

logo

NTDB id 130020 AW20 RS08895 WP 000058556.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



85
85

logo

NTDB id 130020 AW20 RS08895 WP 000058556.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



170
169

logo

NTDB id 130020 AW20 RS08895 WP 000058556.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



255
254

logo

NTDB id 130020 AW20 RS08895 WP 000058556.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



339
339

logo

NTDB id 130020 AW20 RS08895 WP 000058556.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



424
424

logo

NTDB id 130020 AW20 RS08895 WP 000058556.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



509
508

logo

NTDB id 130020 AW20 RS08895 WP 000058556.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



594
593

[illegible][illegible]

Sequence logo

RYLSSLFYRNRAQKLSSTSRDLGDKIKGSVSRMELFNRCAYAHFAQHGLSLRERDIFFKLDAPDIGELFHAAALKRIADRLREN
KLFFSSLFRNEVKQLERSVSRLYGERIQGSVSRMETFNACPSHFAHQHGLHLKERQFFKLAPDIGQLFHHSSLLKLTSDRLREQK
** * * * * ** * * *

RTWRDLTKKQCELFSYWVVERLAPLRQKLLSSNRHYVVKQLQKQVTRVSIILSEHAKSSGFVPVGLLEVFPFGMGGTGSLPPLMT
LDIADLSIEEHFADAIETAPLRKEILLSSNRHFYVKQLQKQITFRTSIILREHAKSSGFVPVDLEVFPFGMGGTGSLPPLME
LDWRDLTKKECLFYSYDAVERLAPLRKEILLSSNRHYVVKQLQKQIVTRVSGILSEHAKASGFVPIGLLE..GFGGKGGLPLPLT

FSLPNQVTKMEVGRIDRVDKAESSENGLTFLRIYDYKSSSKGLDLTEVYYGLALQLMLTYLDWTTNSQTWMKMTATPAGVLVFHIL
FSLPNQVKMEVGRIDRVDKAEDENGTLFLRIDYKSSSKALDLTEVYYGLALQLMLTYLDVVTSNAAQTWMKKGGTASPAGVLVFHIL
FQLKNGCTMELVGRIDRVDKAESSEKGLLRLIVDYKSSDKGLDLAEVYYGLALQLMLTYLDLSITHSADWLGL..MRATPAGVLVFHIL

 non conserved
 similar
 $\geq 50\%$ conserved