

command line values:  
 configuration infos  
 source  
 filter  
 gca list  
 negative gca list  
 custom genomes  
 SSU rRNAs  
 dist metric  
 dist threshold  
 clustering mode  
 dividing scheme  
 kmer engine  
 kmer size  
 kmer canonical  
 priority gca list  
 ranking formula  
 stop conditions  
 other technical infos

