region.data

site-specific

@

@

GENOME

multi-locus
scale

splitting.data
sliding.window.transform
MS/MSMS
create.PopGenome.method
extract.region.as.fasta
.
.
.
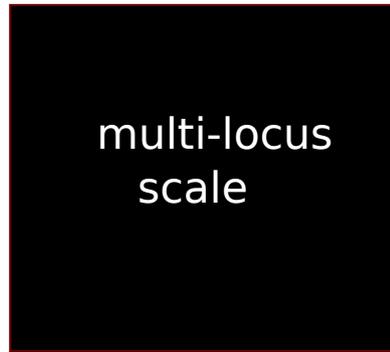
typeof::list()

biallelic.sites [[region]]
biallelic.matrix [[region]]
transitions [[region]]
synonymous [[region]]
.
.
.

@

typeof::vector()

n.biallelic.sites [region]
n.unknowns [region]
n.gaps [region]
Tajima.D [region]
FST [region]
.
.
.

@

region.stats

site-specific

@
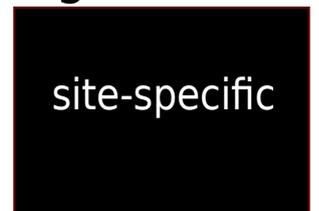
typeof::list()

minor.allele.freqs [[region]]
haplotype.counts [[region]]
.
.
.