Figure S18. Region plot for the top novel locus identified through the G x Sex interaction test compared with other results. Querying the chr7:107127037 region in G x Sex GWAS (top row) shows that this variant is in LD with nearby variants. Comparing it with the following in order: primary GWAS, sex-stratified GWAS for males, sex-stratified GWAS for females, and HGI7no shows that there is a sex-effect for this locus in HostSeq, where males \( N = 3,646 \) have an association with hospitalization. Plots were generated using myLocusZoom.