a

Somatic Analysis - heuristically prioritize features if in...



Decrease priority if common in ExAC



Increase priority if observed in orthogonal sequencing or gene mutated in germline

b

Germline Analysis - annotate with ExAC and ClinVar and report...

Incidental findings from ACMG v2

→ Variants in genes related to hereditary cancers

Variants in genes related to somatic cancers