

SignalP-6.0: between pos. 21 and 22

TargetP-2.0: cleavage site pos. 21-22

DeepSig: signal peptide 1-21

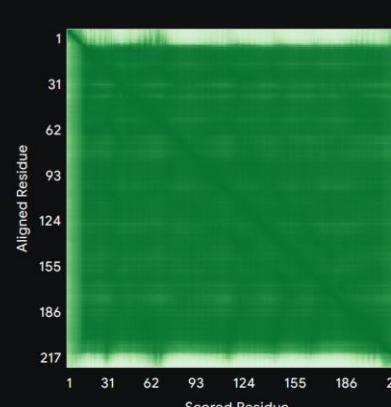
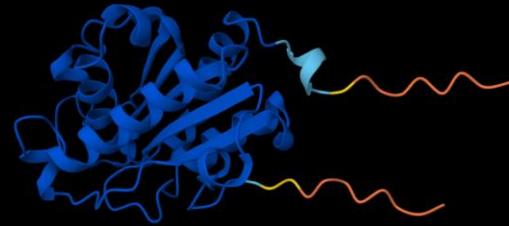
PrediSi: [no result]

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Very high (pIDDT > 90) Confident (90 > pIDDT > 70) Low (70 > pIDDT > 50) Very low (pIDDT < 50)

ipTM = - pTM = 0.9 learn more

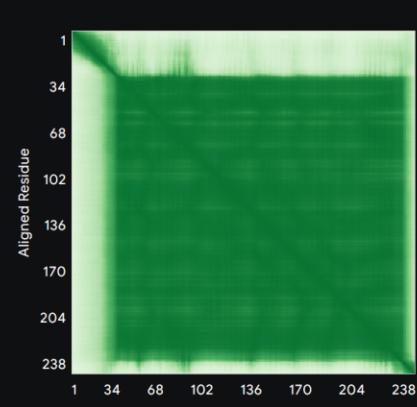
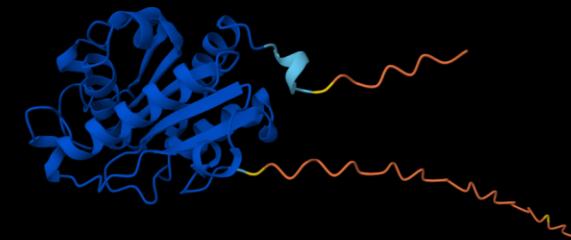


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Very high (pIDDT > 90) Confident (90 > pIDDT > 70) Low (70 > pIDDT > 50) Very low (pIDDT < 50)

ipTM = - pTM = 0.82 learn more



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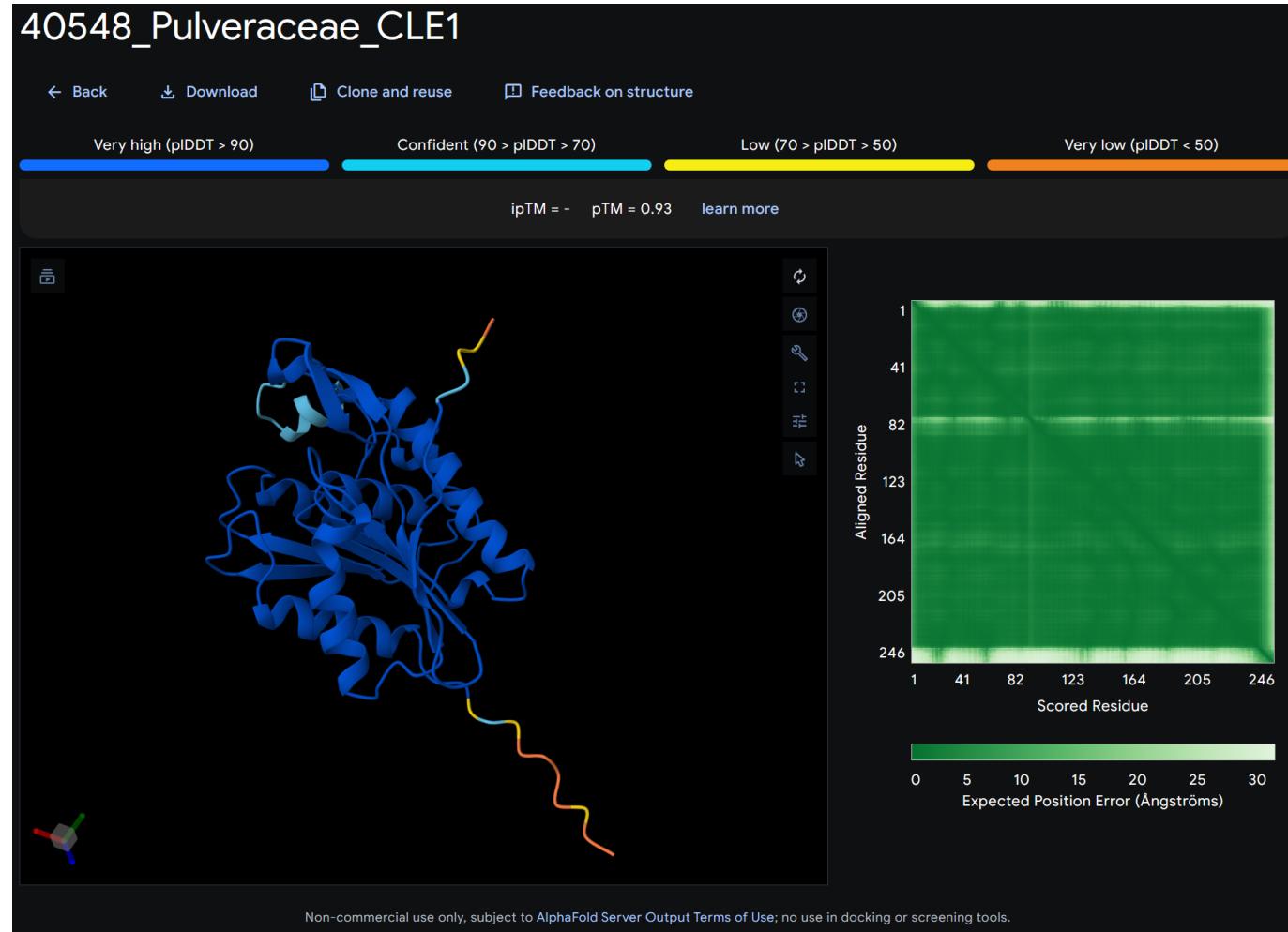
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[SignalP-6.0](#): no signal, other

[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)



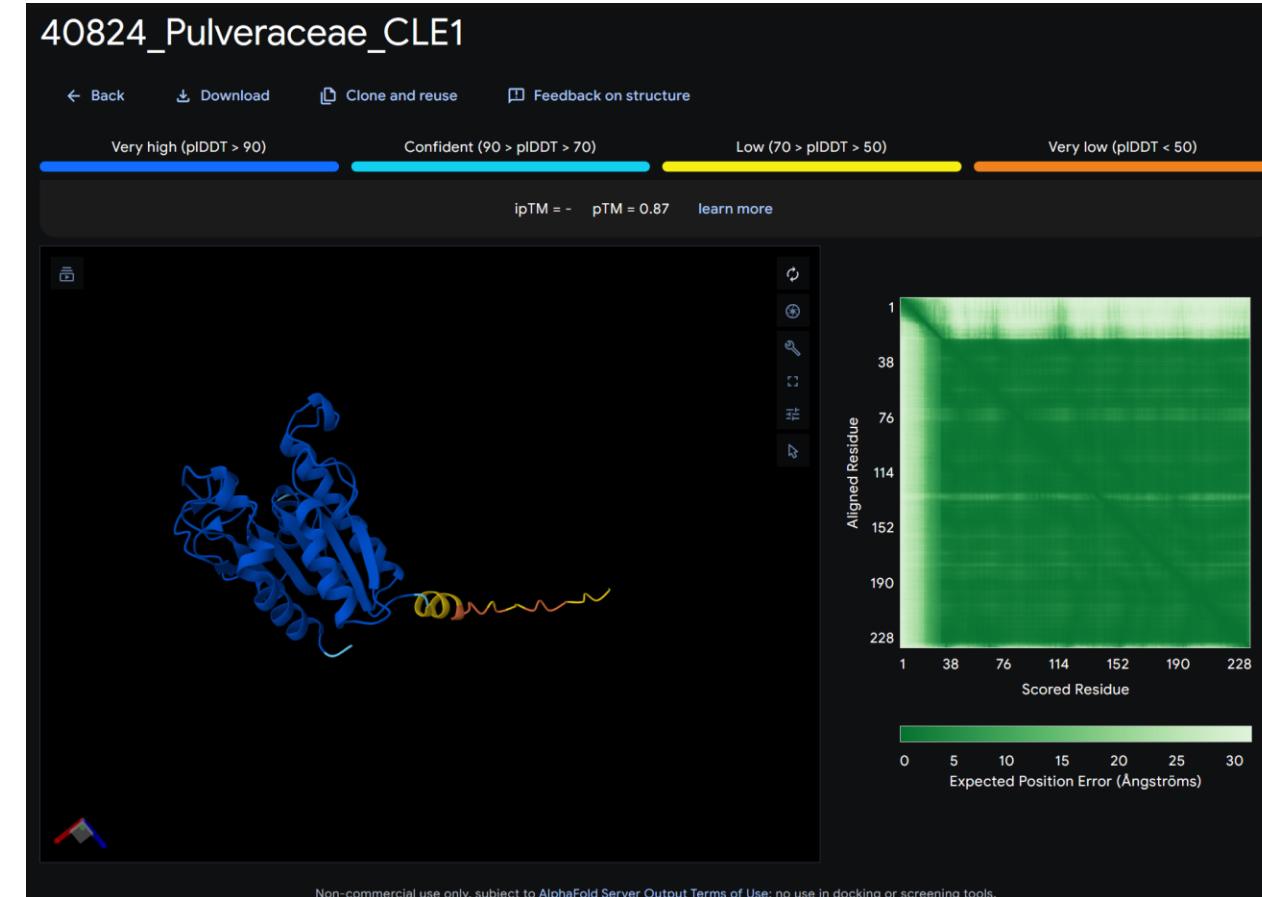
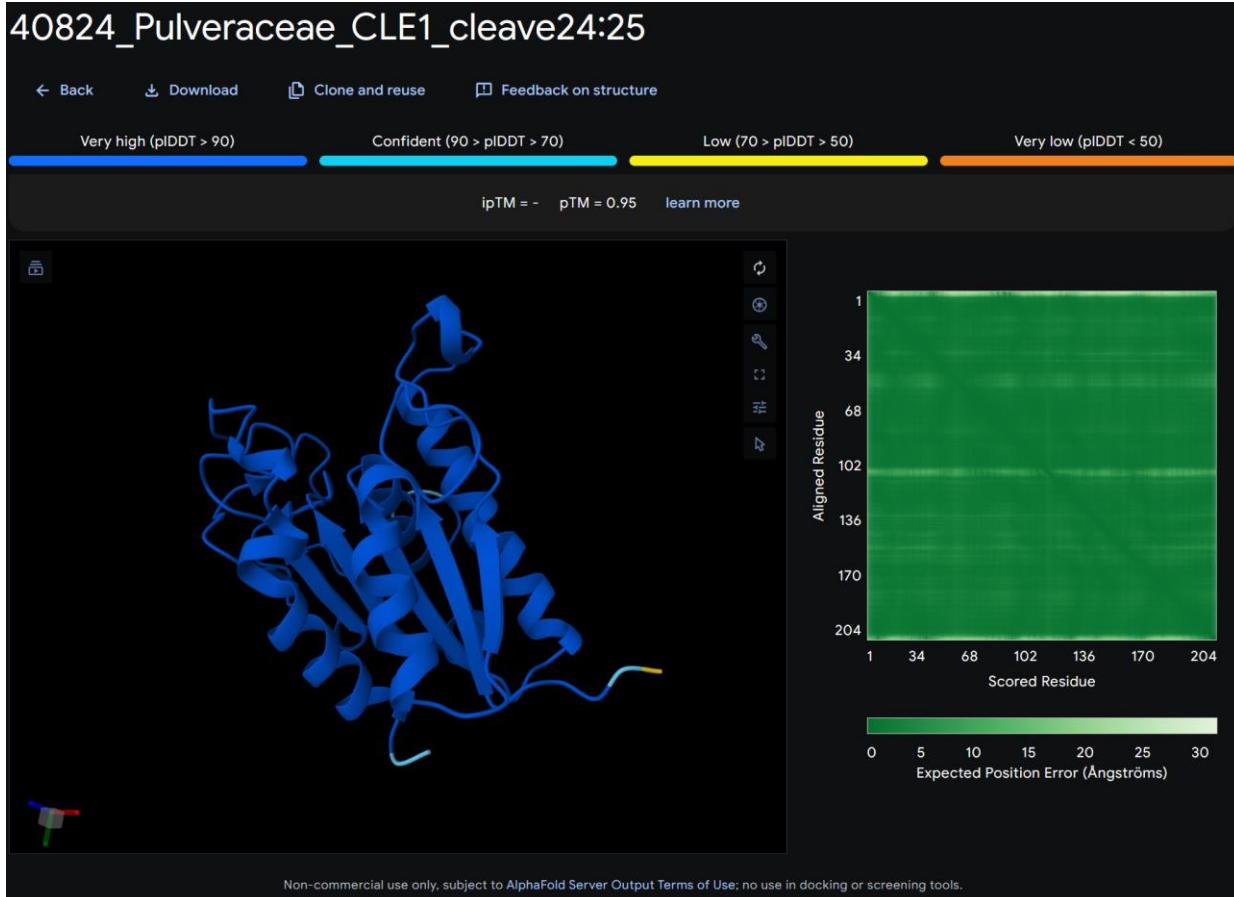
SignalP-6.0: between pos. 24 and 25

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

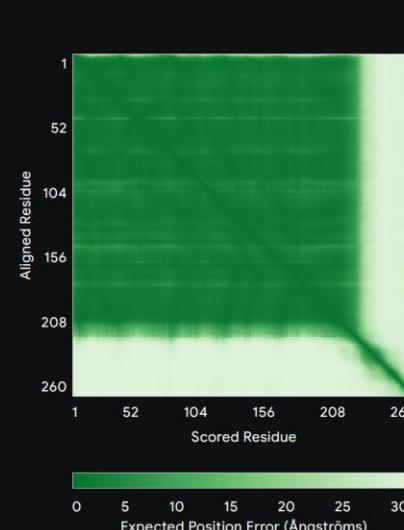
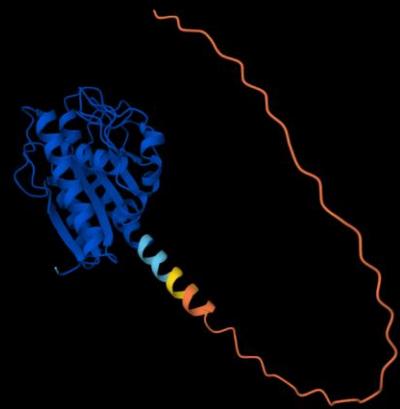
PrediSi: signal peptide, cleavage site pos. 22

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Very high (pIDDT > 90) Confident (90 > pIDDT > 70) Low (70 > pIDDT > 50) Very low (pIDDT < 50)

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

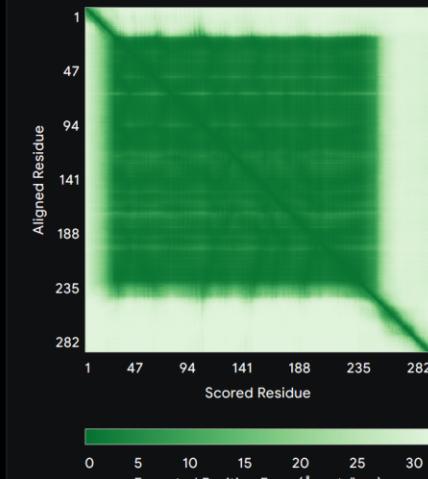
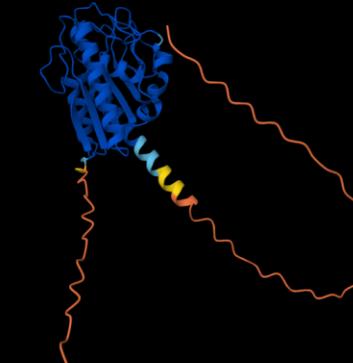


43484_Pulveraceae_CLE1

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Very high (pIDDT > 90) Confident (90 > pIDDT > 70) Low (70 > pIDDT > 50) Very low (pIDDT < 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

