



PhyloWGS	DPClust
Assigns CNAs to clusters	Does not assign CNAs to clusters
Estimates multiplicity of mutations during Markov chain Monte-Carlo (MCMC)	Uses most likely multiplicity of each mutation during MCMC
Reconstructs phylogenies during	Reconstructs phylogenies after MCMC
Estimates purity from SNVs and CNAs	Estimates purity from CNAs

