

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

seqname=

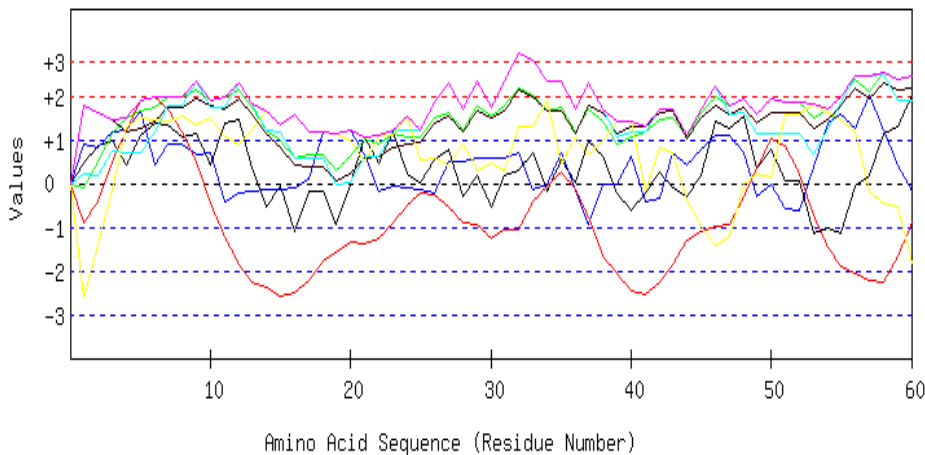
Seq=MTTPSHLSDRYELGEILGFGGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYLRFRREA
QNAAALNHPAIVAVYDTGEAETPAGPLPYIVMEYVDGVTLRDIVHTEGPMPKRAIEVIADACQALNFSHQNGIIH
RDVKPANIMISATNAVKVMDFGIARAIADSGNSVTQTAAVIGTAQYLSPEQARGDSVDARSVDVYSLGCVLYEVLG
EPPFTGDSPVSVAYQHVREDPIPPSARHEGLSADLDAVVLKALAKNPENRYQTAAEMRADLVRVHNGEPPEAPKV
LTDAERTSLLSSAAGNLSGPRTDPLPRQDLDDTDRDRSIGSVGRWVAVVAVLAVLTVVVTIAINTFGGITRDVQVPD
VRGQSSADAIATLQNRGFKIRTLQKPDSTIPPDHVIGTDPAANTSVSAGDEITVNVSTGPEQREIPDVSTLTYAEAVK
KLTAAGFGRFKQANSPSTPELVGKVIGTNPPANQTSAITNVVIIIIVGSGPATKDIPDVAGQTVDVAQKNLNVYGFTK
FSQASVDSPPAGEVTGTNPPAGTTVPVDSVIELQVSKGNQFVMPDLSGMFWVDAEPRLRALGWTGMLDKGAD
VDAGGSQHNRVVYQNPPAGTGVNRDGIITLRFQQ

Length=626

GRAPHICAL

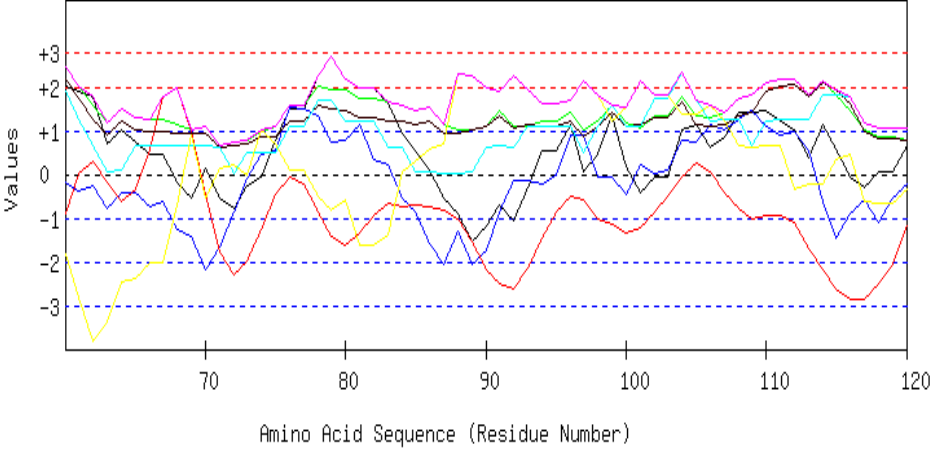
RESULT

GRAPHICAL RESULT :: SEQ 1 to 60



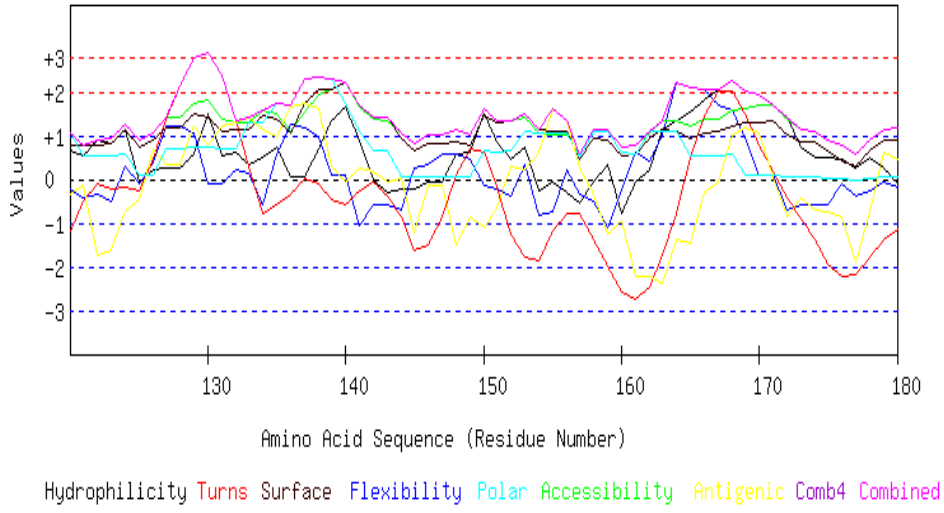
Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 61 to 120

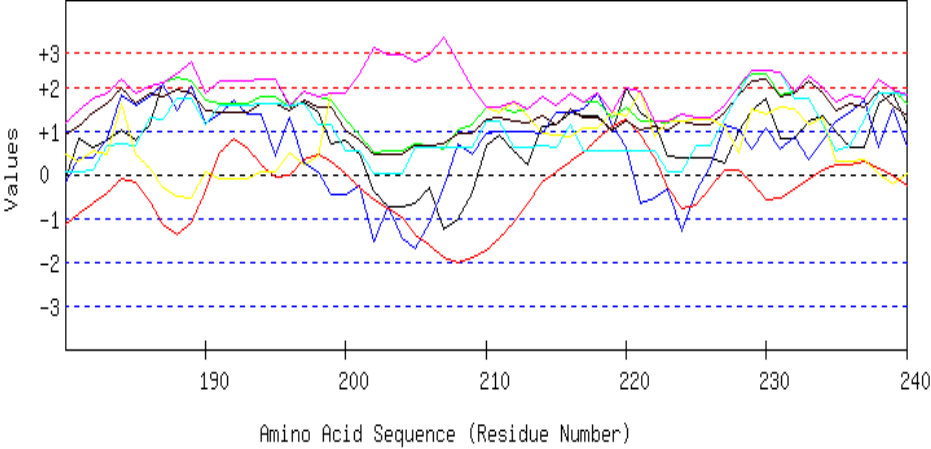


Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 121 to 180

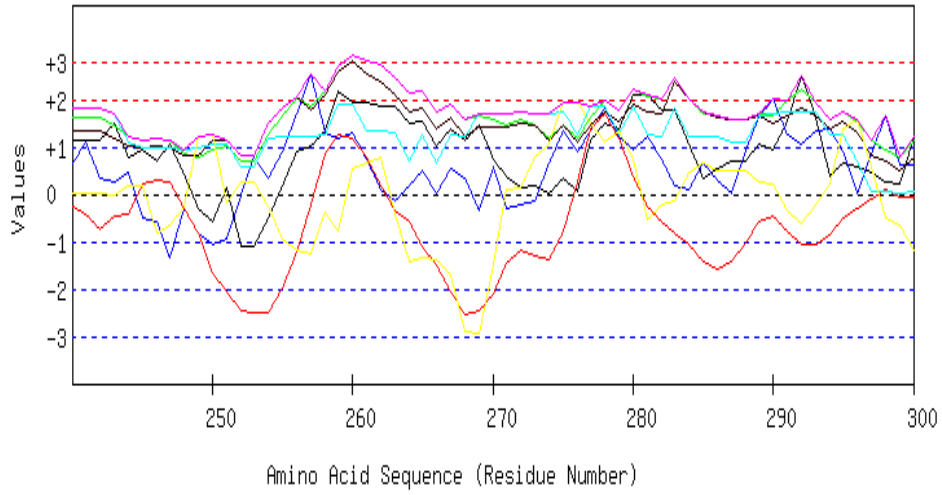


GRAPHICAL RESULT :: SEQ 181 to 240



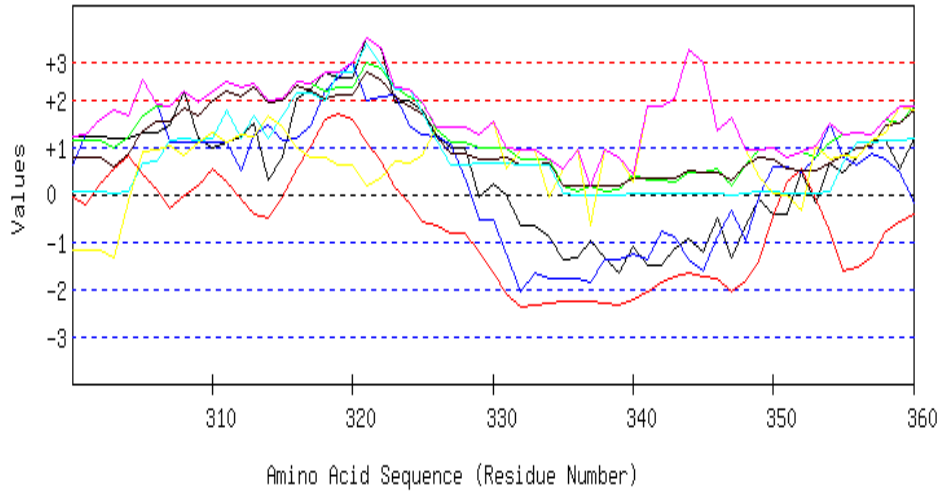
Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 241 to 300



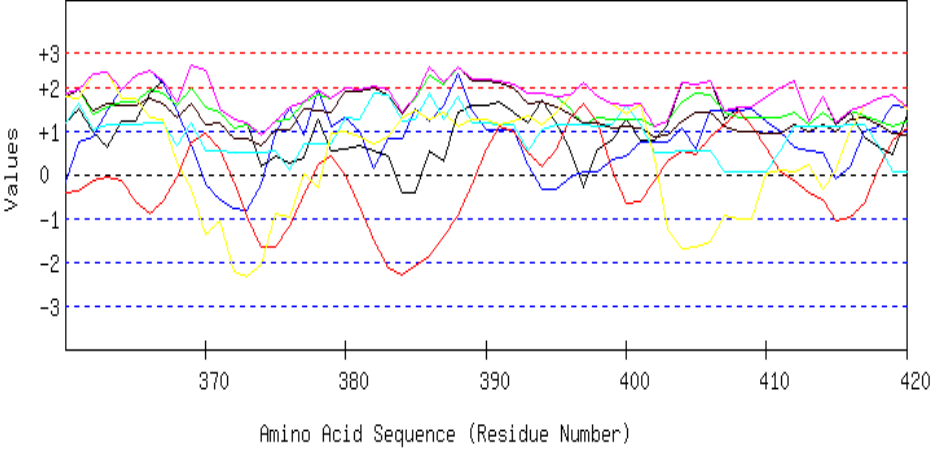
Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 301 to 360



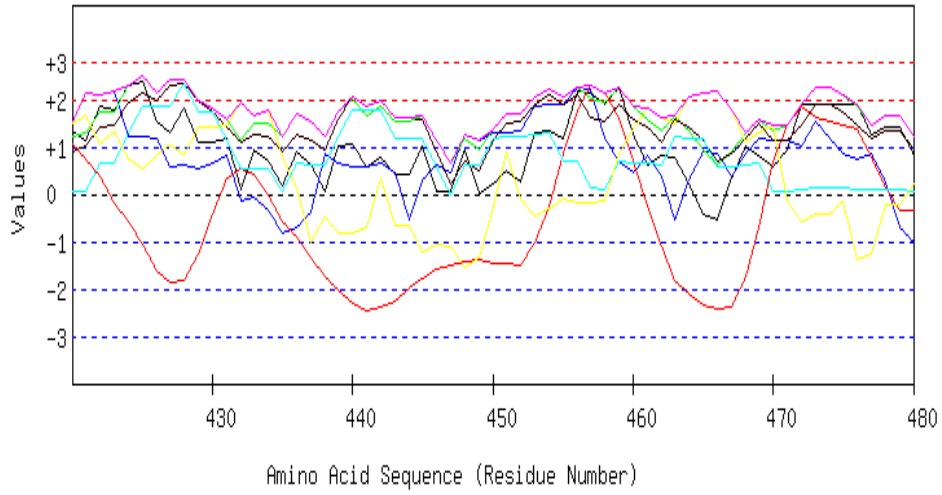
Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 361 to 420



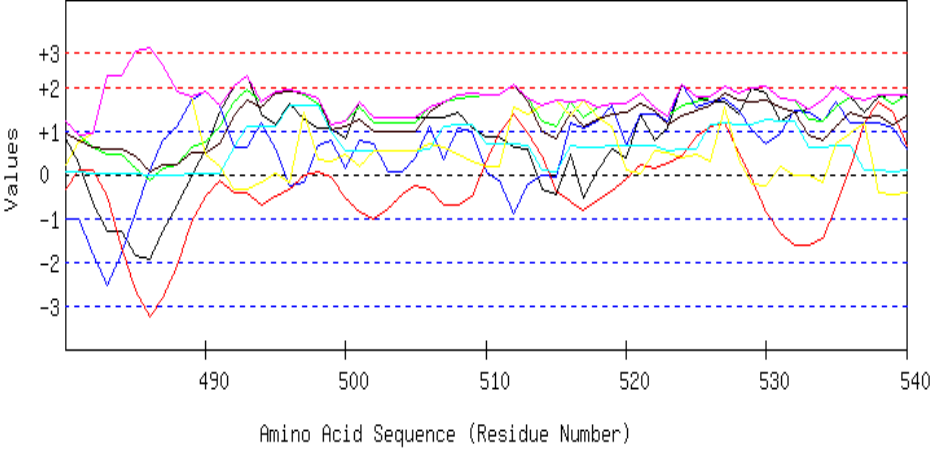
Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 421 to 480



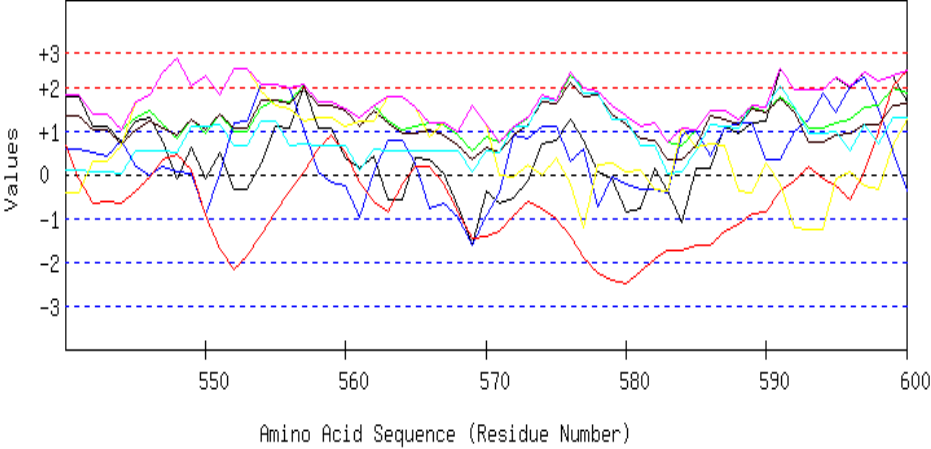
Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 481 to 540



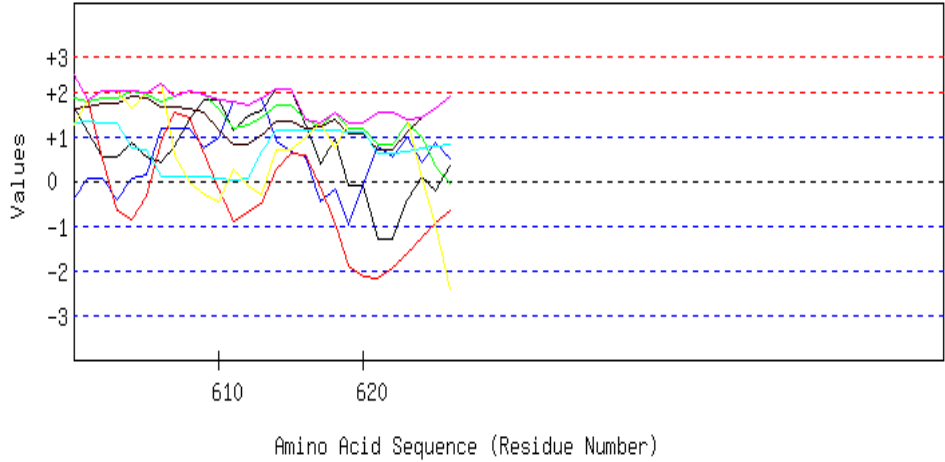
Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 541 to 600



Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

GRAPHICAL RESULT :: SEQ 601 to 660



Hydrophilicity Turns Surface Flexibility Polar Accessibility Antigenic Comb4 Combined

[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

MTTPSHLSDRYELGEILGFQGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYLRFRREA
QNAAALNHPAIVAVYDTGEAETPAGPLPYIVMEYVDGVTLLRDIVHTEGPMTPKRAIEVIA
DACQALNFHQNGIIHRDVKPANIMISATNAVKVMDFGIARAIADSGNSVTQTAAVIGTA
QYLSPEQARGDSVDARSDVYSLGCVLYEVLGTGEPFPTGDSPVSVAYQHVREDPIPPSARH
EGLSADLDAVVLKALAKNPENRYQTAAEMRADLVRVHNGEPPEAPKVLTDARTSLLSSA
AGNLSGPRTDPLPRQDLDDTDRDRSIGSVGRWVAVVAVLAVLTVVVTIAINTFGGITRDV
QVPDVRGQSSADAIAATLQNRGFKIRTLOKPDSTIPPDHVIGTDPAAANTSVSAGDEITVNV
STGPEQREIPDVSTLTLYAEAVKQLTAAGFGRFKQANSPTPELVGKVIGTNPPANQTSAI
TNVVIIVGSGPATKDI PDVAGQTVDVAAQKNLNVYGF TKFSQASVDS PRPAGEVTGTNPP
AGTTVPVDSVIELQVSKGNQFVMPDL SGMFWVDAE PRLRALGWTGMLDKGADV DAGGSQH
NRVYQNPPAGTGVNRDGIITLRFQ

Length=626

A.A.	Parameter							Combined	
	Hydro	Flexi	Access	Turns	Surface	Polar	AntiPro	MAX	MIN
AVG									
1 M -0.033	0.446	0.898	-0.102	-0.881	1.786	0.226	-2.602	1.786	-2.602
2 T 0.301	0.857	0.808	0.505	-0.436	1.622	0.186	-1.432	1.622	-1.432
3 T 0.836	0.990	1.173	1.122	0.501	1.476	0.746	-0.155	1.476	-0.155
4 P 1.070	0.408	1.221	1.496	1.217	1.166	0.691	1.290	1.496	0.408
5 S 1.396	1.084	1.676	1.655	1.883	1.276	0.694	1.505	1.883	0.694
6 H 1.367	1.388	0.413	1.730	1.977	1.440	1.163	1.456	1.977	0.413
7 L 1.545	1.325	0.898	1.963	1.685	1.759	1.768	1.414	1.963	0.898
8 S 1.450	1.072	0.898	1.973	1.194	1.722	1.768	1.526	1.973	0.898
9 D 1.440	1.154	0.670	2.150	0.485	1.932	2.348	1.340	2.348	0.485
10 R 1.084	0.440	0.706	1.907	-0.477	1.768	1.733	1.508	1.907	-0.477
11 Y 0.890	1.382	-0.432	1.982	-1.215	1.713	1.728	1.073	1.982	-1.215
12 E 0.953	1.464	-0.230	2.160	-1.842	1.923	2.307	0.887	2.307	-1.842
13 L 0.608	0.326	-0.178	1.748	-2.260	1.586	1.820	1.217	1.820	-2.260
14 G 0.307	-0.521	-0.148	1.234	-2.393	1.121	1.201	1.653	1.653	-2.393
15 E 0.222	-0.041	-0.148	0.973	-2.570	0.838	1.182	1.321	1.321	-2.570
16 I -0.074	-1.116	-0.096	0.580	-2.501	0.428	0.586	1.599	1.599	-2.501
17 L 0.070	-0.174	0.115	0.655	-2.224	0.373	0.581	1.164	1.164	-2.224
18 G 0.283	-0.174	1.175	0.655	-1.795	0.373	0.581	1.164	1.175	-1.795
19 F 0.018	-0.932	1.123	0.318	-1.573	0.054	-0.002	1.135	1.135	-1.573
20 G 0.238	-0.016	1.201	0.608	-1.341	0.228	0.017	0.973	1.201	-1.341
21 G 0.439	1.059	0.664	1.019	-1.378	0.583	0.611	0.512	1.059	-1.378
22 M 0.324	0.465	-0.168	0.907	-1.271	0.619	0.612	1.101	1.101	-1.271
23 S 0.651	1.179	-0.054	1.132	-0.881	0.838	1.228	1.117	1.228	-0.881
24 E 0.623	0.237	-0.096	1.057	-0.514	0.893	1.233	1.551	1.551	-0.514
25 V 0.490	0.010	-0.132	1.066	-0.227	0.938	1.233	0.541	1.233	-0.227
26 H	0.541	-0.240	1.505	-0.270	1.367	1.841	0.595	1.841	-0.270

56 F 1.041	-0.073	1.245	2.365	-2.079	2.187	2.472	1.171	2.472	-2.079
57 R 0.904	0.180	2.028	2.113	-2.228	1.950	2.453	-0.171	2.453	-2.228
58 R 0.998	1.141	1.215	2.524	-2.255	2.315	2.490	-0.443	2.524	-2.255
59 E 0.850	1.318	0.401	2.393	-1.664	2.151	1.906	-0.555	2.393	-1.664
60 A 0.813	2.033	-0.174	2.459	-0.909	2.196	1.902	-1.816	2.459	-1.816
61 Q 0.537	1.900	-0.378	2.029	0.032	1.722	1.277	-2.825	2.029	-2.825
62 N 0.211	1.767	-0.260	1.599	0.308	1.248	0.652	-3.835	1.767	-3.835
63 A -0.213	0.692	-0.779	1.188	-0.166	0.893	0.058	-3.373	1.188	-3.373
64 A 0.040	1.002	-0.420	1.487	-0.616	1.203	0.099	-2.476	1.487	-2.476
65 A 0.087	0.756	-0.420	1.318	-0.354	1.002	0.676	-2.372	1.318	-2.372
66 L 0.162	0.446	-0.745	1.262	0.592	0.966	0.655	-2.039	1.262	-2.039
67 N 0.346	0.446	-0.637	1.262	1.771	0.966	0.655	-2.039	1.771	-2.039
68 H 0.365	-0.193	-1.246	1.122	1.975	0.948	0.657	-0.707	1.975	-1.246
69 P 0.356	-0.559	-1.432	1.001	0.998	0.938	0.658	0.891	1.001	-1.432
70 A -0.056	0.155	-2.197	1.085	-0.465	0.929	0.653	-0.554	1.085	-2.197
71 I -0.276	-0.521	-1.658	0.664	-1.786	0.610	0.614	0.148	0.664	-1.786
72 V -0.324	-0.774	-0.843	0.758	-2.306	0.674	0.013	0.213	0.758	-2.306
73 A -0.055	-0.275	-0.120	0.786	-1.964	0.720	0.483	-0.016	0.786	-1.964
74 V 0.360	-0.079	0.455	0.982	-1.253	0.875	0.503	1.035	1.035	-1.253
75 Y 0.582	0.787	0.552	1.113	-0.442	0.847	0.501	0.713	1.113	-0.442
76 D 0.996	1.514	1.533	1.561	-0.053	1.221	1.099	0.098	1.561	-0.053
77 T 0.966	1.514	1.485	1.561	-0.221	1.221	1.099	0.098	1.561	-0.221
78 G 1.072	2.241	1.353	2.010	-0.879	1.595	1.697	-0.517	2.241	-0.879
79 E 0.906	2.690	0.726	1.954	-1.431	1.513	1.698	-0.808	2.690	-1.431
80 A 0.770	2.191	0.778	1.926	-1.621	1.467	1.228	-0.579	2.191	-1.621
81 E 0.629	1.995	1.137	1.730	-1.350	1.312	1.208	-1.630	1.995	-1.630
82 T 0.573	1.995	0.357	1.730	-0.957	1.312	1.208	-1.630	1.995	-1.630
83 P 0.471	1.634	0.225	1.646	-0.675	1.221	0.627	-1.384	1.646	-1.384
84 A 0.447	0.920	-0.540	1.561	-0.739	1.230	0.633	0.061	1.561	-0.739
85 G	0.560	-0.865	1.477	-0.718	1.139	0.052	0.308	1.477	-0.865

115A	0.566	-1.444	1.832	-2.617	1.923	1.811	0.340	1.923	-2.617
0.344									
116I	-0.073	-0.905	1.449	-2.861	1.631	1.794	0.442	1.794	-2.861
0.211									
117E	-0.300	-0.581	1.001	-2.846	0.993	1.199	-0.634	1.199	-2.846
-0.167									
118V	0.067	-1.119	0.842	-2.492	0.838	1.064	-0.642	1.064	-2.492
-0.206									
119I	0.067	-0.532	0.842	-2.047	0.838	1.064	-0.642	1.064	-2.047
-0.059									
120A	0.661	-0.208	0.767	-1.190	0.765	1.079	-0.341	1.079	-1.190
0.219									
121D	0.547	-0.412	0.767	-0.488	0.774	0.522	-0.151	0.774	-0.488
0.223									
122A	0.914	-0.342	0.889	-0.099	0.784	0.521	-1.750	0.914	-1.750
0.131									
123C	0.838	-0.516	0.945	-0.219	0.811	0.524	-1.637	0.945	-1.637
0.106									
124Q	1.148	0.303	1.244	-0.197	1.121	0.565	-0.740	1.244	-0.740
0.492									
125A	-0.066	-0.098	0.907	-0.247	0.756	0.080	-0.480	0.907	-0.480
0.122									
126L	0.212	0.393	1.057	0.480	0.911	0.100	0.690	1.057	0.100
0.549									
127N	0.256	1.207	1.431	1.370	1.175	0.702	0.335	1.431	0.256
0.925									
128F	0.256	1.225	1.431	2.127	1.175	0.702	0.335	2.127	0.256
1.036									
129S	0.566	1.074	1.730	2.764	1.485	0.743	1.232	2.764	0.566
1.371									
130H	1.508	-0.106	1.804	2.898	1.431	0.738	0.798	2.898	-0.106
1.296									
131Q	0.560	-0.106	1.365	2.374	1.103	0.699	1.232	2.374	-0.106
1.032									
132N	0.636	0.217	1.290	1.333	1.130	0.696	1.303	1.333	0.217
0.944									
133G	0.357	0.147	1.300	0.286	1.148	1.296	1.410	1.410	0.147
0.849									
134I	0.490	-0.576	1.571	-0.781	1.449	1.301	1.142	1.571	-0.781
0.656									
135I	0.743	0.580	1.515	-0.591	1.394	1.748	0.970	1.748	-0.591
0.908									
136H	0.067	1.263	1.094	-0.358	1.075	1.708	1.672	1.708	-0.358
0.931									
137R	0.067	1.173	1.552	0.037	1.759	2.303	1.736	2.303	0.037
1.232									
138D	0.705	0.968	1.935	-0.088	2.050	2.320	1.635	2.320	-0.088
1.361									
139V	1.344	0.105	2.075	-0.450	2.069	2.318	0.303	2.318	-0.450
1.109									
140K	1.653	0.087	2.216	-0.561	2.205	1.739	-0.077	2.216	-0.561
1.037									
141P	0.882	-1.069	1.646	-0.274	1.713	1.116	0.245	1.713	-1.069
0.608									
142A	-0.016	-0.572	1.365	-0.068	1.440	0.645	0.199	1.440	-0.572
0.427									
143N	-0.288	-0.572	1.346	-0.365	1.431	0.645	-0.068	1.431	-0.572
0.304									
144I	-0.237	-0.691	1.047	-0.863	0.948	0.070	0.027	1.047	-0.863

174A	0.484	-0.576	1.103	-1.393	0.856	0.056	-0.683	1.103	-1.393
-0.022									
175A	0.515	-0.576	0.898	-1.957	0.656	0.036	-0.724	0.898	-1.957
-0.165									
176V	0.465	-0.086	0.767	-2.206	0.437	0.013	-0.846	0.767	-2.206
-0.208									
177I	0.269	-0.396	0.571	-2.182	0.282	-0.007	-1.897	0.571	-2.182
-0.480									
178G	0.515	-0.276	0.898	-1.778	0.656	0.036	-0.724	0.898	-1.778
-0.096									
179T	0.263	-0.048	1.150	-1.369	0.893	0.055	0.619	1.150	-1.369
0.223									
180A	-0.085	-0.180	1.188	-1.156	0.911	0.059	0.465	1.188	-1.156
0.172									
181Q	0.832	0.395	1.477	-0.912	1.084	0.077	0.303	1.477	-0.912
0.465									
182Y	0.604	0.395	1.730	-0.676	1.403	0.096	0.523	1.730	-0.676
0.582									
183L	0.768	0.802	1.860	-0.419	1.613	0.676	0.456	1.860	-0.419
0.822									
184S	1.015	1.820	2.188	-0.100	1.987	0.718	1.629	2.188	-0.100
1.322									
185P	0.768	1.591	1.860	-0.188	1.613	0.676	0.456	1.860	-0.188
0.968									
186E	1.154	1.772	2.038	-0.623	1.850	1.281	0.123	2.038	-0.623
1.085									
187Q	2.096	2.052	2.113	-1.146	1.795	1.276	-0.312	2.113	-1.146
1.125									
188A	2.317	1.465	2.234	-1.398	1.959	1.744	-0.481	2.317	-1.398
1.120									
189R	2.595	2.004	2.141	-1.085	1.841	1.745	-0.541	2.595	-1.085
1.243									
190G	1.868	1.191	1.692	-0.360	1.467	1.147	0.074	1.868	-0.360
1.011									
191D	2.121	1.377	1.636	0.484	1.412	1.594	-0.098	2.121	-0.098
1.218									
192S	2.121	1.694	1.636	0.822	1.412	1.594	-0.098	2.121	-0.098
1.312									
193V	2.121	1.377	1.636	0.623	1.412	1.594	-0.098	2.121	-0.098
1.238									
194D	2.172	1.377	1.795	0.201	1.613	1.614	0.061	2.172	0.061
1.262									
195A	2.172	0.431	1.795	-0.045	1.613	1.614	0.061	2.172	-0.045
1.092									
196R	1.527	1.287	1.524	-0.033	1.449	1.596	0.490	1.596	-0.033
1.120									
197S	1.641	0.269	1.898	0.337	1.695	1.613	0.234	1.898	0.234
1.098									
198D	1.420	0.041	1.776	0.461	1.531	1.144	0.402	1.776	0.041
0.968									
199V	0.705	-0.462	1.692	0.297	1.540	1.150	1.847	1.847	-0.462
0.967									
200Y	0.800	-0.462	1.253	0.001	1.020	0.525	1.849	1.849	-0.462
0.712									
201S	0.477	-0.260	0.889	-0.304	0.774	0.523	2.311	2.311	-0.304
0.630									
202L	-0.389	-1.522	0.496	-0.591	0.446	0.036	2.909	2.909	-1.522
0.198									
203G	-0.736	-0.743	0.533	-0.761	0.465	0.039	2.755	2.755	-0.761

233P	1.205	0.333	2.262	-0.088	2.160	1.742	1.127	2.262	-0.088
1.249									
234I	1.350	0.788	1.982	0.091	1.841	1.137	1.288	1.982	0.091
1.211									
235P	0.990	1.203	1.655	0.221	1.476	0.538	0.304	1.655	0.221
0.912									
236P	0.623	1.419	1.814	0.229	1.631	0.673	0.312	1.814	0.229
0.957									
237S	0.623	1.688	1.730	0.290	1.531	1.274	0.360	1.730	0.290
1.071									
238A	1.622	0.628	2.197	0.137	1.914	1.872	0.012	2.197	0.012
1.197									
239R	1.849	1.483	1.945	-0.026	1.595	1.853	-0.208	1.945	-0.208
1.213									
240H	1.135	0.670	1.617	-0.260	1.330	1.839	0.007	1.839	-0.260
0.906									
241E	1.135	1.119	1.617	-0.431	1.330	1.839	0.007	1.839	-0.431
0.945									
242G	1.135	0.339	1.617	-0.741	1.330	1.839	0.007	1.839	-0.741
0.790									
243L	1.502	0.251	1.459	-0.480	1.175	1.704	-0.001	1.704	-0.480
0.801									
244S	0.787	0.455	1.216	-0.415	1.011	1.089	0.166	1.216	-0.415
0.616									
245A	0.926	-0.496	1.160	0.204	0.966	0.979	0.184	1.160	-0.496
0.560									
246D	0.699	-0.593	1.169	0.285	1.011	0.979	-0.827	1.169	-0.827
0.389									
247L	1.046	-1.336	1.132	0.277	0.993	0.975	-0.673	1.132	-1.336
0.345									
248D	0.402	-0.300	0.860	-0.285	0.829	0.956	-0.244	0.956	-0.300
0.317									
249A	-0.313	-0.839	0.776	-0.827	0.838	0.962	1.201	1.201	-0.839
0.257									
250V	-0.585	-1.043	0.954	-1.657	1.157	1.067	1.275	1.275	-1.657
0.167									
251V	0.130	-0.947	1.038	-2.076	1.148	1.062	-0.170	1.148	-2.076
0.026									
252L	-1.084	-0.019	0.683	-2.465	0.838	0.578	0.274	0.838	-2.465
-0.171									
253K	-1.084	0.794	0.683	-2.490	0.838	0.578	0.274	0.838	-2.490
-0.058									
254A	-0.490	0.321	1.253	-2.494	1.485	1.171	-0.250	1.485	-2.494
0.142									
255L	0.187	0.896	1.674	-1.991	1.804	1.211	-0.951	1.804	-1.991
0.404									
256A	0.901	1.710	2.001	-1.218	2.069	1.224	-1.166	2.069	-1.218
0.789									
257K	1.034	2.523	1.879	-0.210	1.795	1.229	-1.258	2.523	-1.258
0.999									
258N	1.344	1.285	2.178	0.863	2.105	1.270	-0.361	2.178	-0.361
1.240									
259P	2.191	1.167	2.692	1.263	2.570	1.889	-0.797	2.692	-0.797
1.568									
260E	1.938	1.299	2.945	1.197	2.807	1.908	0.546	2.945	0.546
1.806									
261N	1.957	0.724	2.823	0.789	2.543	1.356	0.644	2.823	0.644
1.548									
262R	1.843	0.115	2.720	0.187	2.388	1.335	0.798	2.720	0.115

292E	2.513	1.052	2.216	-1.051	1.823	1.743	-0.623	2.513	-1.051
1.096									
293R	1.603	1.333	1.935	-1.061	1.677	1.728	-0.229	1.935	-1.061
0.998									
294T	0.389	1.375	1.580	-0.868	1.367	1.245	0.215	1.580	-0.868
0.758									
295S	0.667	0.884	1.730	-0.509	1.522	1.265	1.385	1.730	-0.509
0.992									
296L	0.585	0.029	1.552	-0.287	1.312	0.685	1.571	1.571	-0.287
0.778									
297L	0.452	0.860	1.122	-0.084	0.838	0.061	0.562	1.122	-0.084
0.544									
298S	0.256	1.674	0.926	0.106	0.683	0.041	-0.489	1.674	-0.489
0.457									
299S	0.206	0.614	0.767	-0.056	0.483	0.021	-0.649	0.767	-0.649
0.198									
300A	1.230	0.614	1.150	-0.074	0.784	0.056	-1.197	1.230	-1.197
0.366									
301A	1.230	1.241	1.150	-0.202	0.784	0.056	-1.197	1.241	-1.197
0.437									
302G	1.230	1.599	1.150	0.221	0.784	0.056	-1.197	1.599	-1.197
0.549									
303N	1.179	1.786	0.991	0.528	0.583	0.036	-1.356	1.786	-1.356
0.535									
304L	1.179	1.668	1.234	0.820	0.856	0.055	-0.126	1.668	-0.126
0.812									
305S	1.312	2.411	1.664	0.424	1.330	0.680	0.883	2.411	0.424
1.244									
306G	1.280	1.914	1.870	0.095	1.531	0.700	0.924	1.914	0.095
1.188									
307P	1.470	1.082	1.842	-0.287	1.540	1.148	1.028	1.842	-0.287
1.118									
308R	2.185	1.082	2.169	-0.007	1.804	1.161	0.813	2.185	-0.007
1.315									
309T	1.192	1.082	1.935	0.185	1.658	1.147	1.088	1.935	0.185
1.184									
310D	0.964	1.082	2.188	0.545	1.977	1.166	1.308	2.188	0.545
1.319									
311P	1.097	1.082	2.374	0.260	2.178	1.771	1.087	2.374	0.260
1.407									
312L	1.211	0.519	2.272	-0.118	2.078	1.189	1.251	2.272	-0.118
1.200									
313P	1.514	1.263	2.346	-0.409	2.242	1.658	1.201	2.346	-0.409
1.402									
314R	0.300	1.443	1.991	-0.499	1.932	1.175	1.645	1.991	-0.499
1.141									
315Q	0.800	1.121	2.019	-0.080	1.977	1.645	1.416	2.019	-0.080
1.271									
316D	2.014	1.169	2.374	0.543	2.287	2.128	0.972	2.374	0.543
1.641									
317L	2.210	1.443	2.328	1.021	2.169	2.129	0.793	2.328	0.793
1.728									
318D	2.577	2.186	2.169	1.590	2.014	1.993	0.785	2.577	0.785
1.902									
319D	2.463	2.461	2.272	1.688	2.114	2.576	0.621	2.576	0.621
2.028									
320T	2.463	2.778	2.272	1.564	2.114	2.576	0.621	2.778	0.621
2.055									
321D	3.310	1.962	2.786	1.094	2.579	3.195	0.185	3.310	0.185

351N	-0.408	0.582	0.786	0.262	0.537	0.038	-0.006	0.786	-0.408
0.256									
352T	0.459	0.463	0.917	0.557	0.510	0.036	-0.327	0.917	-0.327
0.374									
353F	-0.180	0.786	0.776	-0.099	0.492	0.038	1.005	1.005	-0.180
0.403									
354G	0.655	1.499	1.113	-0.767	0.665	0.056	0.724	1.499	-0.767
0.564									
355G	0.477	0.776	1.244	-1.621	0.829	0.640	0.836	1.244	-1.621
0.455									
356I	0.781	0.640	1.318	-1.558	0.993	1.109	0.786	1.318	-1.558
0.581									
357T	1.129	0.868	1.262	-1.341	1.030	1.107	1.124	1.262	-1.341
0.740									
358R	1.148	0.736	1.599	-0.775	1.449	1.149	1.287	1.599	-0.775
0.942									
359D	0.553	0.461	1.487	-0.576	1.485	1.151	1.875	1.875	-0.576
0.920									
360V	1.192	-0.174	1.870	-0.405	1.777	1.168	1.773	1.870	-0.405
1.029									
361Q	1.495	0.736	1.945	-0.389	1.941	1.637	1.724	1.945	-0.389
1.298									
362V	0.996	0.872	1.393	-0.128	1.458	1.014	2.313	2.313	-0.128
1.131									
363P	0.629	1.459	1.552	-0.060	1.613	1.150	2.321	2.321	-0.060
1.238									
364D	1.224	1.956	1.664	-0.138	1.576	1.148	1.733	1.956	-0.138
1.309									
365V	1.224	2.273	1.664	-0.636	1.576	1.148	1.733	2.273	-0.636
1.283									
366R	1.868	2.369	1.935	-0.906	1.741	1.166	1.304	2.369	-0.906
1.354									
367G	2.147	2.094	1.842	-0.634	1.622	1.168	1.244	2.147	-0.634
1.355									
368Q	1.647	1.467	1.571	-0.072	1.303	0.679	0.243	1.647	-0.072
0.977									
369S	2.513	0.652	1.963	0.754	1.631	1.166	-0.355	2.513	-0.355
1.189									
370S	2.381	-0.204	1.533	0.944	1.157	0.541	-1.364	2.381	-1.364
0.713									
371A	1.514	-0.568	1.403	0.590	1.185	0.543	-1.043	1.514	-1.043
0.518									
372D	1.268	-0.773	1.075	-0.193	0.811	0.501	-2.216	1.268	-2.216
0.068									
373A	1.186	-0.821	1.122	-0.971	0.811	0.501	-2.335	1.186	-2.335
-0.073									
374I	0.193	-0.212	0.889	-1.644	0.665	0.486	-2.060	0.889	-2.060
-0.240									
375A	0.440	0.926	1.216	-1.660	1.039	0.528	-0.887	1.216	-1.660
0.229									
376T	0.250	1.553	1.244	-1.174	1.030	0.080	-0.991	1.553	-1.174
0.285									
377L	0.383	0.888	1.674	-0.458	1.504	0.705	0.019	1.674	-0.458
0.673									
378Q	1.249	1.924	1.804	0.236	1.476	0.703	-0.303	1.924	-0.303
1.013									
379N	0.534	1.109	1.739	0.423	1.431	0.707	0.958	1.739	0.423
0.986									
380R	0.566	1.313	1.991	-0.005	1.914	1.282	0.983	1.991	-0.005

410V 0.846	1.774	1.227	1.309	0.615	0.920	0.072	0.008	1.774	0.008
411S 0.838	1.963	0.998	1.281	0.063	0.929	0.520	0.112	1.963	0.063
412A 0.902	2.128	0.634	1.412	-0.142	1.139	1.100	0.045	2.128	-0.142
413G 0.670	1.211	0.538	1.122	-0.432	0.966	1.082	0.207	1.211	-0.432
414D 0.720	1.774	0.519	1.440	-0.582	1.130	1.100	-0.341	1.774	-0.582
415E 0.463	1.129	-0.116	1.169	-1.078	0.966	1.082	0.088	1.169	-1.078
416I 0.784	1.438	0.165	1.468	-0.968	1.276	1.122	0.985	1.468	-0.968
417T 0.934	0.844	0.980	1.356	-0.650	1.312	1.124	1.573	1.573	-0.650
418V 0.952	0.623	1.117	1.234	0.142	1.148	0.655	1.742	1.742	0.142
419N 0.963	0.459	1.571	1.103	0.784	0.938	0.075	1.809	1.809	0.075
420V 1.089	1.325	1.537	1.234	1.051	0.911	0.074	1.488	1.537	0.074
421S 1.152	1.129	2.124	1.281	0.758	1.030	0.073	1.667	2.124	0.073
422T 1.309	1.856	2.082	1.730	0.372	1.403	0.671	1.052	2.082	0.372
423G 1.283	1.793	2.166	1.758	-0.200	1.467	0.672	1.328	2.166	-0.200
424P 1.319	2.292	1.215	2.309	-0.570	1.950	1.295	0.739	2.309	-0.570
425E 1.374	2.374	1.215	2.487	-1.047	2.160	1.875	0.553	2.487	-1.047
426Q 1.135	1.540	1.179	2.150	-1.604	1.987	1.857	0.834	2.150	-1.604
427R 1.095	1.312	0.592	2.403	-1.873	2.306	1.876	1.053	2.403	-1.873
428E 1.225	1.812	0.634	2.431	-1.824	2.351	2.346	0.824	2.431	-1.824
429I 1.072	1.084	0.550	1.982	-1.274	1.977	1.748	1.439	1.982	-1.274
430P 1.158	1.116	0.670	1.804	-0.402	1.759	1.725	1.436	1.804	-0.402
431D 1.131	1.179	0.802	1.571	0.324	1.440	1.121	1.477	1.571	0.324
432V 0.744	0.104	-0.144	1.160	0.535	1.084	0.527	1.939	1.939	-0.144
433S 0.893	0.939	-0.048	1.496	0.401	1.257	0.545	1.658	1.658	-0.048
434T 0.766	0.686	-0.328	1.505	-0.035	1.221	0.545	1.770	1.770	-0.328
435L 0.252	0.187	-0.819	1.234	-0.566	0.902	0.056	0.769	1.234	-0.819
436T 0.437	0.914	-0.711	1.683	-0.912	1.276	0.654	0.154	1.683	-0.912
437Y 0.169	0.636	-0.370	1.533	-1.354	1.121	0.634	-1.016	1.533	-1.354
438A 0.218	0.073	0.868	1.216	-1.734	0.957	0.616	-0.468	1.216	-1.734
439E	1.015	0.664	1.748	-2.014	1.586	1.205	-0.838	1.748	-2.014

469G	0.806	1.171	1.561	-0.708	1.504	0.647	1.305	1.561	-0.708
0.898									
470T	0.579	1.153	1.356	0.682	1.139	0.072	1.460	1.460	0.072
0.920									
471N	0.945	1.153	1.477	1.447	1.148	0.070	-0.139	1.477	-0.139
0.872									
472P	1.894	1.034	1.917	1.870	1.476	0.109	-0.573	1.917	-0.573
1.104									
473P	1.913	1.531	2.253	1.604	1.895	0.151	-0.410	2.253	-0.410
1.277									
474A	1.913	1.173	2.253	1.535	1.895	0.151	-0.410	2.253	-0.410
1.216									
475N	1.881	0.848	2.103	1.452	1.741	0.131	-0.138	2.103	-0.138
1.145									
476Q	1.881	0.730	1.860	1.375	1.467	0.112	-1.368	1.881	-1.368
0.865									
477T	1.242	0.848	1.477	0.788	1.175	0.095	-1.266	1.477	-1.266
0.623									
478S	1.438	0.261	1.674	0.146	1.330	0.115	-0.215	1.674	-0.215
0.678									
479A	1.438	-0.691	1.674	-0.347	1.330	0.115	-0.215	1.674	-0.691
0.472									
480I	0.825	-1.015	1.225	-0.324	0.948	0.074	0.210	1.225	-1.015
0.278									
481T	0.263	-1.015	0.907	0.105	0.784	0.055	0.758	0.907	-1.015
0.265									
482N	-0.654	-1.831	0.618	0.086	0.610	0.037	0.920	0.920	-1.831
-0.031									
483V	-1.293	-2.536	0.477	-0.508	0.592	0.039	2.252	2.252	-2.536
-0.139									
484V	-1.293	-1.813	0.477	-1.674	0.592	0.039	2.252	2.252	-1.813
-0.203									
485I	-1.856	-0.861	0.160	-2.676	0.428	0.021	2.800	2.800	-2.676
-0.283									
486I	-1.938	0.091	-0.149	-3.255	0.073	-0.020	2.913	2.913	-3.255
-0.326									
487I	-1.293	0.774	0.122	-2.805	0.237	-0.002	2.484	2.484	-2.805
-0.069									
488V	-0.698	1.099	0.234	-2.097	0.200	-0.003	1.896	1.896	-2.097
0.090									
489G	-0.060	1.686	0.618	-1.092	0.492	0.014	1.794	1.794	-1.092
0.493									
490S	0.579	1.890	0.758	-0.482	0.510	0.012	0.462	1.890	-0.482
0.533									
491G	1.413	1.573	1.094	-0.159	0.683	0.030	0.182	1.573	-0.159
0.688									
492P	2.008	0.622	1.664	-0.401	1.330	0.623	-0.342	2.008	-0.401
0.786									
493A	2.279	0.622	1.945	-0.430	1.695	1.112	-0.351	2.279	-0.430
0.982									
494T	1.363	1.161	1.655	-0.716	1.522	1.094	-0.190	1.655	-0.716
0.841									
495K	1.135	0.574	1.907	-0.505	1.841	1.113	0.030	1.907	-0.505
0.871									
496D	1.634	-0.258	1.935	-0.321	1.886	1.583	-0.199	1.935	-0.321
0.895									
497I	1.268	-0.170	1.814	-0.018	1.877	1.585	1.400	1.877	-0.170
1.108									
498P	1.072	0.646	1.617	0.045	1.722	1.565	0.349	1.722	0.045

528P	1.394	1.459	1.860	0.507	1.686	1.143	0.350	1.860	0.350
1.200									
529R	1.989	1.004	1.973	-0.095	1.649	1.141	-0.238	1.989	-0.238
1.061									
530P	1.849	0.682	2.029	-0.847	1.695	1.252	-0.255	2.029	-0.847
0.915									
531A	1.205	0.950	1.758	-1.336	1.531	1.234	0.173	1.758	-1.336
0.788									
532G	1.401	1.441	1.711	-1.626	1.412	1.235	-0.006	1.711	-1.626
0.795									
533E	1.495	1.423	1.272	-1.640	0.893	0.610	-0.005	1.495	-1.640
0.578									
534V	1.691	1.207	1.225	-1.479	0.774	0.611	-0.184	1.691	-1.479
0.549									
535T	2.001	1.662	1.524	-0.731	1.084	0.651	0.713	2.001	-0.731
0.986									
536G	1.774	1.171	1.776	0.158	1.403	0.670	0.933	1.776	0.158
1.126									
537T	1.413	1.171	1.692	1.173	1.312	0.090	1.179	1.692	0.090
1.147									
538N	1.780	1.171	1.814	1.671	1.321	0.088	-0.420	1.814	-0.420
1.061									
539P	1.812	1.052	1.608	1.430	1.121	0.068	-0.460	1.812	-0.460
0.947									
540P	1.780	0.598	1.814	0.655	1.321	0.088	-0.420	1.814	-0.420
0.834									
541A	1.780	0.598	1.814	-0.094	1.321	0.088	-0.420	1.814	-0.420
0.727									
542G	1.103	0.501	1.393	-0.679	1.002	0.049	0.282	1.393	-0.679
0.522									
543T	1.103	0.413	1.393	-0.625	1.002	0.049	0.282	1.393	-0.625
0.517									
544T	0.737	0.778	1.029	-0.648	0.720	0.032	0.651	1.029	-0.648
0.471									
545V	1.236	0.191	1.300	-0.394	1.039	0.521	1.652	1.652	-0.394
0.792									
546P	1.287	-0.038	1.459	-0.064	1.239	0.541	1.812	1.812	-0.064
0.891									
547V	0.724	0.179	1.141	0.342	1.075	0.522	2.360	2.360	0.179
0.906									
548D	-0.111	0.071	0.804	0.442	0.902	0.504	2.640	2.640	-0.111
0.751									
549S	0.617	0.023	1.253	0.085	1.276	1.102	2.025	2.025	0.023
0.912									
550V	-0.098	-0.929	0.926	-0.925	1.011	1.089	2.240	2.240	-0.929
0.473									
551I	0.515	0.023	1.375	-1.705	1.394	1.130	1.815	1.815	-1.705
0.649									
552E	-0.351	1.179	0.982	-2.186	1.066	0.642	2.413	2.413	-2.186
0.535									
553L	-0.351	1.231	0.982	-1.867	1.066	0.642	2.413	2.413	-1.867
0.588									
554Q	0.244	2.044	1.552	-1.368	1.713	1.235	1.889	2.044	-1.368
1.044									
555V	1.110	2.044	1.683	-0.871	1.686	1.233	1.568	2.044	-0.871
1.208									
556S	1.059	1.966	1.655	-0.387	1.631	0.675	1.481	1.966	-0.387
1.154									
557K	2.020	1.014	2.066	0.059	1.996	0.712	1.209	2.066	0.059

587L	1.116	1.058	1.449	-1.299	1.276	1.096	0.659	1.449	-1.299
0.765									
588D	0.920	1.167	1.253	-1.150	1.121	1.076	-0.392	1.253	-1.150
0.571									
589K	1.192	1.167	1.533	-0.887	1.485	1.565	-0.401	1.565	-0.887
0.808									
590G	1.224	0.335	1.421	-0.858	1.431	1.549	0.243	1.549	-0.858
0.763									
591A	2.437	0.335	1.776	-0.379	1.741	2.033	-0.201	2.437	-0.379
1.106									
592D	1.938	0.962	1.505	-0.152	1.422	1.544	-1.202	1.938	-1.202
0.859									
593V	1.938	1.279	1.047	0.175	0.738	0.949	-1.267	1.938	-1.267
0.694									
594D	1.938	1.866	1.047	-0.089	0.738	0.949	-1.267	1.938	-1.267
0.740									
595A	2.216	1.417	1.197	-0.278	0.893	0.969	-0.097	2.216	-0.278
0.902									
596G	1.963	2.026	1.253	-0.564	0.948	0.523	0.075	2.026	-0.564
0.889									
597G	2.330	2.213	1.533	0.052	1.130	1.141	-0.246	2.330	-0.246
1.165									
598S	2.140	1.489	1.561	0.991	1.121	0.693	-0.350	2.140	-0.350
1.092									
599Q	2.273	0.538	1.991	2.002	1.595	1.317	0.659	2.273	0.538
1.482									
600H	1.679	-0.360	1.879	2.389	1.631	1.319	1.247	2.389	-0.360
1.398									
601N	1.084	0.041	1.767	1.814	1.668	1.321	1.836	1.836	0.041
1.361									
602R	0.553	0.041	1.870	0.491	1.750	1.320	2.008	2.008	0.041
1.147									
603V	0.553	-0.414	1.870	-0.669	1.750	1.320	2.008	2.008	-0.669
0.917									
604V	0.863	0.041	2.010	-0.874	1.886	0.741	1.628	2.010	-0.874
0.899									
605Y	0.553	0.137	1.954	-0.323	1.850	0.719	1.961	1.961	-0.323
0.979									
606Q	0.421	1.171	1.767	0.908	1.649	0.113	2.182	2.182	0.113
1.173									
607N	0.787	1.171	1.889	1.530	1.658	0.112	0.583	1.889	0.112
1.104									
608P	1.382	1.189	2.001	1.418	1.622	0.110	-0.006	2.001	-0.006
1.102									
609P	1.830	0.734	1.945	0.635	1.540	0.111	-0.297	1.945	-0.297
0.928									
610A	1.812	0.984	1.608	-0.196	1.121	0.068	-0.460	1.812	-0.460
0.705									
611G	1.135	1.798	1.188	-0.903	0.802	0.029	0.242	1.798	-0.903
0.613									
612T	1.445	1.710	1.244	-0.718	0.838	0.051	-0.091	1.710	-0.718
0.640									
613G	1.578	1.846	1.431	-0.491	1.039	0.657	-0.312	1.846	-0.491
0.821									
614V	2.077	0.894	1.702	0.244	1.358	1.146	0.689	2.077	0.244
1.158									
615N	2.077	0.666	1.702	0.615	1.358	1.146	0.689	2.077	0.615
1.179									
616R	1.242	0.548	1.365	0.575	1.185	1.127	0.970	1.365	0.548

1.002									
617D	0.376	-0.470	1.234	-0.146	1.212	1.129	1.291	1.291	-0.470
0.661									
618G	0.939	-0.196	1.552	-0.923	1.376	1.148	0.743	1.552	-0.923
0.663									
619I	-0.085	-0.997	1.169	-1.913	1.075	1.112	1.291	1.291	-1.913
0.236									
620I	-0.085	-0.046	1.169	-2.146	1.075	1.112	1.291	1.291	-2.146
0.339									
621T	-1.299	0.770	0.832	-2.181	0.711	0.628	1.551	1.551	-2.181
0.145									
622L	-1.299	0.534	0.832	-1.946	0.711	0.628	1.551	1.551	-1.946
0.144									
623R	-0.414	0.992	1.300	-1.625	1.103	0.668	1.393	1.393	-1.625
0.488									
624F	0.092	0.433	0.982	-1.269	1.440	0.727	0.061	1.440	-1.269
0.352									
625G	-0.237	0.862	0.328	-0.956	1.604	0.767	-0.990	1.604	-0.990
0.197									
626Q	0.345	0.489	-0.046	-0.647	1.914	0.821	-2.435	1.914	-2.435
0.063									

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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 MTTPSHLSDRYELGEILGFGGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYLRFRREAQNAAALNH DEITVNVSTGPEQREIPDVSTLT ^{YAEAVK} KLTAAGFGRF <u>KQANS</u> PSTPELVGKVIGTNPPANQTSAITNVVI
Hydrophilicity	¹ MTTPSHLSDRYELGEILGFGGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYLRFR <u>REAQNAA</u> ALNH DEITVNV <u>STGPEQRE</u> IPDVSTLT ^{YAEAVK} KLTAAGFGRF <u>KQANS</u> PSTPELVGKVIGTNPPANQTSAITNVVI
Flexibility	¹ MTTPSHLSDRYELGEILGFGGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYLRFR <u>REAQNAA</u> ALNH DEITVNVSTGPEQREIPDVSTLT ^{YAEAVK} KLTAAGFGRF <u>KQANS</u> PSTPELVGKVIGTNPPANQTSAITNVVI
Accessibility	¹ MTT <u>PSHLSDRYELGEILGFGGMSEVHLARDLRLHRD</u> VAVKVL <u>RADLARDPSFYLRFRREAQNAA</u> ALNH DEITVNV <u>STGPEQREIPDV</u> STLT ^{YAEAVK} KLTAAGFGRF <u>KQANS</u> PSTPELVGKVIGTNPPANQTSAITNVVI
Turns	¹ MTTPSHLSDRYELGEILGFGGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYLRFRREAQNAAALNH

	DEITVNVSTGPEQREIPDVSTLTYAEAVKKLTAAGFGRFKQANSPSTPELVGKVIGTNPPANQTSAITNVVI
Exposed Surface	¹ MTTPSHLSDRYELGEILGFGGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYL RFRREAQ NAAALNH DEITVNVSTG PEQREIPD VSTLTYAEAVKKLTAAGFGRFKQANSPSTPELVGKVIGTNPPANQTSAITNVVI
Polarity	¹ MTTPS HLSDRYELGEI LGFGGM SEVHLARDLRLHRDVAVKVL RADLARDPSFYL RFRREAQ NAAALNH DEITVNVST TGPEQREIPD VSTLTYAEAVKKLTAAGFGRFKQANSPSTPELVGKVIGTNPPANQTSAITNVVI
Antigenic Propensity	¹ MTTPSHLSDRYELGEILGFGGMSEVHLARDLRLHRDVAVKVLRADLARDPSFYL RFRREAQ NAAALNH DEITVNVSTGPEQRE IPDVSTL TYAEAVKKLTAAGFGRFKQANSPST PELVGKVIG TNPPANQTS AITNVVI

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