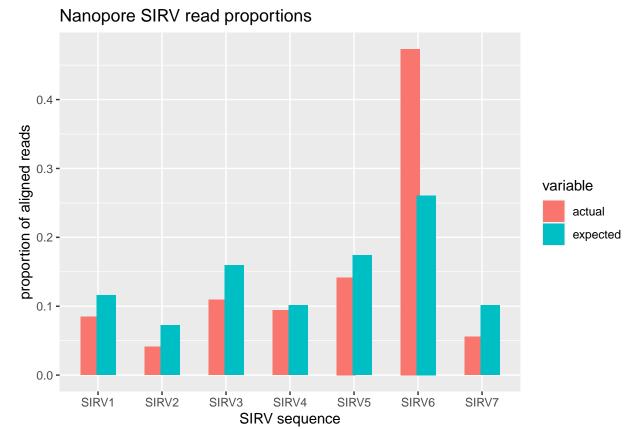
SIRV analysis

Quinn 7/9/2020

this markdown takes in two files of sam RNAMEs, along with relevant abundance information from stringtie output $\frac{1}{2}$

plotting expected vs actual nanopore SIRV read proportions

Warning: Ignoring unknown parameters: stat



plotting expected vs actual illumina SIRV read proportions

Warning: Ignoring unknown parameters: stat

Illumina SIRV read proportions

