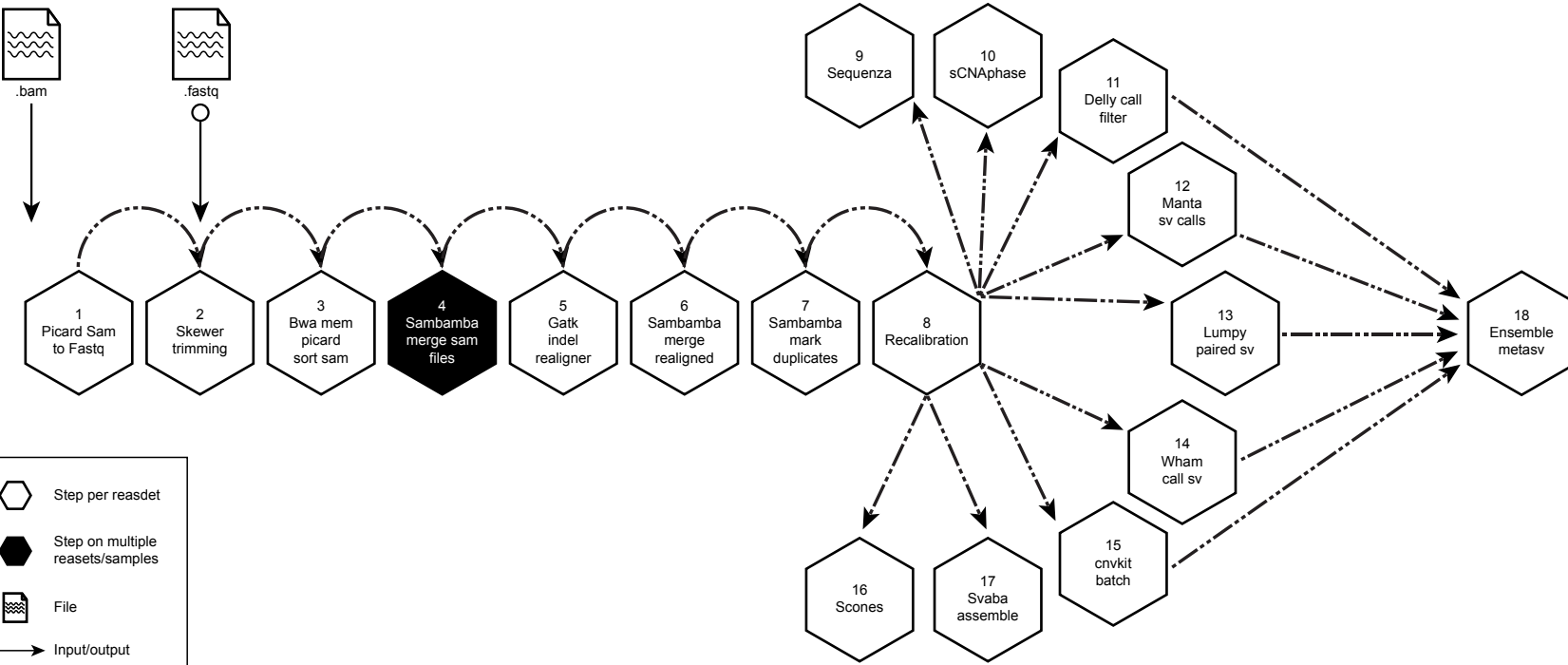


tumor_pair.py -t sv



Step per readset

Step on multiple readsets/samples

File

Input/output

Alternative entry

Step dependency