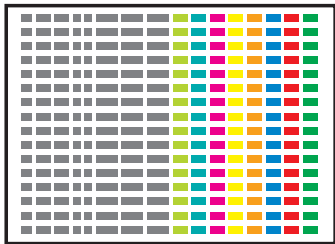


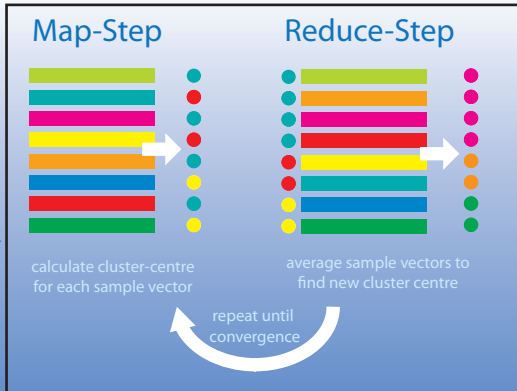


vcf file



sample
genotypes

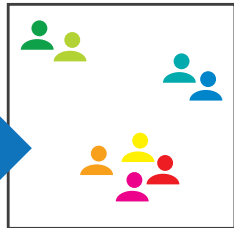
Interface: read in
genomic variants



k-means clustering

...

other machine
learning methods



Interface: visualize
genotype-based
clusters