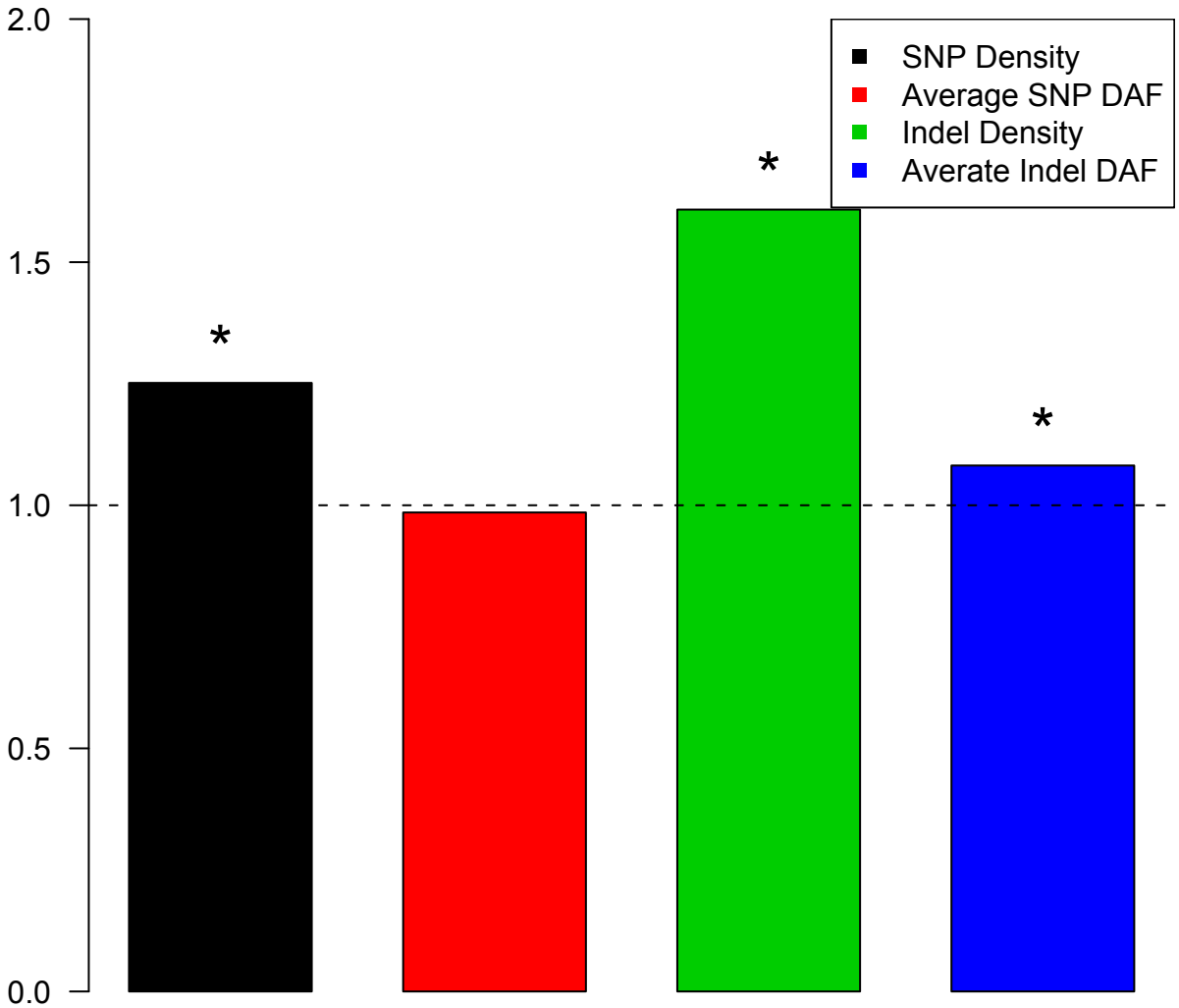
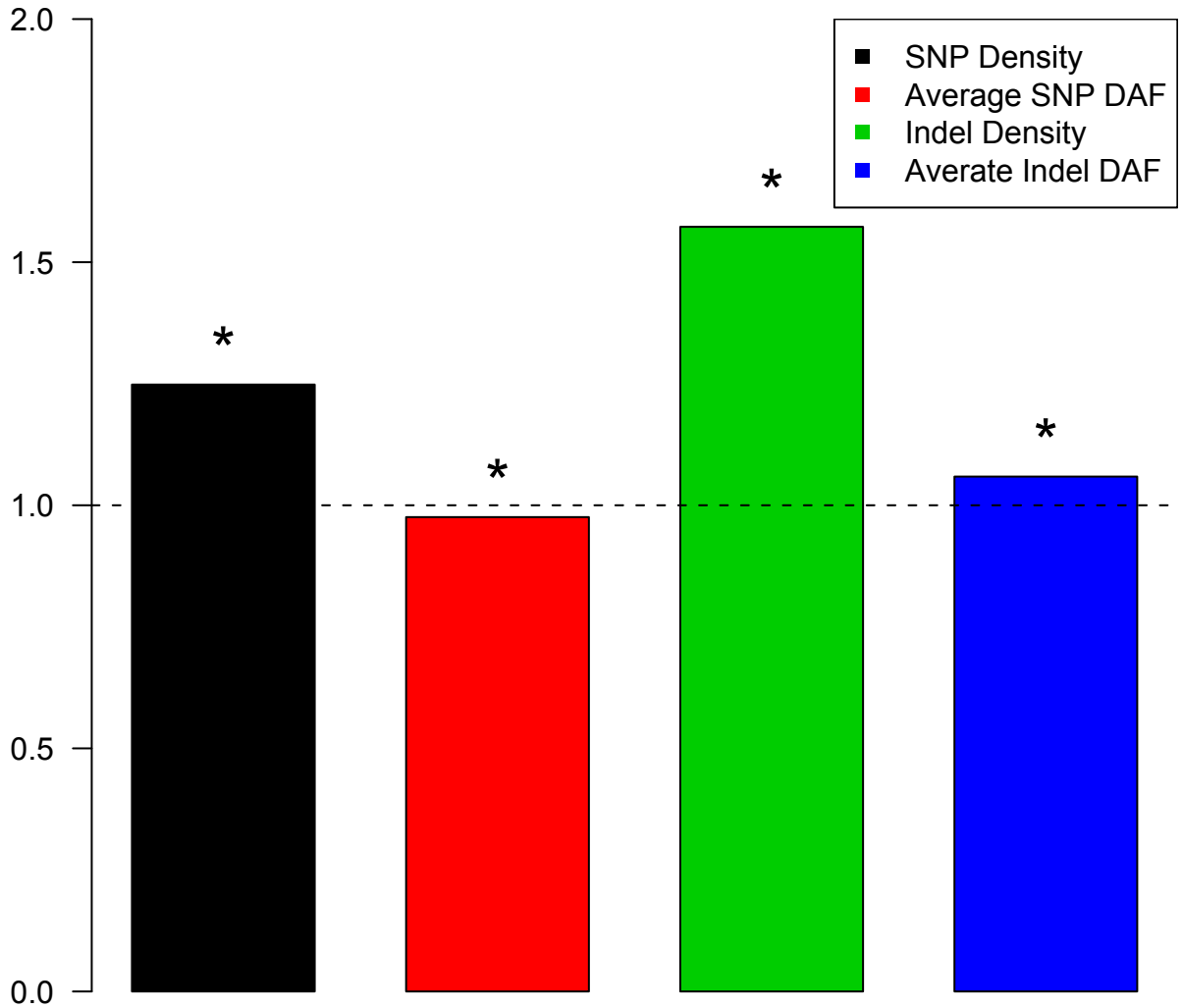


# AlleleSeq.peaks.hg19



# AlleleSeq.peaks.hg19.2kb.end.all\_transcript



# AlleleSeq.peaks.hg19.2kb.0

