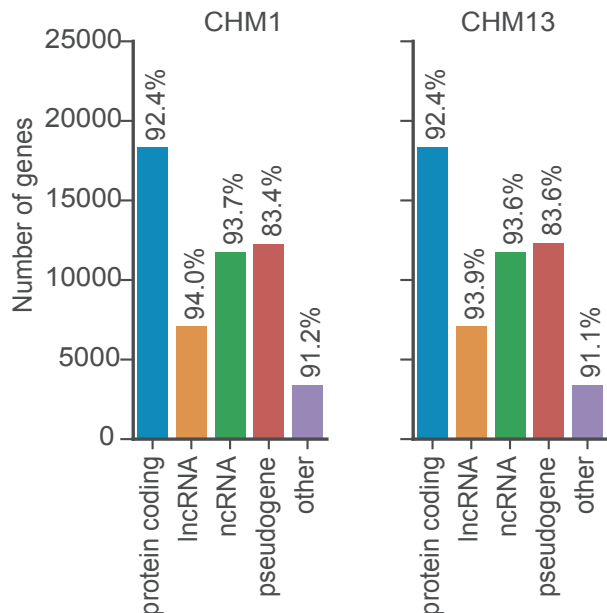
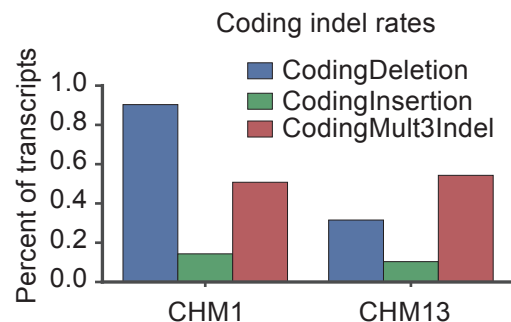


## A) Completeness of comparative annotation



## B) Comparative assembly metrics

	CHM1	CHM13
Number of intra-contig split genes	2	2
Number of contig split genes	39	141



## C) CHM1 specific deletion inactivates TRIB3

hg38 chr20:362,280-400,250 (37,971 bp)

