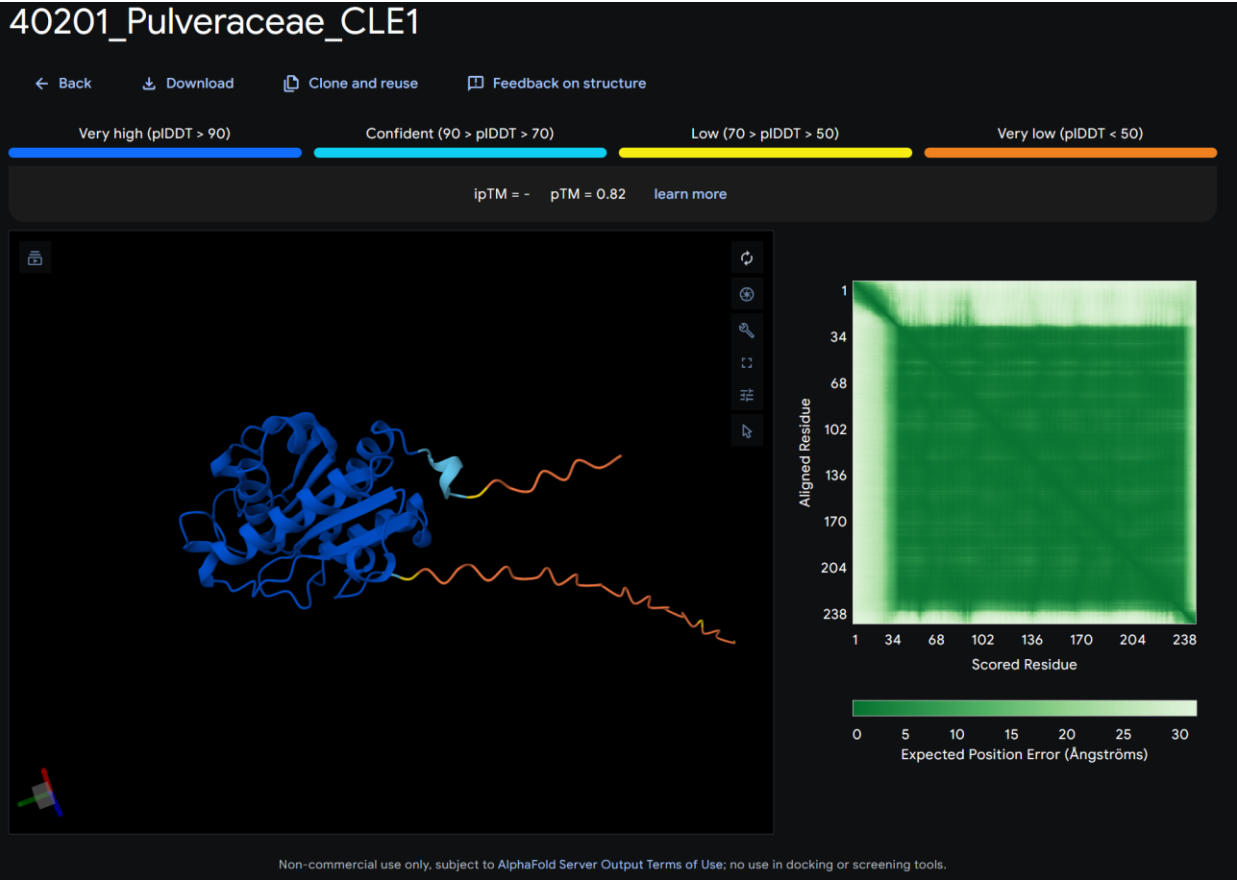
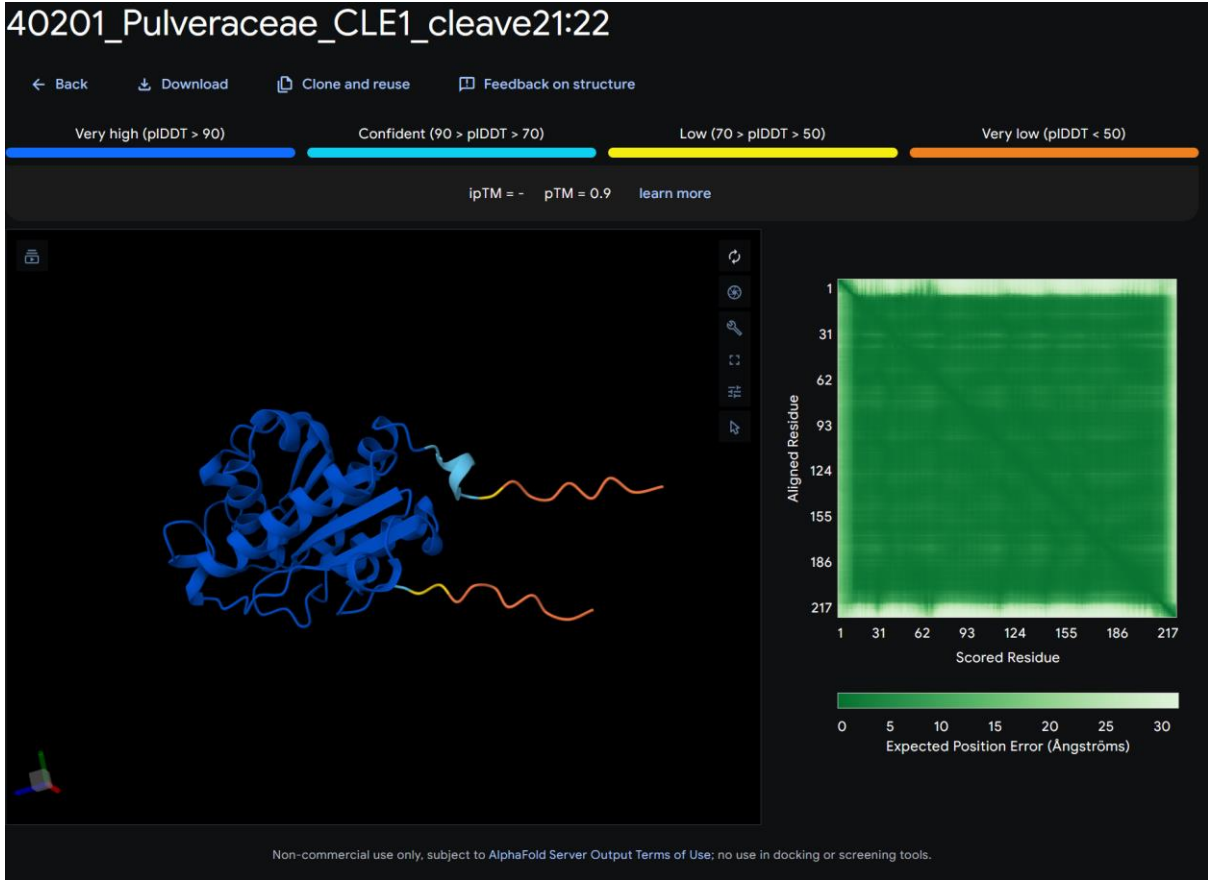


SignalP-6.0: between pos. 21 and 22  
TargetP-2.0: cleavage site pos. 21-22  
DeepSig: signal peptide 1-21  
PrediSi: [no result]

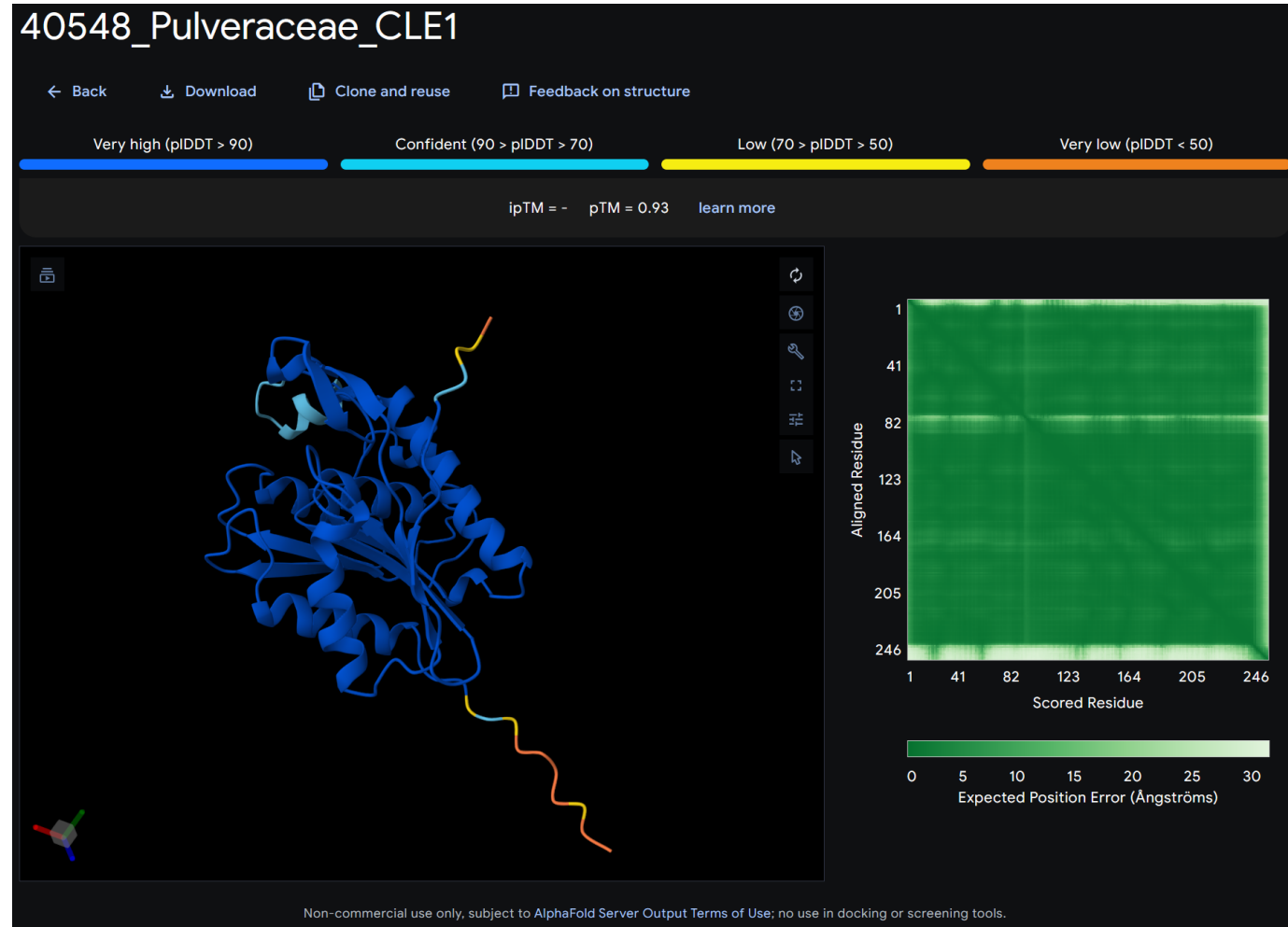


[SignalP-6.0](#): no signal, other

[TargetP-2.0](#): no signal, other

[DeepSig](#): no signal peptide, likely localization somewhere within cytoplasm

[PrediSi](#): no signal peptide, incorrect cleavage position at 37 (low score 0.2564)



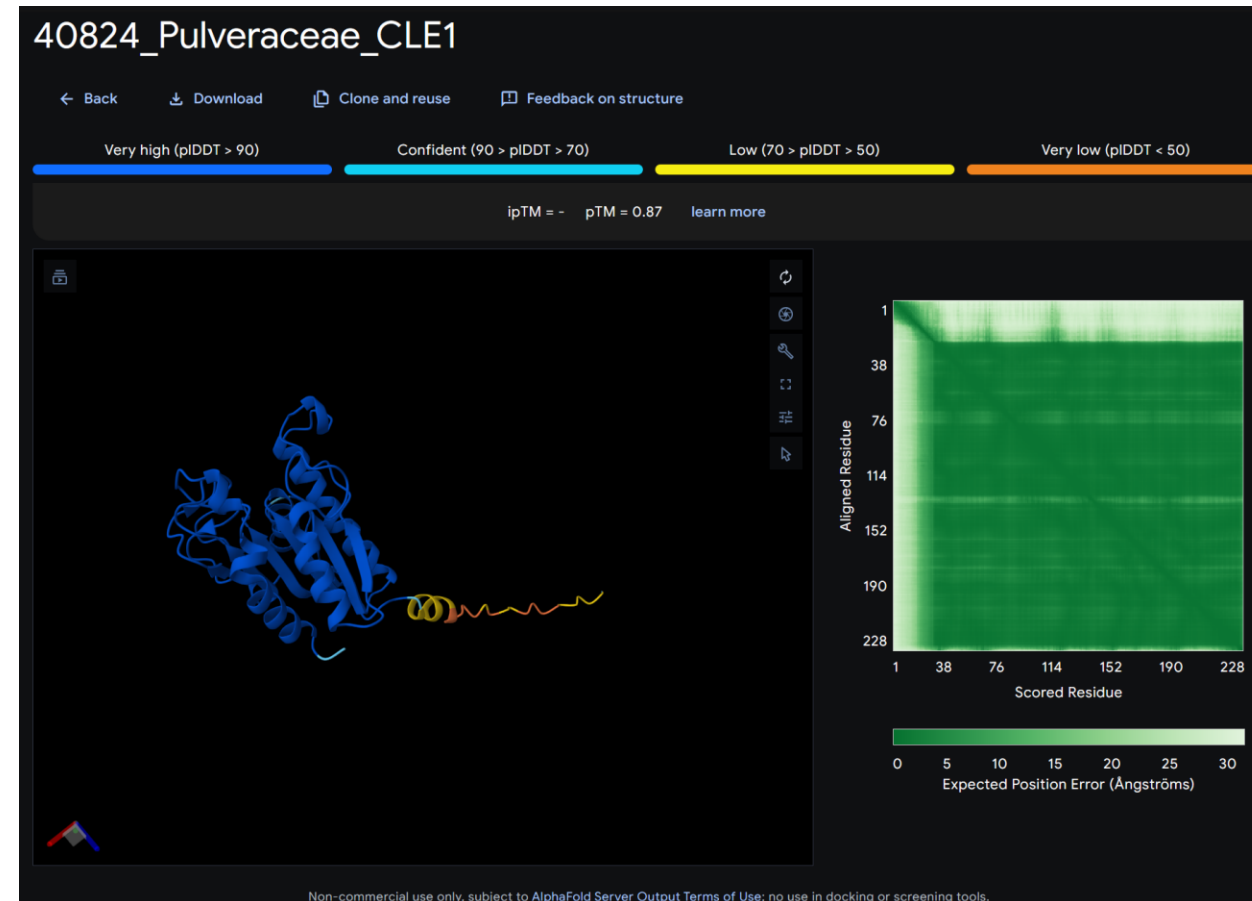
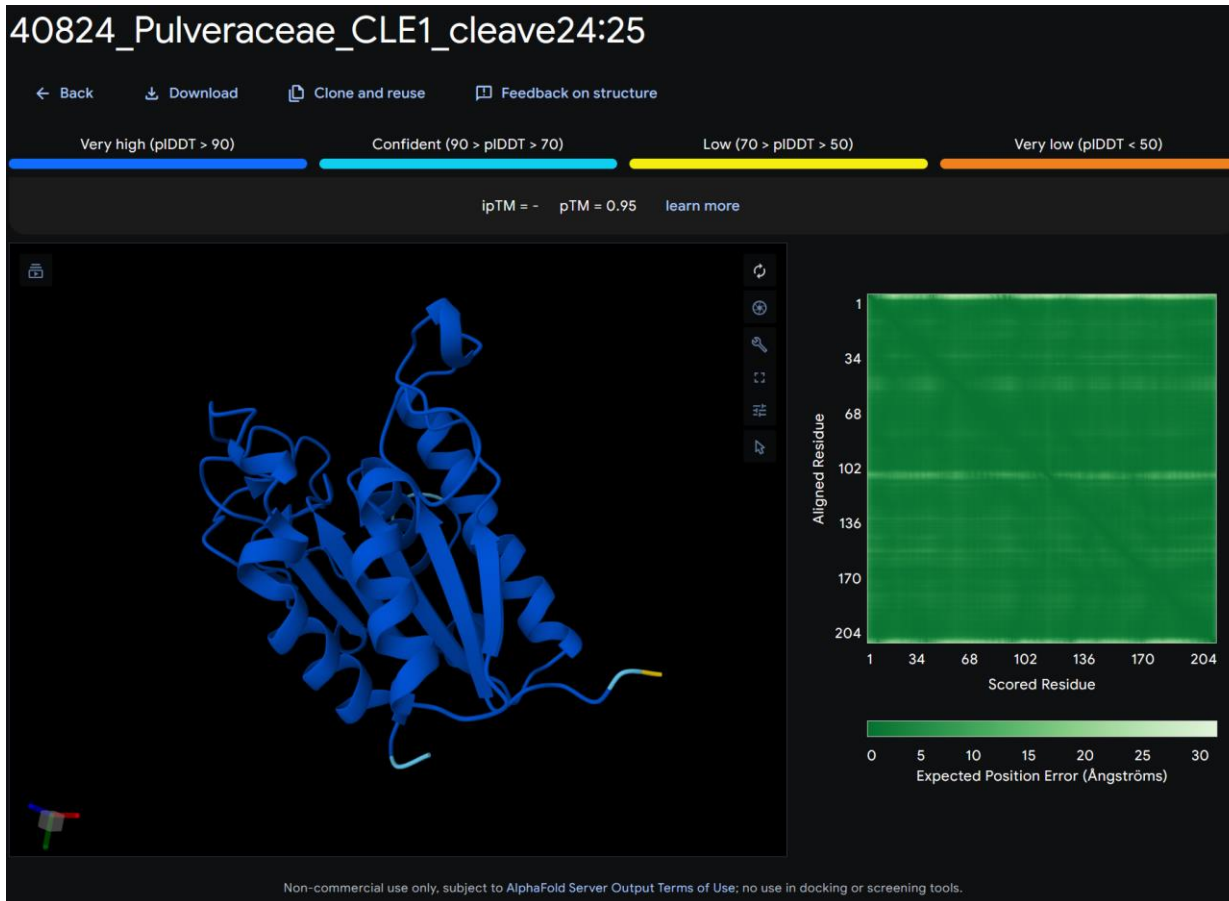
SignalP-6.0: between pos. 24 and 25

TargetP-2.0: cleavage site pos. 16-17

DeepSig: no signal peptide, likely localization somewhere within cytoplasm

PrediSi: signal peptide, cleavage site pos. 21

Cleavage below done at SignalP-6.0 site



[SignalP-6.0](#): between pos. 22 and 23

[TargetP-2.0](#): cleavage site pos. 22-23

[DeepSig](#): signal peptide 1-22

[PrediSi](#): signal peptide, cleavage site pos. 22

### 43484\_Pulveraceae\_CLE1\_cleave22:23

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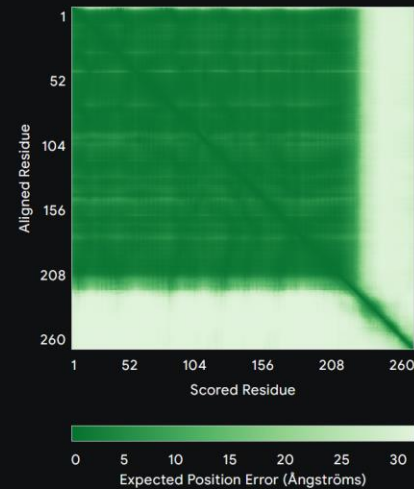
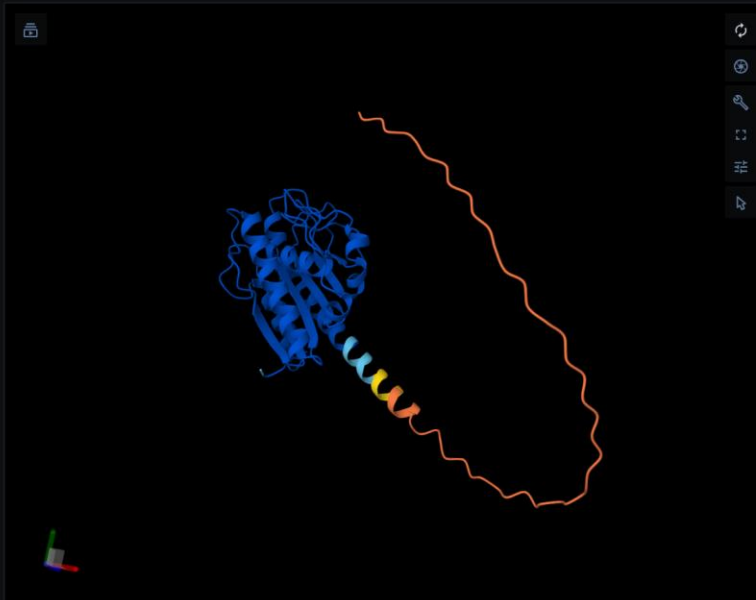
Very high (pLDDT > 90)

Confident (90 > pLDDT > 70)

Low (70 > pLDDT > 50)

Very low (pLDDT < 50)

ipTM = - pTM = 0.8 [learn more](#)



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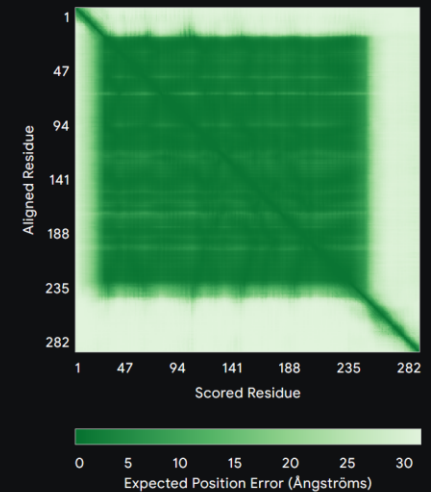
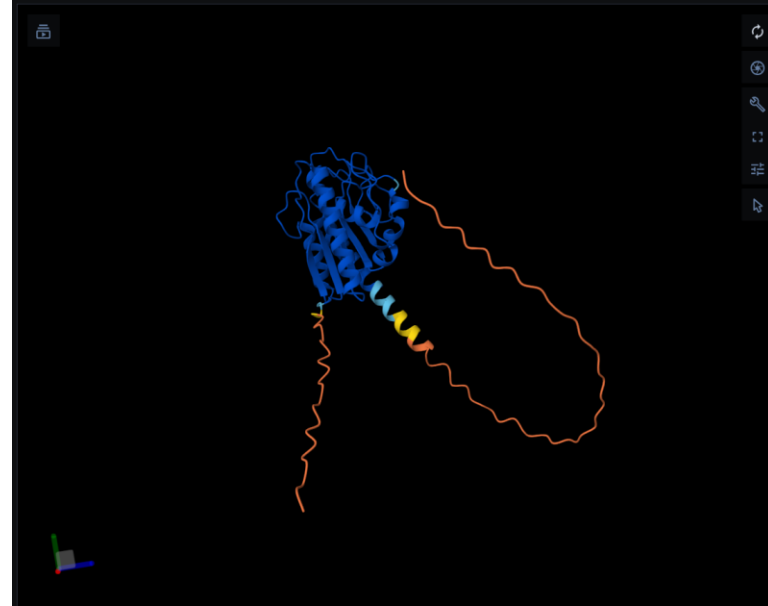
Very high (pLDDT > 90)

Confident (90 > pLDDT > 70)

Low (70 > pLDDT > 50)

Very low (pLDDT < 50)

ipTM = - pTM = 0.75 [learn more](#)



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SignalP-6.0: between pos. 21 and 22  
TargetP-2.0: cleavage site pos. 21-22  
DeepSig: signal peptide 1-21  
PrediSi: signal peptide, cleavage site pos. 21

