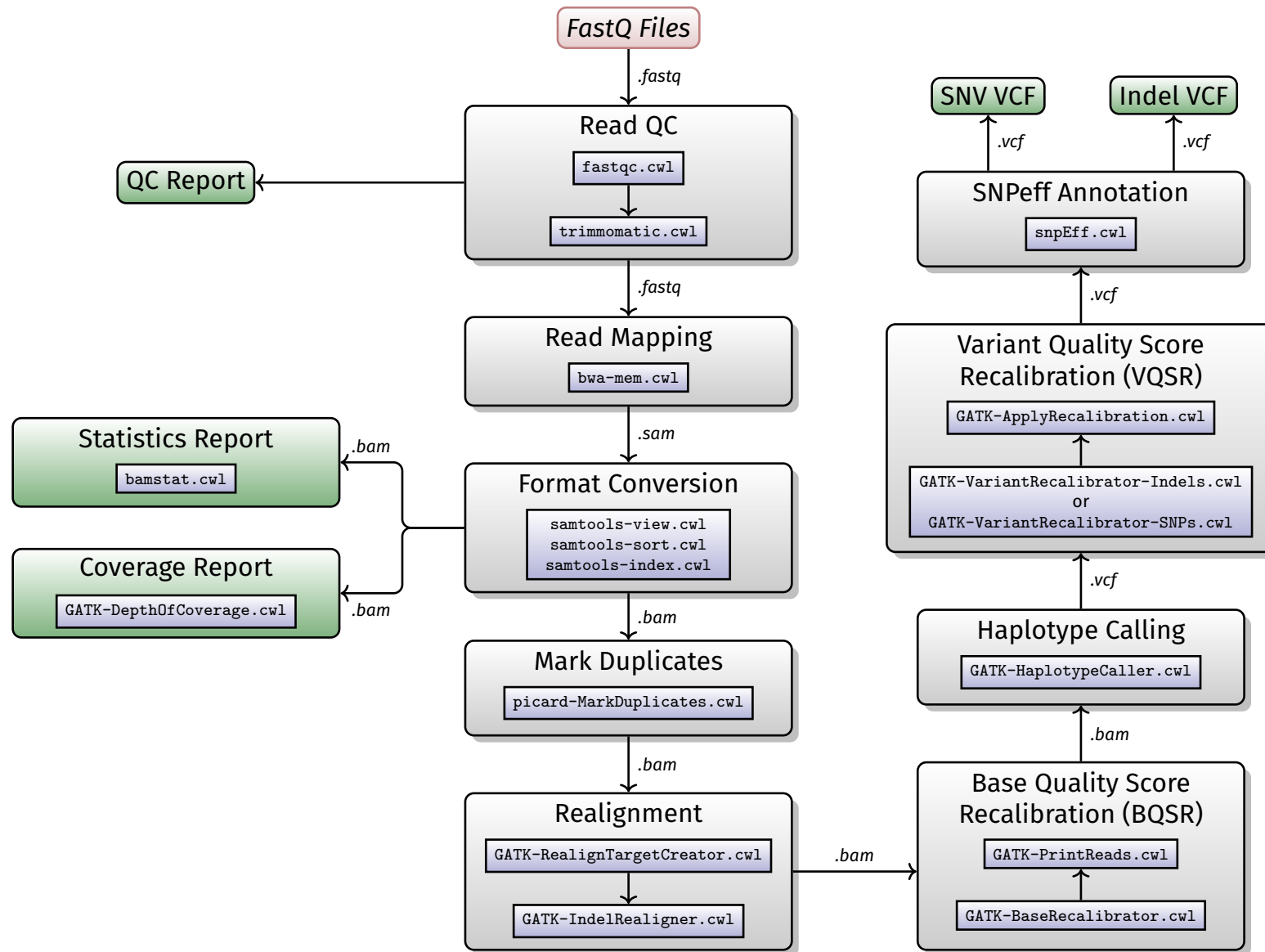
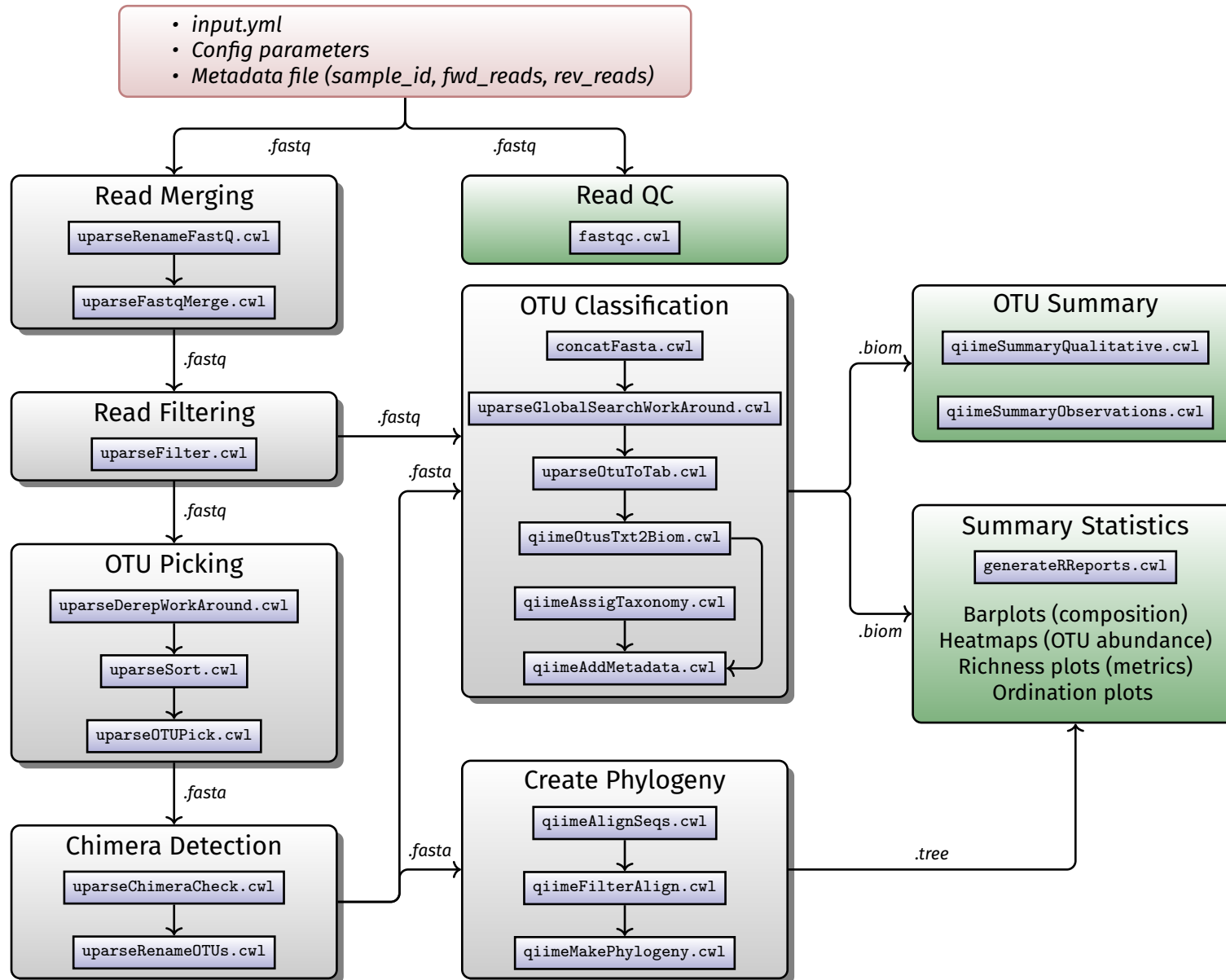


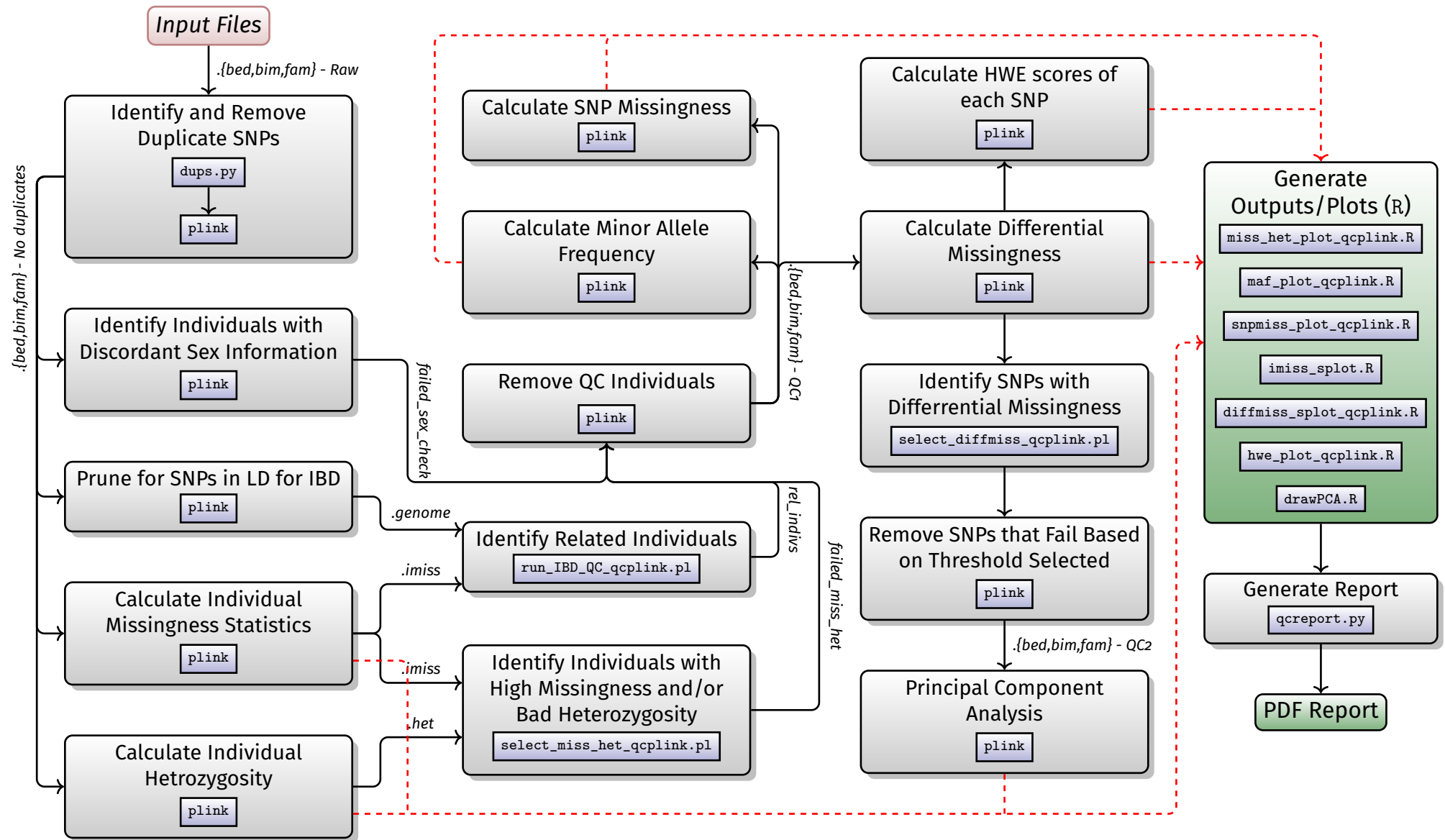
1 Workflow A: Whole Genome/Exome NGS Data Analysis



2 Workflow B: 16S rRNA Diversity Analysis



3 Workflow C: Genome Wide-association studies



4 Workflow D: SNPs Imputation

