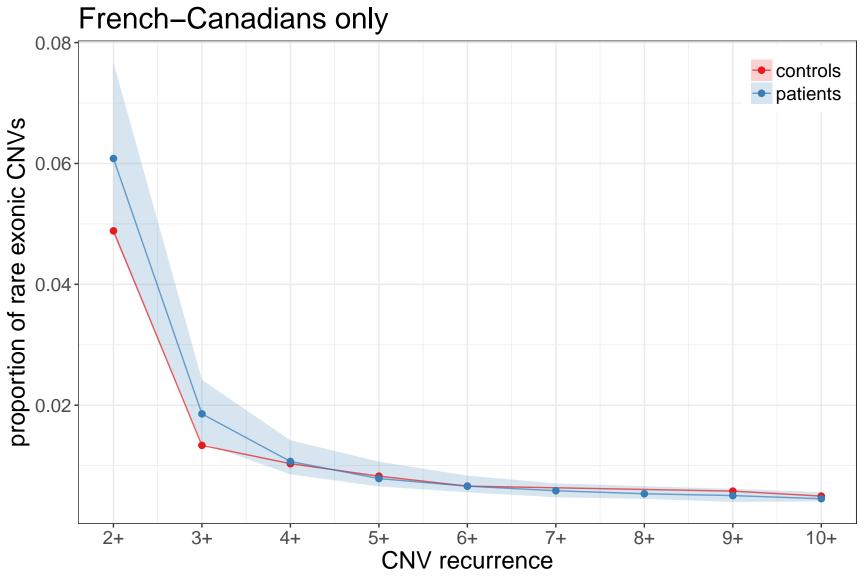
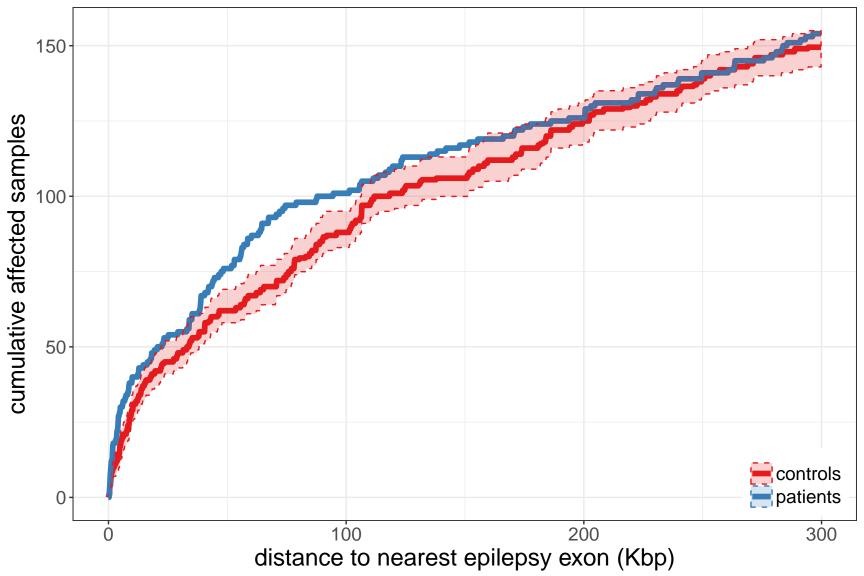
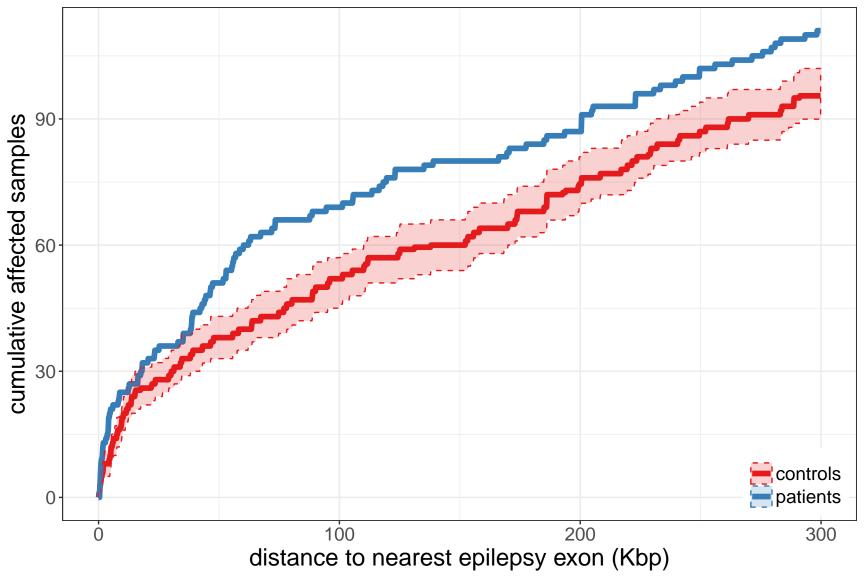
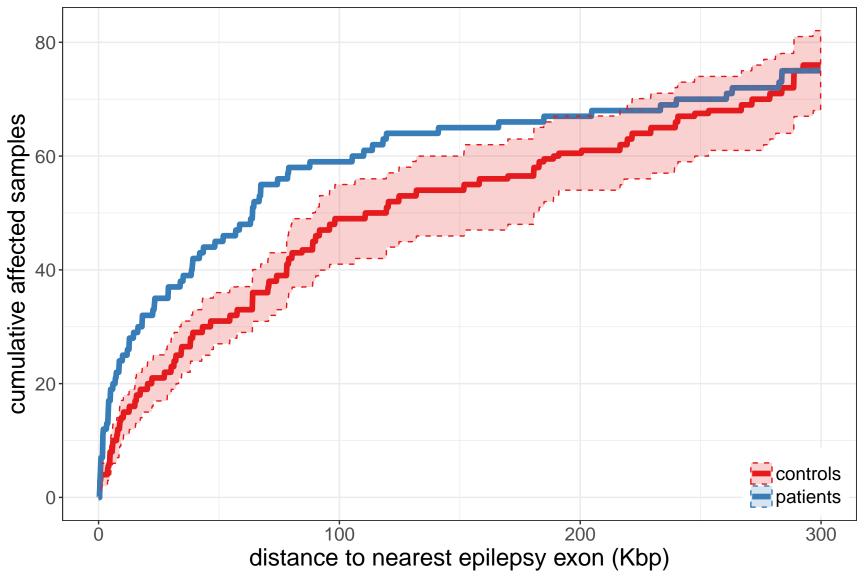


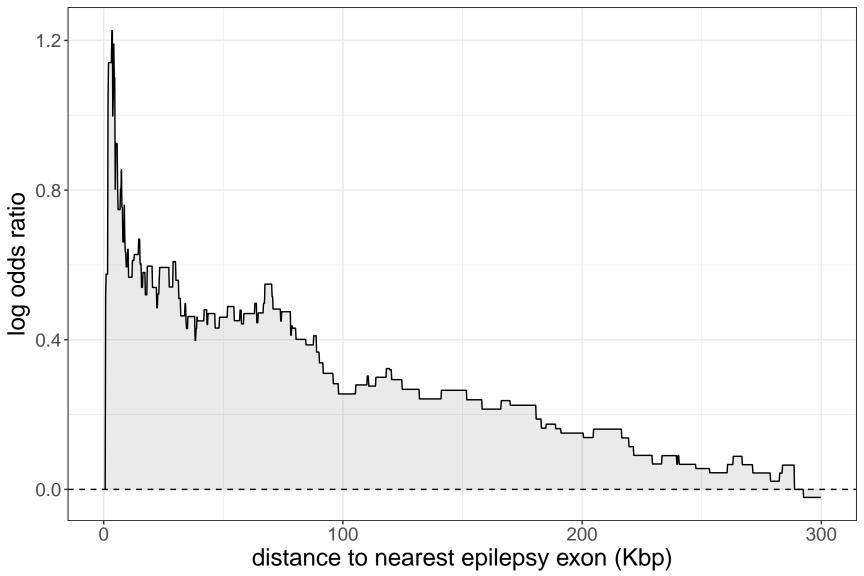
Top 20 most extreme samples removed 0.06 controls patients proportion of rare exonic CNVs 2+ 3+ 4+ 5+ 6+ 7+ 8+ 9+ 10+ **CNV** recurrence

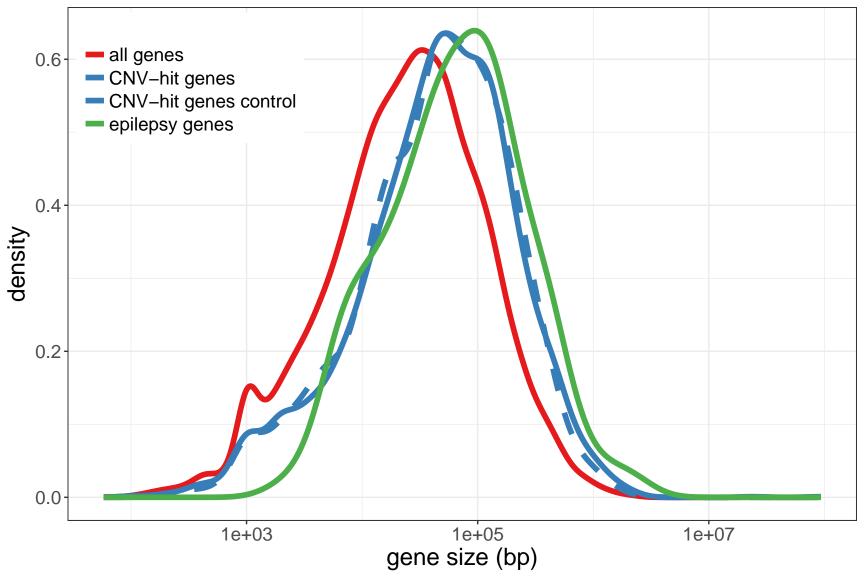












Genes hit deletions never seen in public databases

