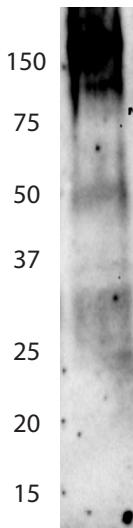


Rhesus



child1>head>body>style @type="http://w3.org/ns/hbox-wo/; break-word; white-space: pre-wrap;">#CLDRD (1 1 2 4) multiple sequence alignment	
Phe	MEELPSKGVVPLVCAZGEEVLLILFVPLVGSVTPDASRASLILPFTASQGNNYHQQEE
antigen	60
Phe	SEELSLQGVVPPQVNBLNLLIIGPFLMLTFLXGGCAGLNLLNPGSLLVQHNPAGPQGLUN
antigen	120
Phe	TANQAGAQNVEHVEIPVGAVTLEPOTCPANLISLQAGLKVTPFLVLLAUVEEVKE
antigen	180
Phe	AMENEGVQVQHAAAGCCTPFIWVQVATLAAVQVQVQVQVQVQVQVQVQVQVQVQVQVQ
antigen	240
Phe	WHEEYVPEAQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQVQ
antigen	50
Phe	SIFVTVNGSPLAVVTEBTATITVIALCGVVAFLPQALQDSGRSCHLWHRQ
antigen	100
Phe	TLAQGQVQGAGAAAGCQGGALLIACQACQLCVQSNLPGVQVQVQVQVQVQVQVQVQ
antigen	160
Phe	MHQGAGACAGAGAGVLTATESYGGDQHQSFSNSPAQGQGQGQGQGQGQGQGQGQ
antigen	220
Phe	MGLAEIATPCINTQZQFLQFSPVNECEEDQDIEHNGEJLQJDFL
antigen	460

```

<html><head><style>*{font-family: arial;}</style></head><body><pre style="word-wrap: break-word; white-space: pre-wrap;">CLUSTAL_OU(1.2.4) multiple sequence alignment

antigen
Nv_Bhl
MADDTGTCRGCFCUULLPQQLLILYVVFVVSQDVQGQNLNTIISFFPVVQIICVLL
0

antigen
L1VGHGLMTTAYLFCGSGLQWNLCAIACVNNATAGQGTTTFSRNNLQIGDLSKKGOLA
0
120

antigen
AVTTCAGCAGGCTTGGCGGTTTGTATTTTGTATTTTGTTGTTGTTGTTGTTGTTGTTG
31
140

antigen
GLAVAGLIVLGSL
50
55

Nv_Bhl
GIVAVVLYHSSVVAEAEKARREKTFDLFATGTLFLWVTFSFSLVNLVQQLSRAVNT
240

antigen
TYTAAASCVTTAATSSLTAAKSLNVEHQHTLAGQVANIIAMIBPGAILLNG
50
300

antigen
GIISTQKYYTWPEIMHLTHDVGCGHSLGKQGCPGQIJAASIAISVSSCLAGQPGVTSLYT
50
360

Nv_Bhl
VPPASRPVPTVQYIYLQYPTVPAAGLQGATPATQDGQVQAHVPLHHSRGGLITPFI
50
410

```

Rhesus



NKA

