BcePred Prediction Server

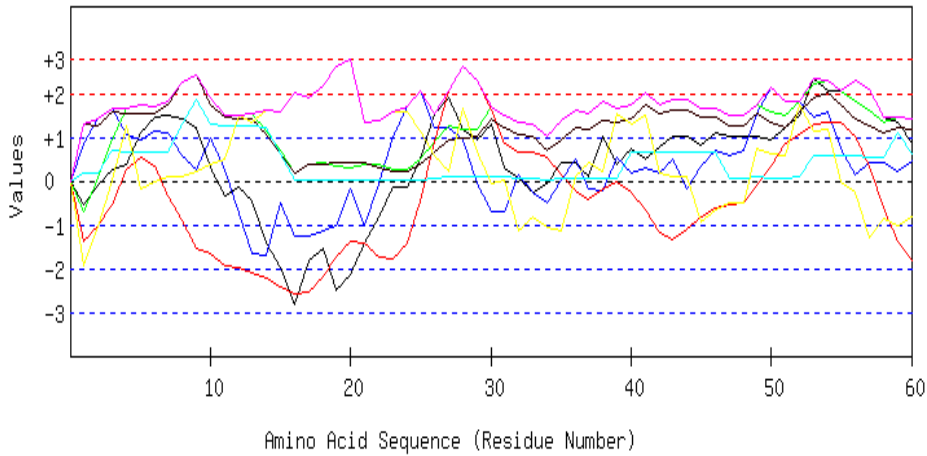The server displays 1.GRAPHICAL RESULT 2.TABULAR RESULT   3.Overlap Display

seqname=
Seq=LLTGNKPAVQRRFIGLLMLSVLVAGCSSNPLANFAPGYPPTIEPAQPAVSPPTSQDPAGAVRPLSGHPRAALF
DNGTRQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSDDHGTAFLAARGGYFVADLSSGHTARVNV
ADAAHTDFTAIARRSDGKLVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFARVDALVTQGNTTVVLDRGQTSV
TTIGADGHAQQALRAGQGATTMAADPLGRVLIADTRGGQLLVYGVDPLILRQAYPVRQAPYGLAGSRELAWVSQ
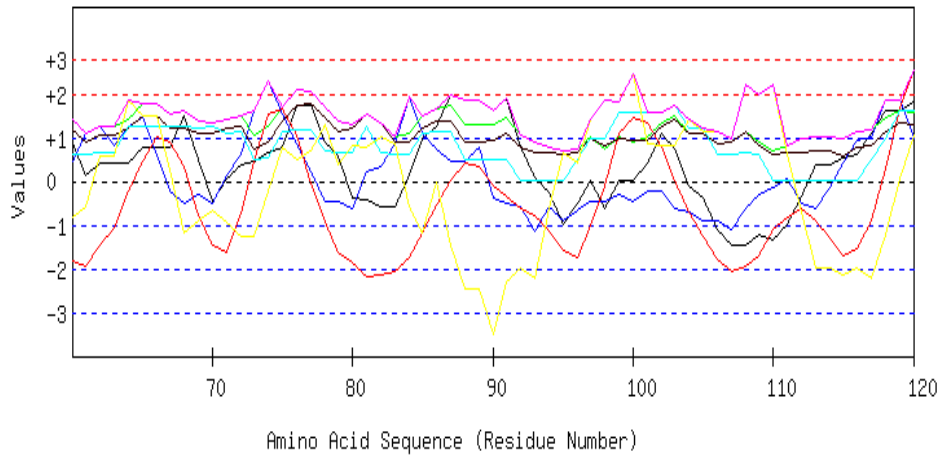TASNTVIGYDLTTGIPVEKVRYPTVQQPNSLAFDETSDTLYVVSGSGAGVQVIEHAAGTR

Length=358

GRAPHICAL RESULT

GRAPHICAL RESULT :: SEQ 1 to 60

Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 61 to 120

Values

+3
+2
+1
0
-1
-2
-3

70   80   90   100   110   120

Amino Acid Sequence (Residue Number)

Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 121 to 180

Values

+3
+2
+1
0
-1
-2
-3

130   140   150   160   170   180

Amino Acid Sequence (Residue Number)

Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 181 to 240

Amino Acid Sequence (Residue Number)

Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

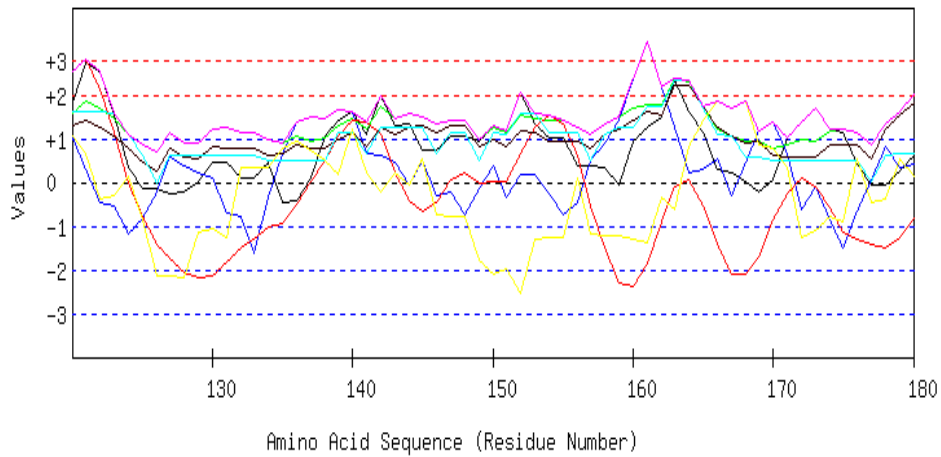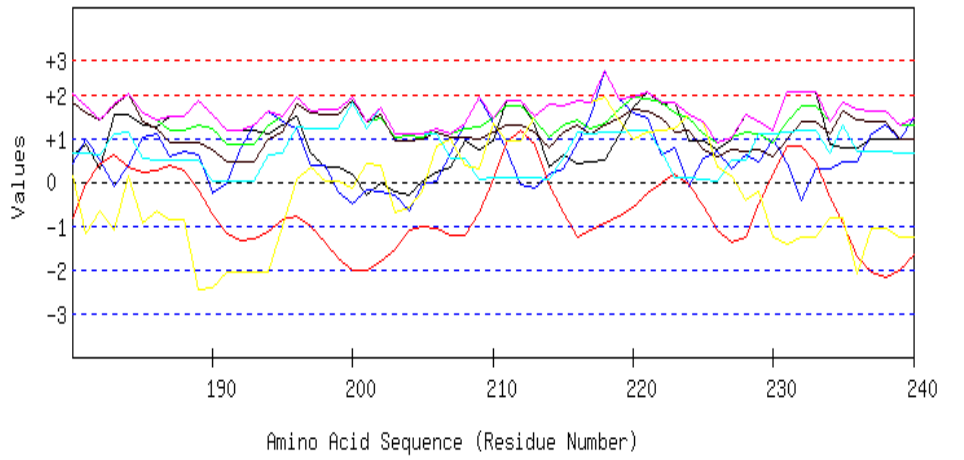GRAPHICAL RESULT :: SEQ 241 to 300

Amino Acid Sequence (Residue Number)

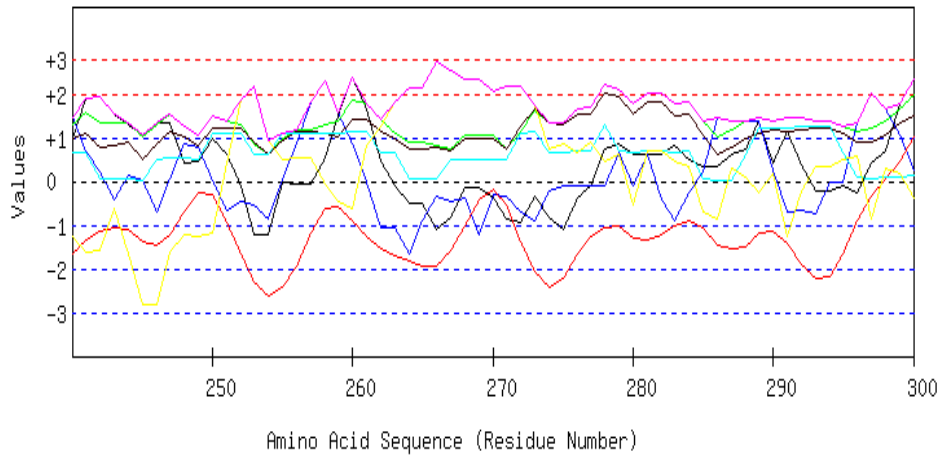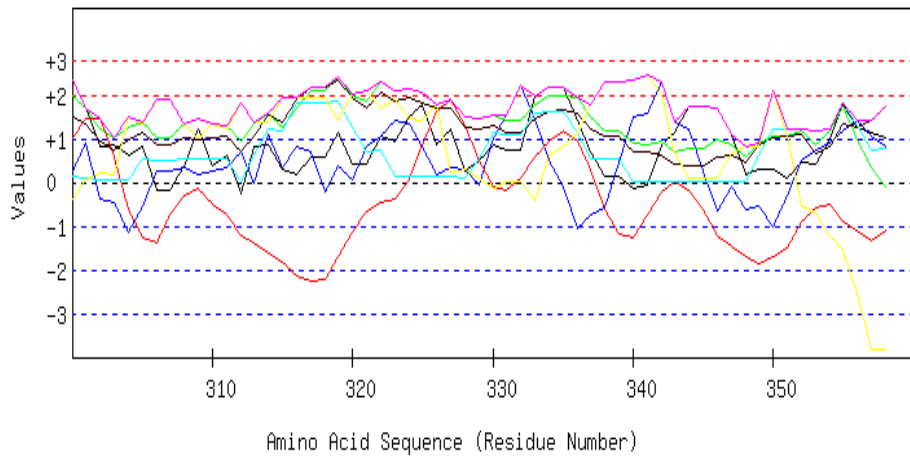Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 301 to 360

Amino Acid Sequence (Residue Number)

Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

TABULAR RESULT

Selected Programs: hydro flexi access turns surface polar antipro

Respective Threshold: 1.9 2 1.9 2.4 2.3 1.8 1.9

LLTGNKPAVQRRFIGLLMLSVLVAGCSSNPLANFAPGYPPTIEPAQPAVSPPTSQDPAGA
VRPLSGHPRAALFDNGTRQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSD
DHGTAFLAARGGYFVADLSSGHTARVNVADAAHTDFTAIARRSDGKLVLGSADGAVYTLA
KNPAVDPASGAATVASRTKIFARVDALVTQGNTTVVLDRGQTSVTTIGADGHAQQALRAG
QGATTMAADPLGRVLIADTRGGQLLVYGVDPLILRQAYPVRQAPYGLAGSRELAWVSQTA
SNTVIGYDLTTGIPVEKVRYPTVQQPNSLAFDETSDTLYVVSGSGAGVQVIEHAAGTR

Length=358

A.A.					**Parameter**					**Combined**	
	Hydro	Flexi	Access	Turns	Surface	Polar	AntiPro	MAX	MIN	AVG
1 L	-0.553	0.876	-0.710	-1.376	1.285	0.181	-1.938	1.285	-1.938	-0.319
2 L	-0.111	1.439	0.047	-1.032	1.276	0.161	-1.040	1.439	-1.040	0.106
3 T	0.250	1.644	0.954	-0.530	1.595	0.696	0.035	1.644	-0.530	0.663
4 G	0.383	1.056	1.655	0.276	1.549	0.655	1.265	1.655	0.276	0.977
5 N	1.097	0.920	1.739	0.548	1.540	0.649	-0.180	1.739	-0.180	0.902
6 K	1.445	1.125	1.702	0.345	1.522	0.646	-0.027	1.702	-0.027	0.965
7 P	1.495	1.107	1.832	-0.344	1.741	0.668	0.096	1.832	-0.344	0.942
8 A	1.401	0.574	2.272	-0.910	2.260	1.293	0.095	2.272	-0.910	0.998

9 V	1.224	0.249	2.403	-1.545	2.424	1.877	0.207	2.424	-1.545	0.977
10 Q	0.281	0.972	1.889	-1.657	1.741	1.286	0.393	1.889	-1.657	0.701
11 R	-0.357	0.277	1.505	-1.938	1.449	1.269	0.495	1.505	-1.938	0.386
12 R	-0.129	-0.741	1.496	-1.997	1.403	1.269	1.505	1.505	-1.997	0.401
13 F	-0.477	-1.668	1.533	-2.094	1.422	1.273	1.351	1.533	-2.094	0.191
14 I	-1.438	-1.699	1.122	-2.220	1.057	1.236	1.623	1.623	-2.220	-0.046
15 G	-1.969	-0.518	0.683	-2.404	0.629	0.629	1.569	1.569	-2.404	-0.198
16 L	-2.817	-1.242	0.169	-2.564	0.164	0.009	2.005	2.005	-2.817	-0.611
17 L	-1.824	-1.242	0.384	-2.528	0.364	0.025	1.913	1.913	-2.528	-0.415
18 M	-1.552	-1.133	0.403	-2.195	0.373	0.025	2.180	2.180	-2.195	-0.271
19 L	-2.494	-1.019	0.328	-1.724	0.428	0.030	2.615	2.615	-2.494	-0.262
20 S	-2.146	-0.188	0.290	-1.397	0.410	0.026	2.769	2.769	-2.146	-0.034
21 V	-1.432	-1.007	0.375	-1.436	0.401	0.021	1.324	1.324	-1.436	-0.251
22 L	-0.806	-0.056	0.375	-1.734	0.310	0.004	1.379	1.379	-1.734	-0.075
23 V	-0.136	1.004	0.244	-1.767	0.209	0.016	1.566	1.566	-1.767	0.162
24 A	-0.136	1.710	0.244	-1.427	0.209	0.016	1.566	1.710	-1.427	0.312
25 G	0.509	2.068	0.515	-0.460	0.373	0.035	1.138	2.068	-0.460	0.597
26 C	1.533	1.237	0.898	0.888	0.674	0.070	0.590	1.533	0.070	0.841
27 S	1.900	1.201	1.262	2.048	0.957	0.087	0.221	2.048	0.087	1.097
28 S	1.186	0.954	1.178	2.630	0.966	0.093	1.666	2.630	0.093	1.239
29 N	0.958	-0.076	1.188	2.310	1.011	0.093	0.656	2.310	-0.076	0.877
30 P	1.312	-0.685	1.702	1.619	1.412	0.116	-0.080	1.702	-0.685	0.771
31 L	0.319	-0.685	1.487	0.845	1.212	0.100	0.012	1.487	-0.685	0.470
32 A	0.041	0.147	1.337	0.644	1.057	0.080	-1.158	1.337	-1.158	0.307
33 N	-0.269	-0.260	1.281	0.656	1.020	0.058	-0.825	1.281	-0.825	0.237
34 F	-0.041	-0.510	1.029	0.544	0.701	0.039	-1.045	1.029	-1.045	0.102
35 A	0.421	0.023	1.365	0.129	0.929	0.053	-1.148	1.365	-1.148	0.253
36 P	0.421	0.513	1.608	-0.220	1.203	0.072	0.082	1.608	-0.220	0.526
37 G	0.111	-0.170	1.552	-0.431	1.166	0.050	0.415	1.552	-0.431	0.385
38 Y	1.021	-0.222	1.814	-0.186	1.367	0.066	0.205	1.814	-0.222	0.581
39 P	0.383	0.544	1.674	-0.027	1.349	0.068	1.536	1.674	-0.027	0.789
40 P	0.743	0.185	1.758	-0.255	1.440	0.648	1.290	1.758	-0.255	0.830
41 T	0.515	0.317	2.010	-0.659	1.759	0.667	1.510	2.010	-0.659	0.874
42 I	0.768	0.185	1.758	-1.166	1.522	0.648	0.168	1.758	-1.166	0.555
43 E	1.015	0.509	1.842	-1.327	1.622	0.672	0.111	1.842	-1.327	0.635
44 P	1.015	-0.162	1.842	-1.090	1.622	0.672	0.111	1.842	-1.090	0.573
45 A	0.819	0.335	1.646	-0.802	1.467	0.652	-0.940	1.646	-0.940	0.454
46 Q	1.091	0.694	1.664	-0.630	1.476	0.651	-0.673	1.664	-0.673	0.610
47 P	1.009	0.562	1.487	-0.529	1.267	0.072	-0.487	1.487	-0.529	0.483
48 A	1.009	0.694	1.487	-0.504	1.267	0.072	-0.487	1.487	-0.504	0.505
49 V	1.009	1.549	1.730	-0.110	1.540	0.091	0.743	1.730	-0.110	0.936
50 S	0.958	2.136	1.599	0.357	1.321	0.068	0.621	2.136	0.068	1.009
51 P	1.236	1.820	1.505	0.842	1.203	0.069	0.561	1.820	0.069	1.034
52 P	1.483	1.820	1.832	1.066	1.576	0.112	1.734	1.832	0.112	1.375
53 T	2.349	1.461	2.225	1.302	1.905	0.599	1.136	2.349	0.599	1.568
54 S	2.071	1.597	2.318	1.340	2.023	0.598	1.197	2.318	0.598	1.592
55 Q	2.071	0.742	2.075	1.344	1.750	0.579	-0.034	2.075	-0.034	1.218
56 D	2.298	0.155	1.823	1.012	1.431	0.560	-0.253	2.298	-0.253	1.004
57 P	2.102	0.429	1.627	0.343	1.276	0.540	-1.304	2.102	-1.304	0.716
58 A	1.457	0.429	1.356	-0.606	1.112	0.522	-0.875	1.457	-0.875	0.485
59 G	1.344	0.225	1.459	-1.398	1.212	1.104	-1.039	1.459	-1.398	0.415
60 A	0.844	0.453	1.431	-1.814	1.166	0.634	-0.810	1.431	-1.814	0.272
61 V	0.130	1.080	1.103	-1.926	0.902	0.620	-0.595	1.103	-1.926	0.188
62 R	0.408	1.267	1.253	-1.467	1.057	0.641	0.575	1.267	-1.467	0.533
63 P	0.408	0.812	1.253	-1.051	1.057	0.641	0.575	1.253	-1.051	0.528
64 L	0.408	1.267	1.412	-0.221	1.230	1.260	1.852	1.852	-0.221	1.030
65 S	0.775	1.471	1.776	0.452	1.513	1.278	1.483	1.776	0.452	1.250
66 G	0.775	0.616	1.776	1.020	1.513	1.278	1.483	1.776	0.616	1.209
67 H	0.775	-0.216	1.533	0.913	1.239	1.259	0.253	1.533	-0.216	0.822

	1	2	3	4	5	6	7	8	9	10
68 P	1.489	-0.480	1.617	0.358	1.230	1.253	-1.192	1.617	-1.192	0.611
69 R	0.496	-0.300	1.384	-0.770	1.084	1.239	-0.917	1.384	-0.917	0.317
70 A	-0.446	-0.504	1.328	-1.471	1.084	1.243	-0.666	1.328	-1.471	0.081
71 A	0.054	0.123	1.440	-1.624	1.230	1.112	-0.942	1.440	-1.624	0.199
72 L	0.364	0.614	1.496	-0.736	1.267	1.134	-1.275	1.496	-1.275	0.409
73 F	0.459	1.631	1.057	0.479	0.747	0.509	-1.274	1.631	-1.274	0.515
74 D	0.655	2.297	1.253	1.554	0.902	0.529	-0.223	2.297	-0.223	0.995
75 N	0.787	1.553	1.683	1.656	1.376	1.154	0.786	1.683	0.786	1.285
76 G	1.748	0.848	2.094	1.059	1.741	1.191	0.514	2.094	0.514	1.314
77 T	1.748	0.221	2.075	-0.080	1.795	1.192	0.698	2.075	-0.080	1.093
78 R	0.882	-0.474	1.683	-0.959	1.467	0.704	1.296	1.683	-0.959	0.657
79 Q	0.572	-0.474	1.384	-1.632	1.157	0.664	0.399	1.384	-1.632	0.296
80 L	-0.370	-0.607	1.309	-1.868	1.212	0.669	0.834	1.309	-1.868	0.168
81 V	-0.433	0.225	1.543	-2.191	1.531	1.274	0.792	1.543	-2.191	0.392
82 A	-0.566	0.321	1.356	-2.154	1.330	0.668	1.013	1.356	-2.154	0.281
83 L	-0.585	0.860	1.019	-2.069	0.911	0.626	0.850	1.019	-2.069	0.230
84 R	0.130	1.920	1.103	-1.726	0.902	0.620	-0.595	1.920	-1.726	0.336
85 P	0.996	1.107	1.496	-1.194	1.230	1.108	-1.193	1.496	-1.194	0.507
86 G	1.274	0.748	1.646	-0.552	1.385	1.128	-0.023	1.646	-0.552	0.801
87 A	1.989	0.479	1.730	0.026	1.376	1.122	-1.468	1.989	-1.468	0.751
88 D	1.856	0.479	1.300	0.413	0.902	0.498	-2.477	1.856	-2.477	0.424
89 S	1.856	0.796	1.300	0.352	0.902	0.498	-2.477	1.856	-2.477	0.461
90 A	1.628	-0.384	1.309	-0.110	0.948	0.498	-3.488	1.628	-3.488	0.057
91 A	1.906	-0.498	1.459	-0.348	1.103	0.518	-2.318	1.906	-2.318	0.260
92 P	0.768	-0.595	1.047	-0.635	0.765	0.031	-1.987	1.047	-1.987	-0.086
93 A	0.092	-1.127	0.889	-0.776	0.656	0.028	-2.202	0.889	-2.202	-0.349
94 S	-0.275	-0.589	0.767	-1.152	0.647	0.029	-0.603	0.767	-1.152	-0.168
95 I	-0.989	-0.905	0.702	-1.598	0.601	0.034	0.658	0.702	-1.598	-0.214
96 M	-0.490	-0.677	0.730	-1.754	0.647	0.504	0.429	0.730	-1.754	-0.087
97 V	0.010	-0.472	1.001	-1.016	0.966	0.993	1.430	1.430	-1.016	0.416
98 F	-0.635	-0.472	0.730	-0.116	0.802	0.974	1.859	1.859	-0.635	0.449
99 D	0.003	-0.298	1.029	1.090	0.993	1.592	1.805	1.805	-0.298	0.888
100D	0.035	-0.478	0.917	1.474	0.938	1.576	2.449	2.449	-0.478	0.987
101V	0.402	-0.204	1.038	1.324	0.948	1.575	0.850	1.575	-0.204	0.847
102H	1.116	-0.204	1.346	0.777	1.267	1.590	0.819	1.590	-0.204	0.959
103V	0.749	-0.619	1.505	0.033	1.422	1.725	0.827	1.725	-0.619	0.806
104A	-0.117	-0.697	1.113	-0.701	1.093	1.238	1.424	1.424	-0.701	0.479
105P	-0.389	-0.901	1.094	-1.249	1.084	1.238	1.157	1.238	-1.249	0.291
106R	-1.103	-0.901	0.870	-1.787	0.866	0.623	1.141	1.141	-1.787	-0.042
107V	-1.451	-1.087	0.907	-2.071	0.884	0.626	0.987	0.987	-2.071	-0.172
108I	-1.451	-0.633	1.150	-1.926	1.157	0.645	2.217	2.217	-1.926	0.166
109F	-1.223	-0.308	0.898	-1.715	0.838	0.626	1.998	1.998	-1.715	0.159
110L	-1.356	-0.134	0.711	-1.110	0.638	0.021	2.219	2.219	-1.356	0.141
111P	-0.989	0.071	0.832	-0.805	0.647	0.019	0.620	0.832	-0.989	0.056
112G	-0.351	-0.492	0.973	-0.622	0.665	0.017	-0.712	0.973	-0.712	-0.075
113P	0.364	-0.629	1.038	-0.909	0.711	0.013	-1.973	1.038	-1.973	-0.198
114A	0.364	-0.132	1.038	-1.280	0.711	0.013	-1.973	1.038	-1.973	-0.180
115A	0.560	0.407	0.991	-1.680	0.592	0.014	-2.153	0.991	-2.153	-0.181
116A	0.610	0.946	1.150	-1.543	0.793	0.034	-1.993	1.150	-1.993	-0.000
117L	1.110	1.036	1.178	-0.902	0.838	0.504	-2.222	1.178	-2.222	0.220
118T	1.609	1.868	1.449	0.286	1.157	0.993	-1.221	1.868	-1.221	0.877
119S	1.609	1.868	1.608	1.675	1.330	1.613	0.057	1.868	0.057	1.394
120D	1.837	1.012	1.599	2.534	1.285	1.613	1.067	2.534	1.012	1.564
121D	2.747	0.299	1.879	2.813	1.431	1.628	0.673	2.813	0.299	1.638
122H	2.551	-0.444	1.683	2.152	1.276	1.608	-0.378	2.551	-0.444	1.207
123G	1.559	-0.534	1.468	1.138	1.075	1.592	-0.287	1.592	-0.534	0.859
124T	0.345	-1.162	1.113	-0.016	0.765	1.108	0.157	1.113	-1.162	0.330
125A	-0.155	-0.839	0.842	-0.786	0.446	0.619	-0.844	0.842	-0.844	-0.102
126F	-0.155	-0.212	0.683	-1.421	0.273	-0.001	-2.121	0.683	-2.121	-0.422

127L	-0.250	0.590	1.122	-1.743	0.793	0.624	-2.123	1.122	-2.123	-0.141
128A	-0.218	0.387	0.917	-2.076	0.592	0.604	-2.163	0.917	-2.163	-0.279
129A	0.010	0.213	0.907	-2.181	0.547	0.604	-1.153	0.907	-2.181	-0.150
130R	0.471	0.117	1.225	-2.147	0.829	0.619	-1.072	1.225	-2.147	0.006
131G	0.471	-0.697	1.244	-1.830	0.774	0.618	-1.255	1.244	-1.830	-0.096
132G	0.104	-0.785	1.122	-1.517	0.765	0.620	0.343	1.122	-1.517	0.093
133Y	0.104	-1.616	1.122	-1.305	0.765	0.620	0.343	1.122	-1.616	0.005
134F	0.471	-0.354	0.963	-1.015	0.610	0.484	0.335	0.963	-1.015	0.213
135V	-0.471	0.676	0.889	-0.921	0.665	0.489	0.770	0.889	-0.921	0.299
136A	-0.420	1.399	1.047	-0.500	0.866	0.509	0.930	1.399	-0.500	0.547
137D	0.111	1.489	0.945	0.021	0.784	0.510	0.757	1.489	0.021	0.659
138L	1.053	1.441	1.001	0.480	0.784	0.506	0.506	1.441	0.480	0.824
139S	1.420	1.646	1.281	1.051	0.966	1.124	0.185	1.646	0.185	1.096
140S	1.616	1.603	1.477	1.410	1.121	1.144	1.236	1.616	1.121	1.372
141G	1.116	0.652	1.206	1.399	0.802	0.655	0.235	1.399	0.235	0.866
142H	1.963	0.634	1.720	1.082	1.267	1.274	-0.201	1.963	-0.201	1.106
143T	1.318	0.447	1.449	0.222	1.103	1.256	0.228	1.449	0.222	0.860
144A	1.350	-0.044	1.599	-0.429	1.257	1.276	-0.045	1.599	-0.429	0.709
145R	0.756	0.495	1.487	-0.643	1.294	1.278	0.543	1.487	-0.643	0.744
146V	0.756	-0.318	1.328	-0.446	1.121	0.658	-0.734	1.328	-0.734	0.338
147N	1.059	-0.222	1.403	0.074	1.285	1.127	-0.784	1.403	-0.784	0.563
148V	1.059	-0.741	1.403	0.212	1.285	1.127	-0.784	1.403	-0.784	0.509
149A	0.926	-0.154	0.973	-0.006	0.811	0.503	-1.793	0.973	-1.793	0.180
150D	1.293	0.385	1.253	0.037	0.993	1.121	-2.114	1.293	-2.114	0.424
151A	1.179	-0.328	1.150	0.021	0.838	1.100	-1.961	1.179	-1.961	0.286
152A	2.045	0.163	1.543	0.668	1.166	1.588	-2.558	2.045	-2.558	0.659
153H	1.331	0.163	1.477	1.252	1.121	1.592	-1.297	1.592	-1.297	0.805
154T	1.028	-0.252	1.403	1.542	0.957	1.123	-1.247	1.542	-1.247	0.650
155D	1.028	-0.743	1.403	1.332	0.957	1.123	-1.247	1.403	-1.247	0.550
156F	0.389	-0.468	1.262	0.632	0.938	1.125	0.084	1.262	-0.468	0.566
157T	0.389	0.519	1.103	-0.532	0.765	0.505	-1.193	1.103	-1.193	0.222
158A	0.326	0.884	1.337	-1.491	1.084	1.109	-1.235	1.337	-1.491	0.288
159I	-0.041	1.423	1.496	-2.303	1.239	1.245	-1.227	1.496	-2.303	0.262
160A	0.952	2.375	1.711	-2.386	1.440	1.261	-1.318	2.375	-2.386	0.576
161R	1.255	3.206	1.786	-1.847	1.604	1.730	-1.368	3.206	-1.847	0.909
162R	1.483	2.188	1.776	-0.918	1.558	1.730	-0.357	2.188	-0.918	1.066
163S	2.349	1.279	2.365	-0.082	2.214	2.323	-0.614	2.365	-0.614	1.405
164D	1.634	0.219	2.281	0.048	2.224	2.328	0.831	2.328	0.048	1.366
165G	1.135	0.307	1.730	-0.532	1.741	1.705	1.421	1.741	-0.532	1.072
166K	0.288	0.536	1.216	-1.403	1.276	1.086	1.857	1.857	-1.403	0.693
167L	0.237	-0.296	1.057	-2.085	1.075	1.066	1.697	1.697	-2.085	0.393
168V	0.016	0.447	0.935	-2.100	0.911	0.597	1.866	1.866	-2.100	0.382
169L	-0.212	1.171	0.945	-1.696	0.957	0.597	0.856	1.171	-1.696	0.374
170G	0.060	1.375	0.767	-0.864	0.638	0.491	0.782	1.375	-0.864	0.464
171S	1.002	0.652	0.842	-0.256	0.583	0.486	0.347	1.002	-0.256	0.522
172A	1.369	-0.611	0.963	0.094	0.592	0.484	-1.252	1.369	-1.252	0.234
173D	1.717	-0.120	0.926	-0.114	0.574	0.480	-1.098	1.717	-1.098	0.338
174G	1.236	-0.863	1.188	-0.568	0.856	0.500	-0.766	1.236	-0.863	0.226
175A	1.154	-1.490	1.234	-1.136	0.856	0.500	-0.886	1.234	-1.490	0.033
176V	0.440	-0.659	1.150	-1.318	0.866	0.505	0.560	1.150	-1.318	0.221
177Y	-0.060	0.047	0.879	-1.437	0.547	0.016	-0.442	0.879	-1.437	-0.064
178T	-0.060	0.812	1.337	-1.495	1.230	0.611	-0.377	1.337	-1.495	0.294
179L	0.250	0.321	1.636	-1.254	1.540	0.651	0.520	1.636	-1.254	0.524
180A	0.617	0.429	2.001	-0.830	1.823	0.669	0.151	2.001	-0.830	0.694
181K	0.869	0.968	1.748	-0.062	1.586	0.649	-1.191	1.748	-1.191	0.653
182N	0.307	0.495	1.431	0.420	1.422	0.631	-0.643	1.431	-0.643	0.580
183P	1.521	-0.114	1.786	0.622	1.731	1.115	-1.087	1.786	-1.087	0.796
184A	1.521	0.383	2.029	0.337	2.005	1.134	0.143	2.029	0.143	1.079
185V	1.293	1.010	1.580	0.216	1.367	0.539	-0.932	1.580	-0.932	0.725

186D	1.261	1.107	1.431	0.266	1.212	0.518	-0.659	1.431	-0.659	0.734
187P	1.489	0.568	1.178	0.368	0.893	0.499	-0.879	1.489	-0.879	0.588
188A	1.489	0.700	1.178	0.241	0.893	0.499	-0.879	1.489	-0.879	0.589
189S	1.856	0.604	1.300	-0.197	0.902	0.498	-2.477	1.856	-2.477	0.355
190G	1.552	-0.252	1.225	-0.723	0.738	0.029	-2.428	1.552	-2.428	0.020
191A	1.186	-0.023	0.860	-1.190	0.455	0.011	-2.059	1.186	-2.059	-0.108
192A	1.186	0.790	0.860	-1.350	0.455	0.011	-2.059	1.186	-2.059	-0.015
193T	1.186	1.281	0.860	-1.313	0.455	0.011	-2.059	1.281	-2.059	0.060
194V	1.091	1.621	1.300	-1.151	0.975	0.636	-2.060	1.621	-2.060	0.345
195A	1.287	1.393	1.496	-0.856	1.130	0.656	-1.009	1.496	-1.009	0.585
196S	1.514	1.219	1.945	-0.793	1.768	1.251	0.066	1.945	-0.793	0.996
197R	0.680	0.363	1.608	-1.023	1.595	1.233	0.347	1.608	-1.023	0.686
198T	0.332	0.363	1.664	-1.377	1.558	1.235	0.009	1.664	-1.377	0.541
199K	0.332	-0.224	1.664	-1.724	1.558	1.235	0.009	1.664	-1.724	0.407
200I	0.187	-0.516	1.945	-2.033	1.877	1.840	-0.151	1.945	-2.033	0.450
201F	-0.313	-0.192	1.393	-2.026	1.394	1.217	0.438	1.394	-2.026	0.273
202A	-0.009	-0.222	1.468	-1.818	1.558	1.686	0.388	1.686	-1.818	0.436
203R	-0.237	-0.318	1.019	-1.539	0.920	1.091	-0.687	1.091	-1.539	0.036
204V	-0.313	-0.641	1.075	-1.111	0.948	1.095	-0.573	1.095	-1.111	0.069
205D	0.035	-0.054	1.019	-1.013	0.984	1.092	-0.236	1.092	-1.013	0.261
206A	0.231	0.035	1.216	-1.051	1.139	1.112	0.815	1.216	-1.051	0.499
207L	0.345	0.644	1.113	-1.239	1.039	0.530	0.979	1.113	-1.239	0.487
208V	0.939	1.339	1.225	-1.207	1.002	0.528	0.391	1.339	-1.207	0.602
209T	0.749	1.926	1.253	-0.699	0.993	0.080	0.287	1.926	-0.699	0.656
210Q	0.945	1.339	1.449	0.074	1.148	0.100	1.338	1.449	0.074	0.913
211G	1.856	0.752	1.730	0.889	1.294	0.114	0.943	1.856	0.114	1.083
212N	1.856	-0.080	1.730	1.185	1.294	0.114	0.943	1.856	-0.080	1.006
213T	1.293	-0.150	1.412	0.869	1.130	0.096	1.491	1.491	-0.150	0.877
214T	0.332	0.173	1.001	-0.072	0.765	0.059	1.763	1.763	-0.072	0.574
215V	0.604	0.309	1.281	-0.696	1.130	0.548	1.754	1.754	-0.696	0.704
216V	0.427	0.896	1.412	-1.259	1.294	1.132	1.866	1.866	-1.259	0.824
217L	0.459	1.483	1.206	-1.094	1.093	1.112	1.825	1.825	-1.094	0.869
218D	0.509	2.543	1.337	-0.937	1.312	1.134	1.948	2.543	-0.937	1.121
219R	1.072	1.908	1.655	-0.778	1.476	1.153	1.400	1.908	-0.778	1.126
220G	1.717	1.585	1.926	-0.579	1.640	1.171	0.971	1.926	-0.579	1.204
221Q	2.064	1.449	1.889	-0.272	1.622	1.167	1.125	2.064	-0.272	1.292
222T	1.761	0.634	1.814	-0.009	1.458	0.698	1.175	1.814	-0.009	1.076
223S	1.824	0.770	1.580	0.182	1.139	0.094	1.216	1.824	0.094	0.972
224V	0.958	-0.086	1.449	-0.077	1.166	0.096	1.538	1.538	-0.086	0.721
225T	0.939	0.550	1.113	-0.557	0.747	0.053	1.375	1.375	-0.557	0.603
226T	0.743	0.686	0.917	-1.110	0.592	0.033	0.324	0.917	-1.110	0.312
227I	0.964	0.285	1.038	-1.364	0.756	0.502	0.155	1.038	-1.364	0.334
228G	1.559	0.610	1.150	-1.263	0.720	0.500	-0.434	1.559	-1.263	0.406
229A	1.363	0.473	1.113	-0.489	0.738	1.100	-0.207	1.363	-0.489	0.585
230D	1.167	0.964	0.917	0.170	0.583	1.080	-1.258	1.167	-1.258	0.518
231G	2.052	0.425	1.384	0.839	0.975	1.121	-1.416	2.052	-1.416	0.768
232H	2.071	-0.406	1.720	0.801	1.394	1.163	-1.253	2.071	-1.253	0.784
233A	2.071	0.317	1.720	0.453	1.394	1.163	-1.253	2.071	-1.253	0.838
234Q	0.857	0.317	1.365	-0.332	1.084	0.680	-0.809	1.365	-0.809	0.452
235Q	0.762	0.453	1.804	-0.938	1.604	1.305	-0.810	1.804	-0.938	0.597
236A	0.762	0.453	1.646	-1.697	1.431	0.685	-2.088	1.646	-2.088	0.170
237L	0.990	1.080	1.636	-2.067	1.385	0.685	-1.077	1.636	-2.067	0.376
238R	0.990	1.285	1.636	-2.176	1.385	0.685	-1.077	1.636	-2.176	0.390
239A	0.971	0.962	1.300	-2.005	0.966	0.642	-1.240	1.300	-2.005	0.228
240G	0.971	1.453	1.300	-1.668	0.966	0.642	-1.240	1.453	-1.668	0.346
241Q	1.881	0.712	1.580	-1.358	1.112	0.657	-1.635	1.881	-1.635	0.421
242G	1.944	0.221	1.346	-1.121	0.793	0.052	-1.593	1.944	-1.593	0.235
243A	1.546	-0.406	1.337	-1.071	0.838	0.069	-0.638	1.546	-1.071	0.239
244T	1.318	0.133	1.346	-1.096	0.884	0.069	-1.649	1.346	-1.649	0.144

245T	1.072	0.001	1.019	-1.381	0.510	0.027	-2.822	1.072	-2.822	-0.225
246M	1.344	-0.695	1.300	-1.451	0.875	0.516	-2.831	1.344	-2.831	-0.135
247A	1.344	0.047	1.543	-1.228	1.148	0.535	-1.601	1.543	-1.601	0.255
248A	0.433	0.860	1.262	-0.711	1.002	0.520	-1.207	1.262	-1.207	0.309
249D	0.465	0.764	1.057	-0.275	0.802	0.500	-1.247	1.057	-1.247	0.295
250P	0.996	0.021	1.496	-0.313	1.230	1.108	-1.193	1.496	-1.193	0.478
251L	0.629	-0.663	1.375	-0.928	1.221	1.109	0.406	1.375	-0.928	0.450
252G	-0.085	-0.458	1.290	-1.644	1.230	1.115	1.851	1.851	-1.644	0.471
253R	-1.223	-0.546	0.879	-2.293	0.893	0.628	2.182	2.182	-2.293	0.074
254V	-1.223	-0.869	0.636	-2.617	0.619	0.609	0.951	0.951	-2.617	-0.270
255L	-0.009	0.041	0.991	-2.413	0.929	1.092	0.508	1.092	-2.413	0.163
256I	-0.041	0.872	1.197	-1.927	1.130	1.112	0.548	1.197	-1.927	0.413
257A	-0.041	1.824	1.197	-1.181	1.130	1.112	0.548	1.824	-1.181	0.655
258D	0.553	2.315	1.309	-0.628	1.093	1.111	-0.040	2.315	-0.628	0.816
259T	1.495	1.571	1.384	-0.597	1.039	1.105	-0.475	1.571	-0.597	0.789
260R	2.381	0.876	1.851	-0.899	1.431	1.146	-0.634	2.381	-0.899	0.879
261G	1.666	-0.034	1.767	-1.272	1.440	1.151	0.812	1.767	-1.272	0.790
262G	0.452	-1.067	1.412	-1.543	1.130	0.668	1.256	1.412	-1.543	0.330
263Q	-0.111	-1.067	1.094	-1.700	0.966	0.649	1.803	1.803	-1.700	0.234
264L	-0.496	-1.654	0.917	-1.810	0.729	0.044	2.136	2.136	-1.810	-0.019
265L	-0.496	-0.911	0.917	-1.958	0.729	0.044	2.136	2.136	-1.958	0.066
266V	-1.090	-0.348	0.804	-1.934	0.765	0.046	2.725	2.725	-1.934	0.138
267Y	-0.838	-0.456	0.748	-1.585	0.711	0.492	2.553	2.553	-1.585	0.232
268G	-0.123	-0.374	1.075	-1.004	0.975	0.506	2.338	2.338	-1.004	0.485
269V	-0.123	-1.206	1.075	-0.421	0.975	0.506	2.338	2.338	-1.206	0.449
270D	-0.395	-0.296	1.057	-0.195	0.966	0.506	2.071	2.071	-0.395	0.530
271P	-0.857	-0.344	0.720	-0.541	0.738	0.492	2.173	2.173	-0.857	0.340
272L	-0.951	-0.703	1.160	-1.395	1.257	1.117	2.172	2.172	-1.395	0.379
273I	-0.338	-0.905	1.608	-2.062	1.640	1.157	1.747	1.747	-2.062	0.407
274L	-0.838	-0.222	1.337	-2.433	1.321	0.669	0.746	1.337	-2.433	0.083
275R	-1.090	-0.114	1.346	-2.203	1.285	0.669	0.858	1.346	-2.203	0.107
276Q	-0.376	-0.114	1.674	-1.678	1.549	0.682	0.643	1.674	-1.678	0.340
277A	-0.104	-0.114	1.692	-1.243	1.558	0.682	0.910	1.692	-1.243	0.483
278Y	0.743	-0.114	2.206	-1.069	2.023	1.301	0.474	2.206	-1.069	0.795
279P	0.857	0.652	2.103	-1.037	1.923	0.719	0.638	2.103	-1.037	0.836
280V	0.610	-0.114	1.776	-1.295	1.549	0.677	-0.535	1.776	-1.295	0.381
281R	0.610	0.610	2.019	-1.321	1.823	0.696	0.695	2.019	-1.321	0.733
282Q	0.610	-0.408	2.019	-1.221	1.823	0.696	0.695	2.019	-1.221	0.602
283A	0.838	-0.899	1.767	-1.001	1.504	0.677	0.475	1.767	-1.001	0.480
284P	0.490	-0.272	1.804	-0.906	1.522	0.681	0.321	1.804	-0.906	0.520
285Y	0.357	0.225	1.375	-1.049	1.048	0.056	-0.688	1.375	-1.049	0.189
286G	0.338	1.445	1.038	-1.478	0.629	0.014	-0.851	1.445	-1.478	0.162
287L	0.617	1.393	1.188	-1.525	0.784	0.034	0.319	1.393	-1.525	0.401
288A	0.749	1.393	1.375	-1.489	0.984	0.639	0.098	1.393	-1.489	0.536
289G	1.363	1.393	1.449	-1.171	1.112	1.220	-0.260	1.449	-1.171	0.729
290S	0.421	0.263	1.375	-1.144	1.166	1.225	0.175	1.375	-1.144	0.497
291R	1.135	-0.689	1.459	-1.408	1.157	1.220	-1.270	1.459	-1.408	0.229
292E	0.370	-0.647	1.477	-1.915	1.175	1.245	-0.238	1.477	-1.915	0.210
293L	-0.224	-0.731	1.365	-2.231	1.212	1.246	0.350	1.365	-2.231	0.141
294A	-0.224	-0.036	1.365	-2.178	1.212	1.246	0.350	1.365	-2.178	0.248
295W	-0.111	-0.036	1.262	-1.662	1.112	0.664	0.514	1.262	-1.662	0.249
296V	-0.275	1.323	1.132	-0.913	0.902	0.084	0.582	1.323	-0.913	0.405
297S	0.440	2.028	1.216	-0.352	0.893	0.079	-0.864	2.028	-0.864	0.491
298Q	0.718	1.664	1.365	0.066	1.048	0.099	0.306	1.664	0.066	0.752
299T	1.793	1.076	1.646	0.468	1.339	0.114	0.171	1.793	0.114	0.944
300A	2.355	0.261	1.963	1.016	1.504	0.133	-0.377	2.355	-0.377	0.979
301S	1.710	0.888	1.692	1.472	1.339	0.114	0.052	1.710	0.052	1.038
302N	0.825	-0.374	1.225	1.441	0.948	0.074	0.210	1.441	-0.374	0.621
303T	0.857	-0.444	1.019	0.597	0.747	0.054	0.170	1.019	-0.444	0.428

304V	0.604	-1.140	1.272	-0.602	0.984	0.073	1.512	1.512	-1.140	0.386
305I	0.825	-0.552	1.393	-1.268	1.148	0.542	1.343	1.393	-1.268	0.490
306G	-0.199	0.263	1.010	-1.380	0.847	0.507	1.891	1.891	-1.380	0.420
307Y	-0.199	0.263	1.010	-0.756	0.847	0.507	1.891	1.891	-0.756	0.509
308D	0.364	0.345	1.328	-0.299	1.011	0.525	1.343	1.343	-0.299	0.660
309L	1.230	0.165	1.459	-0.142	0.984	0.523	1.022	1.459	-0.142	0.749
310T	0.364	0.273	1.328	-0.481	1.011	0.525	1.343	1.343	-0.481	0.623
311T	0.617	0.357	1.318	-0.748	1.048	0.525	1.231	1.318	-0.748	0.621
312G	-0.250	0.698	0.926	-1.228	0.720	0.037	1.829	1.829	-1.228	0.390
313I	0.825	-0.026	1.337	-1.380	1.075	0.632	1.368	1.368	-1.380	0.547
314P	0.857	1.113	1.589	-1.616	1.558	1.206	1.392	1.589	-1.616	0.871
315V	0.294	0.347	1.272	-1.817	1.394	1.188	1.940	1.940	-1.817	0.660
316E	0.199	0.802	1.711	-2.131	1.914	1.813	1.939	1.939	-2.131	0.892
317K	0.585	0.718	2.103	-2.254	2.169	1.830	1.949	2.169	-2.254	1.014
318V	0.585	-0.210	2.103	-2.202	2.169	1.830	1.949	2.169	-2.202	0.889
319R	1.148	0.377	2.421	-1.705	2.333	1.849	1.401	2.421	-1.705	1.118
320Y	0.421	0.055	1.973	-1.140	1.959	1.251	2.016	2.016	-1.140	0.933
321P	0.440	0.820	1.851	-0.644	1.695	0.698	2.114	2.114	-0.644	0.996
322T	1.053	1.070	2.300	-0.474	2.078	0.739	1.689	2.300	-0.474	1.208
323V	0.920	1.435	2.113	-0.380	1.877	0.133	1.910	2.113	-0.380	1.144
324Q	1.483	1.327	2.160	0.066	1.950	0.155	1.465	2.160	0.066	1.229
325Q	1.761	0.836	2.066	0.851	1.832	0.156	1.405	2.066	0.156	1.272
326P	0.850	0.171	1.786	1.628	1.686	0.141	1.799	1.799	0.141	1.152
327N	1.217	0.351	1.907	1.885	1.695	0.140	0.200	1.907	0.140	1.056
328S	0.256	0.317	1.515	1.307	1.276	0.102	0.288	1.515	0.102	0.723
329L	0.509	-0.048	1.459	0.501	1.221	0.548	0.116	1.459	-0.048	0.615
330A	0.869	1.012	1.543	-0.108	1.312	1.129	-0.131	1.543	-0.131	0.804
331F	0.756	1.551	1.440	-0.182	1.157	1.108	0.023	1.551	-0.182	0.836
332D	0.756	2.217	1.440	0.094	1.157	1.108	0.023	2.217	0.023	0.971
333E	1.970	1.473	1.795	0.569	1.467	1.591	-0.421	1.970	-0.421	1.206
334T	2.166	0.491	1.991	0.947	1.622	1.611	0.630	2.166	0.491	1.351
335S	2.166	-0.096	1.973	1.184	1.677	1.613	0.814	2.166	-0.096	1.333
336D	1.413	-1.047	1.954	0.953	1.595	1.143	1.155	1.954	-1.047	1.024
337T	0.686	-0.731	1.505	0.312	1.221	0.545	1.770	1.770	-0.731	0.758
338L	0.123	-0.595	1.188	-0.628	1.057	0.527	2.318	2.318	-0.628	0.570
339Y	0.123	0.465	1.188	-1.161	1.057	0.527	2.318	2.318	-1.161	0.645
340V	-0.148	1.499	0.907	-1.279	0.692	0.038	2.327	2.327	-1.279	0.577
341V	-0.066	1.595	0.860	-0.755	0.692	0.038	2.446	2.446	-0.755	0.687
342S	0.876	2.319	0.935	-0.238	0.638	0.032	2.012	2.319	-0.238	0.939
343G	1.129	1.367	0.683	0.030	0.401	0.013	0.669	1.367	0.013	0.613
344S	1.723	1.231	0.795	-0.165	0.364	0.012	0.081	1.723	-0.165	0.577
345G	1.723	0.279	0.795	-0.637	0.364	0.012	0.081	1.723	-0.637	0.374
346A	1.691	-0.673	0.973	-1.224	0.583	0.034	0.084	1.691	-1.224	0.210
347G	1.097	-0.098	0.860	-1.479	0.619	0.035	0.673	1.097	-1.479	0.244
348V	0.180	-0.635	0.571	-1.717	0.446	0.017	0.835	0.835	-1.717	-0.043
349Q	0.313	-0.538	0.907	-1.857	0.856	0.617	0.808	0.907	-1.857	0.158
350V	0.313	-1.029	1.066	-1.704	1.030	1.237	2.086	2.086	-1.704	0.428
351I	0.085	-0.306	1.075	-1.487	1.075	1.237	1.075	1.237	-1.487	0.394
352E	0.452	0.509	1.197	-0.901	1.084	1.235	-0.524	1.235	-0.901	0.436
353H	0.433	0.748	0.860	-0.574	0.665	1.193	-0.687	1.193	-0.687	0.377
354A	0.996	0.912	1.178	-0.518	0.829	1.211	-1.235	1.211	-1.235	0.482
355A	1.767	1.167	1.748	-0.911	1.321	1.834	-1.557	1.834	-1.557	0.767
356G	1.274	1.421	0.963	-1.118	1.276	1.294	-2.541	1.421	-2.541	0.367
357T	1.141	1.048	0.346	-1.325	1.422	0.735	-3.818	1.422	-3.818	-0.064
358R	1.009	0.812	-0.111	-1.104	1.741	0.795	-3.818	1.741	-3.818	-0.097

Overlap Display

Selected Programs: hydro flexi access turns surface polar antipro

Respective Threshold: 1.9 2 1.9 2.4 2.3 1.8 1.9

The predicted B-cell epitopes are shown in blue colour and underlined.

Sequence	[1]LLTGNKPAVQRRFIGLLMLSVLVAGCSSNPLANFAPGYPPTIEPAQPAVSPPTSQDPAGAVRPLSGH PRAALFDNGTRQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSDDHGTAFLAARGGYF VADLSSGHTARVNVADAAHTDFTAIARRSDGKLVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFA RVDALVTQGNTTVVLDRGQTSVTTIGADGHAQQALRAGQGATTMAADPLGRVLIADTRGGQLLVYG VDPLILRQAYPVRQAPYGLAGSRELAWVSQTASNTVIGYDLTTGIPVEKVRYPTVQQPNSLAFDETS DTLYVVSGSGAGVQVIEHAAGTR[358]
Hydrophilic ity	[1]LLTGNKPAVQRRFIGLLMLSVLV<u>AGCSSNP</u>LANFAPGYPPTIEPAQPAV<u>SPPTSQDPAGA</u>VRPLSGH PRAALFDNGTRQLVAL<u>RPGADSAAPAS</u>IMVFDDVHVAPRVIFLPGPAAAL<u>TSDDHGTA</u>FLAARGGYF VADL<u>SSGHTAR</u>VNV<u>ADAAHTD</u>FTAI<u>ARRSDGK</u>LVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFA RVDALVTQGNTTVVL<u>DRGQTSV</u>TTI<u>GADGHAQQA</u>LR<u>AGQGATT</u>MAADPLGRVLI<u>ADTRGGQ</u>LLVYG VDPLILRQAYPVRQAPYGLAGSRELAWV<u>SQTASNT</u>VIGYDLTTGIPVEKVRYPTVQQPNSL<u>AFDETS DTL</u>YVVSGSGAGVQVIEHAAGTR[358]
Flexibility	[1]LLTGNKPAVQRRFIGLLMLSV<u>LVAGCSS</u>NPLANFAPGYPPTIEPAQ<u>PAVSPPT</u>SQDPAGAVRPLSGH PRA<u>ALFDNGT</u>RQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSDDHGTAFLAARGGYF VADLSSGHTARVNVADAAHTDF<u>TAIARRSDG</u>KLVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFA RVDALVTQGNTT<u>VVLDRGQ</u>TSVTTIGADGHAQQALRAGQGATTMAADPLGRV<u>LIADTRG</u>GQLLVYG VDPLILRQAYPVRQAPYGLAGSREL<u>AWVSQTA</u>SNTVIGYDLTTGIPVEKVRYPTVQQPNSL<u>AFDETS</u> DTL<u>YVVSGSG</u>AGVQVIEHAAGTR[358]
Accessibilit y	[1]LLTG<u>NKPAVQRR</u>FIGLLMLSVLVAGCSSNPLANFAPG<u>YPPTIEP</u>AQPAV<u>SPPTSQDPA</u>GAVRPLSGH PRAAL<u>FDNGTRQL</u>VALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSDDHGTAFLAARGGYF VADLSSGHTARVNVADAAHTDFTAI<u>ARRSDGKL</u>VLGSADGAV<u>YTLAKNPAVDP</u>ASGAA<u>TVASRTKIFA</u> RVDALVTQGNTTVVL<u>DRGQTS</u>VTTIGADGHAQQALRAGQGATTMAADPLGRVLIADTRGGQLLVYG VDPLIL<u>RQAYPVRQAPY</u>GLAGSRELAWV<u>SQTASNT</u>VIGYDLTTGI<u>PVEKVRYPTVQQPNSLAFDETS DTLY</u>VVSGSGAGVQVIEHAAGTR[358]
Turns	[1]LLTGNKPAVQRRFIGLLMLSVLVA<u>GCSSNPL</u>ANFAPGYPPTIEPAQPAVSPPTSQDPAGAVRPLSGH PRAALFDNGTRQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAA<u>LTSDDHGT</u>AFLAARGGYF VADLSSGHTARVNVADAAHTDFTAIARRSDGKLVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFA RVDALVTQGNTTVVLDRGQTSVTTIGADGHAQQALRAGQGATTMAADPLGRVLIADTRGGQLLVYG VDPLILRQAYPVRQAPYGLAGSRELAWVSQTASNTVIGYDLTTGIPVEKVRYPTVQQPNSLAFDETS DTLYVVSGSGAGVQVIEHAAGTR[358]

Exposed Surface	[1]LLTGNKPAVQRRFIGLLMLSVLVAGCSSNPLANFAPGYPPTIEPAQPAVSPPTSQDPAGAVRPLSGHPRAALFDNGTRQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSDDHGTAFLAARGGYFVADLSSGHTARVNVADAAHTDFTAIARRSDGKLVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFARVDALVTQGNTTVVLDRGQTSVTTIGADGHAQQALRAGQGATTMAADPLGRVLIADTRGGQLLVYGVDPLILRQAYPVRQAPYGLAGSRELAWVSQTASNTVIGYDLTTGIPVEKVRYPTVQQPNSLAFDETSDTLYVVSGSGAGVQVIEHAAGTR[358]
Polarity	[1]LLTGNKPAVQRRFIGLLMLSVLVAGCSSNPLANFAPGYPPTIEPAQPAVSPPTSQDPAGAVRPLSGHPRAALFDNGTRQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSDDHGTAFLAARGGYFVADLSSGHTARVNVADAAHTDFTAIARRSDGKLVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFARVDALVTQGNTTVVLDRGQTSVTTIGADGHAQQALRAGQGATTMAADPLGRVLIADTRGGQLLVYGVDPLILRQAYPVRQAPYGLAGSRELAWVSQTASNTVIGYDLTTGIPVEKVRYPTVQQPNSLAFDETSDTLYVVSGSGAGVQVIEHAAGTR[358]
Antigenic Propensity	[1]LLTGNKPAVQRRFIGLLMLSVLVAGCSSNPLANFAPGYPPTIEPAQPAVSPPTSQDPAGAVRPLSGHPRAALFDNGTRQLVALRPGADSAAPASIMVFDDVHVAPRVIFLPGPAAALTSDDHGTAFLAARGGYFVADLSSGHTARVNVADAAHTDFTAIARRSDGKLVLGSADGAVYTLAKNPAVDPASGAATVASRTKIFARVDALVTQGNTTVVLDRGQTSVTTIGADGHAQQALRAGQGATTMAADPLGRVLIADTRGGQLLVYGVDPLILRQAYPVRQAPYGLAGSRELAWVSQTASNTVIGYDLTTGIPVEKVRYPTVQQPNSLAFDETSDTLYVVSGSSAGVQVIEHAAGTR[358]

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