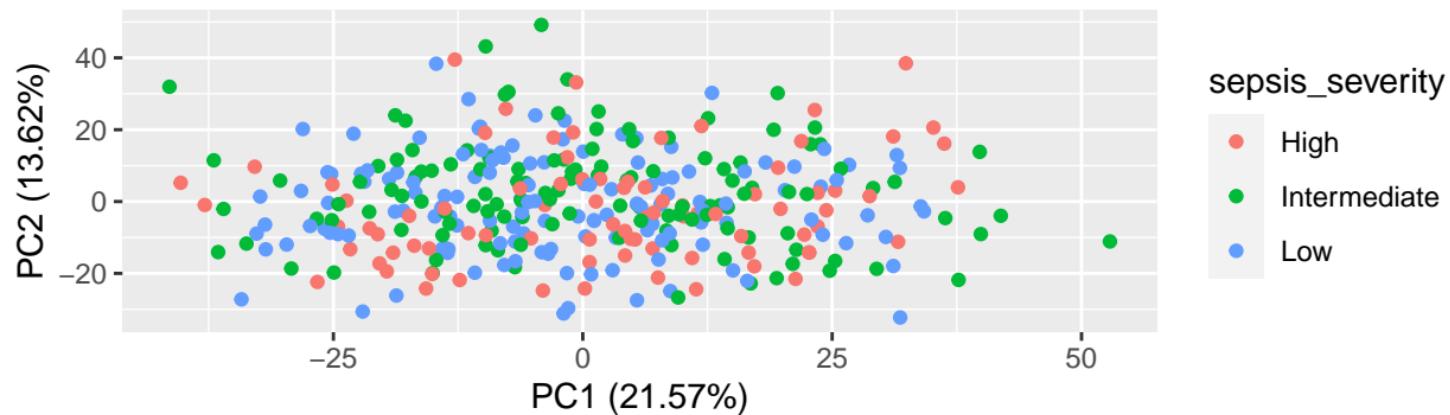


PCA on multiple vars on VST normalized and batch corrected mito–gene counts

PCA regarding sepsis_severity



PCA regarding sample_location

