



Single-allele read: `allele(idx), allele(idx, p), allele(idx, p, ch)`

Single-allele write: `setAllele(allele, idx), setAllele(allele, idx, p), setAllele(allele, idx, p, ch)`

Batch read: `genotype(), genotype(p), genotype(p, ch)`

Batch write: `setGenotype(), setGenotype(p), setGenotype(p, ch)`