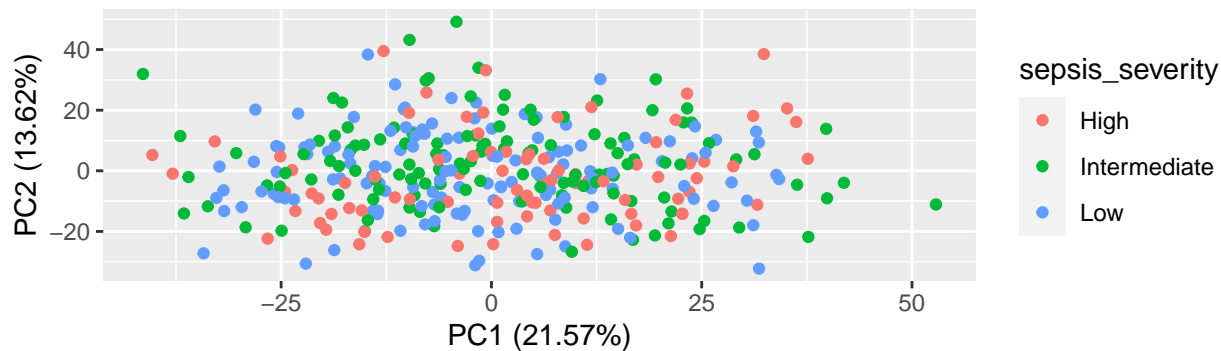


PCA on multiple vars on VST normalized and batch corrected mito-gene counts

PCA regarding sepsis_severity



PCA regarding sample_location

