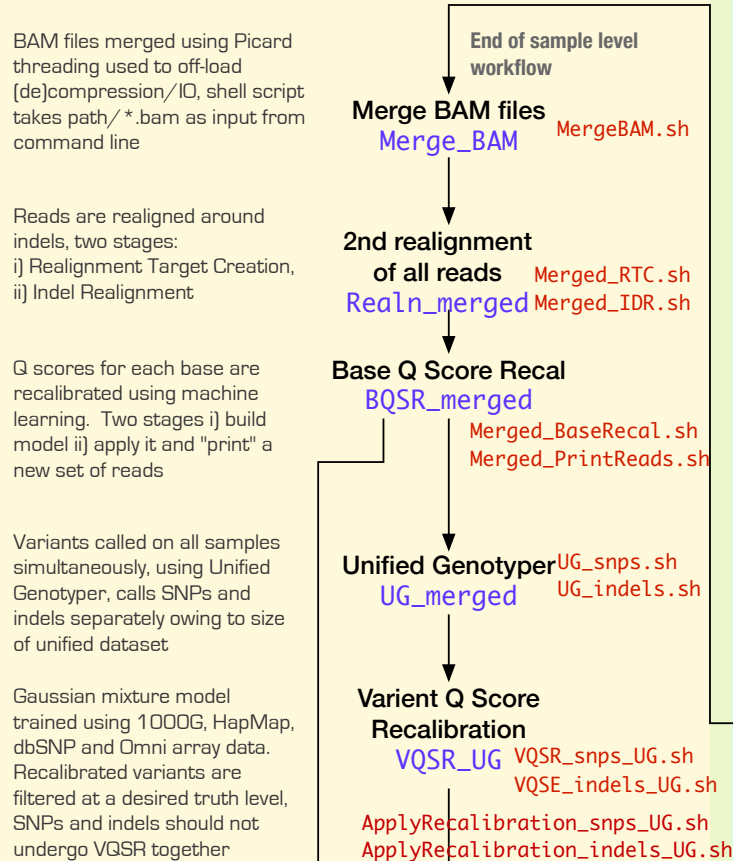


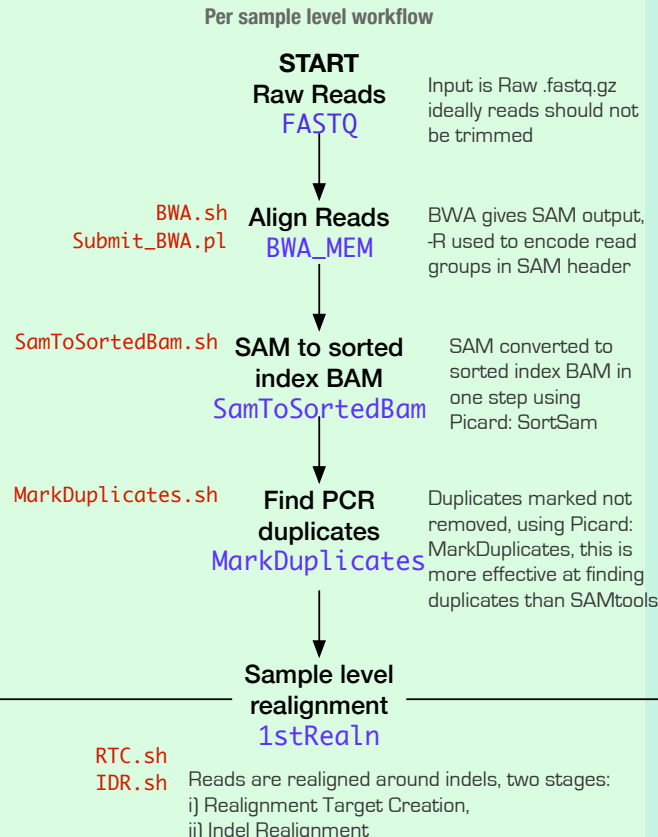
MB-GATK-SGE pipeline

For GATK best practices: classic UG / v3.x HC / MuTect
Matthew Bashton

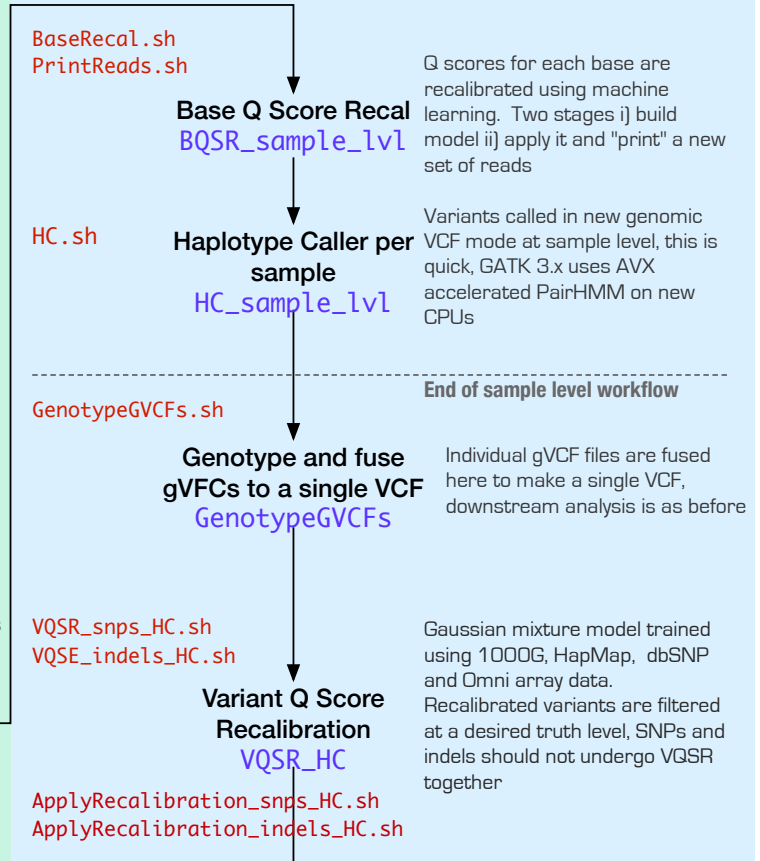
Classic Unified Genotyper workflow



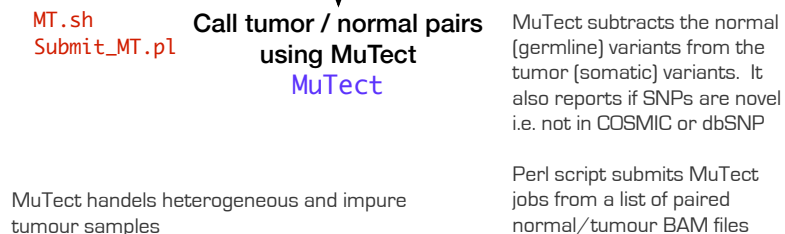
Common per-sample processing



New Haplotype Caller workflow



MuTect 1.x somatic variant calling



Recalibrated variant filtering

