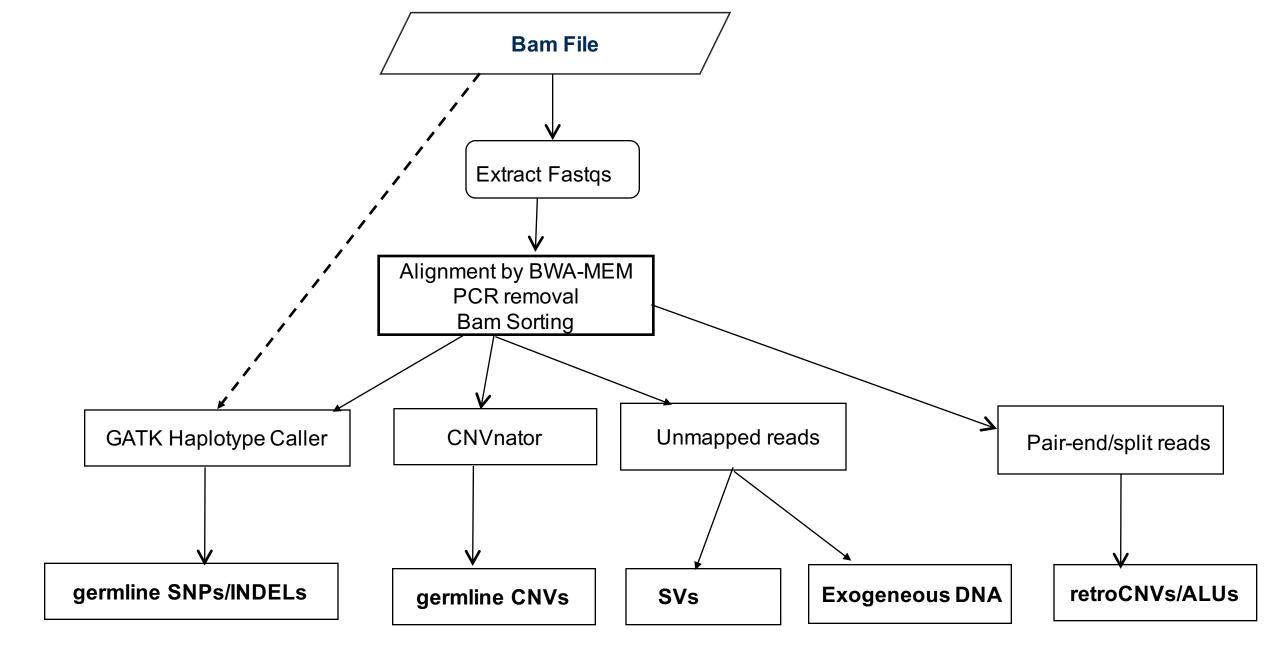
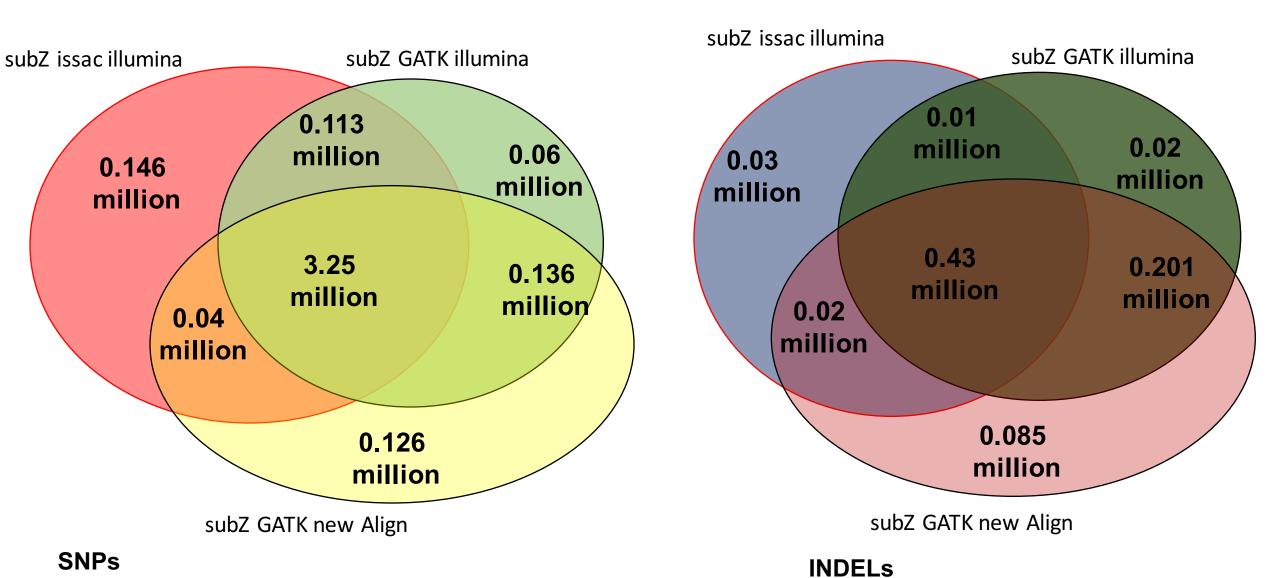
Personal Genome Analysis

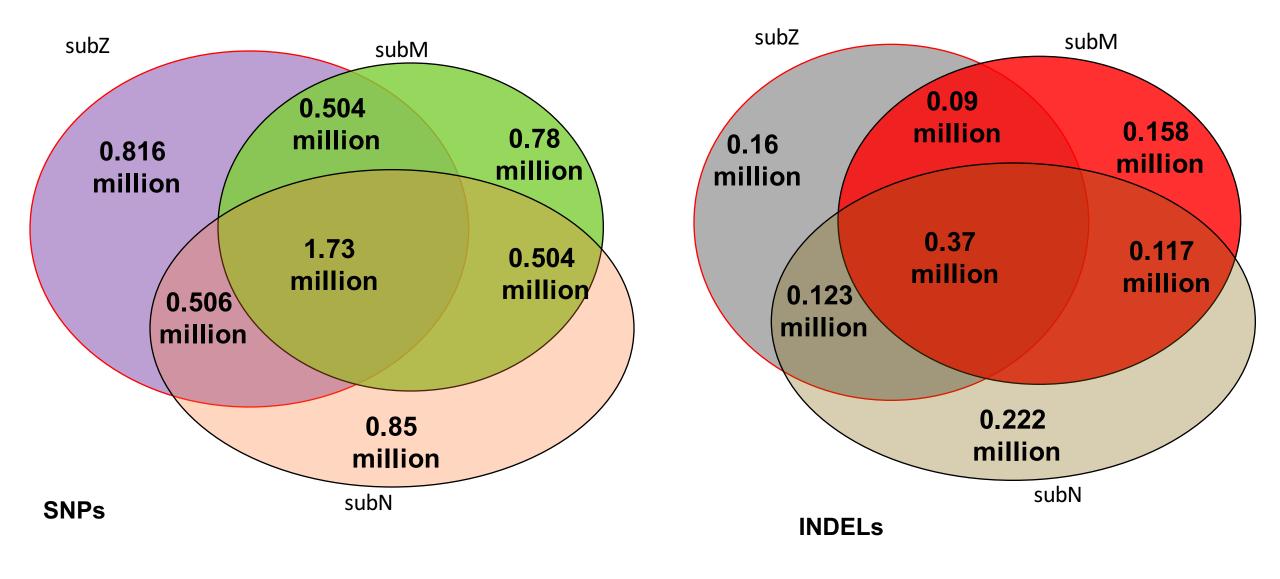


Pipeline to generate variants for personal genome analysis

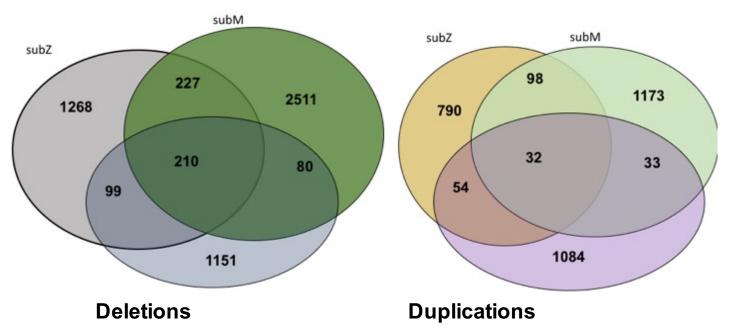
Comparison of variant calls for subject Z



Comparison of SNVs & INDELs across three genomes

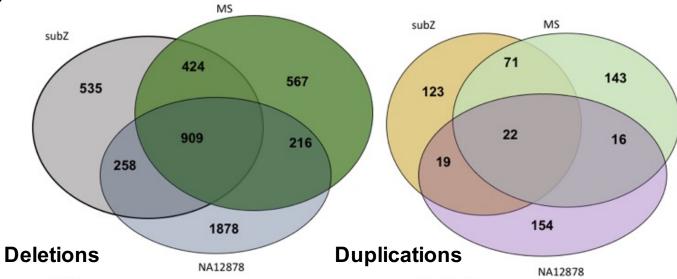


Challenges associated with identifying structural variations



CNVnator assigned large deletions and duplications

CREST assigned large deletions and duplications



Consensus structural variations across three genomes

