

ABCpred Prediction Server

INPUT INFORMATION

Sequence name	
Length of the sequence	265
Number of 16mers from the input sequence	250
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	RGSKDSDWEPVKEAAK	94	0.93	
2	GDIIWAPAIIDGAFVLT	201	0.88	
2	SETSDHGYPIREHLNR	182	0.88	
3	FVLTRGGDFDLQLGT	213	0.85	
4	GYASHDTDTVRLYLQE	233	0.84	
4	TLPEDPREIPDVISQA	141	0.84	
5	SRNEIDDVERGSKDSD	85	0.83	
6	REHLNRLVDGDIWAP	192	0.81	
7	GRRVVVDVSDPGGPVTA	33	0.80	
7	YSAASIEGIRSASSNP	123	0.80	
8	FKRHIAGRVVVDVSDP	27	0.79	

9	KPLVRLRVPFTLSRNE	73	0.78	
10	TGRLIDVKAPTNGVIA	52	0.77	
10	TEAAWAEIELEAARTF	12	0.77	
10	VKEAAKKLAFVEDRTI	104	0.77	
11	SELRLAGVDGPYSVLL	158	0.74	
12	LQETLTFLCYTAEASV	246	0.72	
12	DVYTKVSETSDHGYPI	176	0.72	
13	GVDGPYSVLLSADVYT	164	0.71	
14	LEAARTFKRHIAGR	21	0.70	
14	EGIRSASSNPALTLPE	129	0.70	
14	AFVEDRTIFEGYSAAS	112	0.70	
15	SVLLSADVYT	170	0.67	
16	DVISQALSELRLAGVD	151	0.62	

OVERLAP DISPLAY

MNNLYRDLAPVTEAAWAEIELEAARTFKRHIAGRVVVDSDPGGPVTAAVSTGRLIDVKAPTNGVIAHLRASKPLVRLRVPFT
LSRNEIDDVERGSKDSDWEPVKEAAKKLAFVEDRTIFEGYSAASIEGIRSASSNPALTPEDPREIPDVISQALSELRLAGVD
GPYSVLLSADVYTKVSETSDHGYPIREHNLVDGDIIWAPAIDGAFVLTRGGDFDLQLGTDVAIGYASHDTVRLYQLQET
LTFLCYTAEASVALSH²⁶⁵

RGSKDSDWEPVKEAAK

GDIIWAPAIIDGAFVLT

SETSDHGYPIREHLN

FVLTTTRGGDFDLOLGT-

GYASHDTDTVRLYLYQE

TLPEDPREIPDVISQA-

SRNEIDDVERGSKDSD

REHLNRLVDGDIIWAP

GRVVVDVSDPGGPVTA-

YSAASIEGIRSASSNP

FKRHIAGRVVVDVSDP-

KPLVRLRVPFTLSRNE-

TGRLIDVKAPTNVIA-

TEAAWAEIELEAARTF

VKEAAKKLAFVEDRTI

SELRLAGVDGPYSVLL

LQETLTFLCYTAEASV-

DVYTKVSETSDHGYPI

GV DGP YSV LLSAD VYT

LEAARTFKRHIAGRRV

EGIRSASSNPALTLPE

AFVEDRTIFEGYSAAS

SVLLSADVYTKVSETS

DVISQALSELRLAGVD