

I EDB Analysis Resource

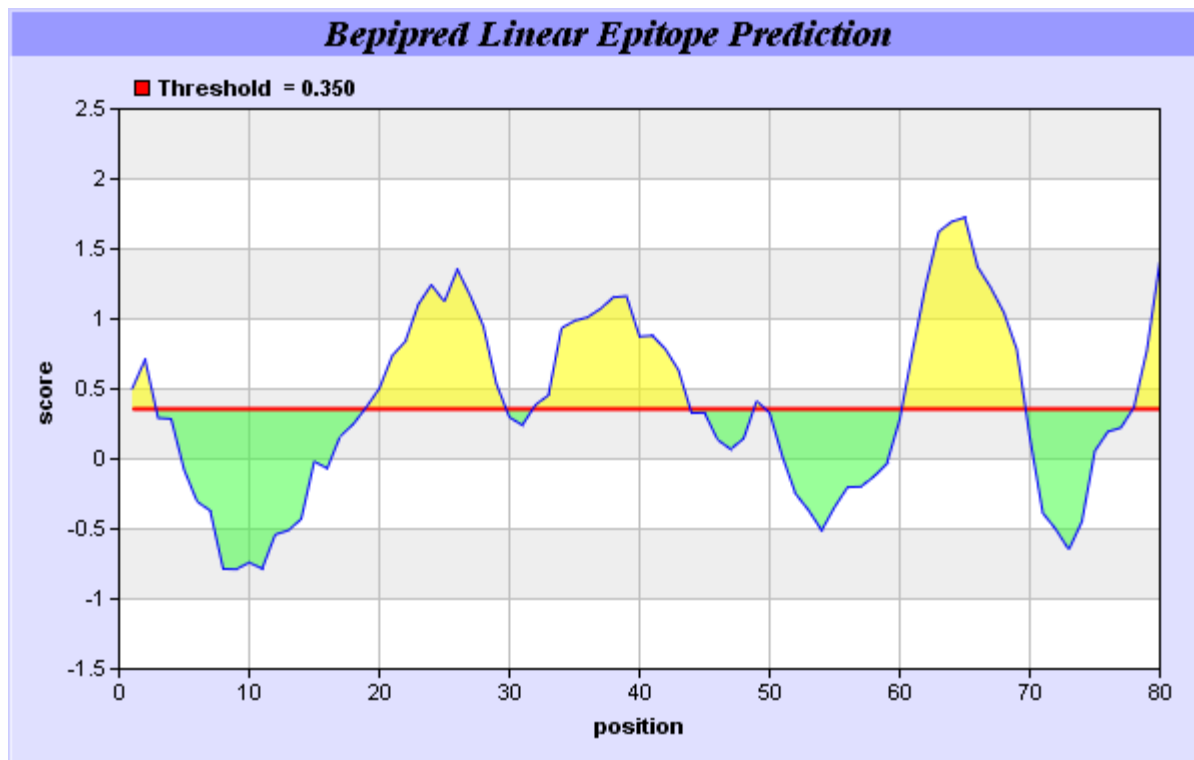
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Sequence:

```

1 MTEHTDFELL ELATPYALNA VSDDERADID RRVAAPSPV AAAFNDEVRA VRETMAVSA
61 ATTAEPFAHL RTAILDATKP EVRRQSRWR TAAFASAAAI AVGLGAFGLG VLTRPSPPT
121 VAEQVLTAPD VRTVSRPLGA GTATVVFSDR RNTGLLMN V APPSRGTV YQMWLLGGAK
181 GPRSAGTMGT AAVTPSTTAT LTDLGASTAL AFTVEPGTGS PQPTGTILAE LPLG

```



Average:0.373 Minimum:-0.788 Maximum:1.725 Threshold: 0.350

[Change](#)

[Click here to view plotted values in table format](#)

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	1	2	MT	2
2	19	29	NAVSDDERADI	11
3	32	43	RVAAAPSPVAAA	12

4	49	49	R	1
5	61	69	ATTAEPPAH	9

Reference: [Jens Erik Pontoppidan Larsen, Ole Lund and Morten Nielsen. Improved method for predicting linear B-cell epitopes. Immunome Res. 2006; 2: 2.](#)