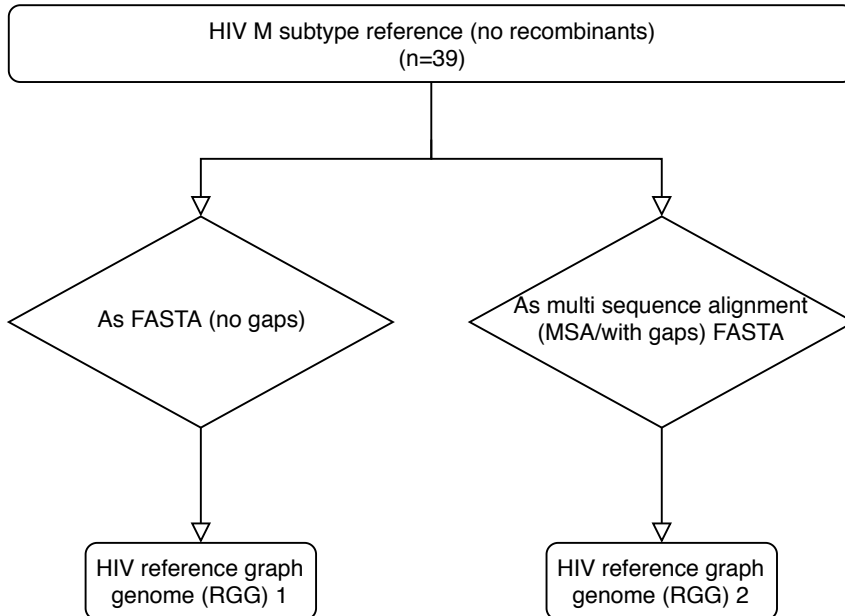## Goal 1 - Make

graph genomes!

Using SWIGG, VG, NovoGraph and
HIV-1 references from LANL HIV

Goal 2 - Use

graph genomes!

Testing performance of single linear reference,
multiple linear references, multiple splice
isoforms as linear references, and graph
genomes for HIV-read capture (with HISAT2).