IEDB Analysis Resource

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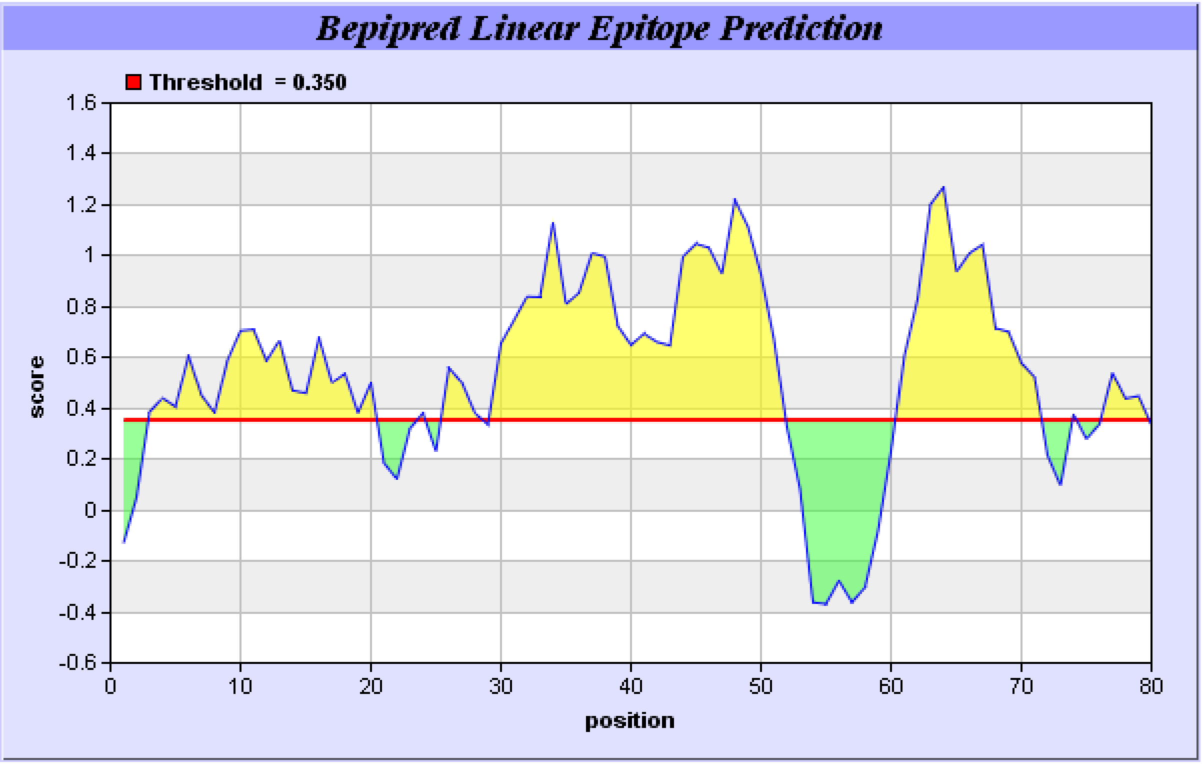
Sequence:

```
 1 MAEMKTDAAT LAQEAGNFER ISGDLKTQID QVESTAGSLQ GQWRGAAGTA AQAAVVRFQE
61 AANKQKQELD EISTNIRQAG  VQYSRADEE QQQALSSQMG F
```



Average:0.539 Minimum:-0.369 Maximum:1.27 Threshold: 0.350 Change

Click here to view plotted values in table format

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	3	20	EMKTDAATLAQEAGNFER	18
2	24	24	D	1
3	26	28	KTQ	3
4	30	51	DQVESTAGSLQGQWRGAAGTAA	22

5	61	71	AANKQKQELDE	11
6	74	74	T	1
7	77	79	RQA	3

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