

Predictions for positions 1 through 100

Threshold for intolerance is 0.05.

Amino acid color code: nonpolar, **uncharged polar**, **basic**, **acidic**.

Capital letters indicate amino acids appearing in the alignment, lower case letters result from prediction.

'Seq Rep' is the fraction of sequences that

contain one of the basic amino acids. A low fraction indicates the position is either severely gapped or unalignable and has little information. Ex

Predict Not Tolerated	Position	Seq Rep	Predict Tolerated
ywvtsrqpnmlkihgfedca	1M	1.00	M
ywvtsrqpnmlkihgfedca	2G	1.00	G
ywvtsrqpnmlkihgfedc	3A	1.00	A
ywvtsrqpnmlkihgfedca	4R	1.00	R
ywvtsrqpnmlkihgfedc	5A	1.00	A
ywvtrqpnmlkihgfedca	6S	1.00	S
hwqdnpr egcksyf mta	7V	1.00	LIIV
ywvtsrqpnmlkihgfedca	8L	1.00	L
w	9R	1.00	cfmyihvlpdgnqetakSR
ywvtsrqpnmlkihgfedca	10G	1.00	G
w	11E	1.00	cmfiyhlpvrtqksnaGdE
cwfmyivhpgldnsta	12K	1.00	erQK
ywvtsrqpnmlkihgfedca	13L	1.00	L
ywvtsrqpnmlkihgfeca	14D	1.00	D
w	15T	1.00	cfymhiplvgndqresATK
yvtsrqpnmlkihgfedca	16W	1.00	W
ywvtsrqpnmlkihgfedca	17E	1.00	E
cwfmdyivgphslnta	18R	1.00	eqKR
ywvtsrqpnmlkihgfedca	19I	1.00	I
cwfmdyivgphslnta	20K	1.00	eqKR
ywvtsrqpnmlkihgfedca	21L	1.00	L
cwfmdyivgphslnta	22K	1.00	eqKR
ywvtsrqpnmlkihgfedca	23P	1.00	P
ywvtsrqpnmlkihgfedca	24G	1.00	G
wmfihclrqvekp	25G	1.00	tdnaSG
ywvtsrqpnmlkihgfedca	26K	1.00	K
ywvtsrqpnmlkihgfedca	27K	1.00	K
	28R	1.00	wCmfipHvygdlnsQaeRK
wvtsrqpnmlkihgfedca	29Y	1.00	Y
cw	30M	1.00	fyidvpghMnlstaeqRK
dhgncewsrkpyqtaf	31L	1.00	viML
ywvtsrqpnmlkihgfedca	32K	1.00	K
ywvtsrqpnmlkihgfedca	33H	1.00	H
dhgncesrkpyqta	34L	1.00	fmvIL
ywvtsrqpnmlkihgfedca	35V	1.00	V
yvtsrqpnmlkihgfedca	36W	1.00	W
ywvtsrqpnmlkihgfedc	37A	1.00	A
ywvtrqpnmlkihgfedca	38S	1.00	S
ywvtsrqpnmlkihgfedca	39R	1.00	R
ywvtsrqpnmlkihgfedca	40E	1.00	E
ywvtsrqpnmlkihgfedca	41L	1.00	L
ywvtsrqpnmlkihgfedca	42E	1.00	E
ywvtsrqpnmlkihgfedca	43R	1.00	R
ywvtsrqpnmlkihgfedca	44F	1.00	F
ywvtsrqpnmlkihgfedc	45A	1.00	A
dhwgncersqpkyta	46L	1.00	fmiVL
wmifcvlyrphqatk	47D	1.00	esgDN
ywvtsrqpnmlkihgfedca	48P	1.00	P
ywvtsrqpnmlkihgfedca	49G	1.00	G
ywvtsrqpnmlkihgfedca	50L	1.00	L
ywvtsrqpnmlkihgfedca	51L	1.00	L
ywvtsrqpnmlkihgfedca	52E	1.00	E
ywvtsrqpnmlkihgfedca	53T	1.00	T
whyfmicrqldnke	54S	1.00	vptgSA
wcfmy	55E	1.00	ihvlpvrtsgqankDE
ywvtsrqpnmlkihgfedca	56G	1.00	G
whdnrqyekfmgps	57V	1.00	tliaVC
wcfym	58N	1.00	ihvpigdtsaenQRK
cwfmyiv	59K	1.00	hpgldntsaerKQ