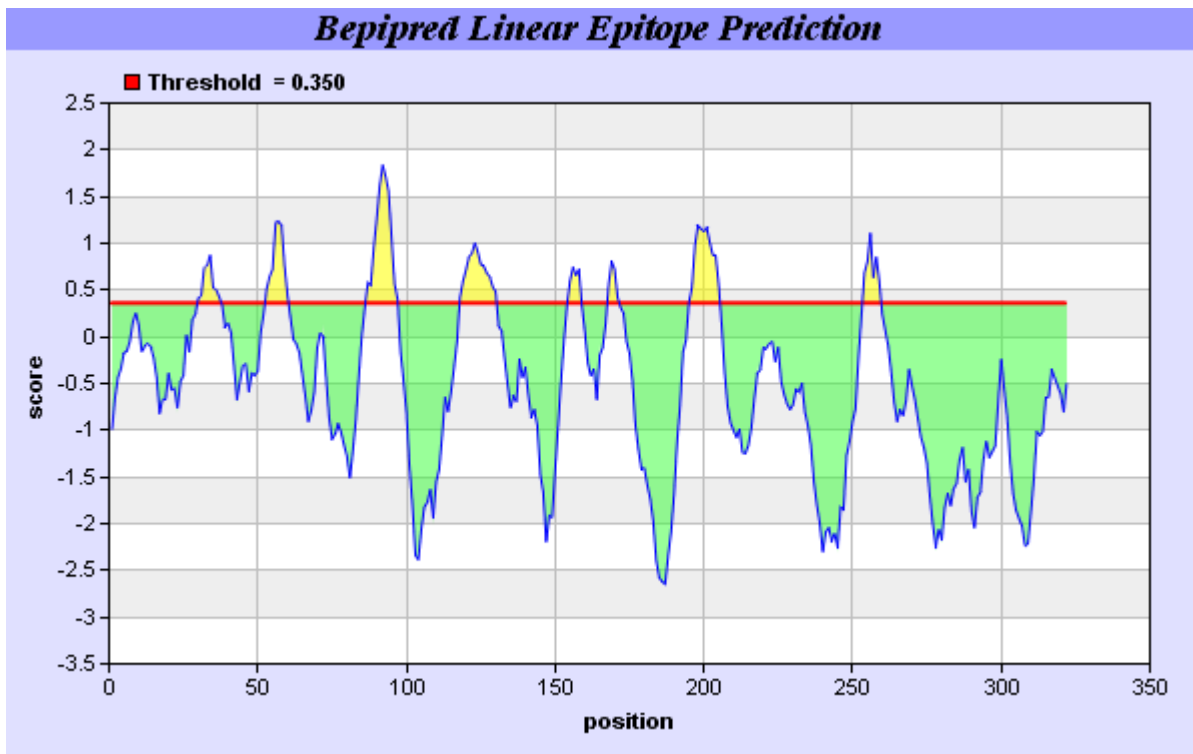


IEDB ANALYSIS RESOURCE

Sequence

```
1 MSYLAGAAQI GGVMVGAPLV IGMTRQVRAR WEGRAGAGLL QPWRDLLKQL GKQQITPAGT
  61 TIVFAAAPVI VAGTLLLIAA IAPLVATGSP LDPSADLFAV VGLLFLGTVA LTLAGIDTGT
 121 SFGGMGASRE ITIAALVEPT ILLAVFALSI PAGESANLGAL VASTIDHPGH VVSLAGVLAF
 181 VALVIVIVAE TGRLPVDNPA THLELTMVHE AMVLEYAGPR LALVEWAAGM RLTVLLALLA
 241 NLFLPWGIAG AAPTALDVLV GVVAVAAKVA ILAVLLATFE VFLAKLRLFR VPELLAGSFL
 301 LALLAVTAAN FFTVGA
```

Average:-0.519 **Minimum:**-2.646 **Maximum:**1.835 **Threshold:**



Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	30	37	RWEGRAGA	8
2	53	60	QQITPAGT	8
3	87	97	TGSPLDPSADL	11
4	118	130	TGTSFGGMGASRE	13
5	155	158	ANLG	4
6	168	171	PGHV	4
7	195	205	PVDNPATHLEL	11
8	254	259	TALDVL	6