

RMSE of  
sequence coverage

0.010  
0.005  
0.000

BCL CU  
BCL UPad  
BT CSU  
HS UTok  
HSA TGen  
MT NIP  
NonT UU  
NonT GU  
NonT NIPS  
OM UHel  
OSA URen  
OSA CSU  
OSA UMis  
OSA UPad  
UC SI