

jobid=69499EA5002FE31FD57F6D82&wait=20 Server Output - DTU Health Tech

Asn-Xaa-Ser/Thr sequons in the sequence output below are highlighted in **blue**.
 Asparagines predicted to be N-glycosylated are highlighted in **red**.

Output for 'YP_003347549.1'

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#####
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Warning: This sequence may not contain a signal peptide!!

Proteins without signal peptides are unlikely to be exposed to the N-glycosylation machinery and thus may not be glycosylated (in vivo) even though they contain potential motifs.

SignalP-NN euk predictions are as follows:

# name	Cmax	pos ?	Ymax	pos ?	Smax	pos ?	Smean ?	D	?	SignalP-noTM
YP_003347549.1	0.113	32	0.112	45	0.149	31	0.111	0.111	N	0.450

SignalP output is explained at <https://services.healthtech.dtu.dk/services/SignalP-4.1/>

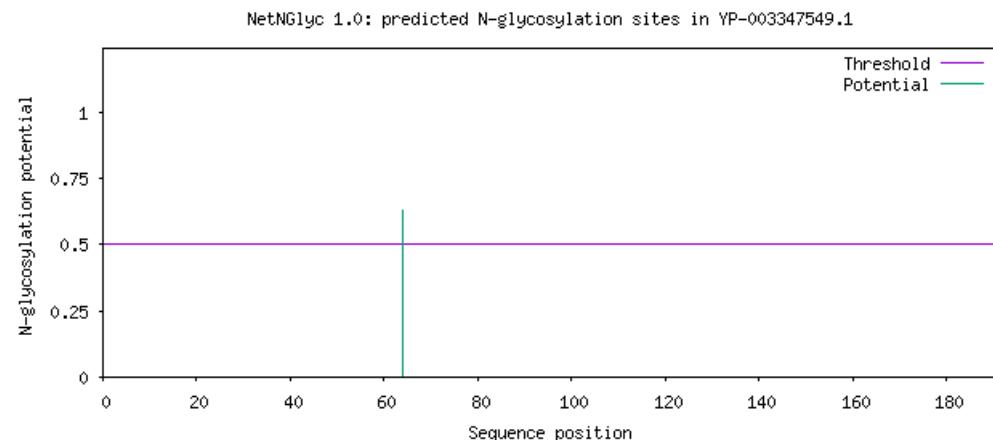
```
#####
```

Name: YP_003347549.1 Length: 192
 MMMQDAYFGSAEELDAVNEMLAAGESPVTTLDEDGSADVANARRILNRINRQIQSKGWFNI**N**ESATLTPDVSTGLIPF 80
 RPAYLSILGGQQVNRGGWVYDKSTGTDFSGPITVTLITLQDYDEMPECFRQWIVTKASRQFNSRFFGAEDVENSLAQEE 160
 MEARMACNEYEMDFGQYNMLDGDAYVQGLIGR 240
N..... 80
 160
 240

(Threshold=0.5)

SeqName	Position	Potential	Jury	N-Glyc
			agreement	result

YP_003347549.1	64	NESA	0.6311	(8/9) +
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