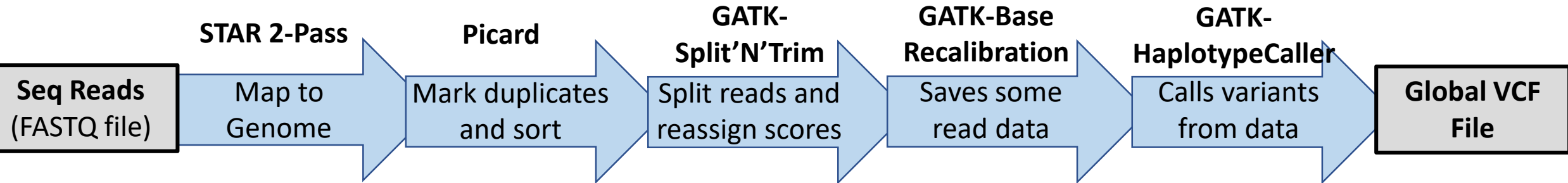
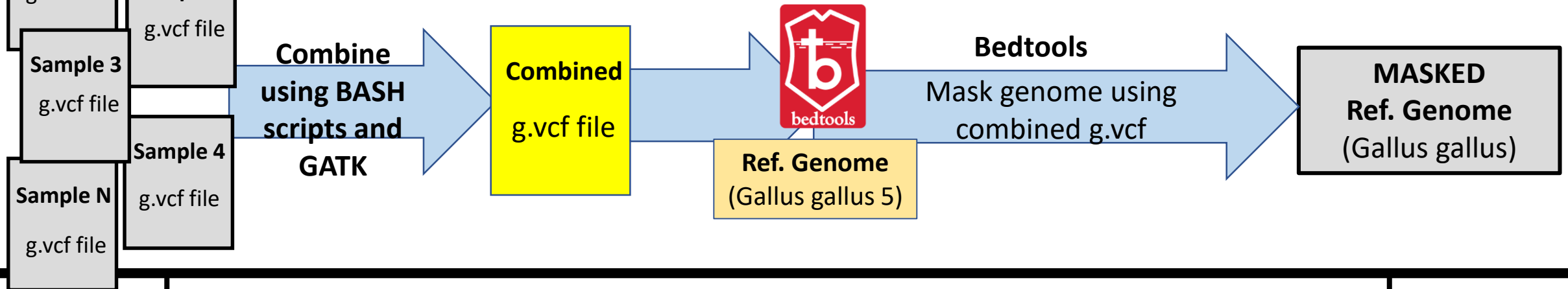


Part 1. Initial Calling of Variants Using GATK



Part 2. Merge All VCFs and Create a Masked Genome



Part 3. Call Variants Using Masked Genome to Reduce Ref. Allele Bias

