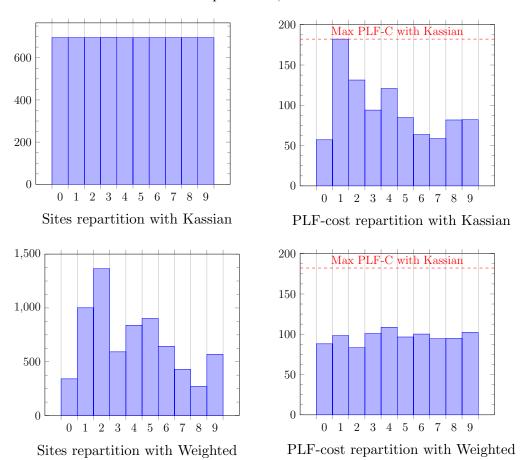
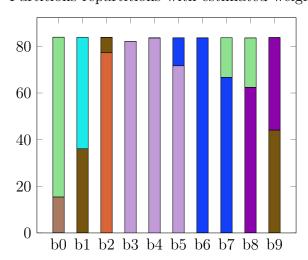
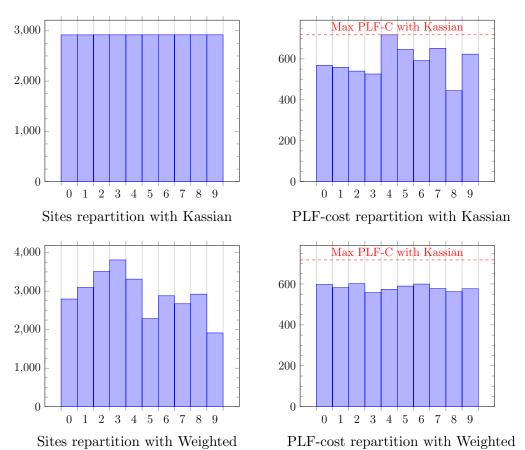
## Sequences 59, 10 CPUS



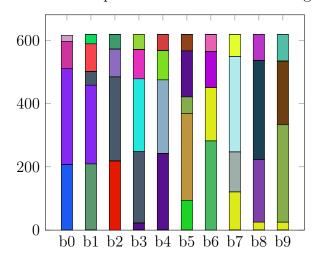
Partitions repartitions with estimated weights



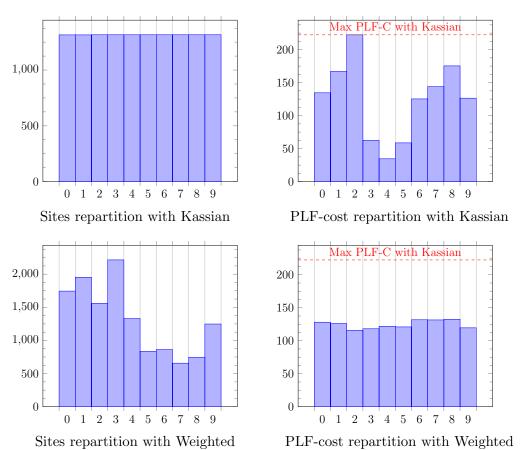
## Sequences 128, 10 CPUS



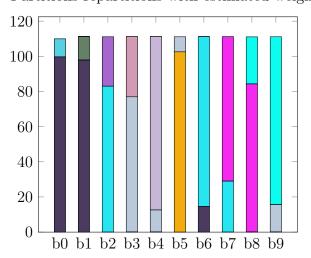
Partitions repartitions with estimated weights



## Sequences 404, 10 CPUS



Partitions repartitions with estimated weights



## Sequences proteins 94, 1487 partitions, 200 CPUS

