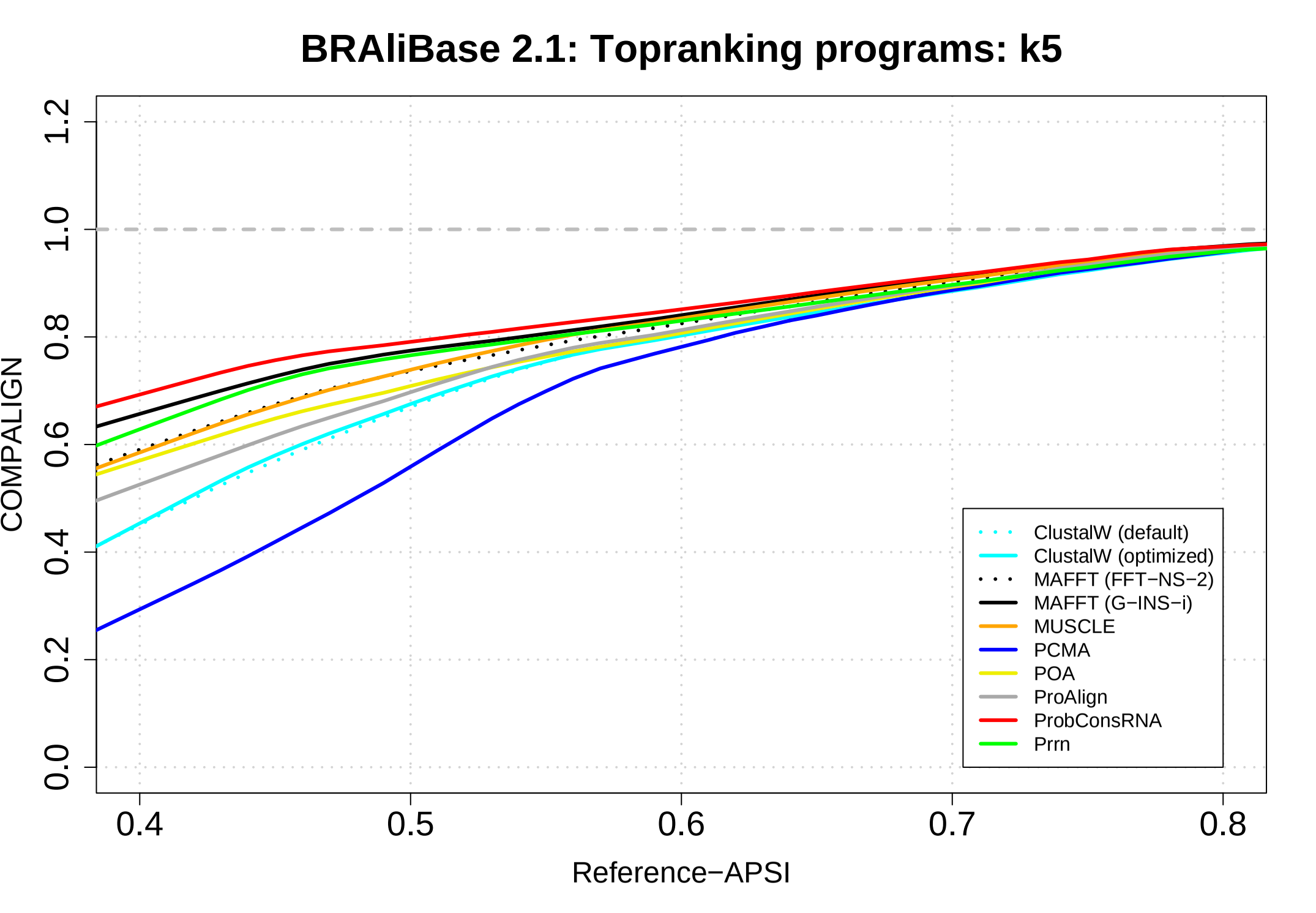
BRAliBase 2.1: Topranking programs: k5
COMPALIGN
Reference−APSI
0.0
0.2
0.4
0.6
0.8
1.0
1.2
0.4
0.5
0.6
0.7
0.8
ClustalW (default)
ClustalW (optimized)
MAFFT (FFT−NS−2)
MAFFT (G−INS−i)
MUSCLE
PCMA
POA
ProAlign
ProbConsRNA
Prrn