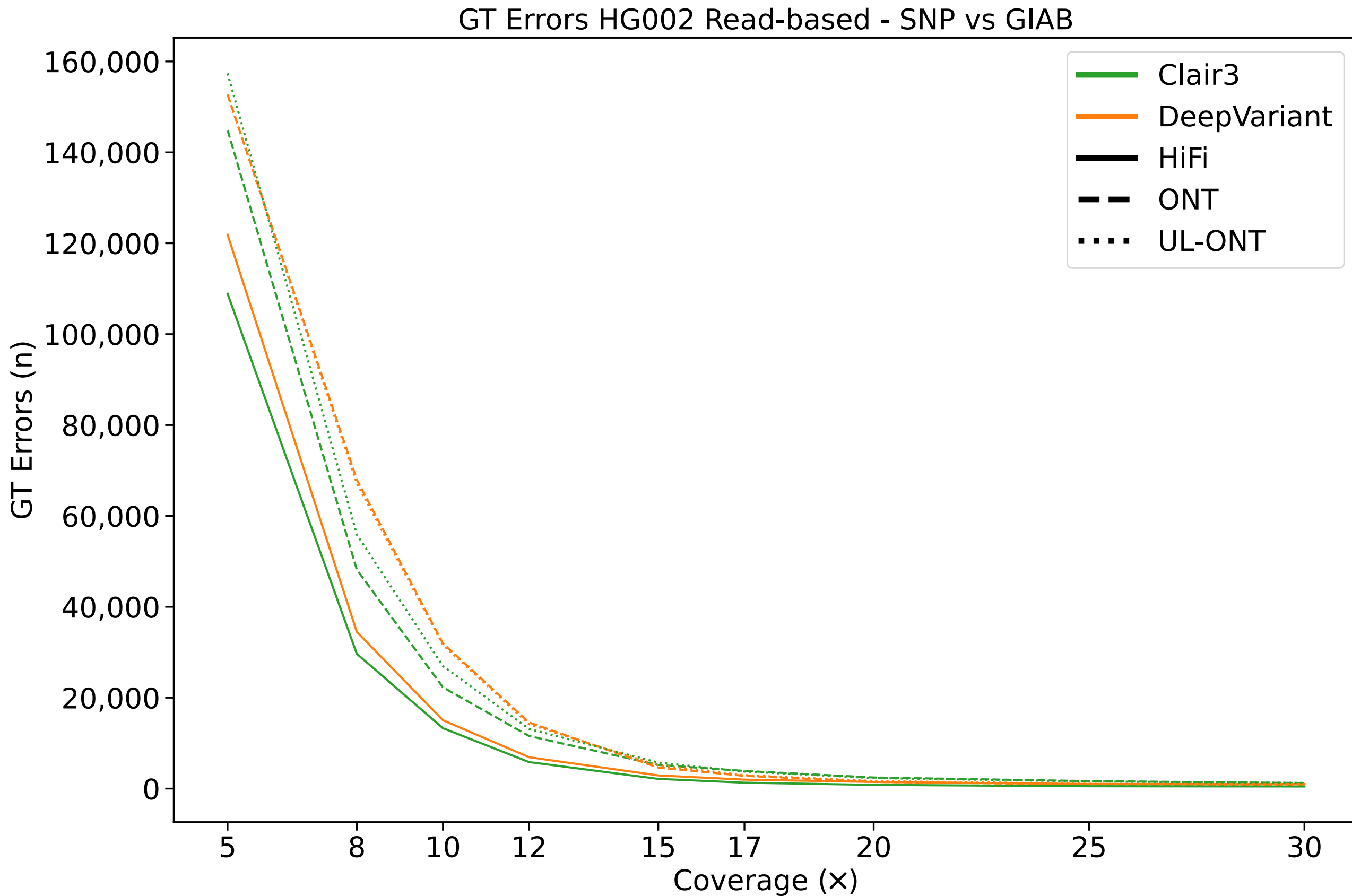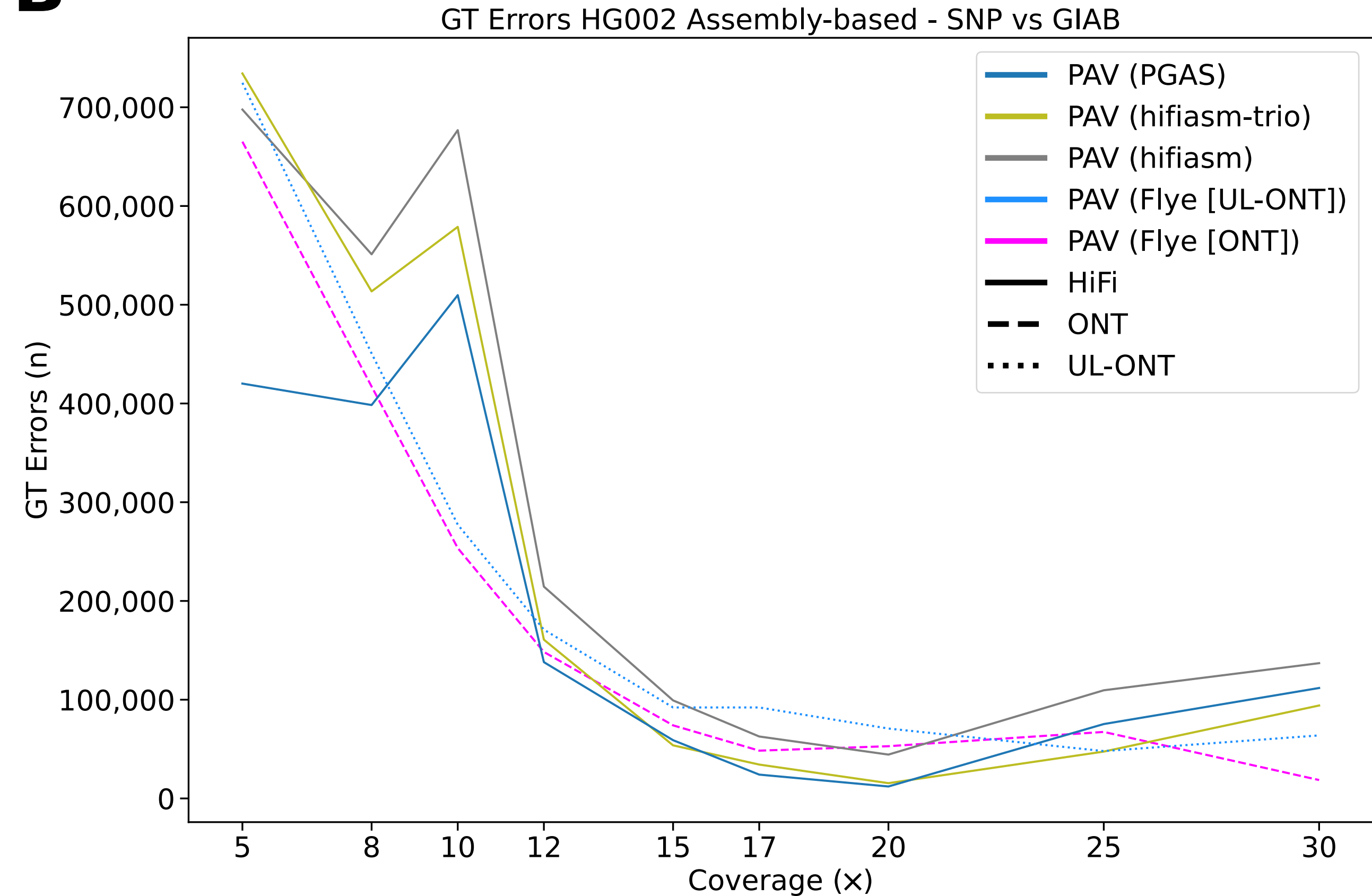**Figure S5. Genotyping errors as a function of coverage in assembly and read-based single-nucleotide polymorphism (SNP) callsets. A)** Genotyping errors in read-based SNP callers show a similar minimum around 12× as caller precision overall saturates. **B)** Assembly-based SNP genotyping error is higher across all coverages compared to read-based methods.