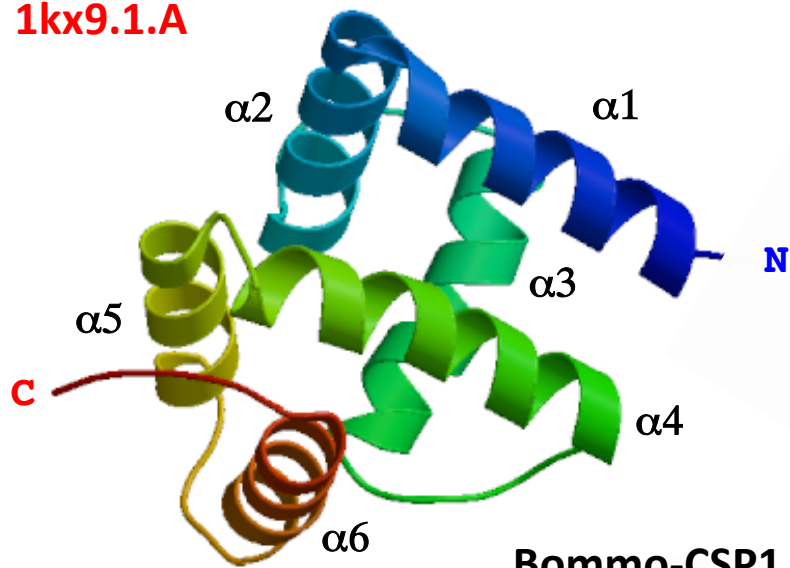
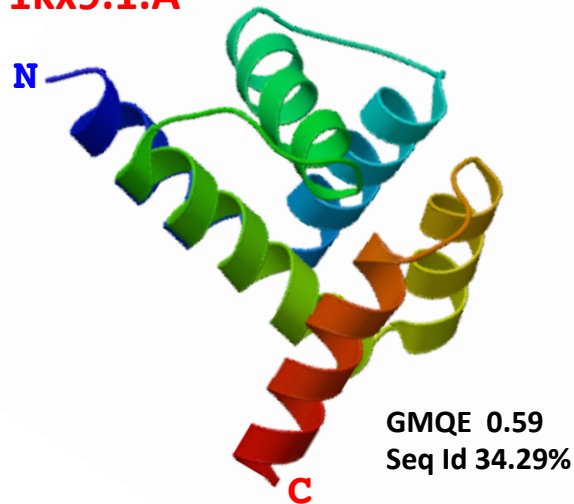


1kx9.1.A



Bommo-CSP1

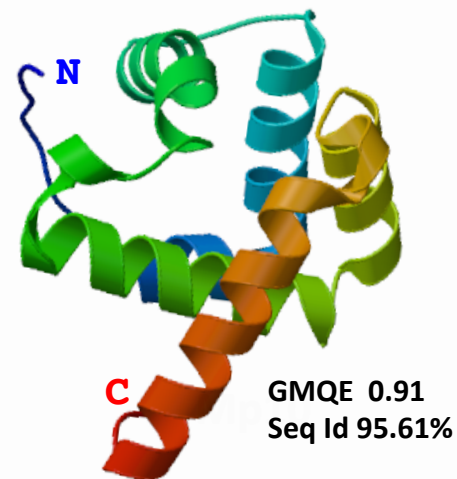
1kx9.1.A



GMQE 0.59
Seq Id 34.29%

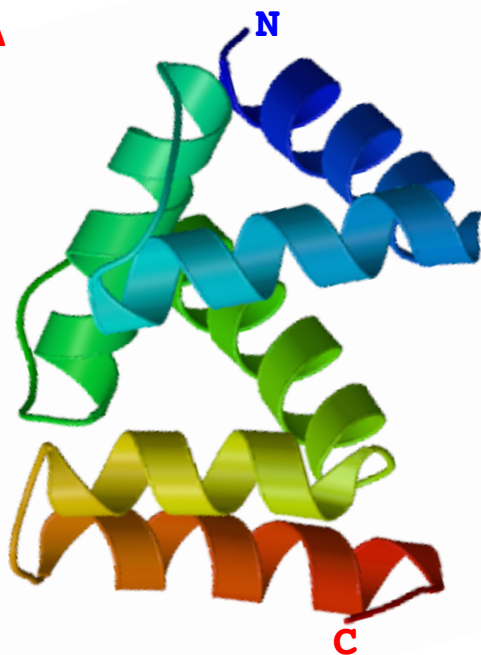
Mp10

A7UL73.1.A



GMQE 0.91
Seq Id 95.61%

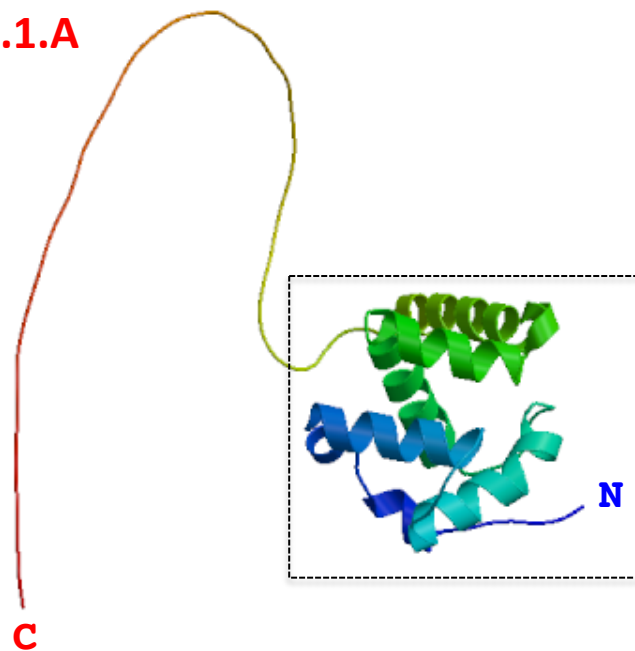
1kx9.1.A



GMQE 0.34
Seq Id 36.11%

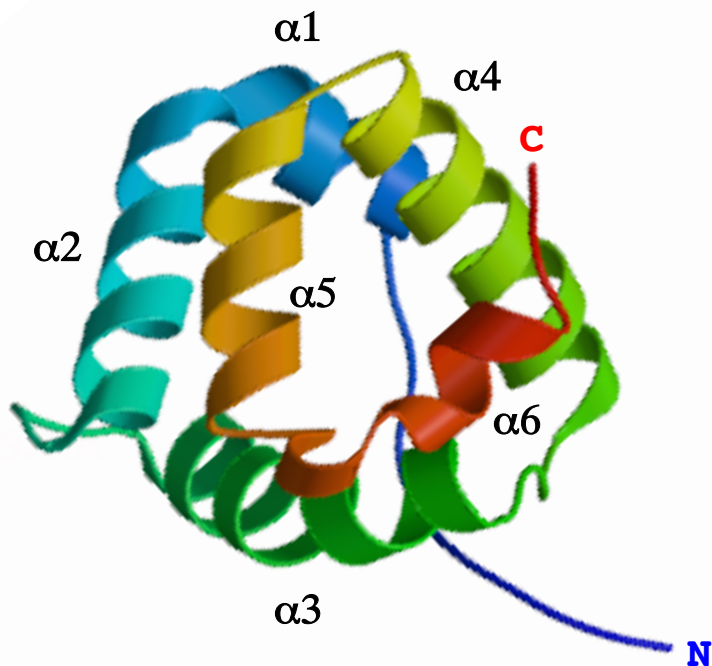
Bommo-CSP10

Q3LB93.1.A



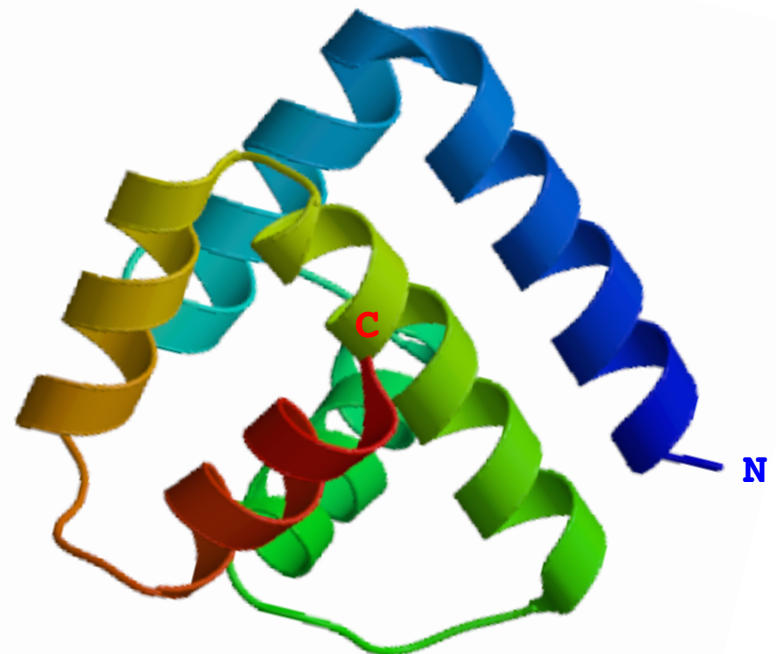
GMQE 0.68
Seq Id 99.38%

Q9W0X2.1.A



GMQE 0.93
Seq Id 79.41%

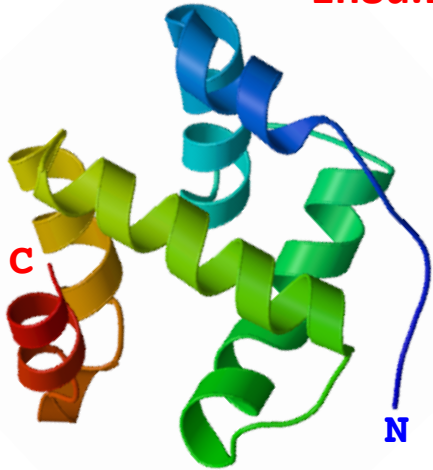
1kx9.1.A



GMQE 0.66
Seq Id 46.53%

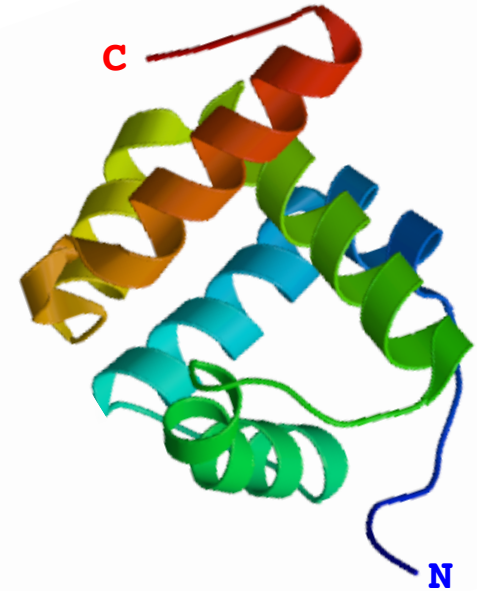
GH22042
Drosophila grimshawi
pherokine (hemolymph protein)

1n8u.1.A



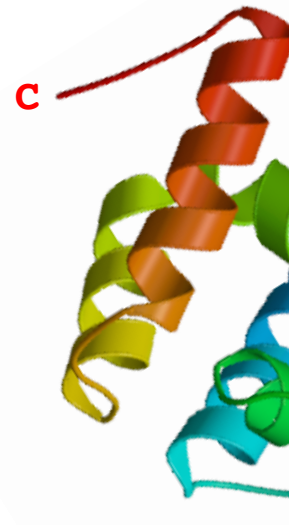
GMQE 0.15
Seq Id 39.18%

A0A835CMW0.1.A



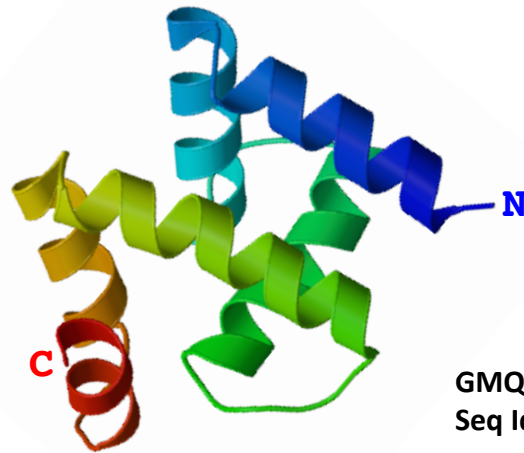
GMQE 0.89
Seq Id 100.00%

1kx9.1.A



GMQE 0.65
Seq Id 46.36%

1kx9.1.A



GMQE 0.15
Seq Id 39.18%

XM_044889794

Coccinella septempunctata

eukaryotic translation initiation factor 4 gamma-like

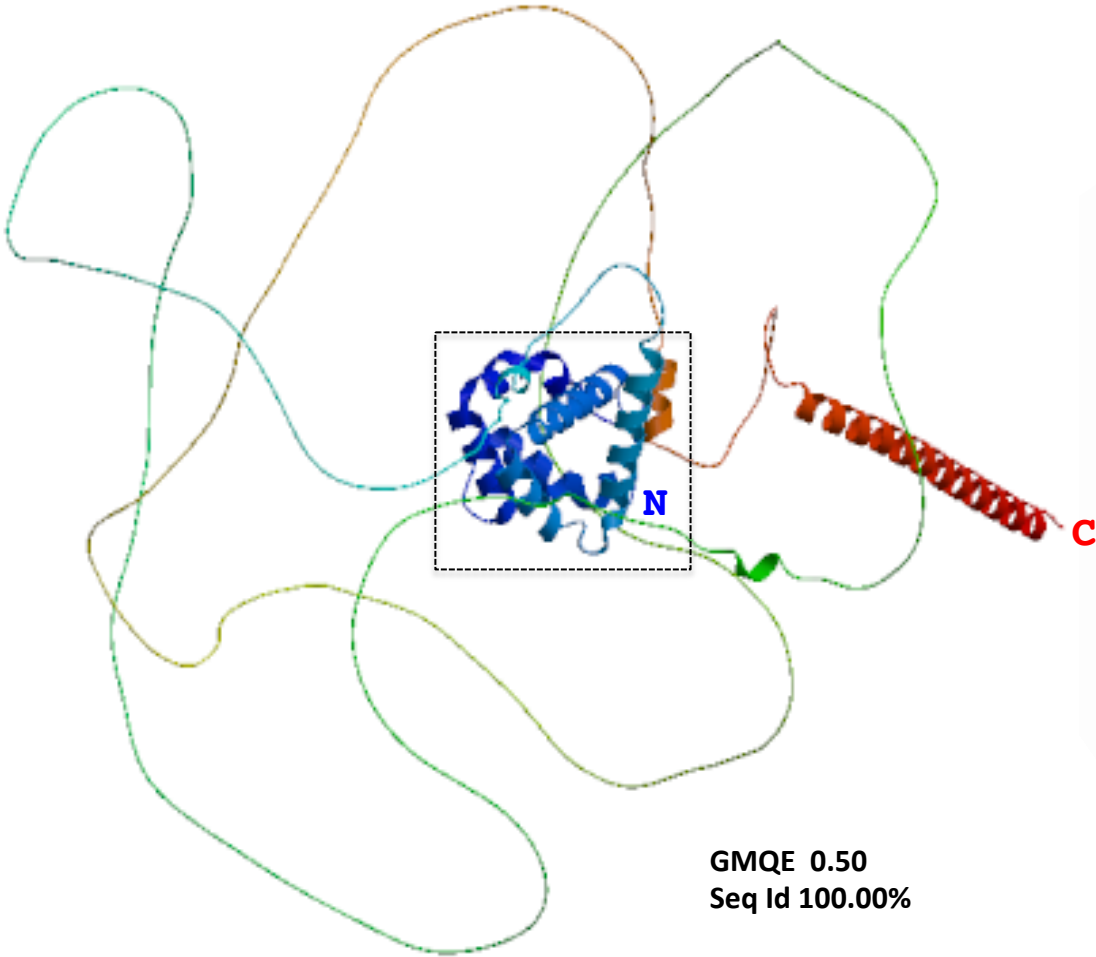
AHX71992 (KJ451426)

Aphis gossypii

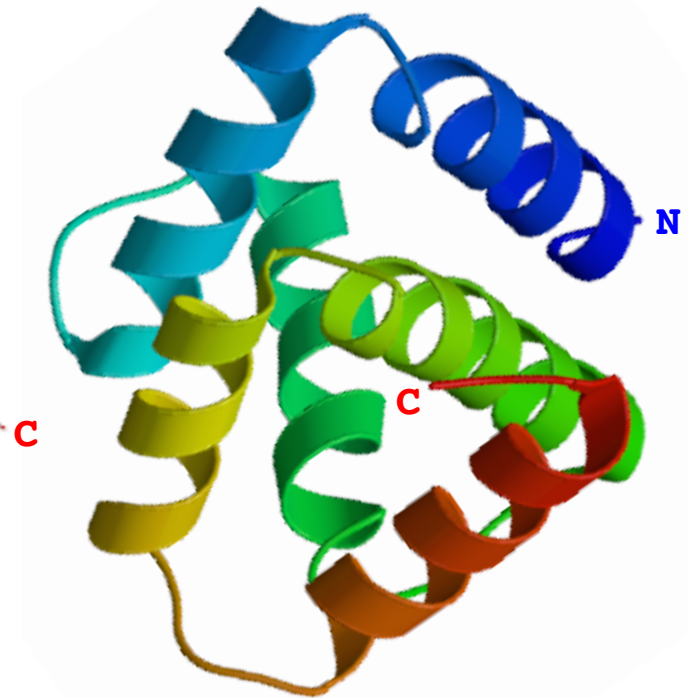
acid trehalase

A0AGI8TVU3.1

1kx9.1.A



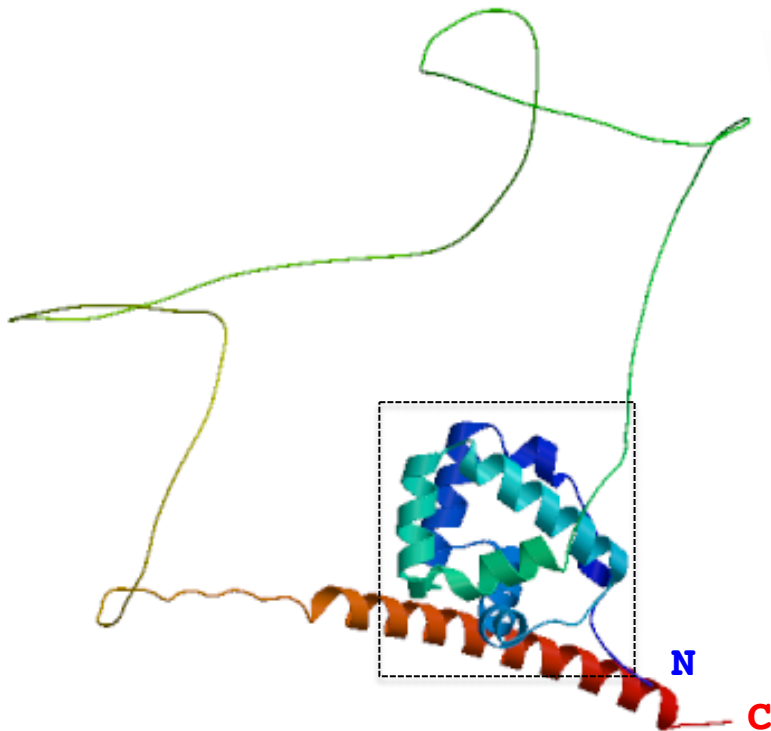
GMQE 0.50
Seq Id 100.00%



GMQE 0.12
Seq Id 30.56%

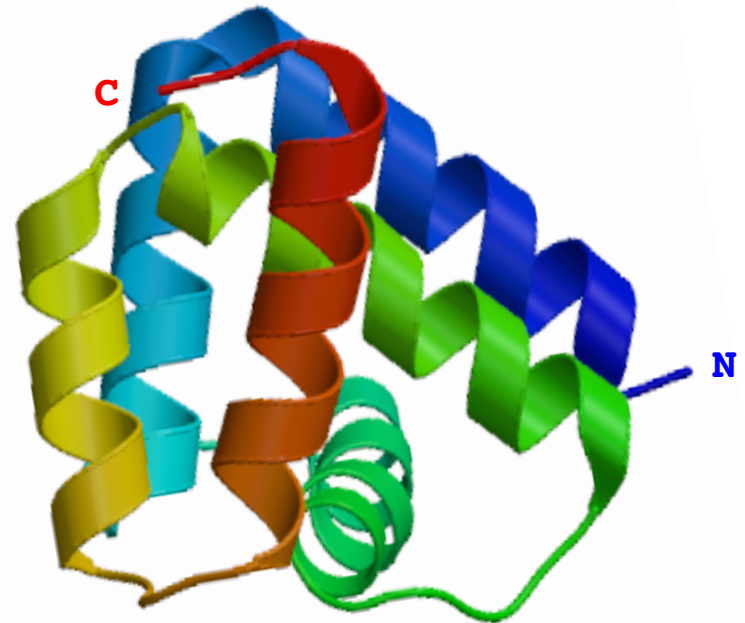
XP_021703245
Aedes aegypti
nuclear pore complex protein
(NPCP)

A0A2H4ZB78.1.A



GMQE 0.62
Seq Id 97.54%

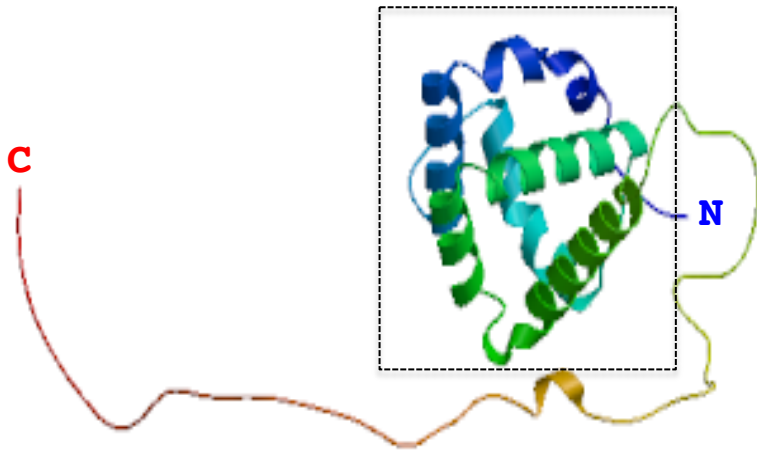
1kx9.1.A



GMQE 0.23
Seq Id 33.64%

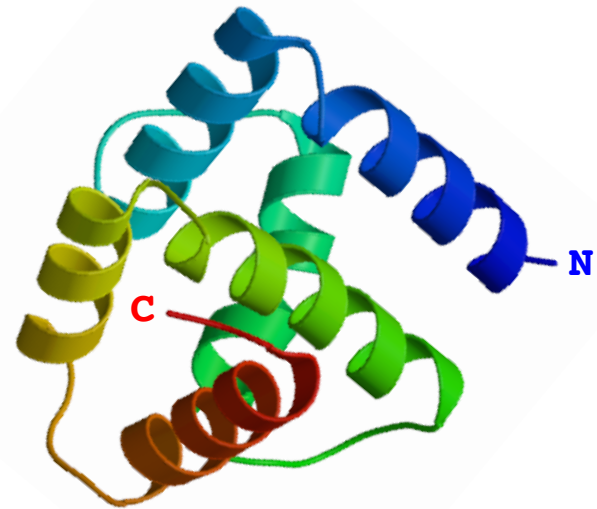
XP_018563025
Anoplophora glabripennis
cell wall integrity and stress response component 3-like
(CWP)

Q6H8Y7.1.A



GMQE 0.37
Seq Id 71.34%

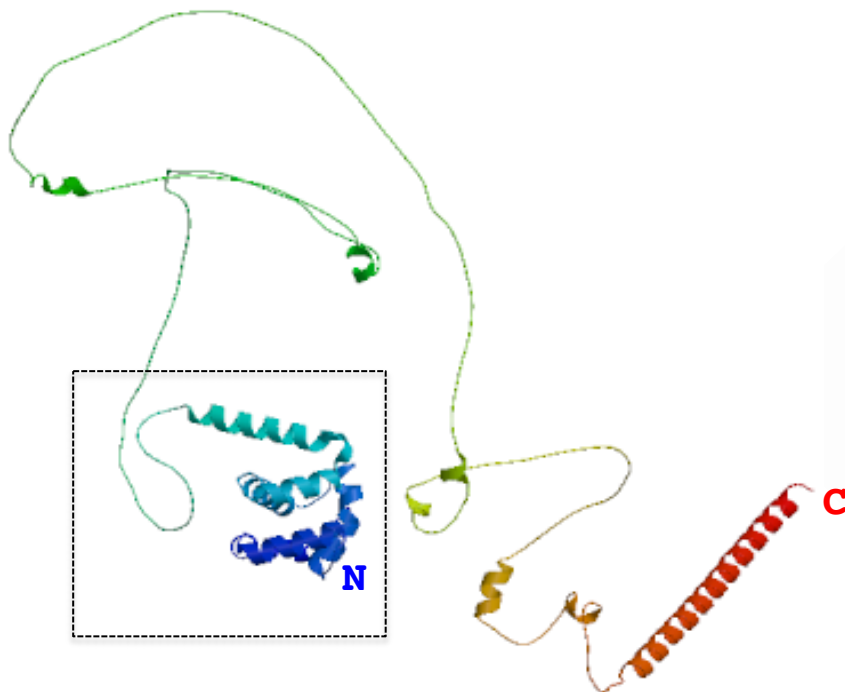
1kx9.1.A



GMQE 0.20
Seq Id 32.41%

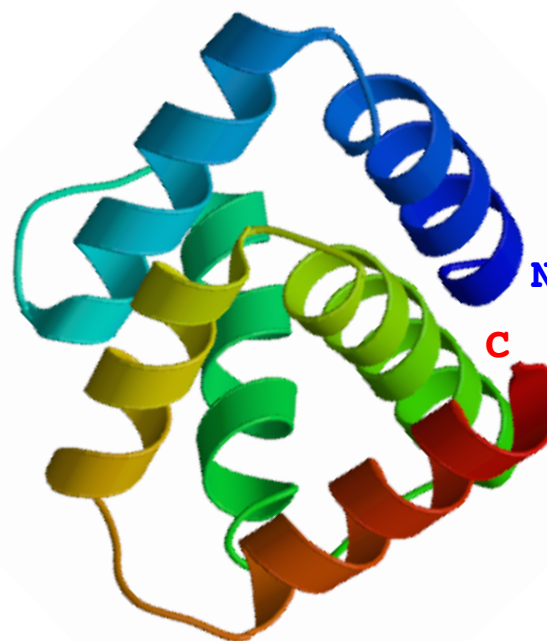
XM_055738352
Uranotaenia lowii
integumentary mucin C.1-like
Translation Initiation Factor (TIF)

A0A6I8TWU8.1.A



GMQE 0.55
Seq Id 89.92%

1kx9.1.A

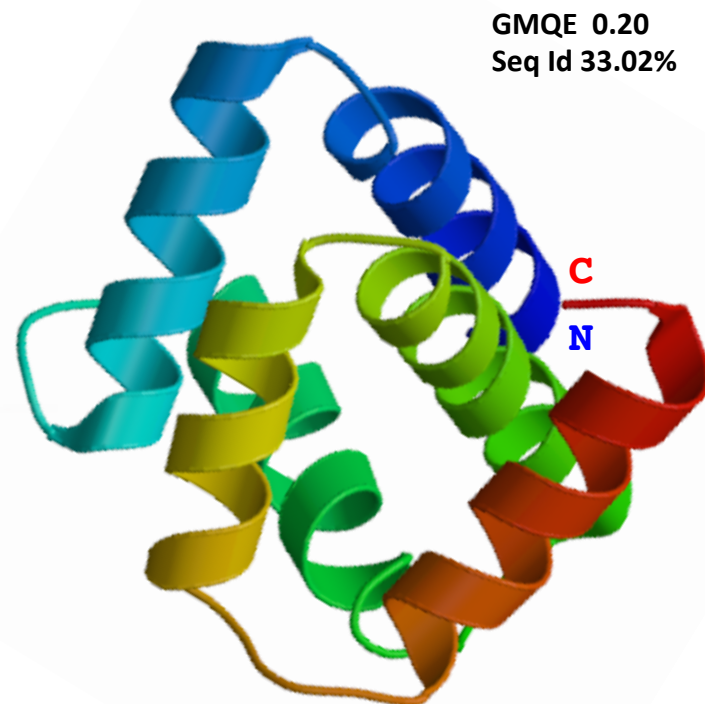
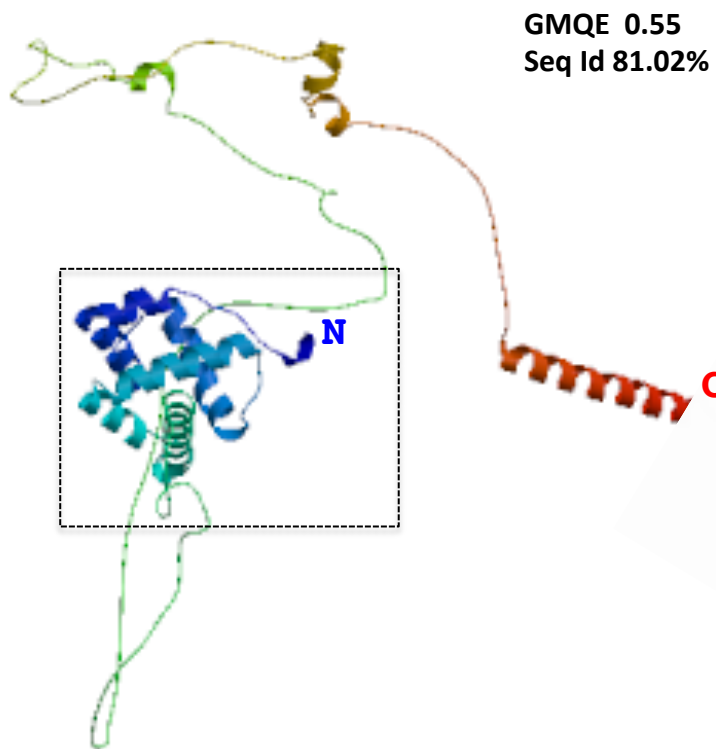


GMQE 0.17
Seq Id 32.08%

XM_029870982
Aedes albopictus
nuclear pore complex protein
(NPCP)

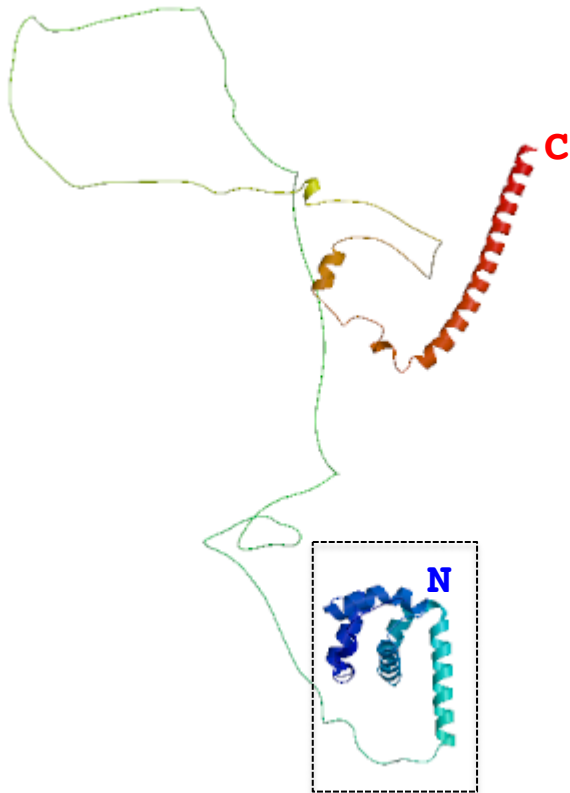
A0A6H0D4B0.1.A

1kx9.1.A



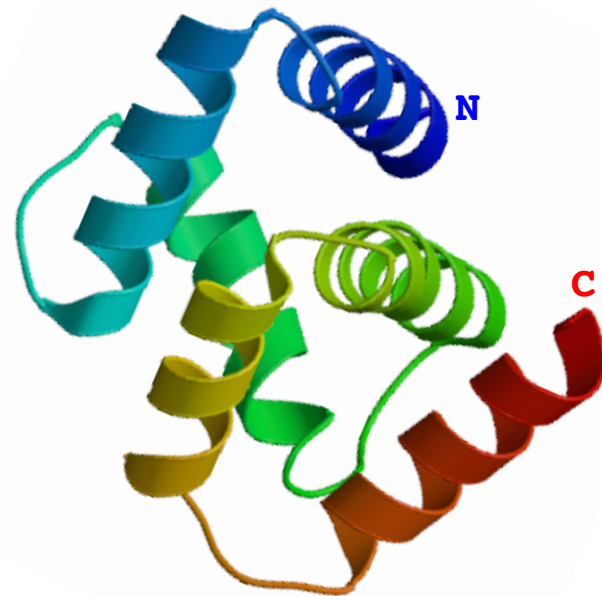
XM_056065396
Eupeodes corollae
rho GTPase-activating protein 17-like
(Rho)

A0A182NU82.1.A



GMQE 0.53
Seq Id 84.12%

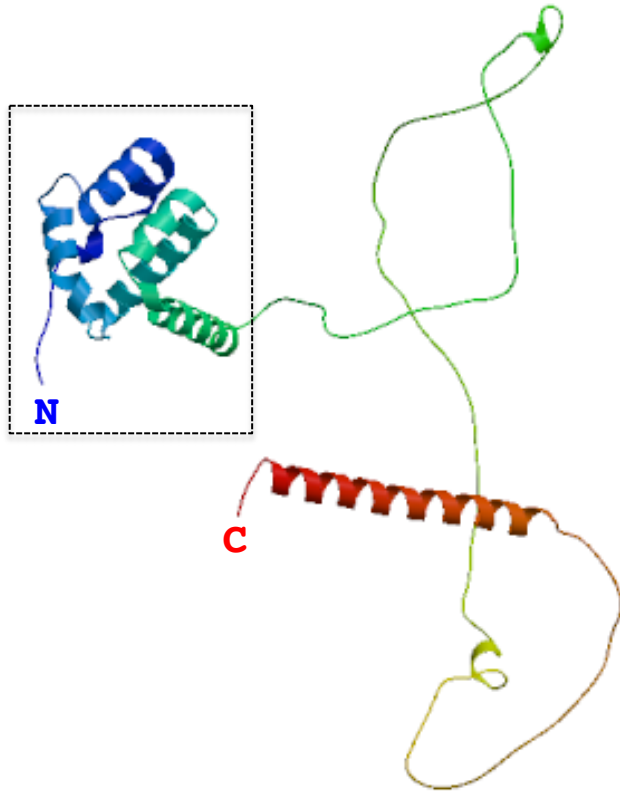
1kx9.1.A



GMQE 0.18
Seq Id 32.38%

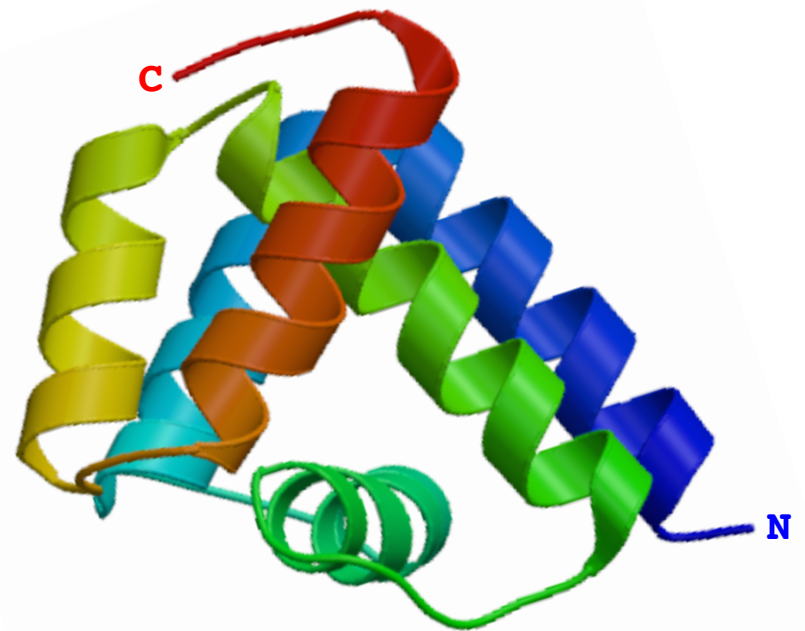
XM_050245628
Anopheles aquasalis
WAS/WASL-interacting protein family
member 1-like isoform X2
(Mucin)

A0A0K8TVC4.1.A



GMQE 0.64
Seq Id 72.48%

1kx9.1.A



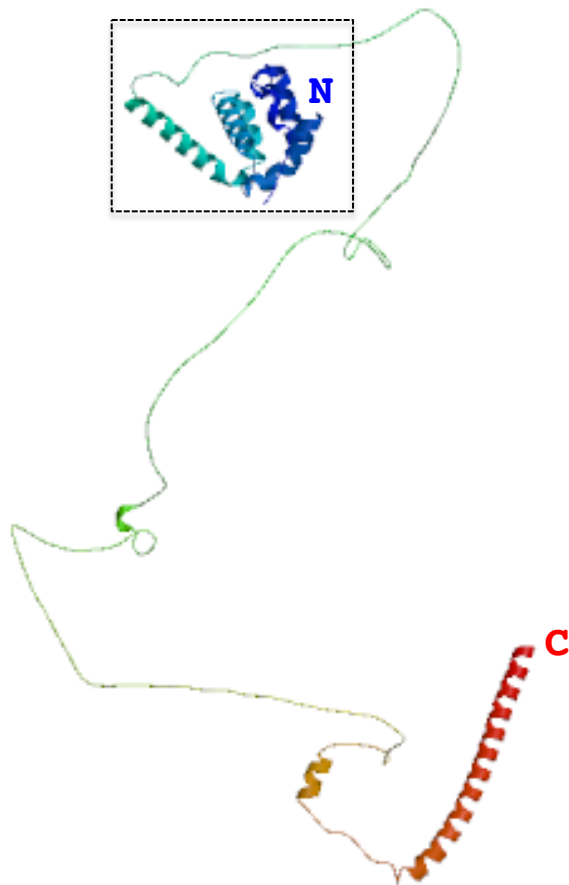
GMQE 0.25
Seq Id 34.26%

XM_047175293

Helicoverpa zea

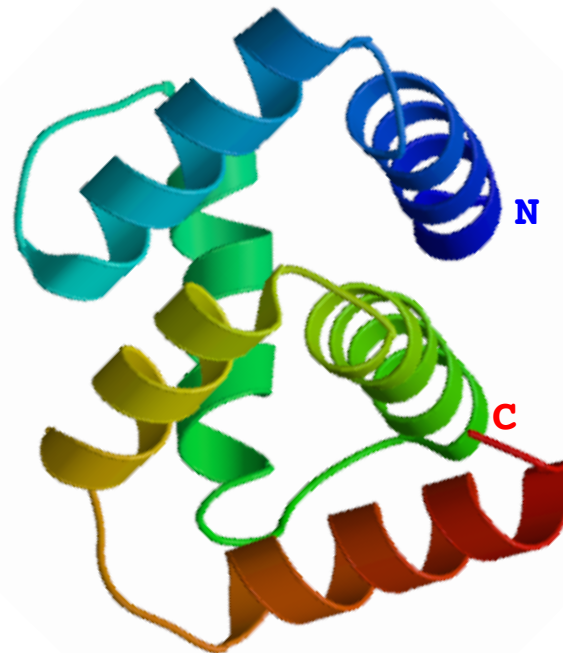
actin cytoskeleton-regulatory complex protein PAN1-like
(ASRP)

A0A182NU82.1.A



GMQE 0.57
Seq Id 88.40%

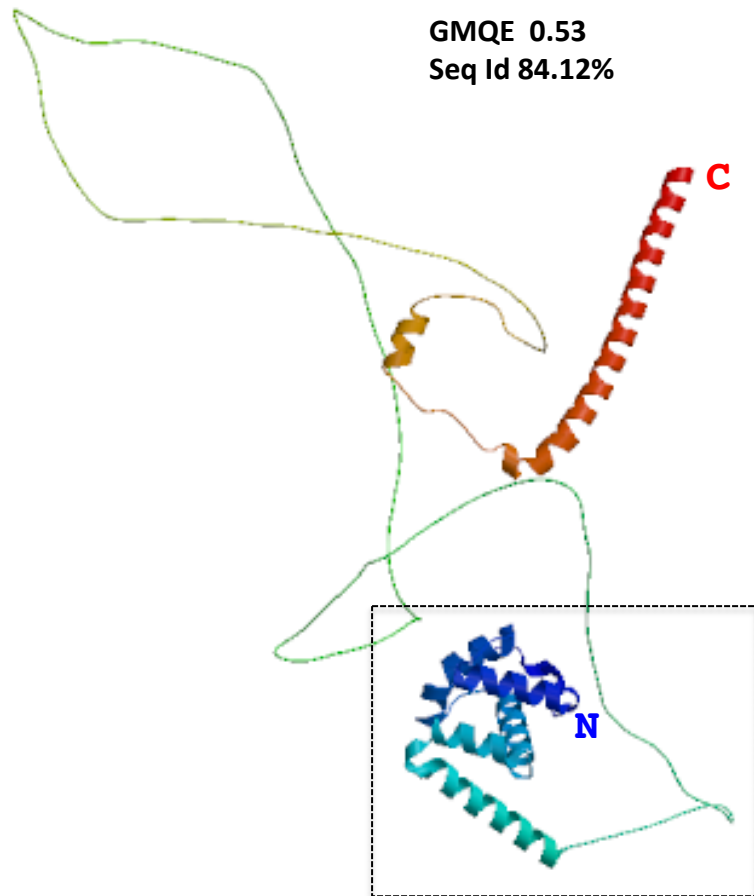
1kx9.1.A



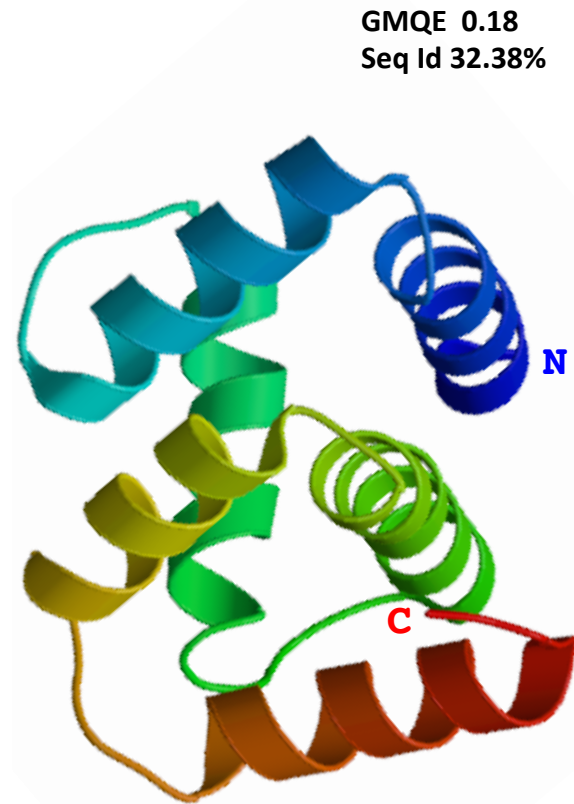
GMQE 0.18
Seq Id 32.08%

XM_058204117
Anopheles bellator
mucin-2-like
Protein UL36

A0A182NU82.1.A

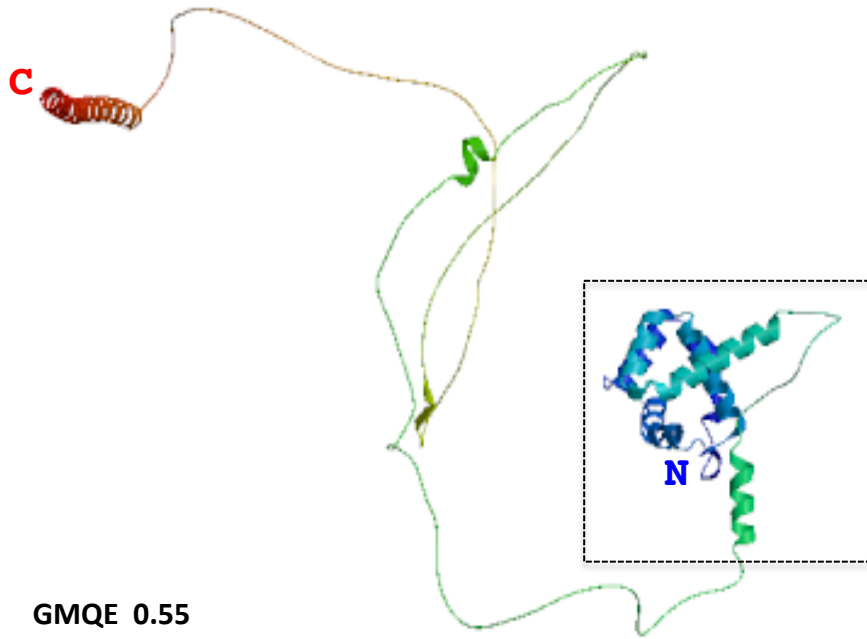


1kx9.1.A



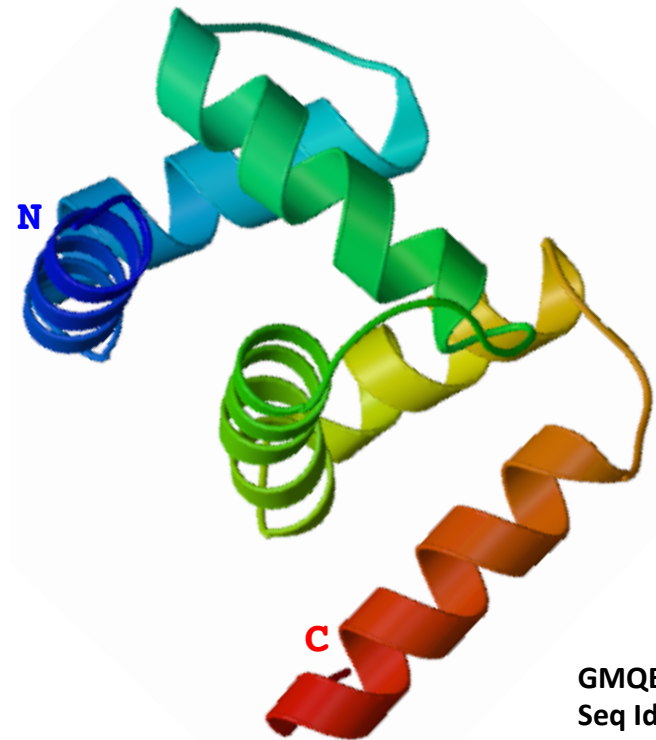
XM_053009541
Anopheles cruzii
WAS/WASL protein
Actin skeleton-regulatory complex (ASRC)

A0A1Q3FP99.1.A



GMQE 0.55
Seq Id 93.30%

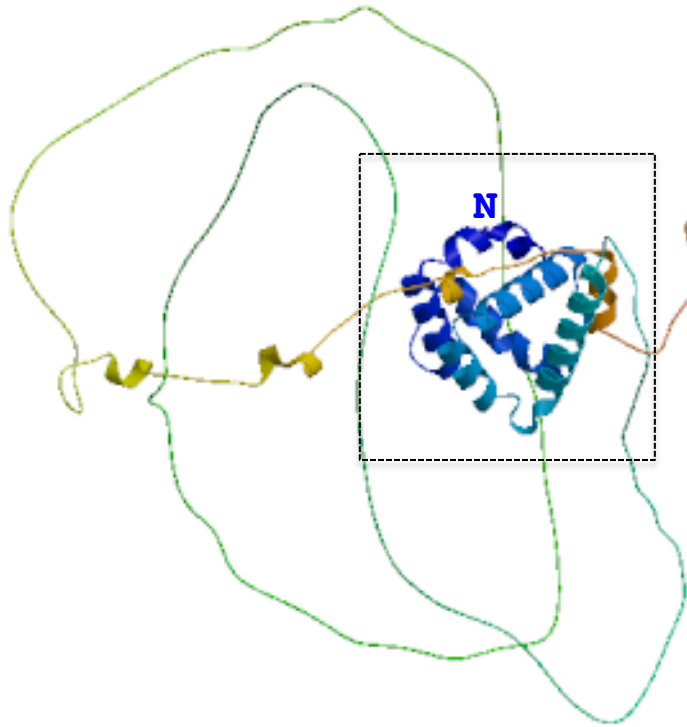
1kx9.1.A



GMQE 0.17
Seq Id 29.91%

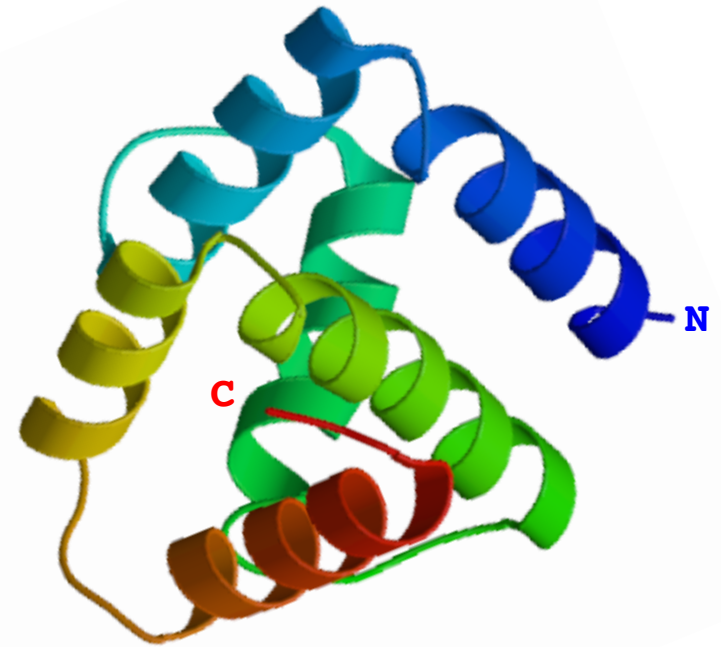
XM_038253956
Culex quinquefasciatus
cell wall protein DAN4
(CWP)

A0A1Q3FQF4.1.A



GMQE 0.52
Seq Id 90.45%

1kx9.1.A

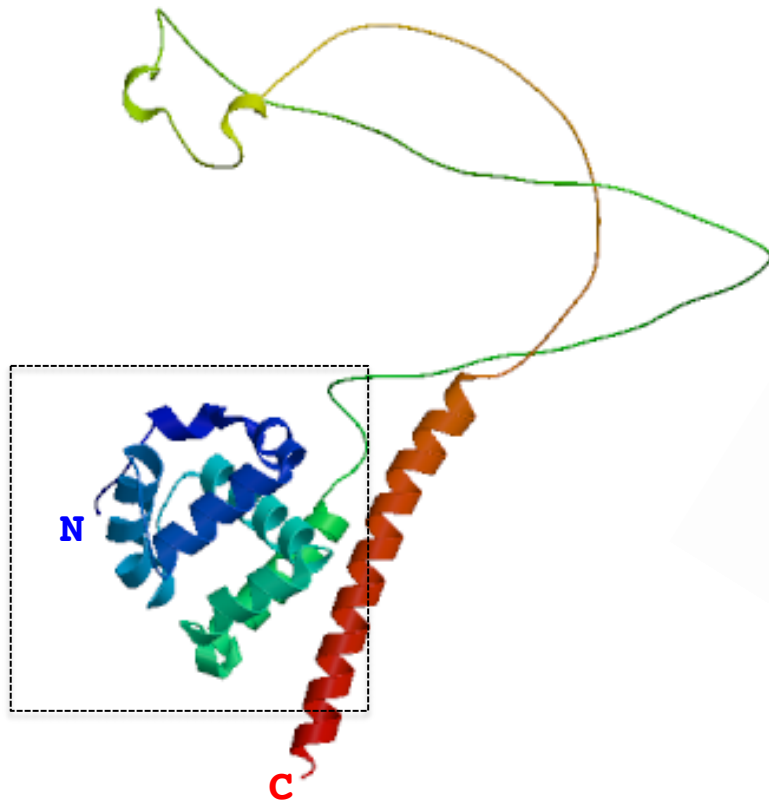


GMQE 0.15
Seq Id 31.43%

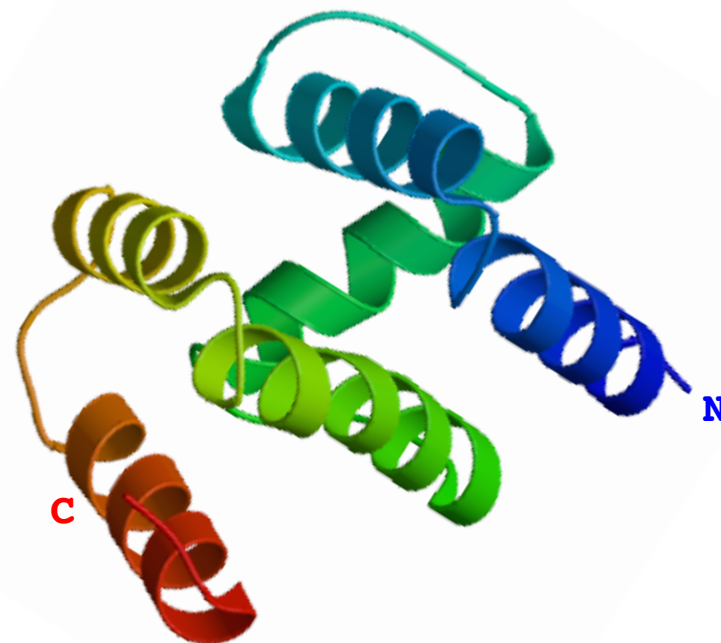
XM_052707633
Culex pipiens pallens
mucin-2-like
(Mucin)

A0A1V1WC08.1.A

1kx9.1.A



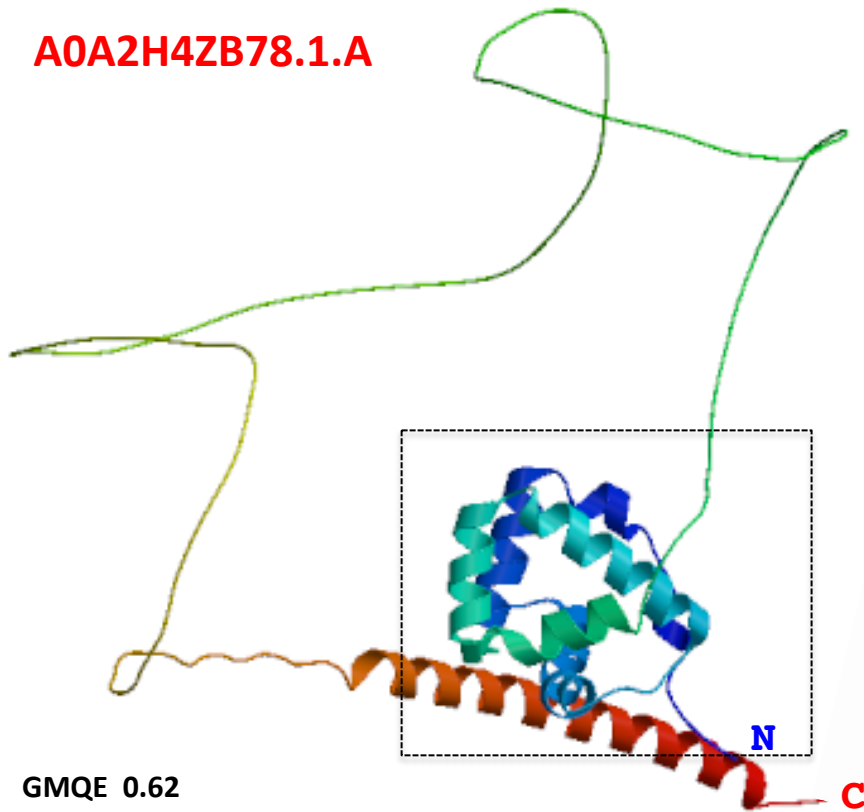
GMQE 0.64
Seq Id 70.31%



GMQE 0.25
Seq Id 34.58%

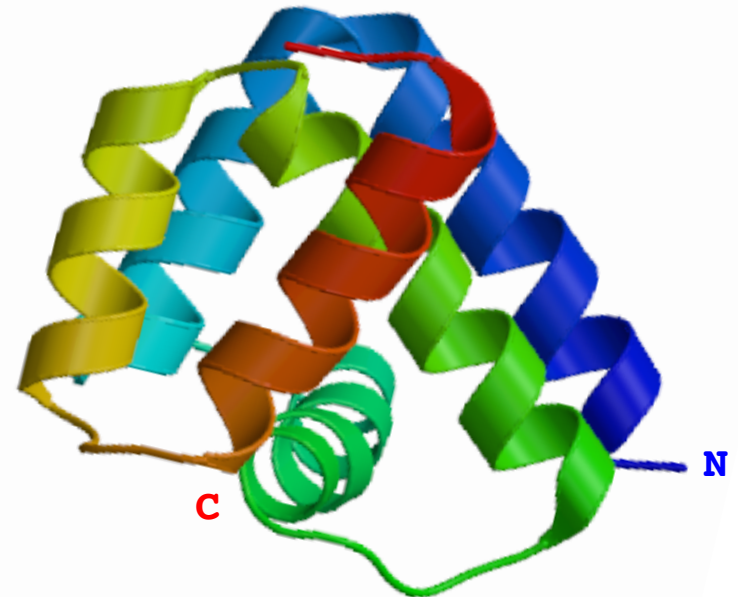
CAH2235359
Pararge aegeria aegeria
Herpes virus major outer envelope glycoprotein
(jg5928)

A0A2H4ZB78.1.A



GMQE 0.62
Seq Id 97.54%

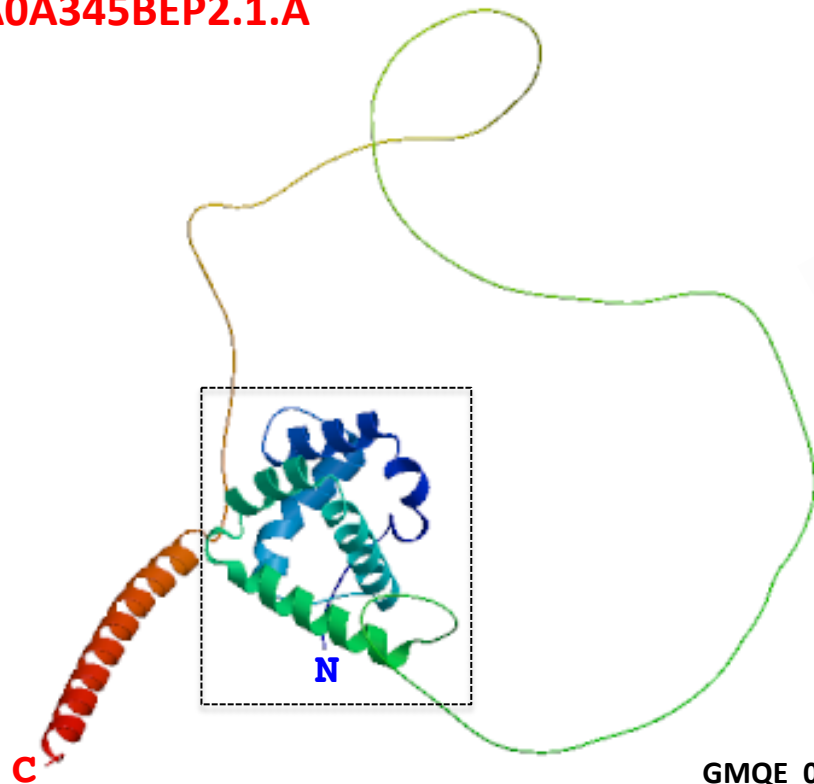
1kx9.1.A



GMQE 0.23
Seq Id 33.64%

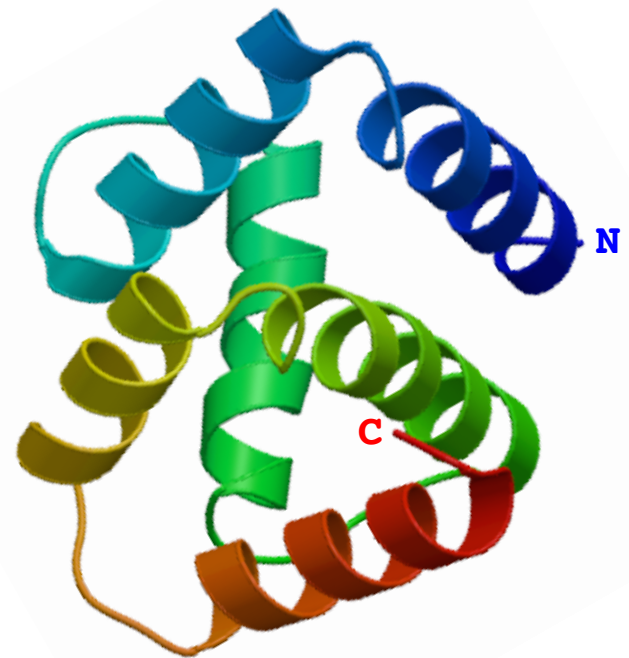
XM_018707509
Anoplophora glabripennis
cell wall integrity and stress response component 3-like
(CWP)

A0A345BEP2.1.A



GMQE 0.64
Seq Id 70.08%

1kx9.1.A

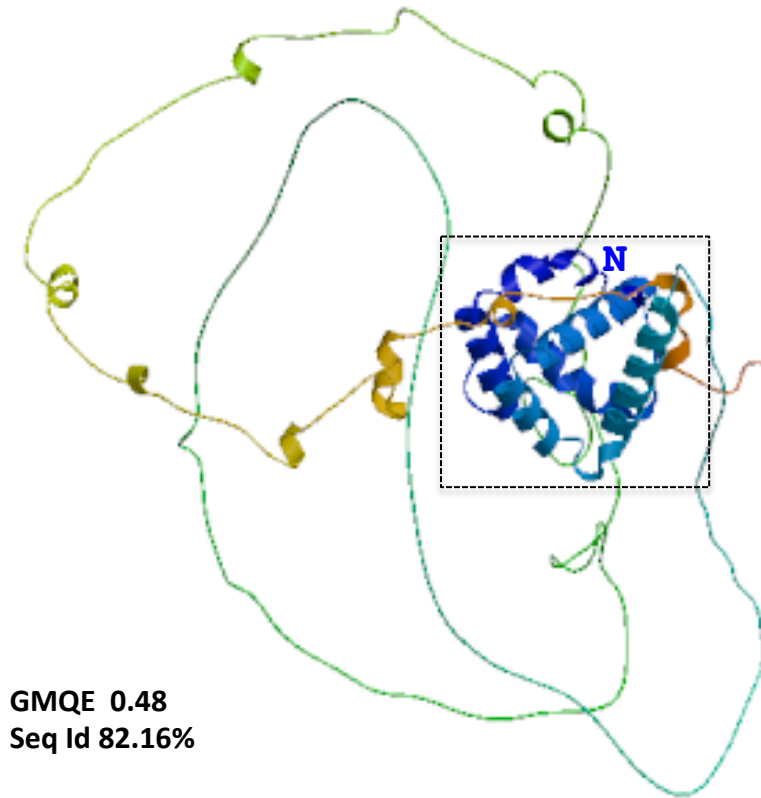


GMQE 0.24
Seq Id 39.25%

XM_048140790
Leguminivora glycinivorella
Complex-activating protein rickA-like isoform X2
mucin-2-like

A0A1Q3FQF4.1.A

1kx9.1.A



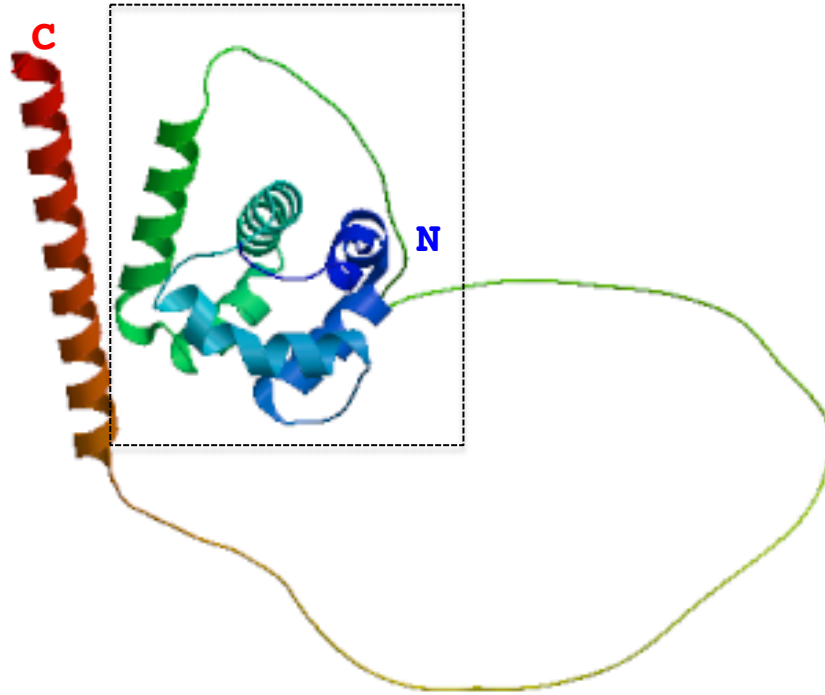
GMQE 0.48
Seq Id 82.16%



GMQE 0.13
Seq Id 30.19%

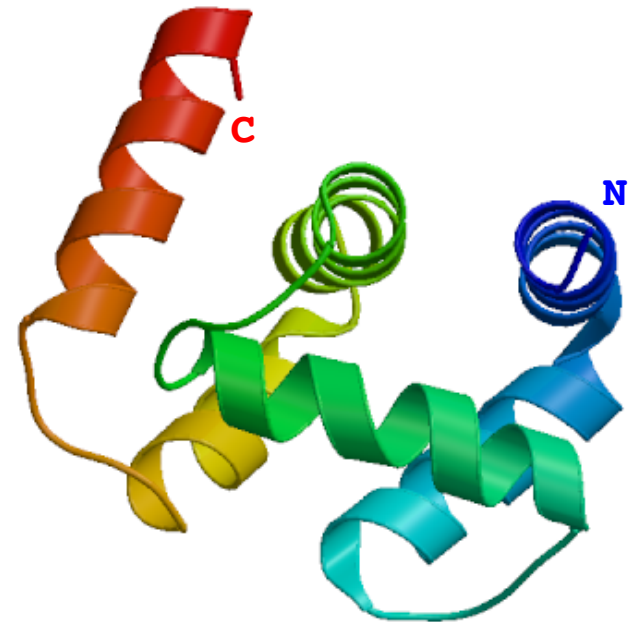
XM_038253951
Culex quinquefasciatus
cell wall protein DAN4
(CWP)

D6WUE8.1.A



GMQE 0.29
Seq Id 33.64%

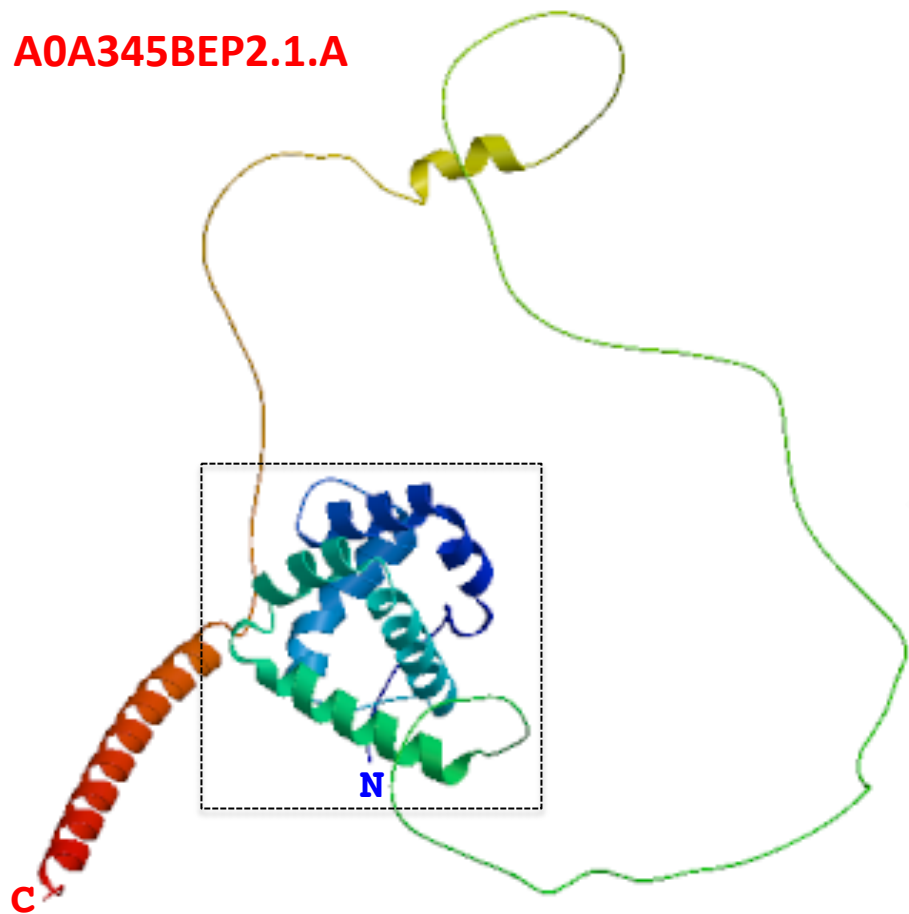
1kx9.1.A



GMQE 0.70
Seq Id 88.79%

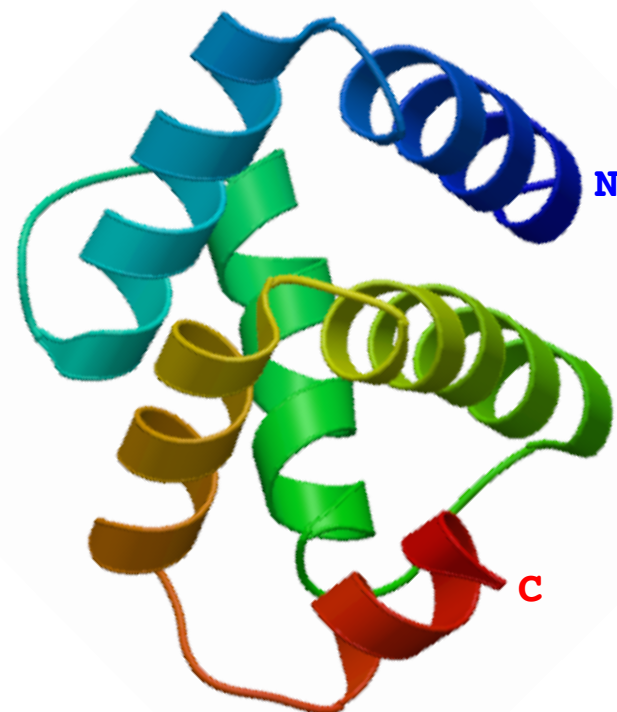
XM_044408485
Tribolium madens
Allergen Tha p 1-like
(IgE-binding protein)

A0A345BEP2.1.A



GMQE 0.63
Seq Id 75.67%

1kx9.1.A

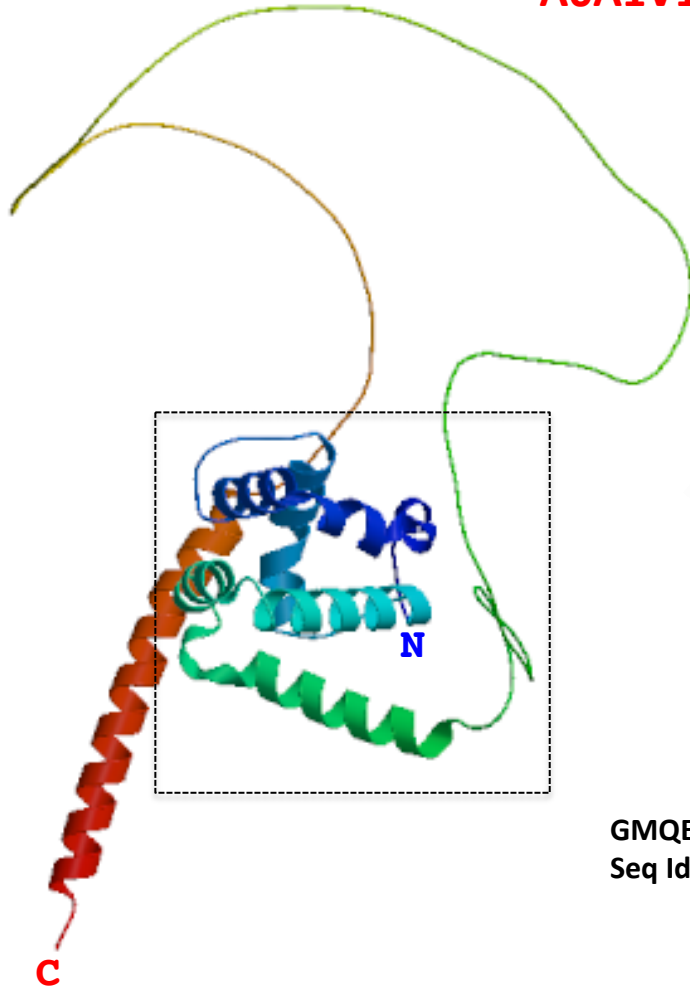


GMQE 0.22
Seq Id 38.78%

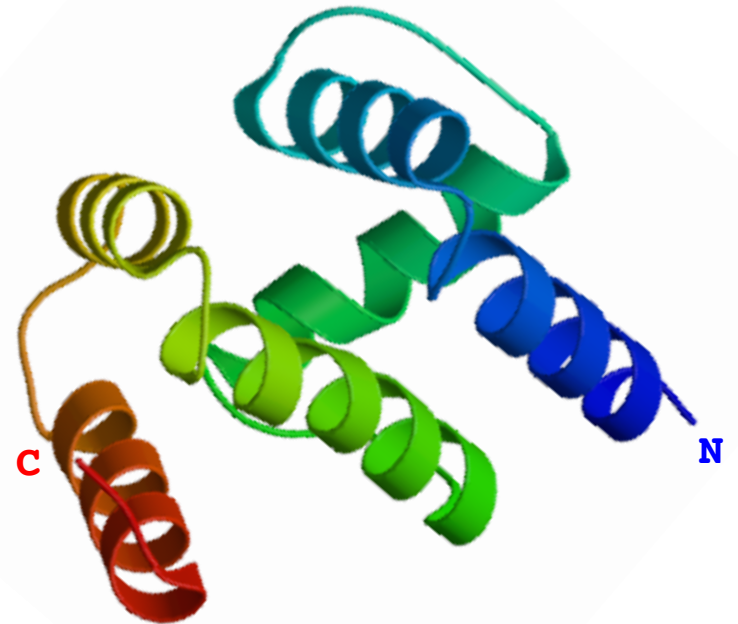
XM_053758324
Plodia interpunctella
Actin microtubule association protein
(Formin-1-like)

A0A1V1WC08.1.A

1kx9.1.A



GMQE 0.65
Seq Id 87.07%

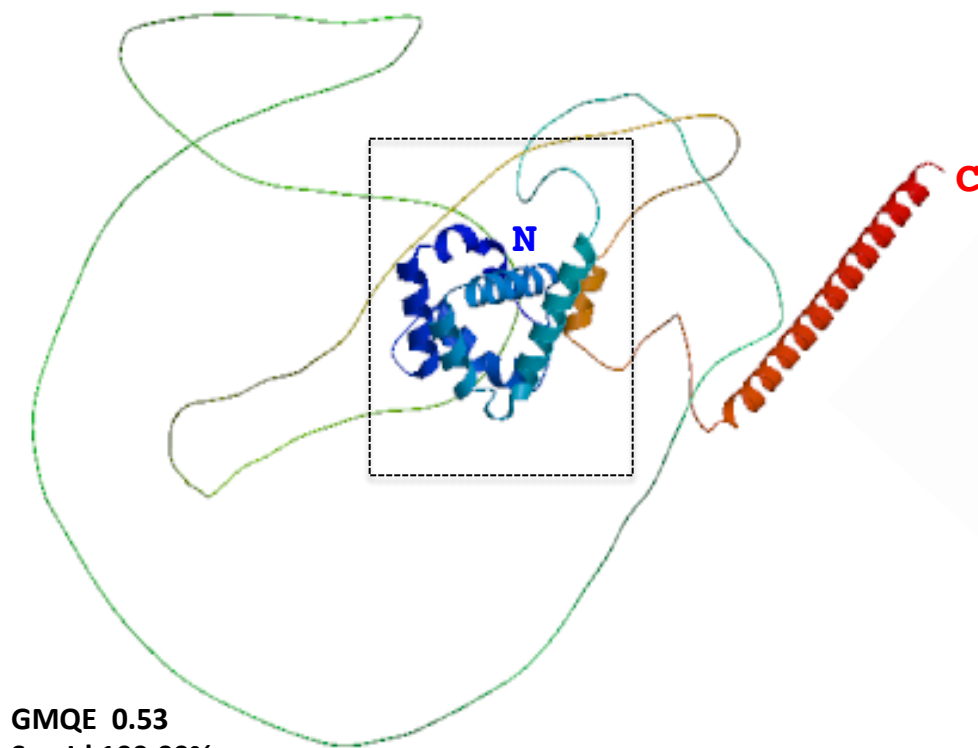


GMQE 0.26
Seq Id 35.51%

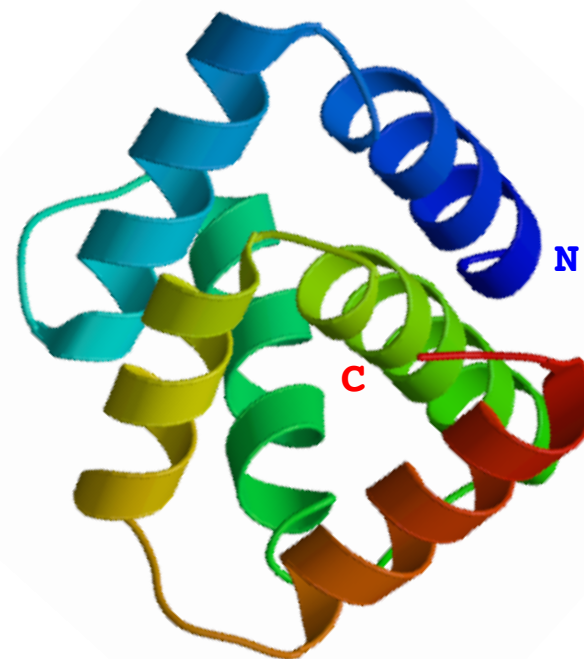
XM_022960083
Spodoptera litura
actin cytoskeleton-regulatory complex protein pan1-like
(AS RP)

AOA6I8TWU8.1.A

1kx9.1.A



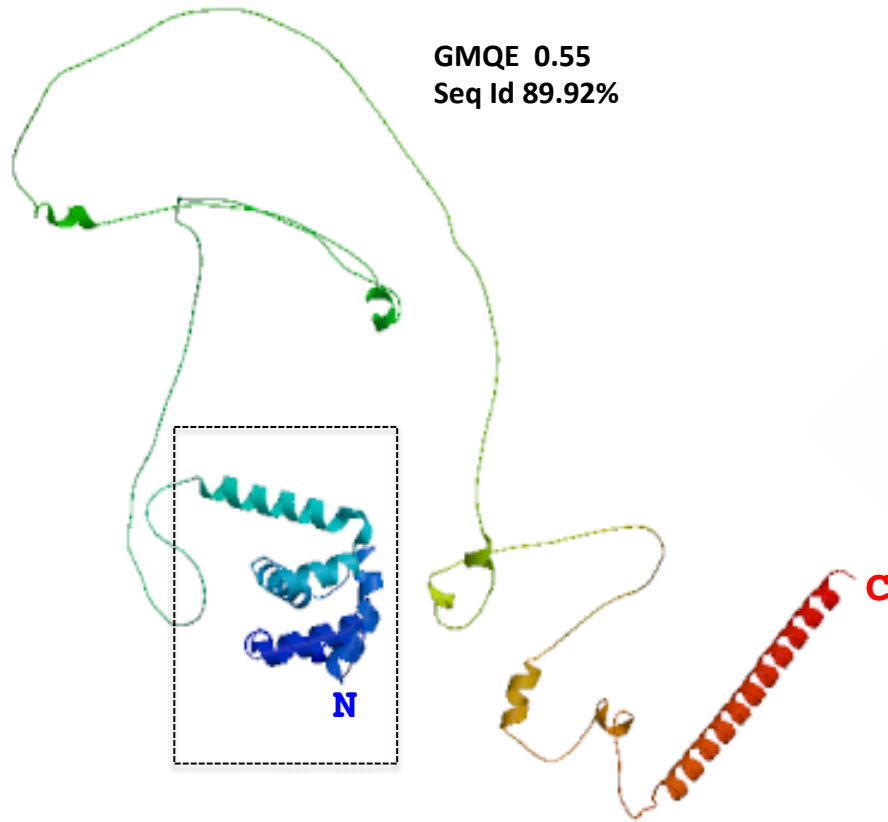
GMQE 0.53
Seq Id 100.00%



GMQE 0.14
Seq Id 30.56%

XM_021847563
Aedes aegypti
nuclear pore complex protein
(NPCP)

A0A8I8TWU8.1.A



1kx9.1.A

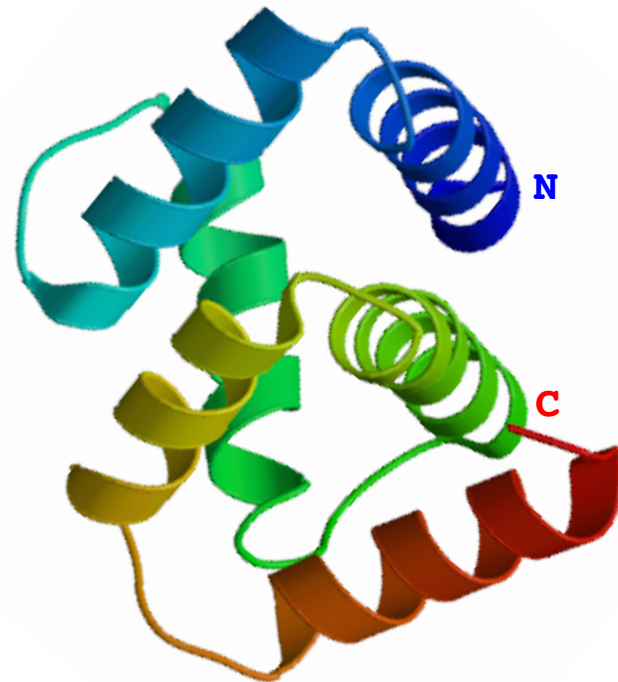
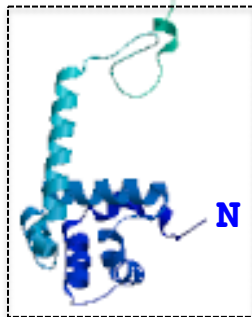


XM_029870982
Aedes albopictus
SamkC nuclear pore complex protein
(NPCP)

AOA182NU82.1.A

1kx9.1.A

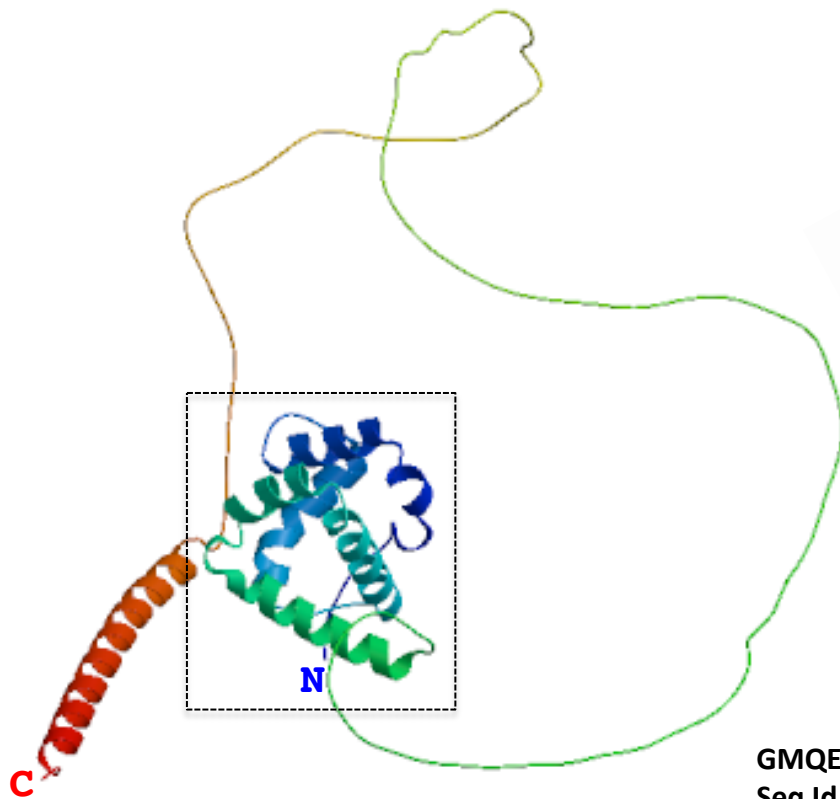
GMQE 0.53
Seq Id 79.39%



GMQE 0.14
Seq Id 32.71%

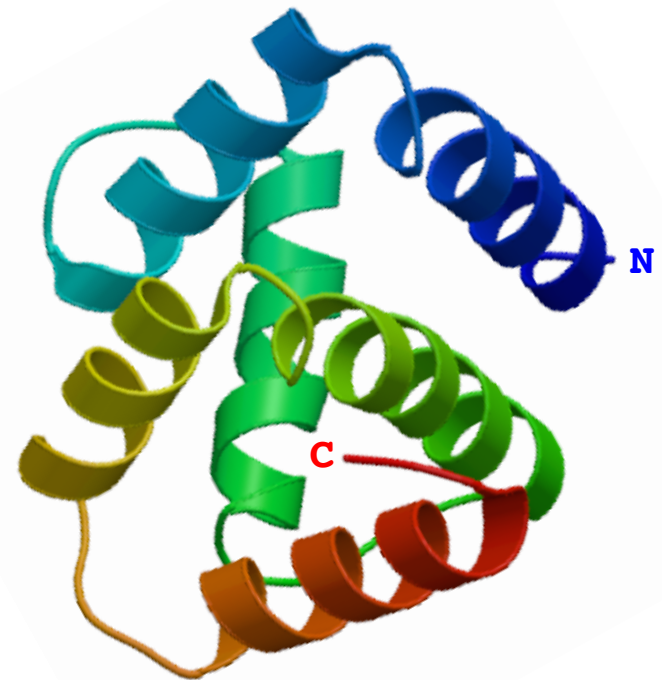
XM_035922754
Anopheles albimanus
extensin-like Leucine-Rich repeat
(LRR)

A0A345BEP2.1.A



GMQE 0.64
Seq Id 70.8%

1kx9.1.A



GMQE 0.24
Seq Id 39.25%

XM_050021059

Pectinophora gossypiella

YLP motif-containing protein 1-like

Nuclear nucleoside kinase (NNK)