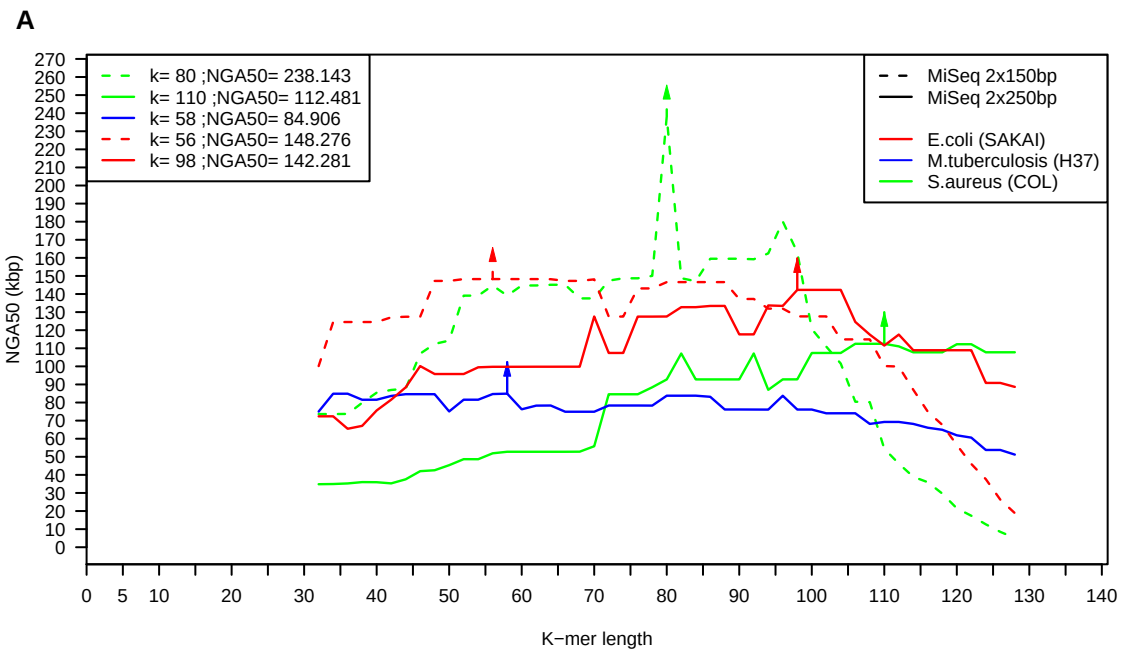
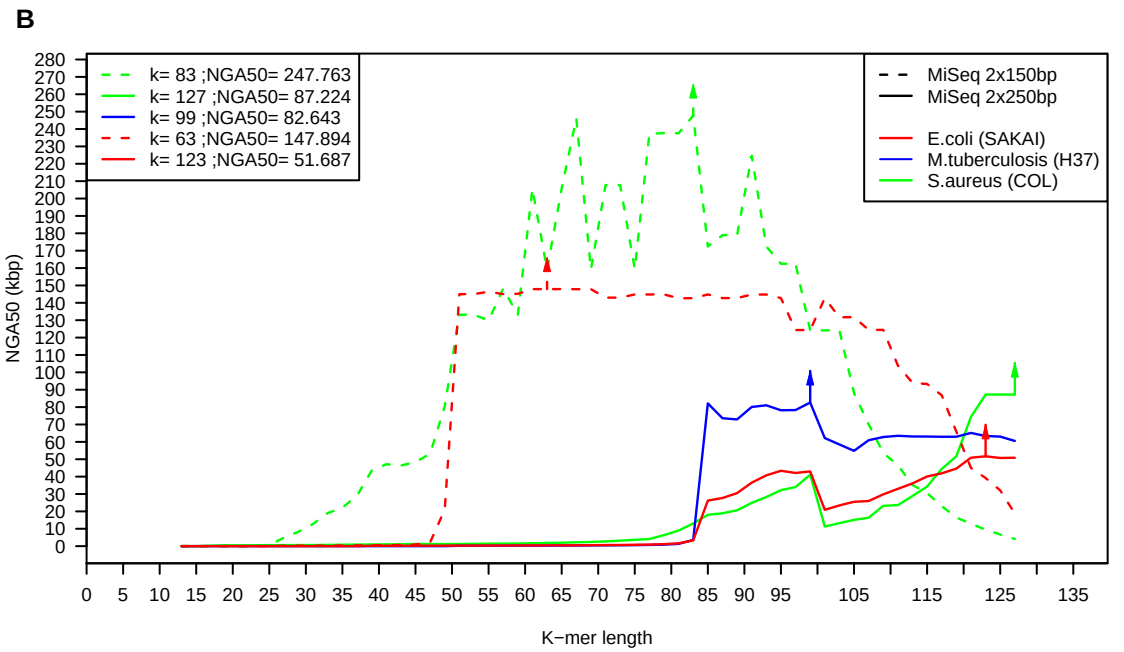
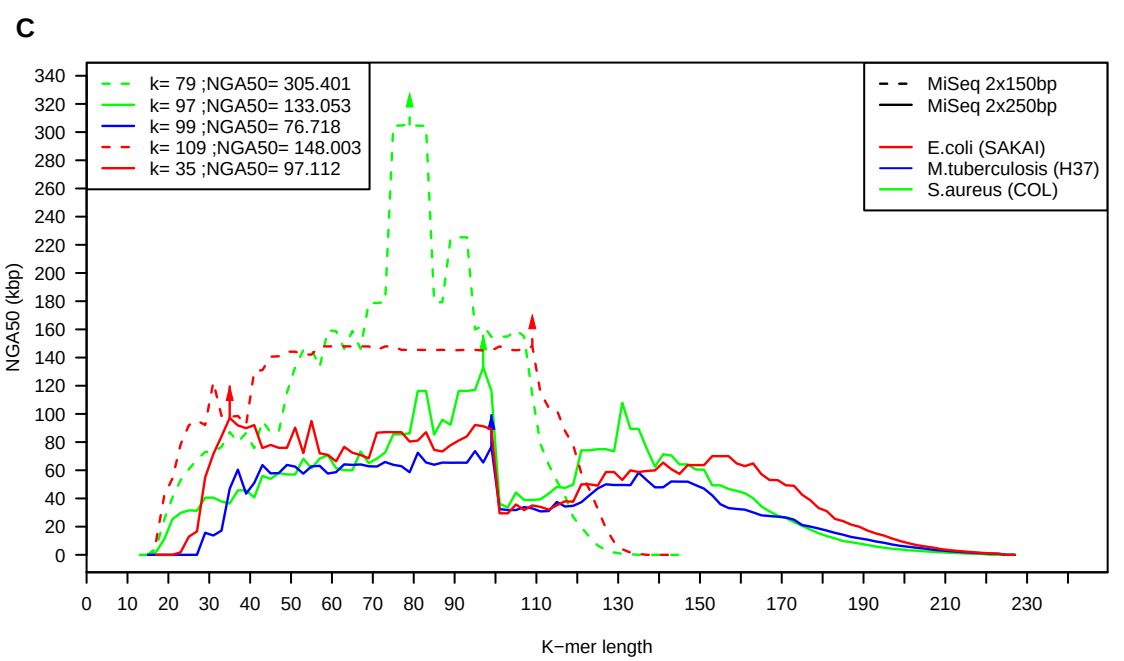
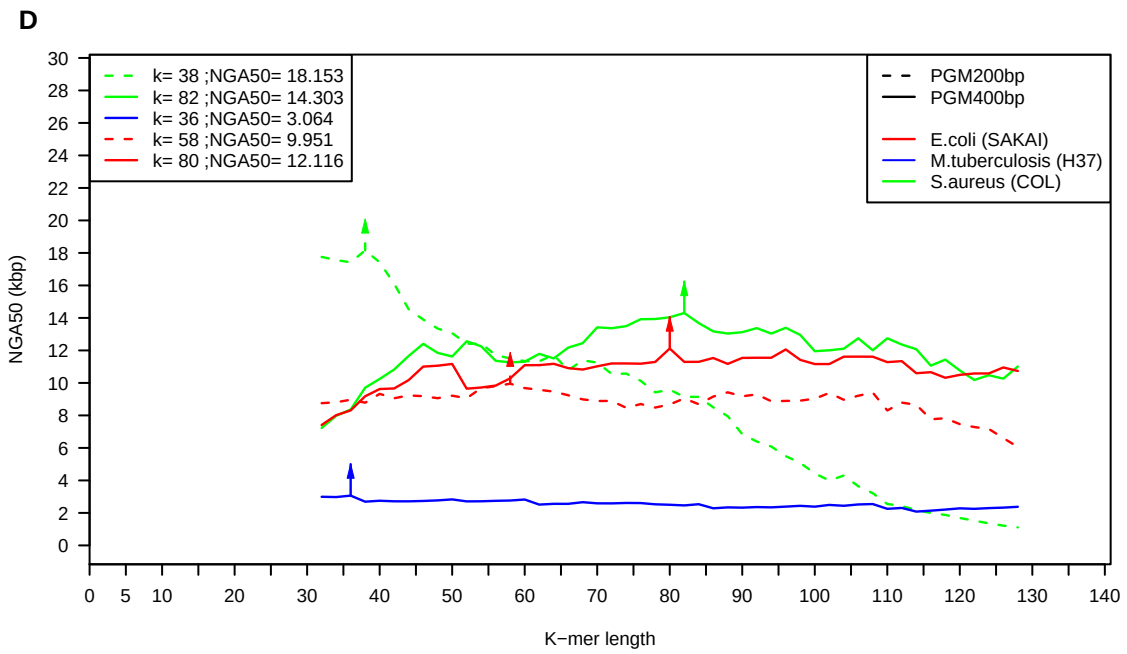
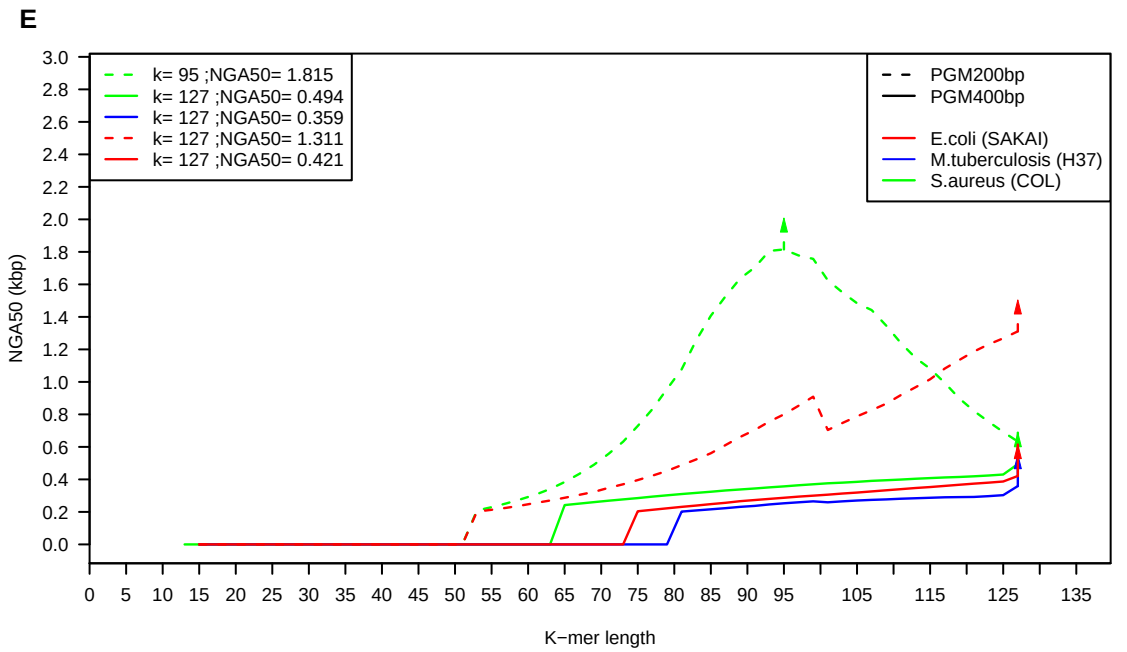
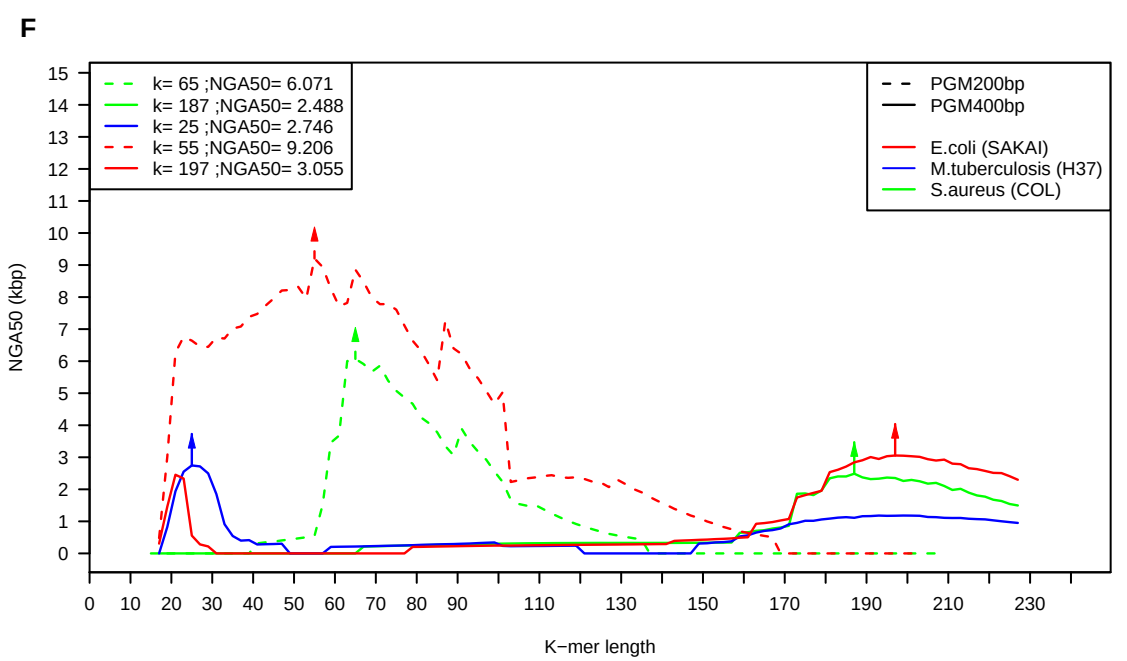

A
k= 80 ;NGA50= 238.143
k= 110 ;NGA50= 112.481
k= 58 ;NGA50= 84.906
k= 56 ;NGA50= 148.276
k= 98 ;NGA50= 142.281
MiSeq 2x150bp
MiSeq 2x250bp
E.coli (SAKAI)
M.tuberculosis (H37)
S.aureus (COL)
NGA50 (kbp)
K−mer length


B
k= 83 ;NGA50= 247.763
k= 127 ;NGA50= 87.224
k= 99 ;NGA50= 82.643
k= 63 ;NGA50= 147.894
k= 123 ;NGA50= 51.687
MiSeq 2x150bp
MiSeq 2x250bp
E.coli (SAKAI)
M.tuberculosis (H37)
S.aureus (COL)
NGA50 (kbp)
K−mer length


C
k= 79 ;NGA50= 305.401
k= 97 ;NGA50= 133.053
k= 99 ;NGA50= 76.718
k= 109 ;NGA50= 148.003
k= 35 ;NGA50= 97.112
MiSeq 2x150bp
MiSeq 2x250bp
E.coli (SAKAI)
M.tuberculosis (H37)
S.aureus (COL)
NGA50 (kbp)
K−mer length


D
k= 38 ;NGA50= 18.153
k= 82 ;NGA50= 14.303
k= 36 ;NGA50= 3.064
k= 58 ;NGA50= 9.951
k= 80 ;NGA50= 12.116
PGM200bp
PGM400bp
E.coli (SAKAI)
M.tuberculosis (H37)
S.aureus (COL)
NGA50 (kbp)
K−mer length


E
k= 95 ;NGA50= 1.815
k= 127 ;NGA50= 0.494
k= 127 ;NGA50= 0.359
k= 127 ;NGA50= 1.311
k= 127 ;NGA50= 0.421
PGM200bp
PGM400bp
E.coli (SAKAI)
M.tuberculosis (H37)
S.aureus (COL)
NGA50 (kbp)
K−mer length


F
k= 65 ;NGA50= 6.071
k= 187 ;NGA50= 2.488
k= 25 ;NGA50= 2.746
k= 55 ;NGA50= 9.206
k= 197 ;NGA50= 3.055
PGM200bp
PGM400bp
E.coli (SAKAI)
M.tuberculosis (H37)
S.aureus (COL)
NGA50 (kbp)
K−mer length