

BcePred Prediction Server

The server displays

1. [GRAPHICAL RESULT](#)
2. [TABULAR RESULT](#)
3. [Overlap Display](#)

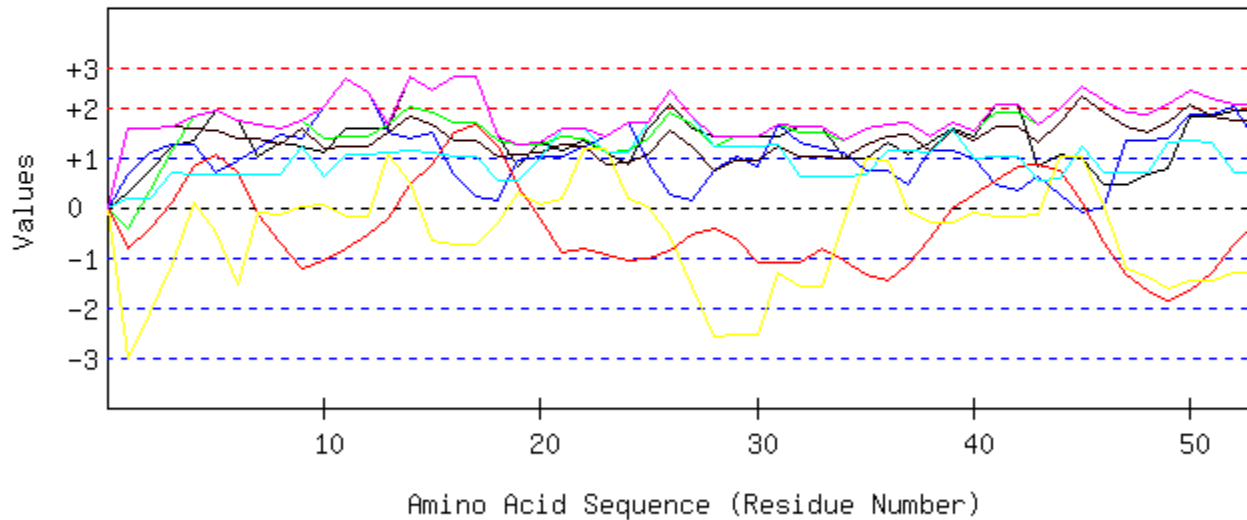
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GCGDGSPA EAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRSITAES
RDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLSVALFFVWMITVAFLYLVGGM
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DLIGGIEVTLADRD
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Length=304

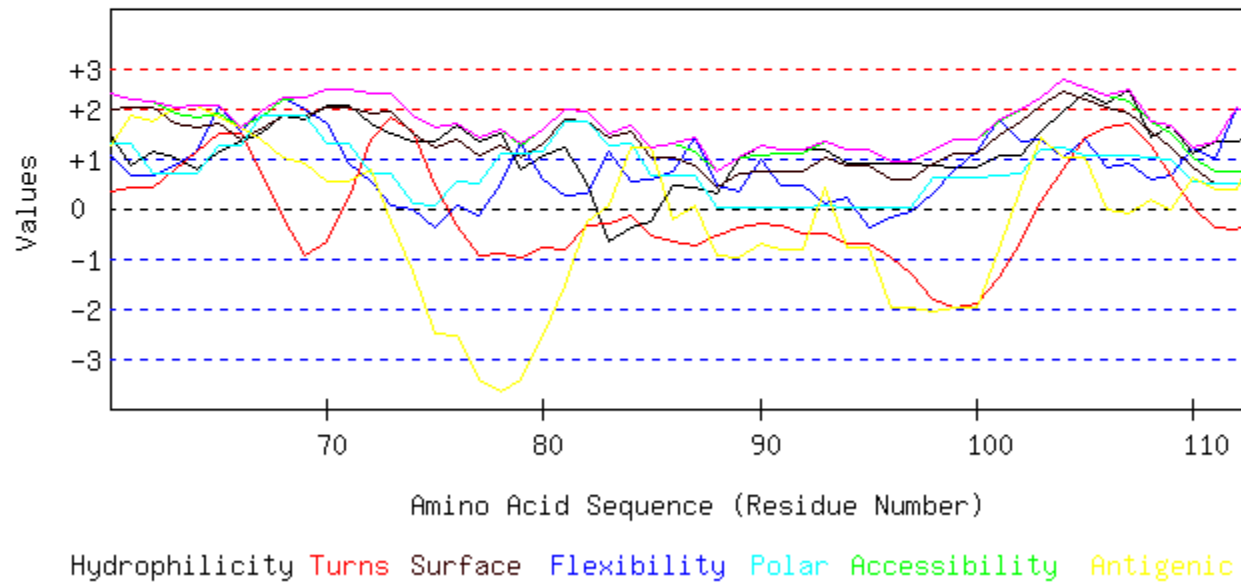
GRAPHICAL RESULT

GRAPHICAL RESULT :: SEQ 1 to 60

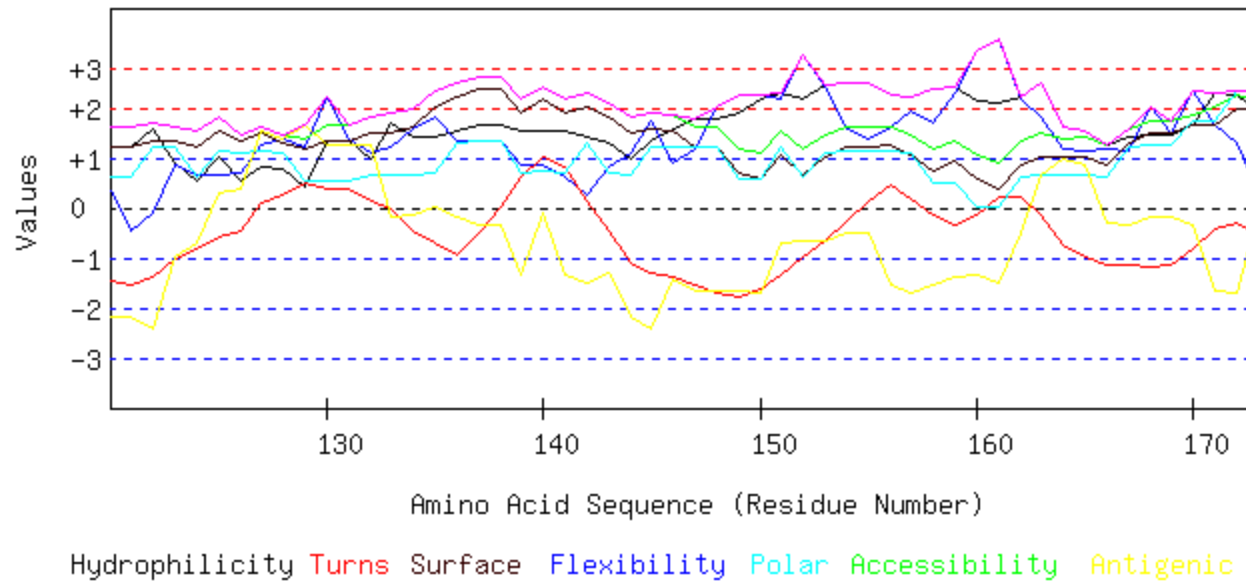


Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic

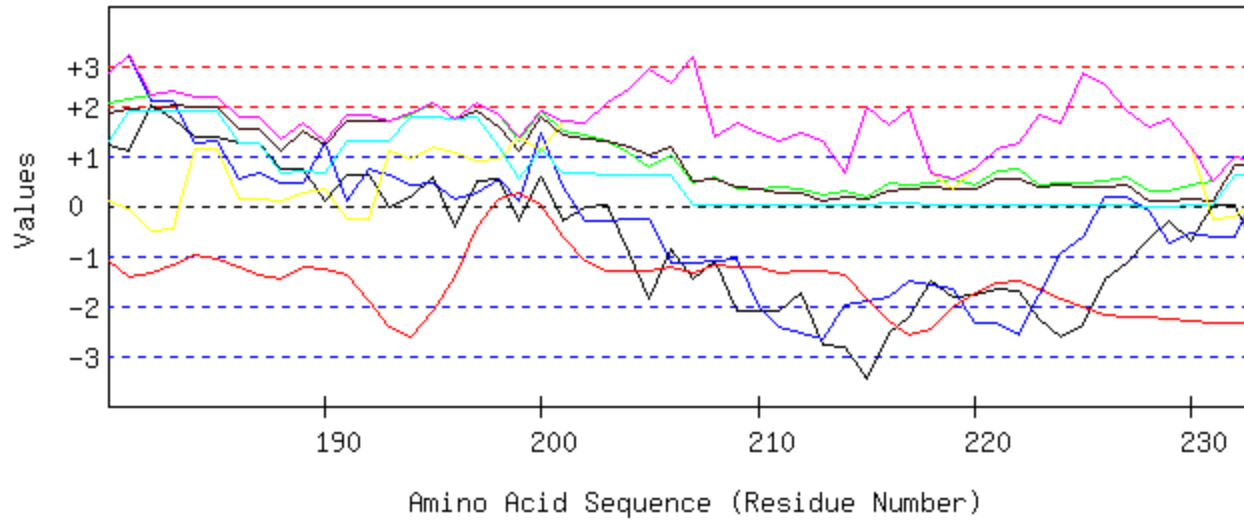
GRAPHICAL RESULT :: SEQ 61 to 120



GRAPHICAL RESULT :: SEQ 121 to 180

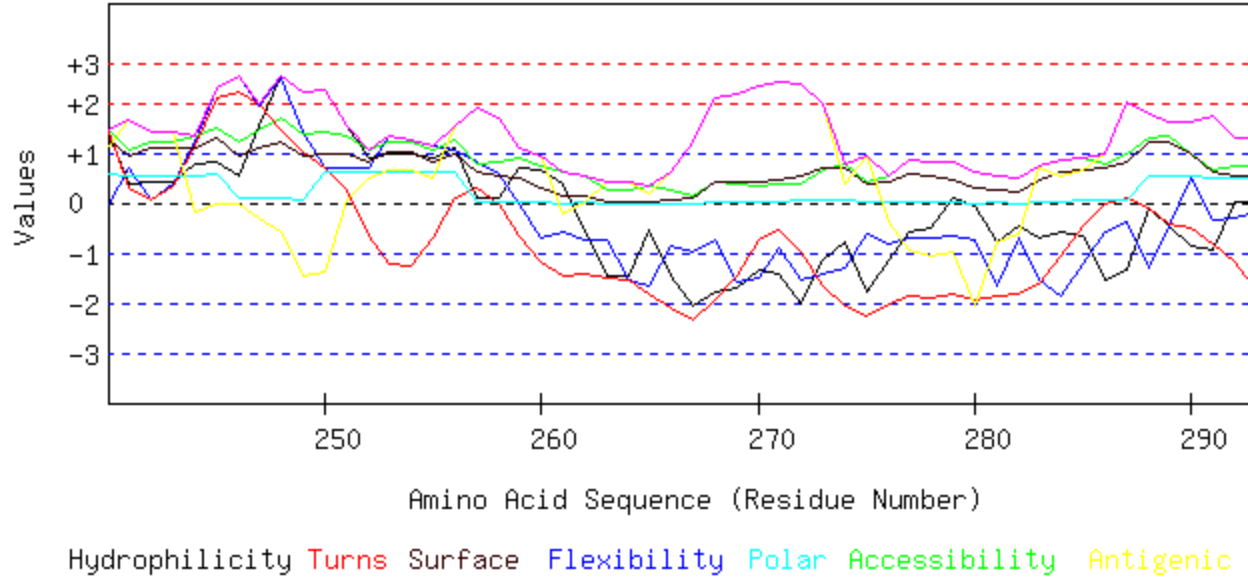


GRAPHICAL RESULT :: SEQ 181 to 240

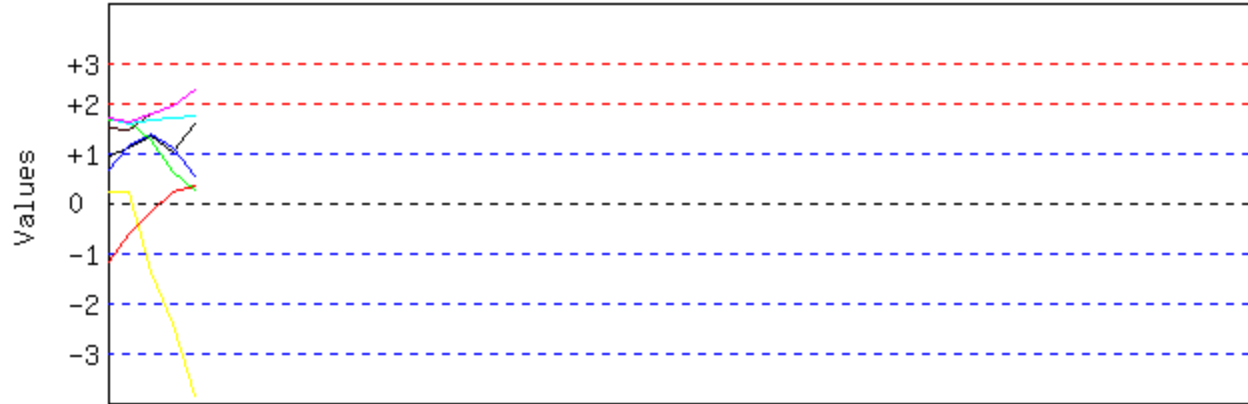


Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic

GRAPHICAL RESULT :: SEQ 241 to 300



GRAPHICAL RESULT :: SEQ 301 to 360



Amino Acid Sequence (Residue Number)

Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic

[TOP](#)

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

VTAPNEPGALSKGDGPNADGLVDRGGAHRAATGPGRIPDAGDPPPWQRAATRQSQAGHRQ
PPPVSHPEGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQDPDASLGCGDGSPA
EAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRSITAESRDARVQLSA
RRSRGPVRASMQIRRIDPWSTLKVSLLLSVALFFVWMITVAFLYLVLGGMGVWAKLNSNV
GDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTALATIGAFVYNLITDLIGGIEVTL

ADRD

Length=304

A.A.

Parameter
Combined

MAX	MIN	Hydro AVG	Flexi	Access	Turns	Surface	Polar	AntiPro
1 V		0.281	0.664	-0.411	-0.819	1.576	0.191	-3.009
1.576	-3.009	-0.218						
2 T		0.724	1.119	0.346	-0.428	1.567	0.171	-2.112
1.567	-2.112	0.198						
3 A		1.217	1.255	1.132	0.116	1.613	0.711	-1.128
1.613	-1.128	0.702						
4 P		1.350	1.255	1.832	0.831	1.567	0.669	0.102
1.832	0.102	1.087						
5 N		1.944	0.692	1.945	1.073	1.531	0.668	-0.487
1.945	-0.487	1.052						
6 E		1.748	0.938	1.748	0.685	1.376	0.648	-1.538
1.748	-1.538	0.801						
7 P		1.034	1.195	1.664	-0.143	1.385	0.653	-0.092
1.664	-0.143	0.814						
8 G		1.312	1.463	1.571	-0.718	1.267	0.654	-0.153
1.571	-0.718	0.771						
9 A		1.230	1.375	1.720	-1.231	1.595	1.208	0.025
1.720	-1.231	0.846						
10 L		1.097	2.002	1.384	-1.079	1.185	0.609	0.052
2.002	-1.079	0.750						
11 S		1.597	2.565	1.412	-0.830	1.230	1.079	-0.177
2.565	-0.830	0.982						
12 K		1.597	2.319	1.412	-0.553	1.230	1.079	-0.177
2.319	-0.553	0.987						
13 G		1.597	1.487	1.655	-0.206	1.504	1.098	1.053
1.655	-0.206	1.170						
14 D		2.621	1.399	2.038	0.452	1.804	1.133	0.505
2.621	0.452	1.422						
15 G		2.343	1.487	1.889	0.863	1.649	1.113	-0.665
2.343	-0.665	1.240						
16 P		2.614	0.656	1.711	1.518	1.330	1.007	-0.739
2.614	-0.739	1.157						
17 N		2.614	0.201	1.711	1.644	1.330	1.007	-0.739
2.614	-0.739	1.110						
18 A		1.401	0.131	1.356	1.232	1.020	0.524	-0.295
1.401	-0.295	0.767						
19 D		0.806	0.944	1.244	0.430	1.057	0.525	0.294
1.244	0.294	0.757						
20 G		1.306	1.032	1.272	-0.221	1.103	0.995	0.065
1.306	-0.221	0.793						
21 L		1.129	1.032	1.403	-0.897	1.267	1.579	0.177
1.579	-0.897	0.813						
22 V		1.356	1.237	1.393	-0.833	1.221	1.579	1.187

1.579	-0.833	1.020						
23 D		1.084	1.423	1.113	-0.949	0.856	1.090	1.197
1.423	-0.949	0.831						
24 R		0.857	1.698	1.122	-1.064	0.902	1.090	0.186
1.698	-1.064	0.684						
25 G		1.571	0.884	1.365	-1.009	1.066	1.705	0.019
1.705	-1.009	0.800						
26 G		2.071	0.257	1.917	-0.844	1.549	2.328	-0.571
2.328	-0.844	0.958						
27 A		1.571	0.121	1.646	-0.549	1.230	1.839	-1.572
1.839	-1.572	0.612						
28 H		1.438	0.748	1.216	-0.412	0.756	1.214	-2.581
1.438	-2.581	0.340						
29 R		1.407	1.016	1.421	-0.623	0.957	1.234	-2.541
1.421	-2.541	0.410						
30 A		1.407	0.830	1.421	-1.105	0.957	1.234	-2.541
1.421	-2.541	0.315						
31 A		1.407	1.644	1.664	-1.107	1.230	1.253	-1.311
1.664	-1.311	0.683						
32 T		1.634	1.319	1.496	-1.097	1.011	0.633	-1.578
1.634	-1.578	0.488						
33 G		1.634	1.187	1.496	-0.835	1.011	0.633	-1.578
1.634	-1.578	0.507						
34 P		0.996	1.099	1.356	-1.052	0.993	0.635	-0.246
1.356	-1.052	0.540						
35 G		0.996	0.740	1.599	-1.349	1.267	0.654	0.984
1.599	-1.349	0.699						
36 R		1.299	0.740	1.674	-1.476	1.431	1.123	0.934
1.674	-1.476	0.818						
37 I		1.072	0.465	1.683	-1.149	1.476	1.123	-0.076
1.683	-1.149	0.656						
38 P		1.299	1.149	1.431	-0.617	1.157	1.104	-0.296
1.431	-0.617	0.747						
39 D		1.571	1.149	1.711	-0.035	1.522	1.593	-0.305
1.711	-0.305	1.029						
40 A		1.438	0.968	1.524	0.242	1.321	0.987	-0.084
1.524	-0.084	0.914						
41 G		2.077	0.465	1.907	0.530	1.613	1.005	-0.186
2.077	-0.186	1.059						
42 D		2.077	0.329	1.907	0.829	1.613	1.005	-0.186
2.077	-0.186	1.082						
43 P		0.813	0.604	1.655	0.879	1.312	0.541	-0.155
1.655	-0.155	0.807						
44 P		1.059	0.245	1.982	0.729	1.686	0.583	1.019
1.982	0.245	1.043						
45 P		0.964	-0.114	2.421	0.106	2.205	1.208	1.017
2.421	-0.114	1.116						
46 W		0.465	0.019	2.150	-0.689	1.886	0.719	0.016
2.150	-0.689	0.652						
47 Q		0.465	1.335	1.907	-1.357	1.613	0.700	-1.214
1.907	-1.357	0.493						
48 R		0.661	1.335	1.860	-1.677	1.494	0.701	-1.393
1.860	-1.677	0.426						
49 A		0.794	1.377	2.047	-1.869	1.695	1.307	-1.614
2.047	-1.869	0.534						
50 A		1.805	1.868	2.356	-1.641	2.050	1.324	-1.473
2.356	-1.641	0.899						

51 T	1.837	1.868	2.178	-1.290	1.832	1.302	-1.476
2.178	-1.476	0.893					
52 R	1.951	2.004	2.075	-0.741	1.731	0.719	-1.312
2.075	-1.312	0.918					
53 Q	1.951	1.281	2.075	-0.299	1.731	0.719	-1.312
2.075	-1.312	0.878					
54 S	2.178	1.603	2.066	-0.086	1.686	0.719	-0.302
2.178	-0.302	1.124					
55 Q	1.982	1.239	2.029	0.015	1.704	1.319	-0.075
2.029	-0.075	1.173					
56 A	1.982	1.107	2.029	-0.063	1.704	1.319	-0.075
2.029	-0.075	1.143					
57 G	1.982	1.465	2.029	0.071	1.704	1.319	-0.075
2.029	-0.075	1.214					
58 H	1.704	1.197	2.122	0.147	1.823	1.318	-0.015
2.122	-0.015	1.185					
59 R	1.457	1.010	2.038	0.281	1.722	1.295	0.042
2.038	0.042	1.121					
60 Q	1.457	1.052	2.281	0.350	1.996	1.314	1.272
2.281	0.350	1.389					
61 P	0.863	0.652	2.169	0.424	2.032	1.315	1.860
2.169	0.424	1.331					
62 P	1.141	0.652	2.160	0.426	2.014	0.715	1.753
2.160	0.426	1.266					
63 P	1.009	0.868	1.889	0.777	1.713	0.711	2.021
2.021	0.711	1.284					
64 V	0.762	1.137	1.804	1.158	1.613	0.687	2.077
2.077	0.687	1.320					
65 S	1.122	2.046	1.889	1.507	1.704	1.268	1.831
2.046	1.122	1.624					
66 H	1.350	1.549	1.636	1.510	1.385	1.249	1.612
1.636	1.249	1.470					
67 P	1.483	1.950	1.823	0.738	1.586	1.855	1.391
1.950	0.738	1.546					
68 E	1.849	2.201	2.188	-0.183	1.868	1.872	1.022
2.201	-0.183	1.545					
69 G	1.767	1.984	2.234	-0.929	1.868	1.872	0.903
2.234	-0.929	1.386					
70 R	2.077	1.716	2.374	-0.665	2.005	1.293	0.523
2.374	-0.665	1.332					
71 P	2.077	0.902	2.374	0.214	2.005	1.293	0.523
2.374	0.214	1.341					
72 T	1.717	0.544	2.290	1.381	1.914	0.712	0.769
2.290	0.544	1.332					
73 N	1.489	0.053	2.300	1.819	1.959	0.712	-0.242
2.300	-0.242	1.156					
74 P	1.356	-0.017	1.870	1.546	1.485	0.087	-1.251
1.870	-1.251	0.725					
75 P	1.356	-0.376	1.627	0.454	1.212	0.068	-2.481
1.627	-2.481	0.266					
76 A	1.660	0.079	1.702	-0.333	1.376	0.537	-2.531
1.702	-2.531	0.356					
77 A	1.350	-0.126	1.403	-0.940	1.066	0.497	-3.428
1.403	-3.428	-0.025					
78 A	1.483	0.483	1.589	-0.890	1.267	1.102	-3.649
1.589	-3.649	0.198					
79 D	0.768	1.297	1.262	-0.964	1.002	1.089	-3.434

1.297	-3.434	0.146						
80 A	1.078	0.584	1.561	-0.776	1.312	1.129	-2.536	
1.561	-2.536	0.336						
81 R	1.211	0.259	1.991	-0.808	1.786	1.754	-1.527	
1.991	-1.527	0.667						
82 L	0.496	0.301	1.926	-0.338	1.741	1.758	-0.266	
1.926	-0.338	0.803						
83 N	-0.642	1.133	1.515	-0.289	1.403	1.271	0.065	
1.515	-0.642	0.637						
84 R	-0.363	0.523	1.664	-0.152	1.558	1.291	1.235	
1.664	-0.363	0.822						
85 F	-0.269	0.566	1.225	-0.528	1.039	0.667	1.236	
1.236	-0.528	0.562						
86 I	0.446	0.740	1.309	-0.664	1.030	0.661	-0.209	
1.309	-0.664	0.473						
87 S	0.414	1.423	1.160	-0.749	0.875	0.641	0.064	
1.423	-0.749	0.547						
88 G	0.281	0.471	0.730	-0.528	0.401	0.016	-0.946	
0.730	-0.946	0.061						
89 A	0.996	0.335	1.038	-0.381	0.720	0.031	-0.977	
1.038	-0.977	0.252						
90 S	1.268	0.962	1.057	-0.289	0.729	0.031	-0.710	
1.268	-0.710	0.435						
91 A	1.186	0.465	1.103	-0.337	0.729	0.030	-0.829	
1.186	-0.829	0.335						
92 P	1.186	0.465	1.103	-0.501	0.729	0.030	-0.829	
1.186	-0.829	0.312						
93 V	1.186	0.107	1.346	-0.501	1.002	0.049	0.401	
1.346	-0.501	0.513						
94 T	0.907	0.203	1.197	-0.706	0.847	0.029	-0.769	
1.197	-0.769	0.244						
95 G	0.907	-0.384	1.197	-0.690	0.847	0.029	-0.769	
1.197	-0.769	0.163						
96 P	0.907	-0.198	0.954	-0.988	0.574	0.010	-1.999	
0.954	-1.999	-0.106						
97 A	0.907	-0.066	0.954	-1.304	0.574	0.010	-1.999	
0.954	-1.999	-0.132						
98 A	0.844	0.293	1.188	-1.807	0.893	0.615	-2.040	
1.188	-2.040	-0.002						
99 A	0.813	0.784	1.393	-1.978	1.093	0.635	-2.000	
1.393	-2.000	0.106						
100V	0.813	1.143	1.393	-1.890	1.093	0.635	-2.000	
1.393	-2.000	0.169						
101R	1.059	1.778	1.720	-1.375	1.467	0.677	-0.827	
1.778	-1.375	0.643						
102T	1.059	1.323	1.963	-0.663	1.741	0.696	0.403	
1.963	-0.663	0.932						
103P	1.559	1.371	2.234	0.137	2.060	1.185	1.405	
2.234	0.137	1.421						
104Q	1.925	1.012	2.599	0.770	2.342	1.203	1.036	
2.599	0.770	1.555						
105P	2.292	1.377	2.440	1.412	2.187	1.067	1.028	
2.440	1.028	1.686						
106D	2.096	0.814	2.244	1.610	2.032	1.047	-0.023	
2.244	-0.023	1.403						
107P	2.374	0.902	2.150	1.687	1.914	1.048	-0.083	
2.374	-0.083	1.427						

108D	1.413	0.580	1.739	1.241	1.549	1.011	0.188
1.739	0.188	1.103					
109A	1.641	0.668	1.487	0.654	1.230	0.992	-0.031
1.641	-0.031	0.949					
110S	1.097	1.207	1.001	0.007	0.820	0.521	0.600
1.207	0.007	0.750					
111L	1.325	0.978	0.748	-0.388	0.501	0.502	0.380
1.325	-0.388	0.578					
112G	1.325	2.038	0.748	-0.418	0.501	0.502	0.380
2.038	-0.418	0.725					
113C	1.552	1.770	0.739	-0.147	0.455	0.502	1.391
1.770	-0.147	0.895					
114G	1.552	1.734	0.739	0.255	0.455	0.502	1.391
1.734	0.255	0.947					
115D	2.267	1.682	1.066	0.631	0.720	0.516	1.176
2.267	0.516	1.151					
116G	2.039	1.143	1.075	0.783	0.765	0.516	0.165
2.039	0.165	0.927					
117S	2.444	0.109	1.617	0.469	1.221	1.097	-0.483
2.444	-0.483	0.925					
118P	2.216	-0.747	1.627	-0.046	1.267	1.097	-1.494
2.216	-1.494	0.560					
119A	1.464	-0.250	1.608	-0.879	1.185	0.628	-1.153
1.608	-1.153	0.372					
120E	1.236	0.325	1.617	-1.448	1.230	0.628	-2.163
1.617	-2.163	0.204					
121A	1.236	-0.454	1.617	-1.552	1.230	0.628	-2.163
1.617	-2.163	0.078					
122Y	1.597	-0.096	1.702	-1.365	1.321	1.208	-2.409
1.702	-2.409	0.280					
123A	0.882	0.850	1.617	-1.004	1.330	1.214	-0.964
1.617	-1.004	0.561					
124S	0.522	0.646	1.533	-0.808	1.239	0.633	-0.718
1.533	-0.808	0.435					
125E	1.021	0.646	1.804	-0.575	1.558	1.122	0.283
1.804	-0.575	0.837					
126L	0.560	0.698	1.468	-0.446	1.330	1.108	0.386
1.468	-0.446	0.729					
127P	0.838	1.261	1.617	0.103	1.485	1.128	1.556
1.617	0.103	1.141					
128D	0.787	1.393	1.459	0.280	1.285	1.108	1.397
1.459	0.280	1.101					
129L	0.427	1.213	1.375	0.483	1.194	0.528	1.643
1.643	0.427	0.980					
130S	1.337	2.231	1.655	0.380	1.339	0.542	1.249
2.231	0.380	1.248					
131G	1.337	1.375	1.655	0.361	1.339	0.542	1.249
1.655	0.361	1.123					
132P	0.971	1.107	1.814	0.136	1.494	0.678	1.257
1.814	0.136	1.065					
133T	1.685	1.239	1.898	-0.000	1.485	0.672	-0.188
1.898	-0.188	0.970					
134P	1.407	1.561	1.991	-0.452	1.604	0.671	-0.128
1.991	-0.452	0.951					
135R	1.426	1.812	2.328	-0.716	2.023	0.714	0.035
2.328	-0.716	1.089					
136A	1.559	1.357	2.515	-0.935	2.224	1.319	-0.186

2.515	-0.935	1.122						
137P		1.672	1.357	2.617	-0.506	2.379	1.340	-0.340
2.617	-0.506	1.217						
138Q		1.672	1.357	2.617	0.003	2.379	1.340	-0.340
2.617	-0.340	1.290						
139R		1.540	0.866	2.188	0.635	1.905	0.715	-1.349
2.188	-1.349	0.928						
140N		1.540	0.866	2.431	1.013	2.178	0.734	-0.119
2.431	-0.119	1.235						
141P		1.540	0.616	2.188	0.807	1.905	0.715	-1.349
2.188	-1.349	0.917						
142A		1.426	0.257	2.290	0.128	2.005	1.298	-1.513
2.290	-1.513	0.841						
143P		1.293	0.832	2.103	-0.447	1.804	0.692	-1.292
2.103	-1.292	0.712						
144A		0.983	1.101	1.804	-1.094	1.494	0.651	-2.190
1.804	-2.190	0.393						
145R		1.344	1.728	1.889	-1.314	1.586	1.232	-2.436
1.889	-2.436	0.575						
146P		1.571	0.914	1.879	-1.382	1.540	1.232	-1.425
1.879	-1.425	0.618						
147A		1.799	1.183	1.627	-1.554	1.221	1.213	-1.645
1.799	-1.645	0.549						
148E		1.799	2.038	1.627	-1.709	1.221	1.213	-1.645
2.038	-1.709	0.649						
149G		1.894	2.277	1.188	-1.772	0.701	0.588	-1.644
2.277	-1.772	0.462						
150G		2.172	2.277	1.094	-1.624	0.583	0.589	-1.704
2.277	-1.704	0.484						
151A		2.305	2.188	1.524	-1.350	1.057	1.214	-0.695
2.305	-1.350	0.892						
152G		2.172	3.044	1.188	-1.000	0.647	0.615	-0.668
3.044	-1.000	0.857						
153S		2.444	2.417	1.468	-0.656	1.011	1.104	-0.677
2.444	-0.677	1.016						
154R		2.494	1.561	1.627	-0.242	1.212	1.124	-0.518
2.494	-0.518	1.037						
155G		2.494	1.375	1.627	0.145	1.212	1.124	-0.518
2.494	-0.518	1.066						
156D		2.267	1.603	1.636	0.454	1.257	1.124	-1.528
2.267	-1.528	0.973						
157S		2.216	1.920	1.477	0.199	1.057	1.104	-1.688
2.216	-1.688	0.898						
158A		2.362	1.692	1.197	-0.138	0.738	0.499	-1.527
2.362	-1.527	0.689						
159A		2.412	2.319	1.356	-0.325	0.938	0.519	-1.368
2.412	-1.368	0.836						
160G		2.140	3.132	1.075	-0.154	0.574	0.030	-1.358
3.132	-1.358	0.777						
161S		2.090	3.361	0.917	0.214	0.373	0.010	-1.518
3.361	-1.518	0.778						
162S		2.222	2.180	1.346	0.208	0.847	0.635	-0.509
2.222	-0.509	0.990						
163G		2.501	1.816	1.496	-0.132	1.002	0.655	0.661
2.501	-0.132	1.143						
164G		1.634	1.189	1.365	-0.748	1.030	0.657	0.983
1.634	-0.748	0.873						

165R		1.552	1.137	1.412	-0.977	1.030	0.656	0.864
1.552	-0.977	0.810						
166S		1.274	1.179	1.262	-1.130	0.875	0.636	-0.306
1.274	-1.130	0.541						
167I		1.407	1.137	1.599	-1.144	1.285	1.236	-0.333
1.599	-1.144	0.741						
168T		1.457	2.000	1.758	-1.167	1.485	1.256	-0.173
2.000	-1.167	0.945						
169A		1.457	1.509	1.758	-1.125	1.485	1.256	-0.173
1.758	-1.125	0.881						
170E		1.679	2.323	1.879	-0.827	1.649	1.725	-0.342
2.323	-0.827	1.155						
171S		2.317	1.652	2.019	-0.432	1.668	1.723	-1.674
2.317	-1.674	1.039						
172R		2.254	1.287	2.253	-0.284	1.987	2.328	-1.716
2.328	-1.716	1.158						
173D		1.887	0.269	2.132	-0.538	1.977	2.329	-0.117
2.329	-0.538	1.134						
174A		1.774	0.586	2.132	-0.915	1.987	1.772	0.073
2.132	-0.915	1.058						
175R		0.781	0.586	1.898	-1.537	1.841	1.757	0.348
1.898	-1.537	0.810						
176V		0.926	0.586	1.617	-1.420	1.522	1.153	0.509
1.617	-1.420	0.699						
177Q		0.427	1.495	1.346	-1.308	1.203	0.664	-0.492
1.495	-1.308	0.476						
178L		0.560	1.860	1.776	-0.952	1.677	1.289	0.517
1.860	-0.952	0.961						
179S		0.560	2.878	1.776	-1.005	1.677	1.289	0.517
2.878	-1.005	1.099						
180A		1.205	2.649	2.047	-1.110	1.841	1.307	0.088
2.649	-1.110	1.147						
181R		1.091	3.008	2.150	-1.437	1.941	1.889	-0.076
3.008	-1.437	1.224						
182R		2.033	2.098	2.225	-1.326	1.886	1.884	-0.511
2.225	-1.326	1.184						
183S		1.755	2.098	2.318	-1.190	2.005	1.883	-0.451
2.318	-1.190	1.203						
184R		1.388	1.243	2.197	-0.994	1.996	1.884	1.148
2.197	-0.994	1.266						
185G		1.388	1.285	2.197	-1.041	1.996	1.884	1.148
2.197	-1.041	1.265						
186P		1.255	0.544	1.767	-1.216	1.522	1.260	0.139
1.767	-1.216	0.753						
187V		1.255	0.676	1.767	-1.392	1.522	1.260	0.139
1.767	-1.392	0.747						
188R		0.724	0.447	1.328	-1.461	1.093	0.652	0.084
1.328	-1.461	0.410						
189A		0.743	0.447	1.664	-1.238	1.513	0.695	0.247
1.664	-1.238	0.582						
190S		0.104	1.261	1.281	-1.274	1.221	0.678	0.349
1.281	-1.274	0.517						
191M		0.604	0.081	1.832	-1.399	1.704	1.301	-0.240
1.832	-1.399	0.555						
192Q		0.604	0.734	1.832	-1.902	1.704	1.301	-0.240
1.832	-1.902	0.576						
193I		-0.035	0.602	1.692	-2.420	1.686	1.302	1.091

1.692	-2.420	0.560						
194R	0.187	0.423	1.814	-2.611	1.850	1.771	0.923	
1.850	-2.611	0.622						
195R	0.585	0.465	2.066	-2.119	2.078	1.773	1.198	
2.078	-2.119	0.864						
196I	-0.427	0.143	1.758	-1.402	1.722	1.756	1.057	
1.758	-1.402	0.658						
197D	0.490	0.263	2.047	-0.428	1.895	1.774	0.895	
2.047	-0.428	0.991						
198P	0.553	0.556	1.814	0.155	1.576	1.169	0.936	
1.814	0.155	0.966						
199W	-0.294	0.101	1.300	0.259	1.112	0.550	1.372	
1.372	-0.294	0.628						
200S	0.572	1.459	1.889	0.020	1.768	1.143	1.116	
1.889	0.020	1.138						
201T	-0.294	0.399	1.496	-0.629	1.440	0.656	1.713	
1.713	-0.629	0.683						
202L	-0.016	-0.296	1.403	-1.092	1.321	0.657	1.653	
1.653	-1.092	0.519						
203K	0.035	-0.296	1.300	-1.319	1.312	0.637	2.066	
2.066	-1.319	0.533						
204V	-0.958	-0.272	1.066	-1.318	1.166	0.622	2.341	
2.341	-1.318	0.378						
205S	-1.868	-0.272	0.786	-1.305	1.020	0.608	2.735	
2.735	-1.868	0.243						
206L	-0.876	-1.127	1.019	-1.217	1.166	0.622	2.460	
2.460	-1.217	0.293						
207L	-1.470	-1.127	0.449	-1.332	0.519	0.029	2.984	
2.984	-1.470	0.008						
208L	-1.103	-1.097	0.571	-1.185	0.528	0.028	1.385	
1.385	-1.185	-0.125						
209S	-2.096	-1.067	0.337	-1.234	0.382	0.013	1.660	
1.660	-2.096	-0.286						
210V	-2.096	-2.019	0.356	-1.206	0.328	0.012	1.476	
1.476	-2.096	-0.450						
211A	-2.096	-2.426	0.375	-1.346	0.273	0.011	1.293	
1.293	-2.426	-0.560						
212L	-1.748	-2.540	0.337	-1.311	0.255	0.007	1.446	
1.446	-2.540	-0.508						
213F	-2.791	-2.660	0.206	-1.315	0.118	0.012	1.309	
1.309	-2.791	-0.732						
214F	-2.823	-1.995	0.318	-1.385	0.173	0.028	0.665	
0.665	-2.823	-0.717						
215V	-3.461	-1.917	0.178	-1.849	0.155	0.029	1.997	
1.997	-3.461	-0.695						
216W	-2.551	-1.821	0.459	-2.284	0.300	0.044	1.602	
1.602	-2.551	-0.607						
217M	-2.203	-1.492	0.403	-2.573	0.337	0.041	1.940	
1.940	-2.573	-0.507						
218I	-1.489	-1.582	0.468	-2.452	0.382	0.037	0.679	
0.679	-2.452	-0.565						
219T	-1.837	-1.664	0.524	-2.039	0.346	0.040	0.341	
0.524	-2.039	-0.613						
220V	-1.786	-2.360	0.421	-1.734	0.337	0.020	0.754	
0.754	-2.360	-0.621						
221A	-1.641	-2.360	0.683	-1.524	0.528	0.022	1.141	
1.141	-2.360	-0.450						

222F	-1.716	-2.564	0.739	-1.506	0.556	0.026	1.254
1.254	-2.564	-0.459					
223L	-2.279	-1.763	0.421	-1.657	0.392	0.007	1.802
1.802	-2.279	-0.439					
224Y	-2.627	-0.931	0.459	-1.866	0.410	0.011	1.649
1.649	-2.627	-0.414					
225L	-2.399	-0.639	0.449	-2.030	0.364	0.011	2.659
2.659	-2.399	-0.226					
226V	-1.457	0.193	0.505	-2.181	0.364	0.007	2.408
2.408	-2.181	-0.023					
227L	-1.141	0.193	0.580	-2.201	0.401	0.019	1.918
1.918	-2.201	-0.033					
228G	-0.661	-0.106	0.318	-2.213	0.118	-0.001	1.586
1.586	-2.213	-0.137					
229G	-0.313	-0.733	0.281	-2.242	0.100	-0.004	1.740
1.740	-2.242	-0.167					
230M	-0.711	-0.528	0.421	-2.302	0.127	0.019	1.173
1.173	-2.302	-0.257					
231G	0.003	-0.619	0.505	-2.325	0.118	0.014	-0.272
0.505	-2.325	-0.368					
232V	0.003	-0.637	0.963	-2.345	0.802	0.608	-0.207
0.963	-2.345	-0.116					
233W	-0.939	0.315	0.889	-2.399	0.856	0.614	0.228
0.889	-2.399	-0.062					
234A	-0.231	1.427	1.197	-1.945	1.121	0.637	0.170
1.427	-1.945	0.339					
235K	-0.180	1.331	1.356	-1.168	1.321	0.657	0.330
1.356	-1.168	0.521					
236L	0.496	1.127	1.776	0.408	1.640	0.696	-0.372
1.776	-0.372	0.825					
237N	0.895	1.870	1.636	1.766	1.613	0.673	0.194
1.870	0.194	1.235					
238S	1.122	1.056	1.627	2.503	1.567	0.673	1.205
2.503	0.673	1.393					
239N	1.394	-0.003	1.449	2.271	1.248	0.567	1.131
2.271	-0.003	1.151					
240V	1.394	-0.003	1.449	1.410	1.248	0.567	1.131
1.449	-0.003	1.028					
241G	0.370	0.702	1.066	0.316	0.948	0.532	1.679
1.679	0.316	0.802					
242D	0.402	0.075	1.216	0.053	1.103	0.552	1.406
1.406	0.053	0.687					
243L	0.402	0.391	1.216	0.334	1.103	0.552	1.406
1.406	0.334	0.772					
244L	0.768	1.223	1.337	1.134	1.112	0.551	-0.193
1.337	-0.193	0.847					
245N	0.819	2.283	1.496	2.098	1.312	0.571	-0.034
2.283	-0.034	1.221					
246N	0.547	2.529	1.216	2.216	0.948	0.082	-0.024
2.529	-0.024	1.073					
247A	1.540	1.920	1.449	1.964	1.093	0.097	-0.299
1.964	-0.299	1.109					
248S	2.532	2.495	1.683	1.463	1.239	0.111	-0.575
2.532	-0.575	1.278					
249G	2.222	1.435	1.384	1.029	0.929	0.071	-1.472
2.222	-1.472	0.800					
250S	2.273	0.712	1.412	0.684	0.984	0.630	-1.385

2.273	-1.385	0.758						
251S	1.559	0.712	1.328	0.272	0.993	0.635	0.060	
1.559	0.060	0.794						
252A	0.914	0.712	1.057	-0.674	0.829	0.617	0.489	
1.057	-0.674	0.563						
253E	0.964	1.339	1.216	-1.237	1.030	0.637	0.649	
1.339	-1.237	0.657						
254L	0.964	1.255	1.216	-1.250	1.030	0.637	0.649	
1.255	-1.250	0.643						
255V	0.914	1.135	1.057	-0.704	0.829	0.617	0.489	
1.135	-0.704	0.619						
256S	1.110	1.056	1.253	0.111	0.984	0.636	1.540	
1.540	0.111	0.956						
257S	0.111	0.828	0.786	0.282	0.601	0.039	1.888	
1.888	0.039	0.648						
258G	0.111	0.600	0.804	-0.028	0.547	0.037	1.704	
1.704	-0.028	0.539						
259T	0.705	-0.028	0.917	-0.674	0.510	0.036	1.116	
1.116	-0.674	0.369						
260I	0.655	-0.693	0.758	-1.179	0.310	0.016	0.956	
0.956	-1.179	0.117						
261F	0.376	-0.572	0.608	-1.480	0.155	-0.004	-0.214	
0.608	-1.480	-0.162						
262G	-0.566	-0.723	0.552	-1.408	0.155	0.000	0.037	
0.552	-1.408	-0.279						
263G	-1.476	-0.723	0.272	-1.507	0.009	-0.015	0.431	
0.431	-1.507	-0.430						
264A	-1.476	-1.554	0.272	-1.558	0.009	-0.015	0.431	
0.431	-1.558	-0.556						
265F	-0.534	-1.650	0.328	-1.813	0.009	-0.019	0.180	
0.328	-1.813	-0.500						
266L	-1.476	-0.867	0.253	-2.103	0.063	-0.013	0.615	
0.615	-2.103	-0.504						
267I	-2.070	-0.987	0.141	-2.353	0.100	-0.012	1.203	
1.203	-2.353	-0.568						
268G	-1.761	-0.759	0.440	-1.959	0.410	0.029	2.101	
2.101	-1.959	-0.214						
269L	-1.685	-1.590	0.365	-1.469	0.437	0.026	2.171	
2.171	-1.685	-0.249						
270V	-1.337	-1.500	0.328	-0.729	0.419	0.023	2.325	
2.325	-1.500	-0.067						
271N	-1.413	-0.913	0.384	-0.525	0.446	0.026	2.438	
2.438	-1.413	0.063						
272I	-2.039	-1.522	0.384	-0.977	0.537	0.043	2.383	
2.383	-2.039	-0.170						
273V	-1.128	-1.402	0.664	-1.659	0.683	0.058	1.988	
1.988	-1.659	-0.114						
274L	-0.762	-1.306	0.786	-2.065	0.692	0.056	0.390	
0.786	-2.065	-0.315						
275M	-1.786	-0.611	0.403	-2.270	0.392	0.021	0.938	
0.938	-2.270	-0.416						
276T	-1.147	-0.821	0.543	-2.034	0.410	0.019	-0.394	
0.543	-2.034	-0.489						
277A	-0.585	-0.685	0.860	-1.853	0.574	0.038	-0.942	
0.860	-1.853	-0.370						
278L	-0.509	-0.685	0.804	-1.885	0.547	0.034	-1.055	
0.804	-1.885	-0.393						

279A	0.117	-0.655	0.804	-1.810	0.455	0.017	-1.000
0.804	-1.810	-0.296					
280T	-0.079	-0.751	0.608	-1.929	0.300	-0.003	-2.051
0.608	-2.051	-0.558					
281I	-0.793	-1.648	0.543	-1.860	0.255	0.001	-0.790
0.543	-1.860	-0.613					
282G	-0.446	-0.715	0.505	-1.832	0.237	-0.003	-0.636
0.505	-1.832	-0.413					
283A	-0.698	-1.546	0.758	-1.611	0.474	0.017	0.706
0.758	-1.611	-0.272					
284F	-0.585	-1.871	0.860	-1.071	0.629	0.037	0.553
0.860	-1.871	-0.207					
285V	-0.661	-1.206	0.917	-0.452	0.656	0.041	0.666
0.917	-1.206	-0.006					
286Y	-1.527	-0.570	0.786	-0.005	0.683	0.043	0.987
0.987	-1.527	0.057					
287N	-1.331	-0.368	0.982	0.082	0.838	0.063	2.038
2.038	-1.331	0.329					
288L	-0.117	-1.302	1.318	-0.116	1.203	0.547	1.778
1.778	-1.302	0.473					
289I	-0.465	-0.470	1.356	-0.430	1.221	0.551	1.624
1.624	-0.470	0.484					
290T	-0.850	0.481	0.963	-0.519	0.966	0.534	1.614
1.614	-0.850	0.456					
291D	-0.932	-0.334	0.655	-0.831	0.610	0.493	1.727
1.727	-0.932	0.198					
292L	0.010	-0.298	0.730	-1.183	0.556	0.488	1.293
1.293	-1.183	0.228					
293I	0.010	-0.190	0.730	-1.780	0.556	0.488	1.293
1.293	-1.780	0.158					
294G	0.174	0.626	0.860	-2.237	0.765	1.067	1.225
1.225	-2.237	0.354					
295G	-0.692	-0.206	0.468	-2.531	0.437	0.580	1.823
1.823	-2.531	-0.017					
296I	0.218	-0.833	0.748	-2.414	0.583	0.595	1.429
1.429	-2.414	0.047					
297E	0.142	0.031	0.804	-2.238	0.610	0.598	1.542
1.542	-2.238	0.213					
298V	-0.085	0.269	0.814	-1.974	0.656	0.598	0.532
0.814	-1.974	0.116					
299T	0.187	0.904	1.094	-1.528	1.020	1.087	0.523
1.094	-1.528	0.470					
300L	0.958	0.668	1.664	-1.188	1.513	1.710	0.200
1.710	-1.188	0.789					
301A	1.097	1.127	1.608	-0.627	1.467	1.599	0.217
1.608	-0.627	0.927					
302D	1.331	1.381	1.272	-0.165	1.795	1.658	-1.381
1.795	-1.381	0.842					
303R	1.002	1.097	0.618	0.230	1.959	1.698	-2.432
1.959	-2.432	0.596					
304D	1.584	0.538	0.244	0.355	2.269	1.753	-3.877
2.269	-3.877	0.409					

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	<p>¹VTAPNEPGALSKGDGPNADGLVDRGGAHRAATGPGRIPDAGDPPPWQRAATRQS QAGHRQPPPVSHP EGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQPDPDAS LGCGDGSPAEAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRS ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLLSV ALFFVWMITVAFL YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL ATIGAFVYNLITDLIGGIEVTLADR³⁰⁴</p>
Hydrophilicity	<p>¹<u>VTAPNEPGALSKGDGPNADGLVDRGGAHR</u>AATGPGRIPDAGDPPPWQR<u>AATRQS</u> <u>QAGHRQPPPVSHP EGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQPDPDAS</u> <u>LGCGDGSPAEAYASELPDLSGPTPRAPQRNPAPARP</u><u>AEGGAGSRGDSAAGSSGGRS</u> <u>ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLLSV ALFFVWMITVAFL</u> <u>YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL</u> <u>ATIGAFVYNLITDLIGGIEVTLADR³⁰⁴</u></p>
Flexibility	<p>¹VTAPNE<u>PGALSKGDGPNADGLVDRGGAHR</u>AATGPGRIPDAGDPPPWQR<u>AATRQS</u> <u>QAGHRQPPPVSHP EGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQPDPDAS</u> <u>LGCGDGSPAEAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRS</u> <u>ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLLSV ALFFVWMITVAFL</u> <u>YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL</u> <u>ATIGAFVYNLITDLIGGIEVTLADR³⁰⁴</u></p>
Accessibility	<p>¹VTAPNEPGAL<u>SKGDGPNADGLVDRGGAHR</u>AATGPGRIPDAGDPPPWQRAATRQS <u>QAGHRQPPPVSHP EGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQPDPDAS</u> LGCGDGSPAEAYASELPDLS<u>GPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRS</u> <u>ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLLSV ALFFVWMITVAFL</u> <u>YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL</u> <u>ATIGAFVYNLITDLIGGIEVTLADR³⁰⁴</u></p>
Turns	<p>¹VTAPNEPGALSKGDGPNADGLVDRGGAHRAATGPGRIPDAGDPPPWQRAATRQS QAGHRQPPPVSHP EGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQPDPDAS LGCGDGSPAEAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRS ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLLSV ALFFVWMITVAFL YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL ATIGAFVYNLITDLIGGIEVTLADR³⁰⁴</p>

Exposure Surface	¹ VTAPNEPGALSKGDGPNADGLVDRGGAHRAATGPGRIPDAGDPPPWQRAATRQS QAGHRQPPPVSHPEGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQDPDAS LGCGDGSPAAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRS ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLSVALFFVWMITVAFL YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL ATIGAFVYNLITDLIGGIEVTLADR ³⁰⁴
Polarity	¹ VTAPNEPGALSKGDGPNADGLVDRGGAHRAATGPGRIPDAGDPPPWQRAATRQS QAGHRQPPPVSHPEGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQDPDAS LGCGDGSPAAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRS ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLSVALFFVWMITVAFL YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL ATIGAFVYNLITDLIGGIEVTLADR ³⁰⁴
Antigenic Property	¹ VTAPNEPGALSKGDGPNADGLVDRGGAHRAATGPGRIPDAGDPPPWQRAATRQS QAGHRQPPPVSHPEGRPTNPPAAADARLNRFISGASAPVTGPAAAVRTPQDPDAS LGCGDGSPAAYASELPDLSGPTPRAPQRNPAPARPAEGGAGSRGDSAAGSSGGRS ITAESRDARVQLSARRSRGPVRASMQIRRIDPWSTLKVSLLSVALFFVWMITVAFL YLVLGGMGVWAKLNSNVGDLLNNASGSSAELVSSGTIFGGAFLIGLVNIVLMTAL ATIGAFVYNLITDLIGGIEVTLADR ³⁰⁴

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