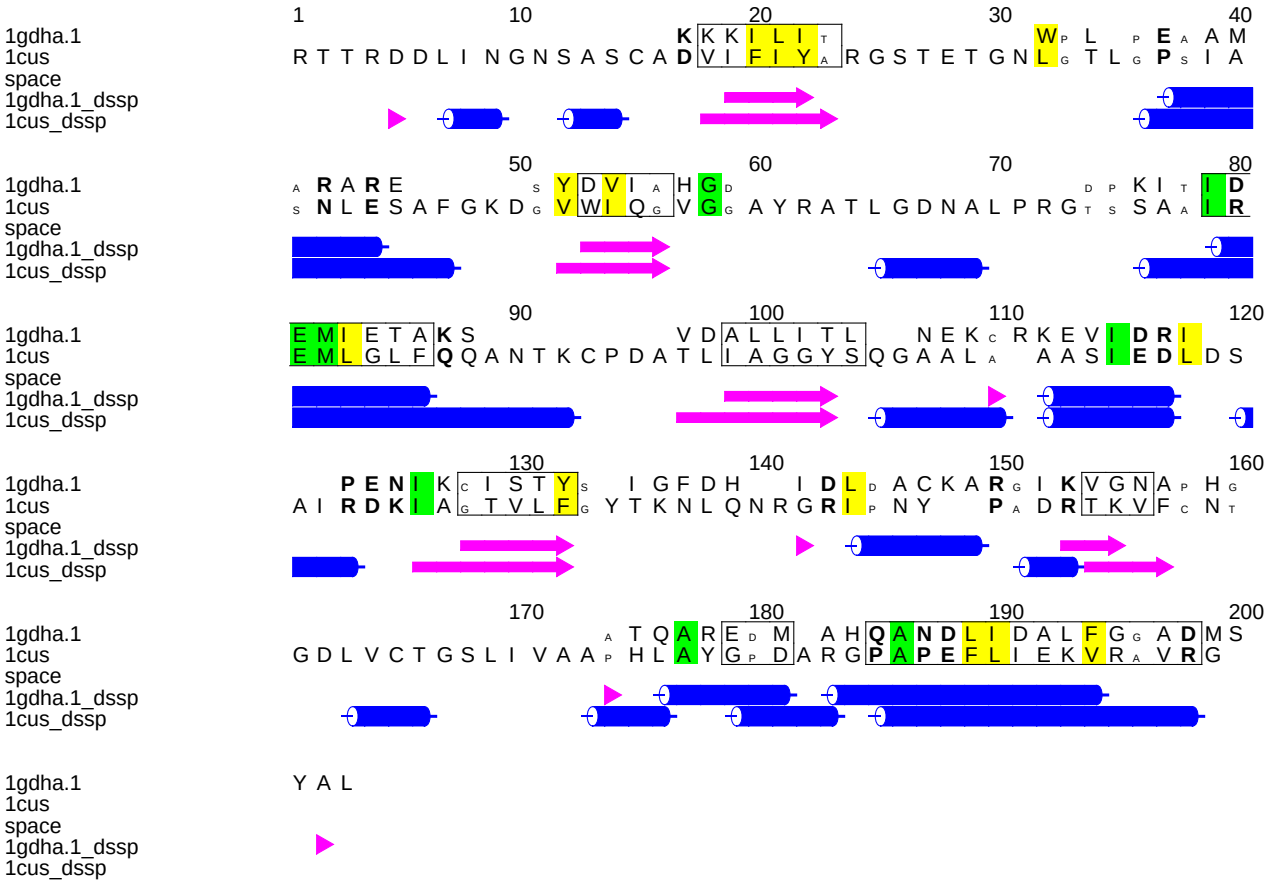
1gdha.1
1cus
space
1gdha.1_dssp
1cus_dssp
1 10 20 30 40
RTTRDDLI NGNSASCA
50 60 70 80
90 100 110 120
130 140 150 160
170 180 190 200
YAL