

ABCpred Prediction Server

INPUT INFORMATION

Sequence name	
Length of the sequence	314
Number of 16mers from the input sequence	299
Threshold setting (Default value is 0.5)	0.51

TABULAR RESULT

Predicted B-cell epitope

The predicted B cell epitopes are ranked according to their score obtained by trained recurrent neural network.

Higher score of the peptide means the higher probability to be as epitope.

All the peptides shown here are above the threshold value chosen.

Rank	Sequence	Start position	Score	
1	CSHIAPANVYGPRQDP	168	0.97	
2	VGGPDDPEFHPPRLGD	259	0.93	
3	VRRSVADPQFDAAVNV	83	0.92	
4	GTPPEYPTPETAPTDP	127	0.91	
5	LLSGKPTRVFGDGTNT	197	0.89	
5	SSGGSIIYGTPEYPTP	120	0.89	
6	DGTNTRDYVVFDDVVD	208	0.88	
7	GWRPQIELADGVRRTV	290	0.86	
7	RHLYGLDCSHIAPANV	161	0.86	
8	SADVGGGLRFNIGTGK	229	0.85	
9	AGFIGSTLVDRLADG	9	0.84	

VGGPDDPEFHPPRLGD

VRRSVADPQFDAAVNV

GTPPEYPTPETAPTDP

LLSGKPTRVFGDGTNT

SSGGSYGTTPPEYPTP

DGTNTRDYVFVDDVVD

GWRPQIELADGVRRTV

RHLYGLDCSHIAPANV

SADVGGGLRFNIGTGK

AGFIGSTLVDRLLDG

TAPTDPASPYAAGKVA

DGVRRTVEYFRHKHTD

RKIVHTSSGGSYGTTP

VRLAEAAARQTGVRKIV

AILEQHRPEVVFHLAA

AGEIYLNTFRHLYGLD

FATGRATNLEHLADNS

FHPPRLGDLKRSCLDI

YGPRQDPHGEGVVAI

SPYAAGKVAGEIYLNT

FNIGTGKETSQRQLHS

HSVGLDNFATGRATN

YVFVDDVVDVAFVRSVA

SCLDIGLAERVLGWRP

VDRLLADGHSVVGLDN

FHLAAQIDVRRSVADP

ALVTGAAGFIGSTLVD

KETSDRQLHSAVAAAV

VFVEADIVTADLHAIL

LEHLADNSAHVFVEAD

LAERVLGWRPQIELAD