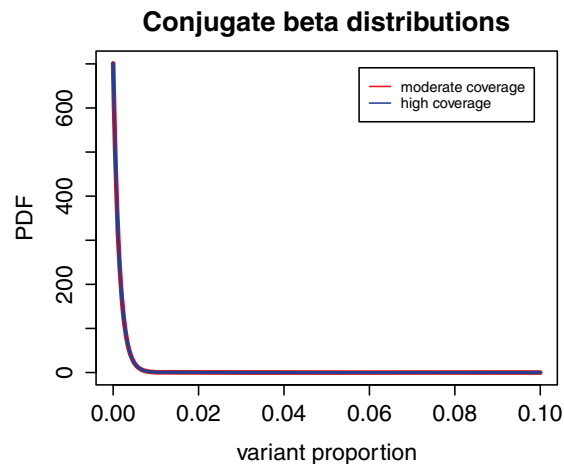
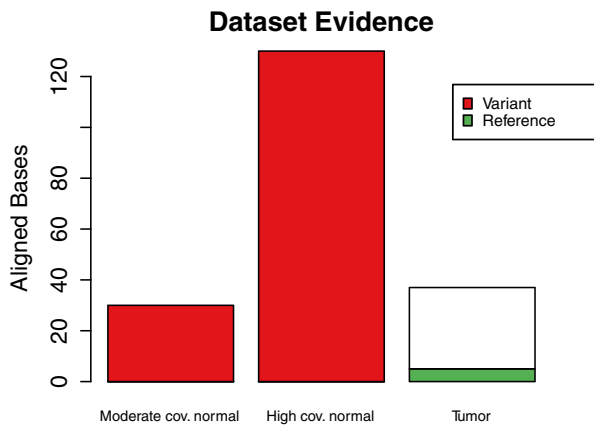


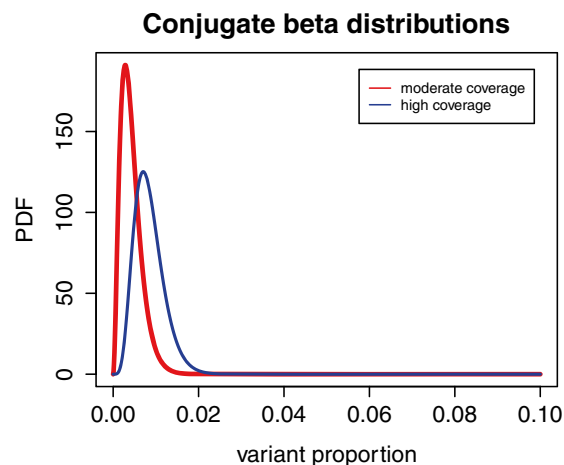
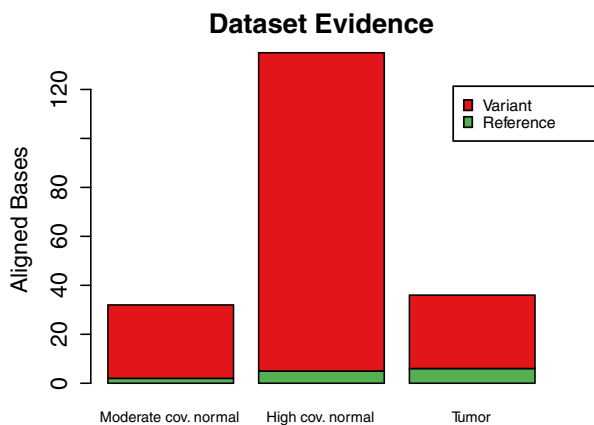
A True Positive (low allelic frequency)

$$Q_{med} = 11.70 \rightarrow Q_{high} = 14.21$$



B False Positive (sequencing/mapping artifact)

$$Q_{med} = 7.02 \rightarrow Q_{high} = 2.02$$



C False Positive (germline variant called somatic)

$$Q_{med} = 1.03 \rightarrow Q_{high} = 1 \times 10^{-7}$$

