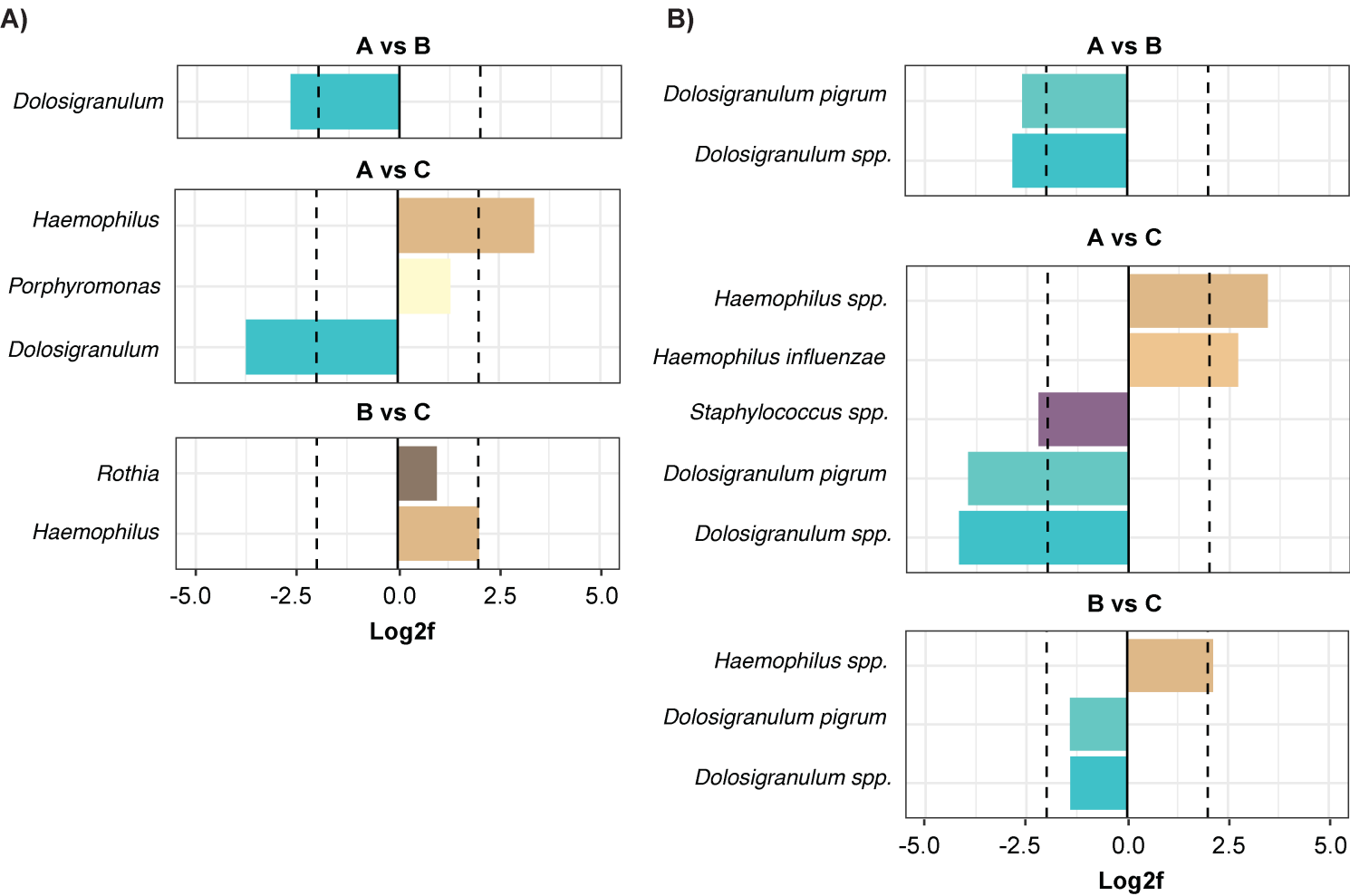


Figure 3. Differentially abundant bacterial taxa by RV/EV infection status and severity.



e-Figure1. Bacterial genera microbiota profiling by subject.

