

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

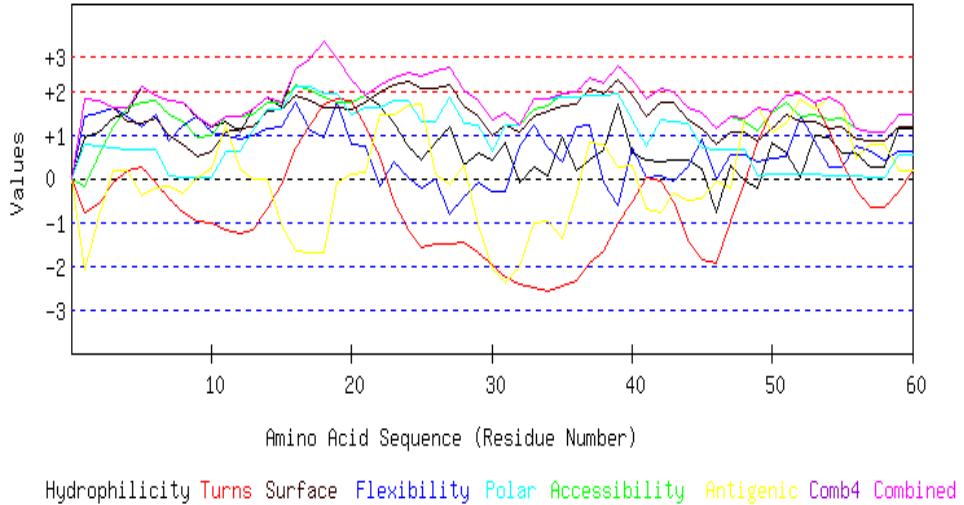
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LPAEQYAVQTGTHPRTRTEANVNFRQLGRLGFHDSRRSFSTTDVFYRWTQWIFLQIYNA
WFDTTANKARPISELVAEFESGARCLDGGRDWAKLTAGERADVIDEYRLVYRADSLVNWCPL
GTVLANEEVTADGRSDRGNFVFRKRLRQWMMRITAYADRLLDDLDVLDWPEQVKTMQRN
WIGRSTGAVALFSARAASDDGFEVDIEVFTTRPDTLFGATYLVLAPEHDLVDELVAASWPAGVN
PLWTYGGGTPGEAIAAYRRAIAAKSDLERQESREKTGVFLGSYAINPANGEPVPIFIADYVLAG
YGTGAIMAVPGHDQRDWDFARAFGLPIVEVIAGGNISESAYTGDGILVNSDYLNGMSVPAAKR
AIVDRLESAGRGRARIEFKLRDWLFARQRYWGEPFPIVYDSDGRPHALDEAALPVELPDVPDY
SPVLFDPPDDADSEPPPLAKATEWVHVLDLGLKPYSRDTNVMPQWAGSSWYELRYTDPH
NSERFCAKEANEAYWMGPRPAEHGPDDPGGVLDLYVGGAEHAVLHLLYSRFWHKVLYDLGHVS
SREPYRRLVNQGYIQAYAYTDARGSYVPAEQVIERGDRFVYPGPGEVEVFQEFGKIGKSLKNS
VSPDEICDAYGADTLRVYEMSMGPLEARPWATKDVVGAYRFLQRVWRLVVDEHTGETRVAD
GVELIDITLRALHRTIVGVSEDFAALRNNTATAKLIETNHLTKKHRDAVPRAAVEPLVQMLAP
LAPHIAEELWLRLGNTTSLAHGPFPKADAAYLVDETVEYPVQVNGKVRGRVVVAADTDEETL
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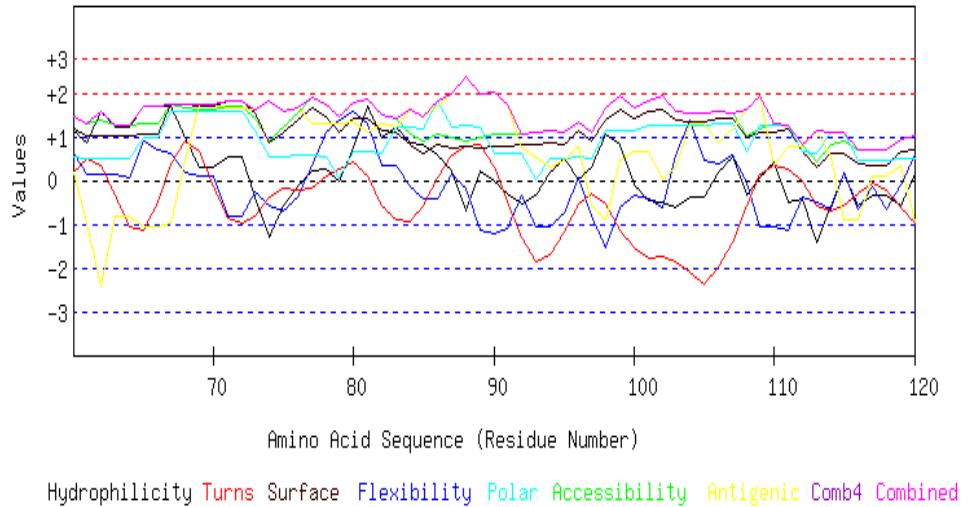
Length=969

GRAPHICAL RESULT

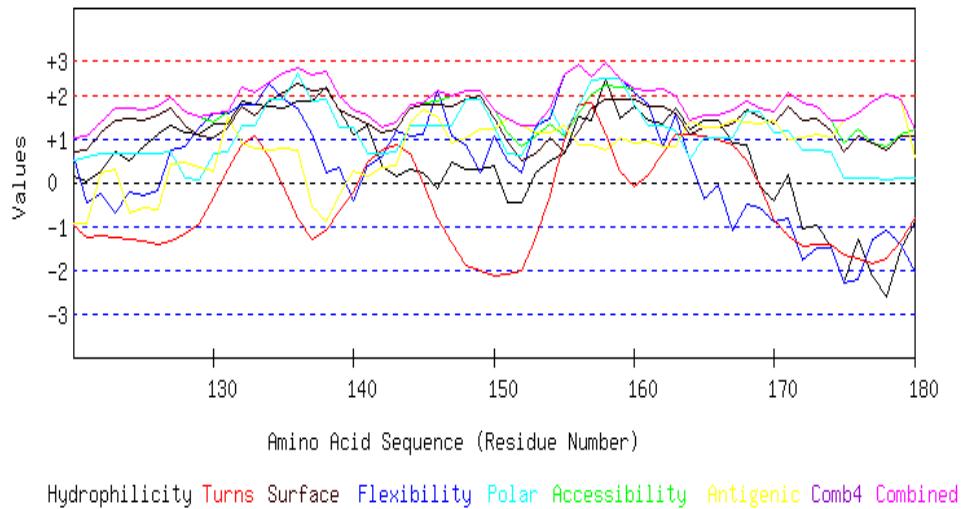
GRAPHICAL RESULT :: SEQ 1 to 60



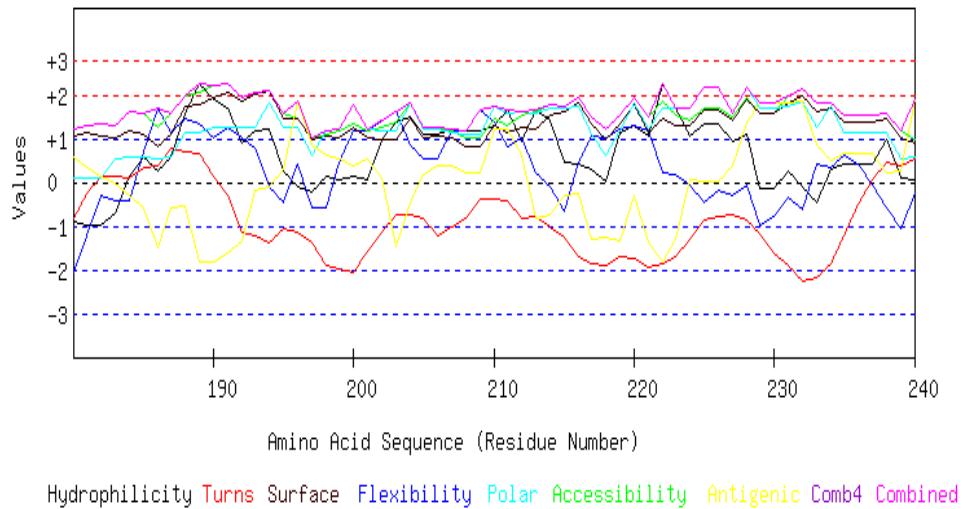
GRAPHICAL RESULT :: SEQ 61 to 120



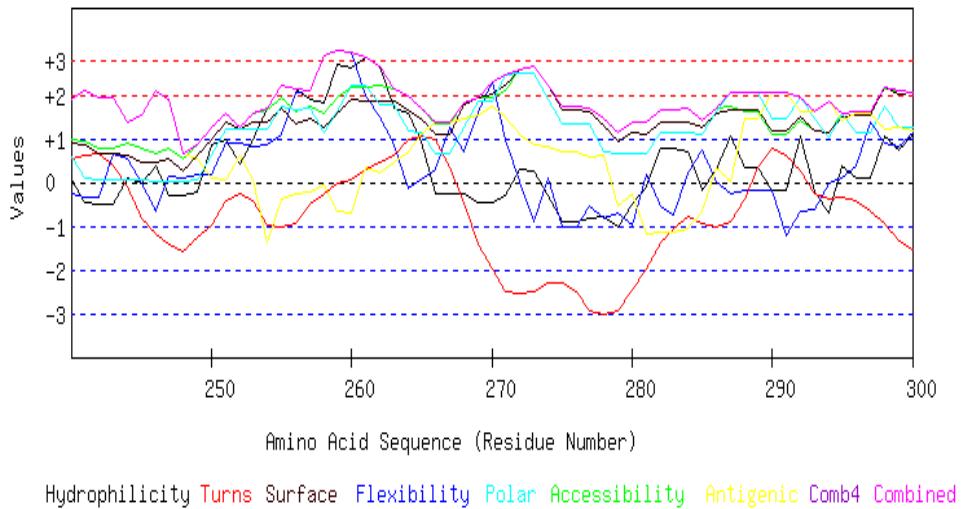
GRAPHICAL RESULT :: SEQ 121 to 180



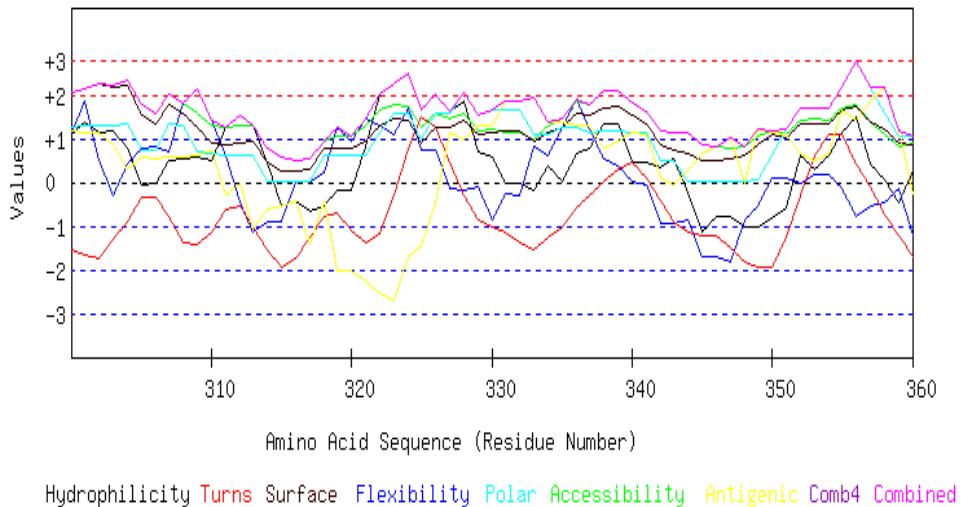
GRAPHICAL RESULT :: SEQ 181 to 240



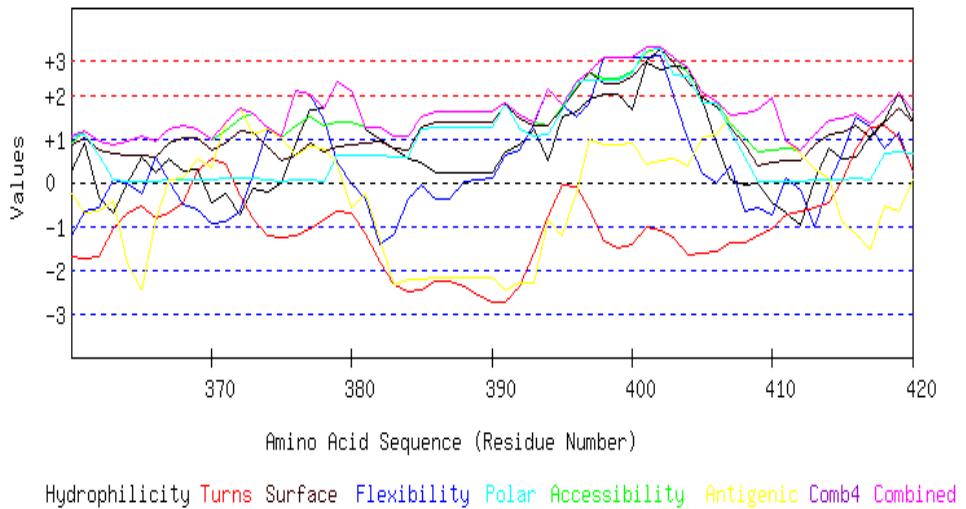
GRAPHICAL RESULT :: SEQ 241 to 300



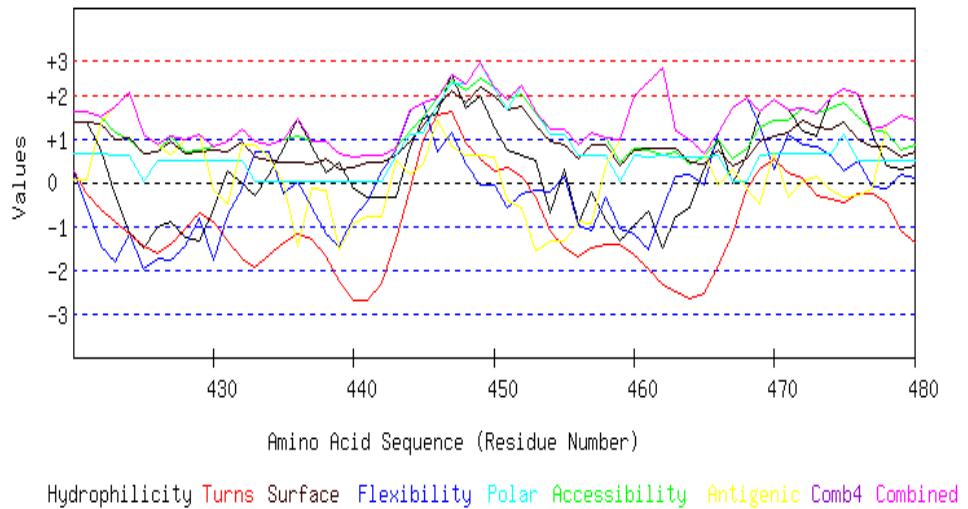
GRAPHICAL RESULT :: SEQ 301 to 360



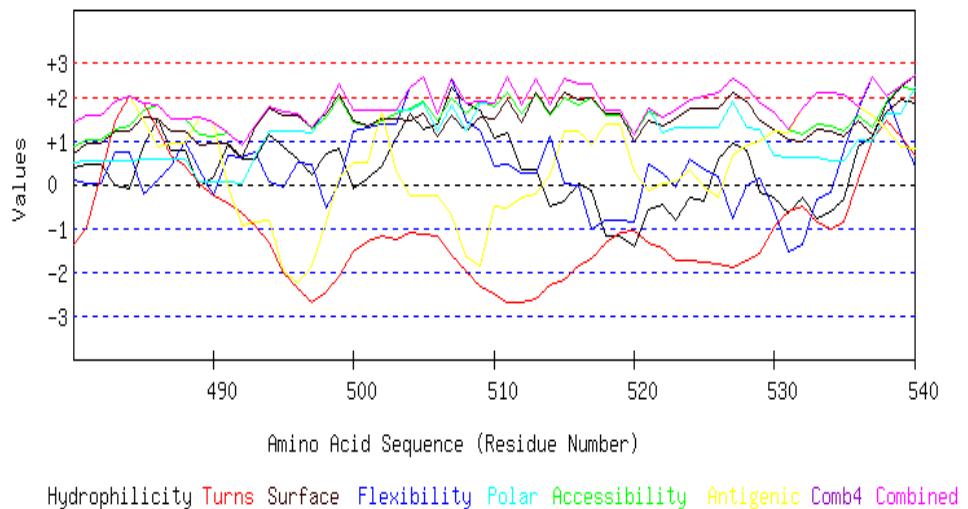
GRAPHICAL RESULT :: SEQ 361 to 420



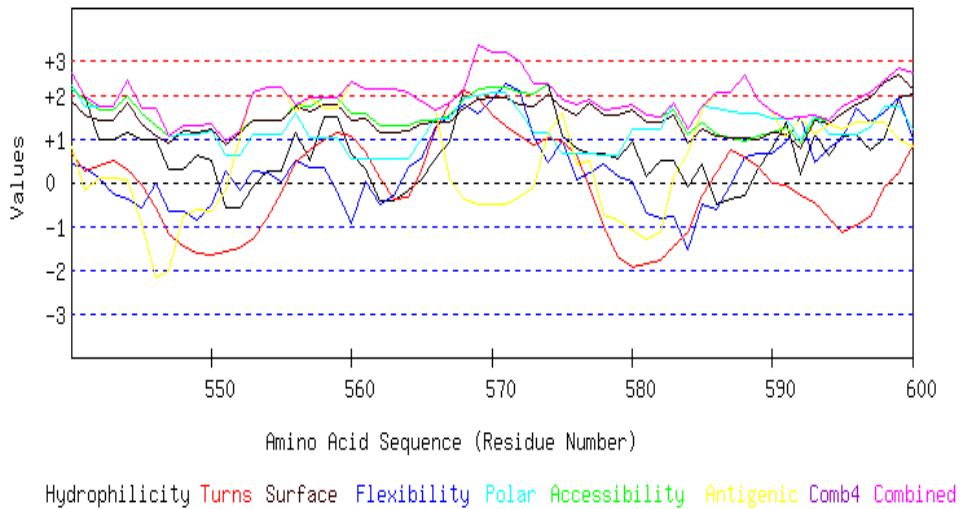
GRAPHICAL RESULT :: SEQ 421 to 480



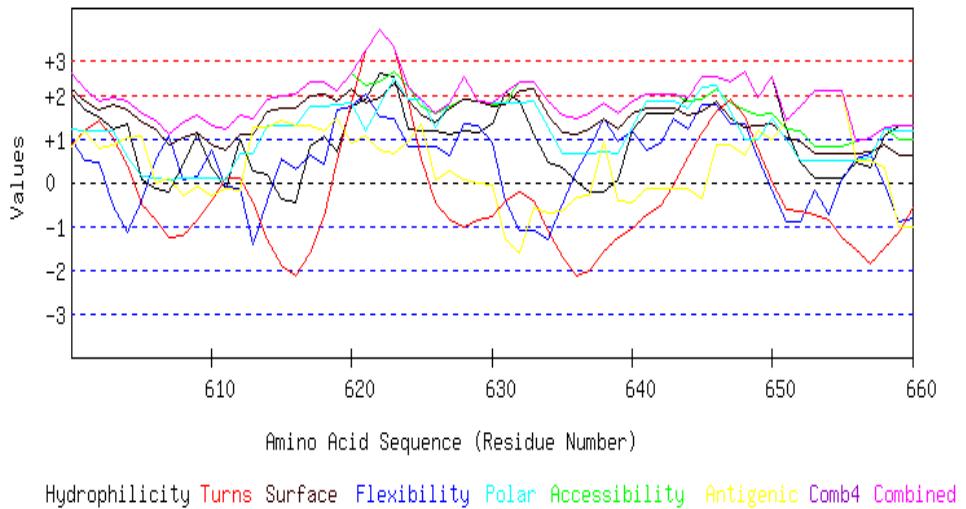
GRAPHICAL RESULT :: SEQ 481 to 540



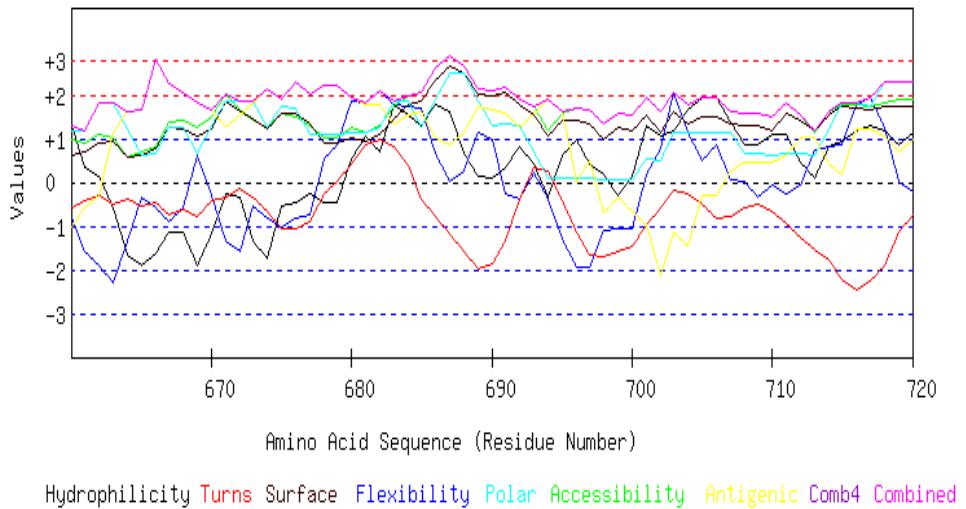
GRAPHICAL RESULT :: SEQ 541 to 600



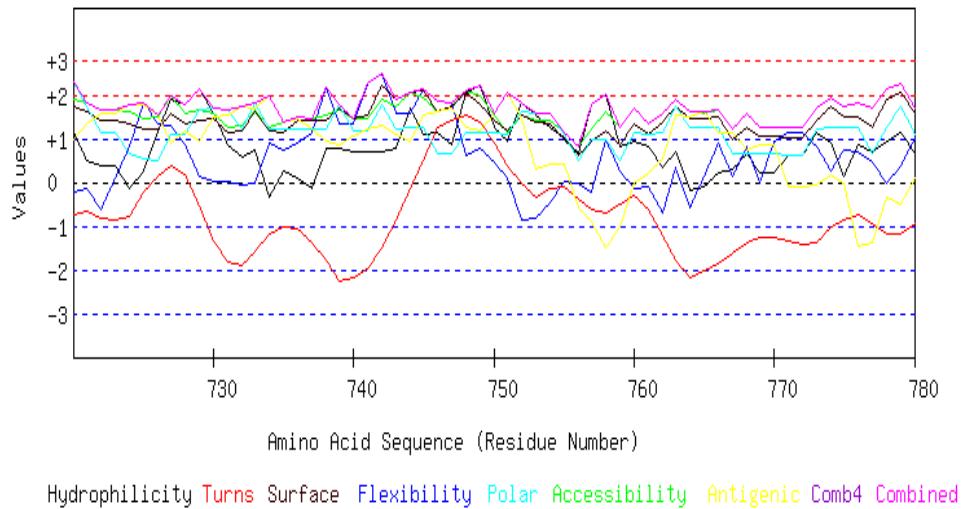
GRAPHICAL RESULT :: SEQ 601 to 660



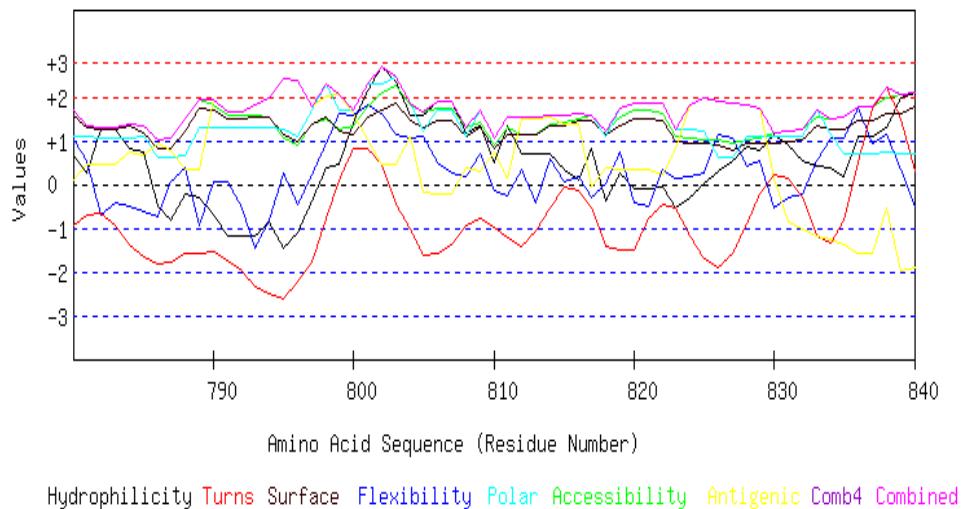
GRAPHICAL RESULT :: SEQ 661 to 720



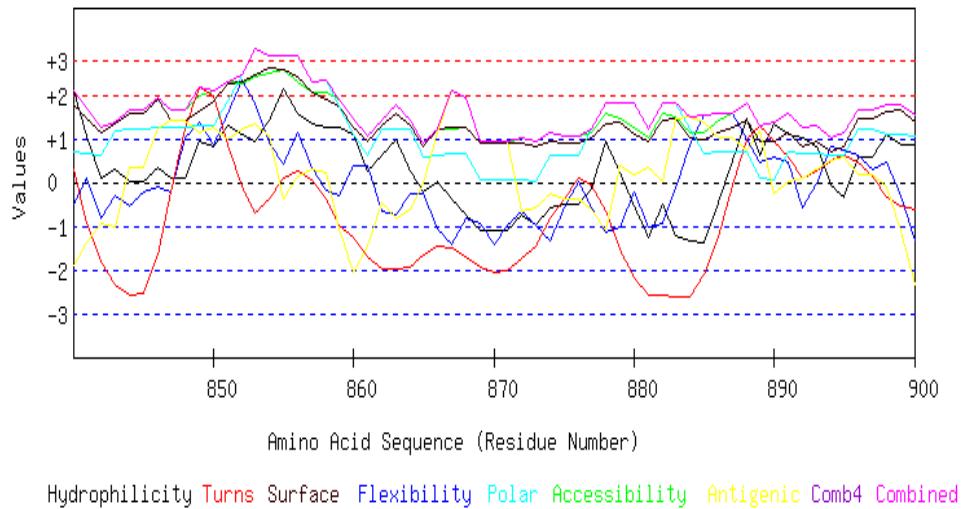
GRAPHICAL RESULT :: SEQ 721 to 780



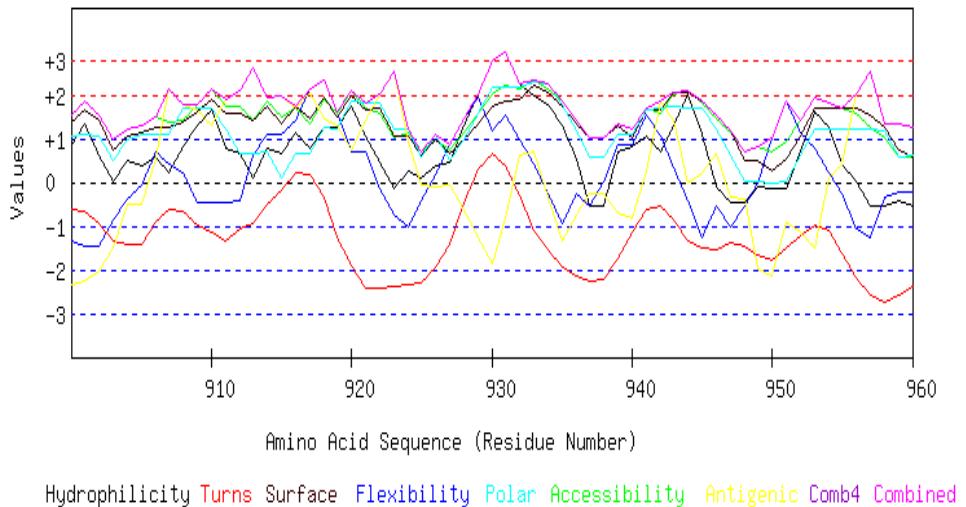
GRAPHICAL RESULT :: SEQ 781 to 840



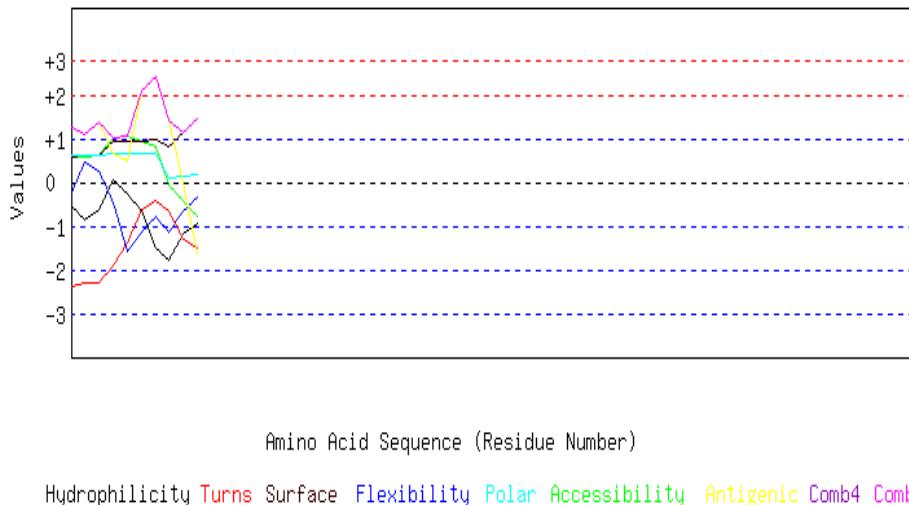
GRAPHICAL RESULT :: SEQ 841 to 900



GRAPHICAL RESULT :: SEQ 901 to 960



GRAPHICAL RESULT :: SEQ 961 to 1020



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

VTESPTAGPGGVPRADDADSDVPRYRYTAELAARLERTWQENWARLGTNFVNPNPGSLAP
PDGAAPDDKLFQDMFPYPSGEGLHVGHPLGYIATDVYARYFRMVGRNVHLALGFDAFG
LPAEQYAVQTGTHPRTTRTEANVNFRQLGRLGFGHSRRSFSTTDVDFYRWTQWIFLQI
YNAWFDTTANKARPISELVAEFESEGARCLDGGRDWAKLTAGERADVIDEYRLVYRADSLV
NWCPGLTVLANEEVTADGRSDRGNFVFRKRLRQWMMRITAYADRLLDDLDVLDWPEQV
KTMQRNWIGRSTGAVALFSARAASDDGFEVDIEVFTTRPDTLFGATYLVLAPEHDLVDEL
VAASWPAGVNPLWTYGGGTGPGEAIAAYRAIAAKSDLERQESREKTGVFLGSYAINPANG
EPVPIFIADYVIAGYGTGAIMAVPGHDQRDWDFAFAGLPIVEVIAGGNISESAYTGDI
LVNSDYLNGMSVPAAKRAIVDRLESAGRGRARIEFKLRLDWLFARQRYWGEPFPIVYDSG
RPHALDEAALPVELPDVDPDYSPVLFDPDDADSEPSPLAKATEWHVHDLDLGDKPYSR
DTNVMPQWAGSSWYELRYTDPHNSERFCAKENEAYWMGPRPAEHGPDDPGGVIDLVGGAE
HAVLHLLYSRFWHKVLYDLGHVSSREPYRRLVNQGYIQAYAYTDARGSYVPAEQVIERGD

RFVYPGPGEVEVFQEFGKIGKSLKNSVSPDEICDAYGADTLRVYEMSMGPLASRPWAT
 KDVVGAYRFLQRVWRLLVVDEHTGETRVADGVELIDTLRALHRTIVGVSEDFAALRNNTA
 TAKLIEYTNHLTKKHRRDAVPRAAVEPLVQMLAPLAPHIAEELWLRLGNNTSLAHGPFPKA
 DAAYLVDETVEYPVQVNGKVRGRVVVAADTDEETLKAAVLTDEKVQAFLAGATPRKVIVV
 AGRLVNLVI

Length=969

A.A.

**Parameter
Combined**

MIN	AVG	Hydro	Flexi	Access	Turns	Surface	Polar	AntiPro	MAX
1 V	0.920	1.401	-0.177	-0.764	1.823	0.791	-2.086	1.823	
-2.086	0.273								
2 T	1.053	1.497	0.524	-0.586	1.777	0.750	-0.855	1.777	
-0.855	0.594								
3 E	1.382	1.633	1.178	-0.116	1.613	0.710	0.195	1.633	
-0.116	0.942								
4 S	1.514	1.417	1.636	0.190	1.294	0.650	0.195	1.636	
0.190	0.985								
5 P	2.109	1.189	1.748	0.250	1.257	0.648	-0.393	2.109	
-0.393	0.973								
6 T	1.913	1.457	1.795	-0.101	1.376	0.647	-0.214	1.913	
-0.214	0.982								
7 A	1.780	0.870	1.459	-0.475	0.966	0.048	-0.187	1.780	
-0.475	0.637								
8 G	1.729	1.229	1.300	-0.795	0.765	0.028	-0.347	1.729	
-0.795	0.558								
9 P	1.363	1.415	0.935	-0.961	0.483	0.010	0.022	1.415	
-0.961	0.467								
10 G	1.167	1.056	0.982	-1.008	0.601	0.009	0.201	1.167	
-1.008	0.430								
11 G	1.299	0.968	1.412	-1.166	1.075	0.634	1.210	1.412	
-1.166	0.776								
12 V	1.072	0.880	1.421	-1.248	1.121	0.634	0.200	1.421	
-1.248	0.583								
13 P	1.571	0.976	1.449	-1.170	1.166	1.104	-0.029	1.571	
-1.170	0.724								
14 R	1.843	1.157	1.730	-0.697	1.531	1.593	-0.038	1.843	
-0.697	1.017								
15 A	1.616	1.199	1.739	-0.132	1.576	1.593	-1.049	1.739	
-1.049	0.935								
16 D	2.482	1.738	2.132	0.699	1.905	2.080	-1.646	2.482	
-1.646	1.341								
17 D	2.760	1.103	2.038	1.202	1.786	2.081	-1.706	2.760	
-1.706	1.323								
18 A	3.127	0.922	1.879	1.686	1.631	1.946	-1.714	3.127	
-1.714	1.354								
19 D	2.760	1.736	1.758	1.803	1.622	1.947	-0.116	2.760	
-0.116	1.644								
20 S	2.260	0.790	1.730	1.772	1.576	1.477	0.113	2.260	
0.113	1.388								
21 D	1.894	0.748	1.889	1.170	1.731	1.613	0.121	1.894	
0.121	1.309								
22 V	1.641	-0.198	2.141	0.493	1.968	1.632	1.464	2.141	
-0.198	1.306								

23 P	1.274	0.389	2.300	-0.543	2.123	1.768	1.472	2.300
-0.543	1.255							
24 R	0.743	0.031	2.403	-1.188	2.205	1.767	1.644	2.403
-1.188	1.086							
25 Y	0.440	-0.208	2.328	-1.570	2.041	1.298	1.694	2.328
-1.570	0.860							
26 R	0.806	-0.005	2.449	-1.515	2.050	1.297	0.095	2.449
-1.515	0.740							
27 Y	1.167	-0.819	2.533	-1.481	2.142	1.877	-0.151	2.533
-1.481	0.753							
28 T	0.319	-0.412	2.019	-1.446	1.677	1.258	0.284	2.019
-1.446	0.528							
29 A	0.572	-0.090	1.767	-1.693	1.440	1.239	-1.058	1.767
-1.693	0.311							
30 E	0.440	-0.294	1.337	-1.989	0.966	0.614	-2.067	1.337
-2.067	-0.142							
31 L	0.825	-0.294	1.515	-2.261	1.203	1.219	-2.400	1.515
-2.400	-0.028							
32 A	-0.085	0.724	1.234	-2.433	1.057	1.205	-2.006	1.234
-2.433	-0.043							
33 A	0.275	1.215	1.561	-2.489	1.422	1.804	-1.022	1.804
-2.489	0.395							
34 R	0.048	0.712	1.664	-2.572	1.531	1.830	-0.997	1.830
-2.572	0.317							
35 L	0.958	0.389	1.945	-2.478	1.677	1.844	-1.391	1.945
-2.478	0.421							
36 E	0.193	1.169	1.963	-2.334	1.695	1.869	-0.358	1.963
-2.334	0.600							
37 R	0.440	1.203	2.290	-1.957	2.069	1.912	0.815	2.290
-1.957	0.967							
38 T	0.667	-0.114	2.188	-1.666	1.959	1.887	0.790	2.188
-1.666	0.816							
39 W	1.691	-0.605	2.571	-1.011	2.260	1.922	0.242	2.571
-1.011	1.010							
40 Q	0.566	0.712	2.262	-0.555	1.914	1.347	0.290	2.262
-0.555	0.934							
41 E	0.433	0.017	1.832	0.015	1.440	0.723	-0.719	1.832
-0.719	0.534							
42 N	0.370	0.069	2.066	-0.059	1.759	1.328	-0.761	2.066
-0.761	0.682							
43 W	0.421	-0.050	1.963	-0.523	1.750	1.308	-0.348	1.963
-0.523	0.646							
44 A	0.402	0.279	1.627	-1.429	1.330	1.265	-0.511	1.627
-1.429	0.423							
45 R	0.237	0.888	1.496	-1.844	1.121	0.686	-0.444	1.496
-1.844	0.306							
46 L	-0.787	-0.021	1.132	-1.949	0.765	0.649	-0.080	1.132
-1.949	-0.042							
47 G	0.288	0.542	1.412	-0.995	1.057	0.665	-0.215	1.412
-0.995	0.393							
48 T	-0.079	0.523	1.290	-0.084	1.048	0.666	1.384	1.384
-0.084	0.678							
49 F	-0.212	0.391	1.103	0.863	0.847	0.061	1.605	1.605
-0.212	0.665							
50 N	0.813	0.469	1.487	1.529	1.148	0.096	1.057	1.529
0.096	0.943							
51 V	0.585	0.487	1.739	1.889	1.467	0.115	1.276	1.889
0.115	1.080							
52 P	0.022	1.439	1.421	1.930	1.303	0.096	1.824	1.930

82 G	0.977	0.345	1.515	-0.593	1.148	0.633	1.303	1.515
-0.593	0.761							
83 E	1.230	0.345	1.421	-0.888	1.084	1.234	1.238	1.421
-0.888	0.809							
84 G	0.863	-0.140	1.057	-0.952	0.802	1.216	1.607	1.607
-0.952	0.636							
85 L	0.813	-0.408	0.898	-0.572	0.601	1.196	1.447	1.447
-0.572	0.568							
86 H	0.585	-0.408	1.066	0.079	0.820	1.816	1.714	1.816
-0.408	0.810							
87 V	0.225	0.129	0.982	0.574	0.729	1.236	1.961	1.961
0.129	0.834							
88 G	-0.717	-0.182	0.907	0.784	0.784	1.241	2.395	2.395
-0.717	0.745							
89 H	0.225	-1.133	0.982	0.833	0.729	1.236	1.961	1.961
-1.133	0.690							
90 P	-0.028	-1.224	1.075	0.287	0.793	0.635	2.025	2.025
-1.224	0.509							
91 L	-0.300	-1.091	1.057	-0.501	0.784	0.635	1.758	1.758
-1.091	0.334							
92 G	-0.528	-0.348	1.066	-1.291	0.829	0.635	0.748	1.066
-1.291	0.159							
93 Y	-0.332	-1.071	1.103	-1.873	0.811	0.035	0.521	1.103
-1.873	-0.115							
94 I	0.168	-1.071	1.132	-1.700	0.856	0.505	0.293	1.132
-1.700	0.026							
95 A	0.515	-0.747	1.094	-1.184	0.838	0.501	0.446	1.094
-1.184	0.209							
96 T	0.035	0.067	1.356	-0.522	1.121	0.521	0.778	1.356
-0.522	0.479							
97 D	0.288	-0.831	1.103	-0.304	0.884	0.501	-0.564	1.103
-0.831	0.154							
98 V	1.059	-1.544	1.674	-0.552	1.376	1.124	-0.887	1.674
-1.544	0.321							
99 Y	0.806	-0.635	1.926	-1.146	1.613	1.144	0.456	1.926
-1.146	0.595							
100A	-0.104	-0.342	1.664	-1.548	1.412	1.128	0.666	1.664
-1.548	0.411							
101R	-0.471	-0.438	1.823	-1.771	1.567	1.264	0.674	1.823
-1.771	0.378							
102Y	-0.503	-0.625	1.935	-1.721	1.622	1.279	0.030	1.935
-1.721	0.288							
103F	-0.616	0.596	1.561	-1.854	1.376	1.261	0.287	1.561
-1.854	0.373							
104R	-0.389	1.379	1.552	-2.111	1.330	1.261	1.297	1.552
-2.111	0.617							
105M	-0.389	0.469	1.552	-2.374	1.330	1.261	1.297	1.552
-2.374	0.450							
106V	0.174	0.379	1.599	-1.997	1.403	1.283	0.852	1.599
-1.997	0.528							
107G	0.522	0.566	1.543	-1.418	1.440	1.280	1.190	1.543
-1.418	0.732							
108R	-0.325	-0.062	1.029	-0.589	0.975	0.661	1.625	1.625
-0.589	0.473							
109N	0.073	-1.079	1.197	0.139	1.103	1.264	1.948	1.948
-1.079	0.663							
110V	0.440	-1.061	1.318	0.348	1.112	1.262	0.349	1.318
-1.061	0.538							
111L	-0.503	-1.140	1.244	0.243	1.166	1.267	0.784	1.267

141N		1.293	0.391	1.524	0.443	1.321	0.674	0.138	1.524
0.138		0.826							
142V		0.383	0.596	1.262	0.724	1.121	0.658	0.348	1.262
0.348		0.727							
143V		0.155	1.183	1.365	0.841	1.230	0.683	0.373	1.365
0.155		0.833							
144N		0.288	1.074	1.795	0.666	1.704	1.308	1.383	1.795
0.288		1.174							
145F		0.225	1.093	1.823	-0.013	1.768	1.310	1.659	1.823
-0.013		1.123							
146R		-0.123	2.080	1.860	-0.818	1.786	1.313	1.505	2.080
-0.818		1.086							
147R		0.471	1.062	1.973	-1.394	1.750	1.312	0.917	1.973
-1.394		0.870							
148Q		0.294	0.876	2.103	-1.896	1.914	1.896	1.029	2.103
-1.896		0.888							
149L		0.294	0.211	2.085	-2.008	1.968	1.897	1.213	2.085
-2.008		0.809							
150G		0.389	1.042	1.646	-2.128	1.449	1.272	1.214	1.646
-2.128		0.698							
151R		-0.458	0.505	1.150	-2.106	0.929	0.652	1.466	1.466
-2.106		0.305							
152L		-0.477	0.231	0.814	-2.001	0.510	0.610	1.303	1.303
-2.001		0.141							
153G		0.237	1.291	1.057	-1.260	0.674	1.224	1.135	1.291
-1.260		0.623							
154F		0.509	1.477	1.337	-0.289	1.039	1.713	1.126	1.713
-0.289		0.987							
155G		0.655	2.465	1.057	0.978	0.720	1.108	1.287	2.465
0.655		1.181							
156H		1.502	2.693	1.571	1.765	1.185	1.728	0.851	2.693
0.851		1.613							
157D		1.407	2.429	2.010	1.800	1.704	2.352	0.850	2.429
0.850		1.793							
158S		2.399	2.745	2.225	1.095	1.905	2.368	0.758	2.745
0.758		1.928							
159R		1.457	2.381	2.169	0.264	1.905	2.372	1.009	2.381
0.264		1.651							
160R		1.736	2.058	2.160	-0.106	1.886	1.772	0.902	2.160
-0.106		1.487							
161S		1.432	1.784	2.085	0.172	1.722	1.304	0.951	2.085
0.172		1.350							
162F		1.350	0.832	2.132	0.658	1.722	1.303	0.832	2.132
0.658		1.261							
163S		1.717	1.545	1.973	1.091	1.567	1.168	0.824	1.973
0.824		1.412							
164T		1.217	0.515	1.421	1.115	1.084	0.545	1.414	1.421
0.515		1.044							
165T		1.438	-0.382	1.543	1.064	1.248	1.013	1.245	1.543
-0.382		1.024							
166D		1.438	-0.060	1.543	0.995	1.248	1.013	1.245	1.543
-0.060		1.060							
167V		0.907	-1.101	1.646	0.869	1.330	1.013	1.417	1.646
-1.101		0.869							
168D		0.844	-0.514	1.879	0.502	1.649	1.617	1.376	1.879
-0.514		1.050							
169F		-0.117	-0.562	1.702	-0.071	1.513	1.623	1.357	1.702
-0.562		0.778							
170Y		-0.420	-0.891	1.627	-0.853	1.349	1.154	1.407	1.627

200A	0.136	1.185	1.356	-2.070	1.248	1.780	0.368	1.780
-2.070	0.572							
201E	0.054	1.185	1.178	-1.585	1.039	1.200	0.554	1.200
-1.585	0.518							
202F	0.996	1.423	1.253	-1.147	0.984	1.195	0.119	1.423
-1.147	0.689							
203E	1.363	1.633	1.375	-0.739	0.993	1.193	-1.480	1.633
-1.480	0.620							
204S	1.495	0.854	1.804	-0.741	1.467	1.818	-0.470	1.818
-0.741	0.890							
205G	1.091	0.538	1.262	-0.839	1.011	1.236	0.178	1.262
-0.839	0.640							
206A	1.091	0.538	1.244	-1.239	1.066	1.237	0.362	1.244
-1.239	0.614							
207R	1.230	1.165	1.188	-1.022	1.020	1.127	0.379	1.230
-1.022	0.727							
208C	1.179	1.165	1.029	-0.824	0.820	1.107	0.220	1.179
-0.824	0.671							
209L	1.179	1.668	1.029	-0.368	0.820	1.107	0.220	1.668
-0.368	0.808							
210D	1.312	1.369	1.459	-0.398	1.294	1.731	1.229	1.731
-0.398	1.142							
211G	1.679	0.830	1.300	-0.417	1.139	1.596	1.221	1.679
-0.417	1.049							
212G	0.958	1.034	1.533	-0.800	1.248	1.603	0.621	1.603
-0.800	0.885							
213R	1.672	0.203	1.617	-0.772	1.239	1.598	-0.824	1.672
-0.824	0.676							
214D	1.401	-0.120	1.795	-1.021	1.558	1.703	-0.750	1.795
-1.021	0.652							
215W	0.459	-0.659	1.720	-1.279	1.613	1.709	-0.316	1.720
-1.279	0.464							
216A	0.427	0.471	1.926	-1.682	1.813	1.729	-0.275	1.926
-1.682	0.630							
217K	0.294	1.046	1.496	-1.871	1.339	1.104	-1.284	1.496
-1.871	0.304							
218L	0.022	1.028	1.216	-1.902	0.975	0.615	-1.275	1.216
-1.902	0.097							
219T	1.148	1.233	1.524	-1.712	1.321	1.189	-1.324	1.524
-1.712	0.483							
220A	1.280	1.281	1.954	-1.723	1.795	1.814	-0.314	1.954
-1.723	0.870							
221G	1.053	1.185	1.505	-1.929	1.157	1.219	-1.390	1.505
-1.929	0.400							
222E	2.267	0.233	1.860	-1.856	1.467	1.703	-1.834	2.267
-1.856	0.549							
223R	1.704	0.197	1.543	-1.694	1.303	1.685	-1.286	1.704
-1.694	0.493							
224A	1.065	-0.042	1.403	-1.290	1.285	1.686	0.046	1.686
-1.290	0.593							
225D	1.337	-0.448	1.683	-0.862	1.649	2.175	0.037	2.175
-0.862	0.796							
226V	1.337	-0.174	1.683	-0.775	1.649	2.175	0.037	2.175
-0.775	0.848							
227I	0.952	-0.282	1.505	-0.738	1.412	1.570	0.370	1.570
-0.738	0.684							
228D	1.084	-0.054	1.935	-0.852	1.886	2.195	1.379	2.195
-0.852	1.082							
229E	-0.129	-0.999	1.580	-1.174	1.576	1.711	1.823	1.823

259G	2.684	3.002	1.954	-0.033	1.576	1.612	-0.646	3.002
-0.646	1.450							
260R	2.621	2.984	2.188	0.049	1.895	2.217	-0.688	2.984
-0.688	1.609							
261S	2.848	1.996	2.178	0.285	1.850	2.217	0.323	2.848
0.285	1.671							
262D	2.659	1.499	2.206	0.476	1.841	1.769	0.219	2.659
0.219	1.524							
263R	1.717	0.864	2.150	0.601	1.841	1.773	0.469	2.150
0.469	1.345							
264G	1.584	-0.124	1.963	0.963	1.640	1.167	0.690	1.963
-0.124	1.126							
265N	0.939	0.063	1.692	1.071	1.476	1.149	1.119	1.692
0.063	1.073							
266F	-0.275	0.285	1.356	0.963	1.112	0.664	1.379	1.379
-0.275	0.783							
267P	-0.275	1.273	1.356	0.346	1.112	0.664	1.379	1.379
-0.275	0.836							
268V	-0.275	0.710	1.814	-0.445	1.795	1.259	1.444	1.814
-0.445	0.900							
269F	-0.452	1.619	1.945	-1.424	1.959	1.843	1.556	1.959
-1.424	1.007							
270R	-0.452	2.285	1.926	-1.982	2.014	1.844	1.740	2.285
-1.982	1.053							
271K	-0.319	0.968	2.113	-2.488	2.214	2.450	1.519	2.450
-2.488	0.922							
272R	0.294	0.023	2.561	-2.549	2.597	2.491	1.094	2.597
-2.549	0.930							
273L	0.244	-0.905	2.646	-2.511	2.661	2.512	0.865	2.661
-2.511	0.787							
274R	-0.288	0.113	2.206	-2.299	2.233	1.904	0.810	2.233
-2.299	0.669							
275Q	-0.913	-1.025	1.748	-2.299	1.640	1.327	0.690	1.748
-2.299	0.167							
276W	-0.913	-1.025	1.748	-2.514	1.640	1.327	0.690	1.748
-2.514	0.136							
277M	-0.838	-0.522	1.692	-2.932	1.613	1.323	0.577	1.692
-2.932	0.130							
278M	-0.774	-0.815	1.459	-3.036	1.294	0.718	0.619	1.459
-3.036	-0.077							
279R	-1.021	-0.701	1.132	-2.950	0.920	0.676	-0.555	1.132
-2.950	-0.357							
280I	-0.509	-0.975	1.365	-2.445	1.139	0.670	-0.245	1.365
-2.445	-0.143							
281T	-0.111	0.163	1.375	-1.980	1.093	0.653	-1.200	1.375
-1.980	-0.001							
282A	0.787	-0.532	1.655	-1.374	1.367	1.125	-1.154	1.655
-1.374	0.268							
283Y	0.787	-0.737	1.655	-1.075	1.367	1.125	-1.154	1.655
-1.154	0.281							
284A	0.711	0.209	1.711	-0.773	1.394	1.128	-1.040	1.711
-1.040	0.477							
285D	-0.199	0.748	1.431	-0.952	1.248	1.114	-0.646	1.431
-0.952	0.392							
286R	0.300	0.005	1.702	-1.005	1.567	1.603	0.355	1.702
-1.005	0.647							
287L	1.053	-0.270	1.720	-0.917	1.649	2.072	0.014	2.072
-0.917	0.760							
288L	0.338	-0.162	1.636	-0.388	1.658	2.078	1.459	2.078

-0.388	0.946								
289D	0.338	-0.162	1.636	0.359	1.658	2.078	1.459	2.078	
-0.162	1.052	-0.161	-0.162	1.085	0.794	1.175	1.454	2.049	2.049
290D	-0.161	-0.161	-1.204	1.085	0.623	1.175	1.454	2.049	2.049
-0.162	0.891	0.717	1.053	-0.641	1.440	0.263	1.485	1.938	1.605
291L	-0.161	1.020	-0.212	-0.605	1.188	-0.246	1.185	1.474	1.636
-1.204	0.717	-0.212	-0.605	1.188	-0.246	1.185	1.474	1.636	1.636
292D	1.053	-0.641	1.440	0.263	1.485	1.938	1.605	1.938	
-0.641	1.020	-0.641	1.440	0.263	1.485	1.938	1.605	1.938	
293V	-0.212	-0.605	1.188	-0.246	1.185	1.474	1.636	1.636	
-0.605	0.631	0.580	0.364	0.091	1.571	-0.348	1.494	1.598	1.403
294L	-0.711	-0.017	1.160	-0.379	1.139	1.004	1.865	1.865	1.598
-0.711	0.580	-0.017	1.160	-0.379	1.139	1.004	1.865	1.865	1.598
295D	0.364	0.091	1.571	-0.348	1.494	1.598	1.403	1.598	
-0.348	0.882	0.111	0.383	1.627	-0.420	1.549	1.152	1.576	1.627
296W	0.882	0.111	1.377	1.627	-0.643	1.549	1.152	1.576	1.627
-0.420	0.854	0.111	1.377	1.627	-0.643	1.549	1.152	1.576	1.627
297P	0.964	1.053	0.904	2.160	-0.920	2.178	1.741	1.206	2.178
-0.643	1.189	0.749	0.820	2.085	-1.320	2.014	1.272	1.256	2.085
298E	0.982	1.116	1.143	2.057	-1.536	2.041	1.264	1.178	2.057
-0.920	1.038	1.363	1.848	2.141	-1.671	2.142	1.288	1.121	2.142
299Q	1.176	1.135	0.513	2.244	-1.729	2.251	1.313	1.147	2.251
-1.320	0.982	0.982	0.982	2.216	-1.316	2.187	1.311	0.871	2.216
300V	1.116	1.143	2.057	-1.536	2.041	1.264	1.178	2.057	
-1.536	1.176	1.135	0.513	2.244	-1.729	2.251	1.313	1.147	2.251
301K	0.982	0.982	0.982	2.216	-1.316	2.187	1.311	0.871	2.216
-1.671	0.982	0.881	0.800	2.356	-0.893	2.214	1.335	0.304	2.356
302T	0.982	0.881	0.800	2.356	-0.893	2.214	1.335	0.304	2.356
-1.729	0.936	0.881	0.800	2.356	-0.893	2.214	1.335	0.304	2.356
303M	0.654	-0.066	0.762	1.767	-0.341	1.558	0.742	0.561	1.767
-0.341	0.654	0.712	0.762	1.767	-0.341	1.558	0.742	0.561	1.767
305R	-0.035	0.804	1.561	-0.355	1.358	0.722	0.520	1.561	
-0.355	0.496	0.686	2.001	-0.783	1.786	1.329	0.575	2.001	
307W	0.870	0.528	1.816	1.823	-1.377	1.567	1.307	0.571	1.823
-0.783	0.528	0.891	0.591	2.140	1.589	-1.411	1.248	0.702	2.140
308I	0.891	0.591	2.140	1.589	-1.411	1.248	0.702	0.613	2.140
-1.377	0.509	1.417	1.281	-1.141	0.893	0.662	0.726	1.417	
309G	0.509	0.621	1.274	0.604	1.262	-0.627	0.875	0.636	-0.306
-1.411	0.509	0.621	1.274	0.604	1.262	-0.627	0.875	0.636	1.274
310R	0.509	0.531	1.546	-0.456	1.281	-0.538	0.884	0.636	-0.039
-1.141	0.509	0.473	1.318	-1.121	1.290	-1.007	0.929	0.636	1.546
311S	0.531	0.473	1.318	-1.121	1.290	-1.007	0.929	0.636	-1.050
-0.627	0.473	0.471	0.471	-0.893	0.776	-1.545	0.465	0.017	1.318
312T	0.471	-0.189	0.471	-0.893	0.776	-1.545	0.465	0.017	-0.614
-0.538	0.471	-0.189	0.471	-0.893	0.776	-1.545	0.465	0.017	0.776
313G	0.142	-0.521	0.142	-0.893	0.561	-1.931	0.264	0.001	-0.523
-1.121	0.142	-0.521	0.142	-0.893	0.561	-1.931	0.264	0.001	0.561
314A	0.471	-0.521	0.471	-0.893	0.561	-1.931	0.264	0.001	-0.403
-1.545	-0.189	-0.434	-0.434	-0.893	0.561	-1.931	0.264	0.001	0.515
315V	-0.189	-0.434	-0.434	-0.893	0.561	-1.931	0.264	0.001	-1.414
-1.931	-0.434	-0.434	-0.434	-0.893	0.561	-1.931	0.264	0.001	0.524
316A	-0.254	-0.439	-0.439	0.017	0.515	-1.729	0.264	0.001	-0.403
-1.729	-0.254	-0.667	-0.667	0.017	0.524	-1.257	0.310	0.001	0.515
317L	-0.667	-0.355	-0.355	0.017	0.524	-1.257	0.310	0.001	-1.414
-1.414	-0.355	-0.355	-0.355	0.017	0.524	-1.257	0.310	0.001	0.524

318F	-0.534	0.221	0.954	-0.782	0.784	0.626	-0.405	0.954
-0.782	0.123							
319S	-0.167	1.251	1.075	-0.703	0.793	0.624	-2.003	1.251
-2.003	0.124							
320A	-0.167	0.934	1.075	-1.095	0.793	0.624	-2.003	1.075
-2.003	0.023							
321R	0.825	1.473	1.309	-1.400	0.938	0.639	-2.279	1.473
-2.279	0.215							
322A	2.039	1.287	1.646	-1.191	1.303	1.124	-2.539	2.039
-2.539	0.524							
323A	2.260	1.113	1.767	-0.294	1.467	1.592	-2.707	2.260
-2.707	0.743							
324S	2.488	1.688	1.758	0.799	1.422	1.592	-1.697	2.488
-1.697	1.150							
325D	1.641	0.736	1.262	1.510	0.902	0.972	-1.445	1.641
-1.445	0.797							
326D	2.001	0.736	1.589	1.266	1.267	1.572	-0.461	2.001
-0.461	1.138							
327G	1.634	-0.128	1.468	0.471	1.257	1.573	1.138	1.634
-0.128	1.059							
328F	1.856	-0.180	1.589	-0.316	1.422	2.042	0.969	2.042
-0.316	1.055							
329E	0.718	-0.102	1.178	-0.847	1.084	1.555	1.299	1.555
-0.847	0.698							
330V	0.579	-0.851	1.234	-1.014	1.130	1.665	1.282	1.665
-1.014	0.575							
331D	-0.016	-0.264	1.122	-1.148	1.166	1.667	1.871	1.871
-1.148	0.628							
332I	-0.016	-0.312	1.122	-1.390	1.166	1.667	1.871	1.871
-1.390	0.587							
333E	-0.180	0.826	0.991	-1.538	0.957	1.087	1.938	1.938
-1.538	0.583							
334V	0.383	0.610	1.309	-1.261	1.121	1.106	1.390	1.390
-1.261	0.665							
335F	0.016	1.245	1.468	-1.029	1.276	1.242	1.398	1.468
-1.029	0.802							
336T	0.655	1.910	1.851	-0.570	1.567	1.259	1.296	1.910
-0.570	1.138							
337T	0.794	1.215	1.795	-0.296	1.522	1.148	1.313	1.795
-0.296	1.070							
338R	1.356	0.550	2.113	0.027	1.686	1.166	0.765	2.113
0.027	1.095							
339P	1.356	0.363	2.094	0.286	1.741	1.168	0.949	2.094
0.286	1.137							
340D	0.446	0.005	1.832	0.442	1.540	1.152	1.160	1.832
0.005	0.939							
341T	0.477	-0.044	1.627	0.108	1.339	1.132	1.119	1.627
-0.044	0.823							
342L	0.345	-0.941	1.197	-0.413	0.866	0.507	0.110	1.197
-0.941	0.239							
343F	0.541	-0.941	1.150	-0.897	0.747	0.508	-0.069	1.150
-0.941	0.148							
344G	-0.212	-0.863	1.132	-1.151	0.665	0.039	0.272	1.132
-1.151	-0.017							
345A	-1.122	-1.695	0.851	-1.203	0.519	0.024	0.666	0.851
-1.695	-0.280							
346T	-0.774	-1.695	0.814	-1.227	0.501	0.020	0.820	0.820
-1.695	-0.220							
347Y	-0.774	-1.827	0.795	-1.499	0.556	0.022	1.003	1.003

377G	1.672	2.032	1.505	-1.065	0.884	0.048	0.816	2.032
-1.065	0.842							
378G	1.704	1.405	1.300	-0.866	0.683	0.028	0.776	1.704
-0.866	0.719							
379T	2.317	0.453	1.375	-0.647	0.811	0.608	0.417	2.317
-0.647	0.762							
380P	2.090	-0.038	1.384	-0.705	0.856	0.608	-0.593	2.090
-0.705	0.515							
381G	1.224	-0.396	1.253	-1.189	0.884	0.610	-0.272	1.253
-1.189	0.302							
382E	0.996	-1.430	1.262	-1.801	0.929	0.610	-1.282	1.262
-1.801	-0.102							
383A	0.800	-1.192	1.066	-2.333	0.774	0.590	-2.333	1.066
-2.333	-0.375							
384I	0.547	-0.378	1.075	-2.516	0.738	0.591	-2.221	1.075
-2.516	-0.309							
385A	0.452	-0.054	1.515	-2.443	1.257	1.215	-2.222	1.515
-2.443	-0.040							
386A	0.225	-0.378	1.617	-2.277	1.367	1.240	-2.197	1.617
-2.277	-0.058							
387Y	0.225	-0.378	1.617	-2.252	1.367	1.240	-2.197	1.617
-2.252	-0.054							
388R	0.225	0.029	1.617	-2.386	1.367	1.240	-2.197	1.617
-2.386	-0.015							
389R	0.225	0.047	1.617	-2.595	1.367	1.240	-2.197	1.617
-2.595	-0.042							
390A	0.225	0.089	1.617	-2.721	1.367	1.240	-2.197	1.617
-2.721	-0.054							
391I	0.705	0.628	1.814	-2.749	1.768	1.816	-2.464	1.816
-2.749	0.217							
392A	0.850	0.748	1.533	-2.372	1.449	1.211	-2.303	1.533
-2.372	0.159							
393A	1.217	1.323	1.375	-1.613	1.294	1.075	-2.311	1.375
-2.311	0.337							
394K	0.503	2.136	1.290	-0.761	1.303	1.081	-0.866	2.136
-0.866	0.669							
395S	1.502	1.796	1.758	-0.052	1.686	1.678	-1.214	1.796
-1.214	1.022							
396D	1.634	1.515	2.188	-0.080	2.160	2.303	-0.205	2.303
-0.205	1.359							
397L	1.881	1.832	2.515	-0.627	2.533	2.346	0.969	2.533
-0.627	1.635							
398E	2.014	2.850	2.393	-1.342	2.260	2.350	0.877	2.850
-1.342	1.629							
399R	2.014	2.850	2.393	-1.519	2.260	2.350	0.877	2.850
-1.519	1.604							
400Q	1.647	2.868	2.552	-1.423	2.415	2.486	0.886	2.868
-1.423	1.633							
401E	2.722	2.868	2.963	-1.028	2.770	3.080	0.424	3.080
-1.028	1.971							
402S	2.589	2.920	3.085	-1.093	3.044	3.075	0.516	3.085
-1.093	2.019							
403R	2.652	1.968	2.851	-1.277	2.725	2.471	0.557	2.851
-1.277	1.707							
404E	2.633	0.980	2.515	-1.641	2.306	2.428	0.394	2.633
-1.641	1.374							
405K	1.906	0.201	2.066	-1.639	1.932	1.830	1.009	2.066
-1.639	1.044							
406T	0.914	-0.003	1.851	-1.563	1.731	1.814	1.101	1.851

436G	1.476	0.033	1.066	-1.162	0.455	0.009	-1.465	1.476
-1.465	0.059							
437T	0.838	-0.595	0.926	-1.280	0.437	0.011	-0.133	0.926
-1.280	0.029							
438G	0.212	-1.182	0.926	-1.691	0.528	0.028	-0.188	0.926
-1.691	-0.195							
439A	0.465	-1.450	0.674	-2.268	0.291	0.009	-1.531	0.674
-2.268	-0.544							
440I	-0.129	-0.823	0.561	-2.701	0.328	0.010	-0.942	0.561
-2.701	-0.528							
441M	-0.325	-0.408	0.608	-2.693	0.446	0.009	-0.763	0.608
-2.693	-0.447							
442A	-0.325	0.245	0.608	-2.289	0.446	0.009	-0.763	0.608
-2.289	-0.296							
443V	-0.325	0.736	0.767	-1.308	0.619	0.629	0.514	0.767
-1.308	0.233							
444P	0.813	1.646	1.178	-0.154	0.957	1.116	0.184	1.646
-0.154	0.820							
445G	1.457	1.826	1.515	0.992	1.285	1.142	0.402	1.826
0.402	1.231							
446H	1.590	0.696	1.945	1.520	1.759	1.766	1.412	1.945
0.696	1.527							
447D	2.456	1.145	2.337	1.625	2.087	2.254	0.814	2.456
0.814	1.817							
448Q	1.691	0.431	2.113	0.902	1.832	2.260	0.616	2.260
0.431	1.406							
449R	1.963	-0.060	2.393	0.556	2.196	2.749	0.607	2.749
-0.060	1.486							
450D	1.249	-0.060	2.169	0.254	1.977	2.133	0.591	2.169
-0.060	1.188							
451W	0.749	-0.599	1.898	0.322	1.658	1.644	-0.410	1.898
-0.599	0.752							
452D	0.636	-0.270	2.001	0.121	1.759	2.226	-0.575	2.226
-0.575	0.843							
453F	0.503	-0.182	1.571	-0.328	1.285	1.602	-1.584	1.602
-1.584	0.409							
454A	-0.711	-0.212	1.234	-1.114	0.920	1.117	-1.324	1.234
-1.324	-0.013							
455R	0.281	0.147	1.206	-1.501	0.856	1.092	-1.346	1.206
-1.501	0.105							
456A	-0.932	-0.991	0.851	-1.719	0.547	0.608	-0.902	0.851
-1.719	-0.363							
457F	-0.218	-1.087	1.160	-1.501	0.866	0.623	-0.933	1.160
-1.501	-0.156							
458G	-0.857	-0.338	1.019	-1.421	0.847	0.625	0.399	1.019
-1.421	0.039							
459L	-1.356	-1.061	0.468	-1.433	0.364	0.002	0.988	0.988
-1.433	-0.290							
460P	-0.996	-1.182	0.795	-1.670	0.729	0.601	1.972	1.972
-1.670	0.036							
461I	-0.648	-1.540	0.739	-1.976	0.765	0.599	2.310	2.310
-1.976	0.036							
462V	-1.514	-0.589	0.608	-2.328	0.793	0.601	2.631	2.631
-2.328	0.029							
463E	-0.800	0.135	0.692	-2.493	0.784	0.595	1.186	1.186
-2.493	0.014							
464V	-0.572	0.169	0.440	-2.664	0.465	0.576	0.966	0.966
-2.664	-0.089							
465I	0.294	-0.060	0.571	-2.529	0.437	0.574	0.645	0.645

495A	0.844	-0.050	1.646	-2.008	1.576	1.210	-1.976	1.646
-2.008	0.177							
496K	0.572	0.489	1.627	-2.398	1.567	1.210	-2.243	1.627
-2.398	0.118							
497R	0.206	0.471	1.262	-2.696	1.285	1.193	-1.874	1.285
-2.696	-0.022							
498A	0.705	-0.546	1.533	-2.464	1.604	1.681	-0.873	1.681
-2.464	0.234							
499I	0.838	0.029	1.963	-2.095	2.078	2.306	0.136	2.306
-2.095	0.751							
500V	-0.104	1.209	1.431	-1.550	1.449	1.717	0.506	1.717
-1.550	0.665							
501D	0.123	1.305	1.328	-1.293	1.339	1.692	0.481	1.692
-1.293	0.711							
502R	0.402	1.393	1.477	-1.181	1.494	1.712	1.651	1.712
-1.181	0.993							
503L	1.040	1.393	1.617	-1.267	1.513	1.710	0.319	1.710
-1.267	0.904							
504E	1.634	2.225	1.730	-1.096	1.476	1.708	-0.269	2.225
-1.096	1.058							
505S	1.268	2.463	1.889	-1.144	1.631	1.844	-0.261	2.463
-1.144	1.098							
506A	1.363	1.607	1.449	-1.176	1.112	1.220	-0.260	1.607
-1.176	0.759							
507G	2.210	2.421	1.963	-1.614	1.576	1.839	-0.696	2.421
-1.614	1.100							
508R	1.849	1.469	1.636	-1.965	1.212	1.239	-1.680	1.849
-1.965	0.537							
509G	1.704	1.231	1.917	-2.348	1.531	1.844	-1.840	1.917
-2.348	0.577							
510R	1.065	0.429	1.776	-2.503	1.513	1.846	-0.509	1.846
-2.503	0.517							
511A	1.198	0.447	2.113	-2.699	1.923	2.445	-0.535	2.445
-2.699	0.699							
512R	0.351	0.243	1.617	-2.690	1.403	1.825	-0.283	1.825
-2.690	0.352							
513I	0.351	0.243	2.075	-2.608	2.087	2.419	-0.218	2.419
-2.608	0.621							
514E	-0.496	1.107	1.561	-2.320	1.622	1.800	0.217	1.800
-2.320	0.499							
515F	-0.363	0.029	1.991	-2.167	2.096	2.425	1.227	2.425
-2.167	0.748							
516K	0.003	-0.001	1.832	-1.873	1.941	2.289	1.219	2.289
-1.873	0.773							
517L	-0.123	-1.007	1.991	-1.682	1.977	2.312	0.919	2.312
-1.682	0.627							
518R	-1.198	-0.803	1.580	-1.337	1.622	1.718	1.380	1.718
-1.337	0.423							
519D	-1.198	-0.803	1.580	-1.113	1.622	1.718	1.380	1.718
-1.198	0.455							
520W	-1.426	-0.851	1.132	-1.071	0.984	1.124	0.305	1.132
-1.426	0.028							
521L	-0.578	0.465	1.646	-1.331	1.449	1.743	-0.131	1.743
-1.331	0.466							
522F	-0.465	0.263	1.543	-1.469	1.349	1.161	0.034	1.543
-1.469	0.345							
523A	-0.831	-0.066	1.702	-1.756	1.504	1.296	0.042	1.702
-1.756	0.270							
524R	-0.319	0.562	1.935	-1.733	1.722	1.291	0.351	1.935

554L	0.263	0.221	1.403	-0.837	1.422	1.105	2.199	2.199
-0.837	0.825							
555P	0.263	0.019	1.403	-0.204	1.422	1.105	2.199	2.199
-0.204	0.886							
556D	1.129	0.515	1.795	0.446	1.750	1.592	1.601	1.795
0.446	1.261							
557V	0.515	0.335	1.720	0.736	1.622	1.012	1.960	1.960
0.335	1.129							
558P	1.508	0.335	1.954	0.998	1.768	1.027	1.684	1.954
0.335	1.325							
559D	1.508	-0.228	1.954	1.134	1.768	1.027	1.684	1.954
-0.228	1.264							
560Y	0.642	-0.941	1.561	1.040	1.440	0.539	2.282	2.282
-0.941	0.938							
561S	0.294	0.005	1.599	0.720	1.458	0.543	2.128	2.128
0.005	0.964							
562P	-0.420	-0.492	1.290	0.108	1.139	0.528	2.160	2.160
-0.492	0.616							
563V	-0.420	-0.312	1.290	-0.398	1.139	0.528	2.160	2.160
-0.420	0.570							
564L	-0.167	0.323	1.281	-0.341	1.175	0.528	2.047	2.047
-0.341	0.692							
565F	0.054	0.527	1.403	0.278	1.339	0.997	1.879	1.879
0.054	0.925							
566D	0.553	1.241	1.431	1.209	1.385	1.467	1.650	1.650
0.553	1.276							
567P	0.920	1.557	1.552	1.825	1.394	1.465	0.051	1.825
0.051	1.252							
568D	2.134	1.774	1.907	2.084	1.704	1.949	-0.393	2.134
-0.393	1.594							
569D	3.127	1.593	2.122	1.942	1.905	1.965	-0.484	3.127
-0.484	1.738							
570A	2.987	1.910	2.178	1.546	1.950	2.075	-0.502	2.987
-0.502	1.735							
571D	2.987	2.269	2.178	1.284	1.950	2.075	-0.502	2.987
-0.502	1.749							
572S	2.766	2.088	2.057	1.048	1.786	1.606	-0.333	2.766
-0.333	1.574							
573E	2.267	1.028	2.029	0.873	1.741	1.136	-0.104	2.267
-0.104	1.281							
574P	2.267	0.453	2.272	1.017	2.014	1.155	1.126	2.272
0.453	1.472							
575S	1.053	0.926	1.917	0.945	1.704	0.672	1.570	1.917
0.672	1.255							
576P	0.775	0.071	1.767	0.614	1.549	0.652	0.400	1.767
0.071	0.832							
577P	0.642	0.203	1.889	-0.084	1.823	0.647	0.491	1.889
-0.084	0.801							
578L	0.642	0.419	1.646	-1.028	1.549	0.628	-0.739	1.646
-1.028	0.445							
579A	0.560	0.121	1.692	-1.697	1.549	0.628	-0.858	1.692
-1.697	0.285							
580K	0.920	0.025	1.776	-1.923	1.640	1.208	-1.104	1.776
-1.923	0.363							
581A	0.155	-0.717	1.552	-1.878	1.385	1.215	-1.302	1.552
-1.878	0.059							
582T	0.503	-0.813	1.515	-1.773	1.367	1.211	-1.148	1.515
-1.773	0.123							
583E	0.503	-0.765	1.674	-1.445	1.540	1.831	0.130	1.831

613W	0.263	-1.406	1.449	-0.467	1.093	0.660	1.265	1.449
-1.406	0.408							
614Y	0.168	-0.412	1.889	-1.289	1.613	1.284	1.264	1.889
-1.289	0.645							
615E	-0.363	0.534	1.991	-1.917	1.695	1.283	1.436	1.991
-1.917	0.666							
616L	-0.446	0.317	2.038	-2.126	1.695	1.283	1.317	2.038
-2.126	0.583							
617R	0.819	0.612	2.290	-1.603	1.996	1.747	1.286	2.290
-1.603	1.021							
618Y	1.072	0.407	2.281	-0.787	2.032	1.747	1.173	2.281
-0.787	1.132							
619T	0.711	1.670	2.113	0.487	1.841	1.767	1.467	2.113
0.487	1.437							
620D	1.736	1.754	2.496	1.824	2.142	1.802	0.919	2.496
0.919	1.810							
621P	1.881	2.028	2.216	3.002	1.823	1.198	1.080	3.002
1.080	1.889							
622H	2.494	1.495	2.290	3.493	1.950	1.778	0.721	3.493
0.721	2.032							
623N	2.431	1.441	2.524	3.106	2.269	2.383	0.680	3.106
0.680	2.119							
624S	1.217	0.832	2.188	1.844	1.905	1.898	0.940	2.188
0.832	1.546							
625E	1.173	0.808	1.730	0.522	1.540	1.897	1.342	1.897
0.522	1.287							
626R	1.173	0.808	1.571	-0.465	1.367	1.277	0.065	1.571
-0.465	0.828							
627F	1.091	0.604	1.720	-0.841	1.695	1.831	0.243	1.831
-0.841	0.906							
628C	1.173	1.353	1.898	-1.024	1.905	2.410	0.057	2.410
-1.024	1.110							
629A	1.122	1.317	1.870	-0.871	1.850	1.851	-0.030	1.870
-0.871	1.016							
630K	1.350	0.910	1.767	-0.790	1.741	1.826	-0.055	1.826
-0.790	0.964							
631E	2.064	-0.424	1.832	-0.407	1.786	1.822	-1.317	2.064
-1.317	0.765							
632N	1.856	-1.113	2.300	-0.227	2.114	1.824	-1.607	2.300
-1.607	0.735							
633E	1.091	-1.095	2.318	-0.401	2.132	1.849	-0.575	2.318
-1.095	0.760							
634A	0.465	-1.312	1.860	-1.105	1.540	1.271	-0.695	1.860
-1.312	0.289							
635Y	0.332	-0.498	1.524	-1.697	1.130	0.672	-0.668	1.524
-1.697	0.113							
636W	0.022	0.267	1.468	-2.120	1.093	0.650	-0.335	1.468
-2.120	0.149							
637M	-0.205	0.770	1.571	-2.024	1.203	0.675	-0.310	1.571
-2.024	0.240							
638G	-0.205	1.459	1.814	-1.585	1.476	0.694	0.920	1.814
-1.585	0.653							
639P	0.048	0.922	1.561	-1.255	1.239	0.675	-0.422	1.561
-1.255	0.396							
640R	1.173	1.191	1.870	-1.044	1.586	1.249	-0.470	1.870
-1.044	0.793							
641P	1.571	0.736	2.038	-0.751	1.713	1.852	-0.148	2.038
-0.751	1.002							
642A	1.571	0.916	2.038	-0.527	1.713	1.852	-0.148	2.038

672W	-0.357	-1.562	1.664	-0.132	1.604	1.860	1.535	1.860
-1.562	0.659							
673H	-1.350	-0.520	1.431	-0.359	1.458	1.845	1.810	1.845
-1.350	0.616							
674K	-1.735	-0.815	1.253	-0.671	1.221	1.240	2.143	2.143
-1.735	0.377							
675V	-0.521	-1.019	1.589	-1.079	1.586	1.725	1.883	1.883
-1.079	0.595							
676L	-0.471	-0.833	1.487	-1.080	1.576	1.705	2.296	2.296
-1.080	0.669							
677Y	-0.243	-0.725	1.318	-0.912	1.358	1.085	2.029	2.029
-0.912	0.559							
678D	-0.471	0.538	1.029	-0.277	0.893	1.110	2.231	2.231
-0.471	0.722							
679L	-0.471	0.854	1.029	0.004	0.893	1.110	2.231	2.231
-0.471	0.807							
680G	0.522	1.872	1.262	0.430	1.039	1.125	1.956	1.956
0.430	1.172							
681H	1.053	1.820	1.160	0.853	0.957	1.126	1.784	1.820
0.853	1.250							
682V	0.686	2.088	1.318	0.979	1.112	1.261	1.792	2.088
0.686	1.320							
683S	1.761	1.778	1.730	0.820	1.467	1.856	1.331	1.856
0.820	1.535							
684S	1.533	1.736	1.982	0.386	1.786	1.875	1.550	1.982
0.386	1.550							
685R	1.280	1.694	2.075	-0.382	1.850	1.274	1.615	2.075
-0.382	1.344							
686E	1.780	0.676	2.627	-0.786	2.333	1.897	1.025	2.627
-0.786	1.365							
687P	1.634	0.005	2.907	-1.179	2.652	2.502	0.865	2.907
-1.179	1.341							
688Y	0.642	0.255	2.674	-1.585	2.506	2.487	1.140	2.674
-1.585	1.160							
689R	0.142	1.153	2.122	-1.978	2.023	1.864	1.730	2.122
-1.978	1.008							
690R	0.092	0.966	2.094	-1.854	1.968	1.305	1.643	2.094
-1.854	0.888							
691L	0.338	-0.254	2.178	-1.356	2.069	1.329	1.586	2.178
-1.356	0.841							
692V	0.819	-0.374	1.917	-0.364	1.786	1.309	1.254	1.917
-0.374	0.907							
693N	0.433	0.213	1.739	0.339	1.549	0.704	1.587	1.739
0.213	0.938							
694Q	-0.338	-0.396	1.169	0.240	1.057	0.081	1.910	1.910
-0.396	0.532							
695G	0.623	-1.294	1.580	-0.427	1.422	0.118	1.638	1.638
-1.294	0.523							
696Y	0.990	-1.921	1.702	-1.160	1.431	0.116	0.039	1.702
-1.921	0.171							
697I	0.427	-1.921	1.655	-1.664	1.358	0.095	0.484	1.655
-1.921	0.062							
698Q	0.180	-1.105	1.328	-1.718	0.984	0.053	-0.689	1.328
-1.718	-0.138							
699A	-0.300	-1.057	1.589	-1.583	1.267	0.072	-0.357	1.589
-1.583	-0.053							
700Y	0.149	-1.057	1.533	-1.472	1.185	0.073	-0.649	1.533
-1.472	-0.034							
701A	1.287	0.163	1.945	-0.990	1.522	0.560	-0.979	1.945

731V	0.850	0.001	1.253	-1.824	1.139	1.665	1.549	1.665
-1.824	0.662							
732E	0.598	-0.078	1.309	-1.895	1.194	1.219	1.721	1.721
-1.895	0.581							
733V	0.730	-0.026	1.646	-1.580	1.604	1.818	1.695	1.818
-1.580	0.841							
734F	-0.344	0.902	1.253	-1.178	1.194	1.223	1.972	1.972
-1.178	0.717							
735Q	0.250	0.752	1.365	-1.002	1.157	1.221	1.384	1.384
-1.002	0.732							
736E	0.117	0.888	1.487	-1.055	1.431	1.216	1.475	1.487
-1.055	0.794							
737F	-0.155	1.145	1.468	-1.375	1.422	1.217	1.208	1.468
-1.375	0.704							
738G	0.787	2.174	1.524	-1.792	1.422	1.212	0.957	2.174
-1.792	0.898							
739K	0.768	1.343	1.646	-2.259	1.686	1.765	0.859	1.765
-2.259	0.830							
740I	0.686	1.343	1.468	-2.198	1.476	1.185	1.045	1.476
-2.198	0.715							
741G	0.686	2.277	1.449	-1.992	1.531	1.186	1.229	2.277
-1.992	0.909							
742K	0.686	2.505	1.907	-1.516	2.214	1.781	1.294	2.505
-1.516	1.267							
743S	0.768	1.577	1.758	-0.855	1.886	1.227	1.116	1.886
-0.855	1.068							
744L	1.685	1.577	2.047	-0.146	2.060	1.245	0.954	2.060
-0.146	1.346							
745K	1.091	2.140	1.935	0.502	2.096	1.247	1.542	2.140
0.502	1.508							
746N	1.141	1.848	1.636	1.245	1.613	0.672	1.637	1.848
0.672	1.399							
747S	0.863	1.814	1.730	1.430	1.731	0.671	1.697	1.814
0.671	1.420							
748V	2.077	0.634	2.085	1.522	2.041	1.155	1.253	2.085
0.634	1.538							
749S	2.210	0.766	1.963	1.361	1.768	1.160	1.162	2.210
0.766	1.484							
750P	1.261	0.449	1.524	0.903	1.440	1.121	1.597	1.597
0.449	1.185							
751D	0.939	0.091	1.160	0.363	1.194	1.119	2.059	2.059
0.091	0.989							
752E	1.805	-0.855	1.552	-0.028	1.522	1.606	1.461	1.805
-0.855	1.009							
753I	1.527	-0.803	1.403	-0.351	1.367	1.586	0.291	1.586
-0.803	0.717							
754C	1.274	-0.478	1.412	-0.145	1.330	1.586	0.403	1.586
-0.478	0.769							
755D	1.002	0.025	1.132	-0.118	0.966	1.097	0.413	1.132
-0.118	0.645							
756A	0.642	-0.023	0.804	-0.362	0.601	0.498	-0.571	0.804
-0.571	0.227							
757Y	1.780	-0.228	1.216	-0.609	0.938	0.985	-0.902	1.780
-0.902	0.454							
758G	2.020	0.992	1.627	-0.698	1.185	0.987	-1.483	2.020
-1.483	0.661							
759A	0.806	0.269	1.272	-0.502	0.875	0.503	-1.039	1.272
-1.039	0.312							
760D	0.939	-0.138	1.702	-0.308	1.349	1.128	-0.030	1.702

790L	-0.686	0.041	1.814	-1.535	1.713	1.292	1.950	1.950
-1.535	0.656							
791Q	-1.198	0.041	1.580	-1.752	1.494	1.298	1.640	1.640
-1.752	0.443							
792R	-1.198	-0.546	1.580	-1.981	1.494	1.298	1.640	1.640
-1.981	0.327							
793V	-1.198	-1.456	1.561	-2.340	1.549	1.299	1.824	1.824
-2.340	0.177							
794W	-0.850	-0.821	1.524	-2.495	1.531	1.295	1.978	1.978
-2.495	0.309							
795R	-1.464	0.257	1.075	-2.613	1.148	1.255	2.403	2.403
-2.613	0.295							
796L	-1.097	-0.466	0.917	-2.228	0.993	1.119	2.395	2.395
-2.228	0.233							
797V	-0.370	0.229	1.365	-1.772	1.367	1.717	1.780	1.780
-1.772	0.617							
798V	0.395	0.952	1.505	-0.772	1.522	2.311	2.025	2.311
-0.772	1.134							
799D	0.459	1.623	1.272	0.102	1.203	1.707	2.067	2.067
0.102	1.205							
800E	1.401	1.575	1.346	0.804	1.148	1.701	1.632	1.701
0.804	1.373							
801H	2.128	1.814	1.795	0.820	1.522	2.299	1.017	2.299
0.820	1.628							
802T	2.690	1.627	2.113	0.477	1.686	2.318	0.469	2.690
0.469	1.626							
803G	2.324	1.137	2.272	-0.445	1.841	2.453	0.477	2.453
-0.445	1.437							
804E	1.597	1.048	1.823	-1.014	1.467	1.855	1.093	1.855
-1.014	1.124							
805T	1.597	1.101	1.664	-1.624	1.294	1.236	-0.185	1.664
-1.624	0.726							
806R	1.900	0.513	1.739	-1.578	1.458	1.705	-0.235	1.900
-1.578	0.786							
807V	1.900	0.275	1.739	-1.384	1.458	1.705	-0.235	1.900
-1.384	0.780							
808A	1.173	0.167	1.290	-0.937	1.084	1.107	0.380	1.290
-0.937	0.609							
809D	1.337	0.706	1.421	-0.799	1.294	1.686	0.313	1.686
-0.799	0.851							
810G	0.490	-0.158	0.907	-0.965	0.829	1.067	0.749	1.067
-0.965	0.417							
811V	1.356	-0.246	1.300	-1.220	1.157	1.554	0.151	1.554
-1.220	0.579							
812E	0.718	0.341	1.160	-1.408	1.139	1.556	1.483	1.556
-1.408	0.713							
813L	0.718	-0.438	1.160	-1.069	1.139	1.556	1.483	1.556
-1.069	0.650							
814D	0.686	0.580	1.365	-0.521	1.339	1.576	1.524	1.576
-0.521	0.936							
815I	0.338	0.041	1.403	-0.075	1.358	1.580	1.370	1.580
-0.075	0.859							
816D	0.111	0.161	1.505	-0.141	1.467	1.605	1.395	1.605
-0.141	0.872							
817T	0.825	-0.288	1.589	-0.531	1.458	1.600	-0.050	1.600
-0.531	0.658							
818L	-0.389	0.035	1.234	-1.403	1.148	1.116	0.394	1.234
-1.403	0.305							
819R	0.250	0.730	1.533	-1.514	1.339	1.734	0.340	1.734

849N	0.945	0.945	1.375	1.991	2.181	1.604	1.295	1.158	2.181
850H	0.813	0.813	0.856	2.113	1.961	1.877	1.290	1.250	2.113
851L	0.860	1.451	1.293	1.579	2.309	0.860	2.278	1.865	0.983
852T	-0.083	1.595	1.097	2.323	2.272	-0.083	2.296	2.465	1.209
853K	-0.707	1.654	0.920	1.832	2.403	-0.707	2.461	3.049	2.309
854K	-0.386	1.611	1.420	0.904	2.515	-0.386	2.606	2.918	1.045
855H	-0.400	1.575	2.134	0.431	2.599	0.097	2.597	2.913	2.918
856R	0.148	1.482	1.571	1.155	2.281	0.261	2.433	2.894	0.148
857D	0.047	1.535	1.344	0.341	2.075	0.047	2.069	2.318	2.318
858A	-0.368	1.214	1.249	-0.198	2.057	-0.368	1.905	2.349	2.349
859V	-1.040	1.033	1.249	-0.294	1.898	-0.988	1.731	1.729	-1.040
860P	-2.050	0.612	1.116	0.377	1.468	-1.269	1.257	1.104	1.898
861R	-1.704	0.286	0.250	0.377	1.075	-1.704	0.929	0.617	-2.050
862A	-1.978	0.013	0.610	-0.641	1.403	-1.978	1.294	1.216	1.468
863A	-1.965	0.205	0.977	-0.737	1.767	-1.965	1.576	1.234	-0.468
864V	-1.929	0.288	0.263	-0.246	1.440	-1.929	1.312	1.220	-0.837
865E	-1.642	0.205	-0.237	-0.264	0.889	-1.642	0.829	0.597	1.440
866P	-1.453	0.020	0.010	-1.043	1.216	-1.453	1.203	0.639	-0.032
867L	-1.502	0.245	-0.389	-1.402	1.206	-1.502	1.248	0.617	1.216
868V	-1.700	0.273	-0.736	-0.839	1.244	-1.700	1.267	0.660	2.096
869Q	-1.954	0.263	-1.097	-0.947	0.917	-1.954	0.902	0.958	0.958
870M	-2.076	-0.166	-1.097	-1.438	0.917	-1.097	0.902	0.958	0.958
871L	-2.003	-0.253	-1.097	-0.965	0.917	-2.003	0.902	0.958	0.958
872A	-1.730	-0.175	-0.730	-0.671	1.038	-0.252	0.911	0.059	0.958
873P	-1.354	-0.252	-0.977	-0.995	0.954	-0.977	0.811	-0.640	0.954
874L	-1.456	-0.316	-0.730	-0.671	1.122	-0.954	0.938	0.036	1.038
875A	-1.354	-0.047	-0.578	-1.354	1.038	-0.954	0.911	-0.584	0.954
876P	-0.574	-0.503	-0.503	0.001	1.122	-0.954	0.911	-0.375	0.954
877H	-0.503	0.111	-0.142	-0.562	1.066	-0.503	0.911	0.635	1.066
878I	-0.621	0.264	0.284	0.933	-1.156	1.150	-0.142	-0.375	1.215
					1.561	-0.052	1.002	1.215	-0.621
					-0.649	1.358	1.810	-1.082	1.215
						1.358	-1.082	1.810	

-1.156	0.396								
879A	0.218	-1.035	1.477	-1.544	1.367	1.815	0.363	1.815	
-1.544	0.380	-0.547	-0.222	1.253	-2.162	1.112	1.821	0.165	1.821
880E	-0.547	-0.222	1.253	-2.162	1.112	1.821	0.165	1.821	
-2.162	0.203	-1.261	-1.001	1.010	-2.563	0.948	1.207	0.333	1.207
881E	-1.261	-1.001	1.010	-2.563	0.948	1.207	0.333	1.207	
-2.563	-0.190	-0.490	-0.949	1.580	-2.564	1.440	1.830	0.010	1.830
882L	-0.490	-0.949	1.580	-2.564	1.440	1.830	0.010	1.830	
-2.564	0.122	-1.204	-0.136	1.496	-2.639	1.449	1.835	1.456	1.835
883W	-1.204	-0.136	1.496	-2.639	1.449	1.835	1.456	1.835	
-2.639	0.322	-1.337	0.858	1.160	-2.638	1.039	1.236	1.482	1.482
884L	-1.337	0.858	1.160	-2.638	1.039	1.236	1.482	1.482	
-2.638	0.257	-1.388	1.553	1.132	-2.101	0.984	0.677	1.395	1.553
885R	-1.388	1.553	1.132	-2.101	0.984	0.677	1.395	1.553	
-2.101	0.322	-0.477	1.595	1.412	-1.250	1.130	0.691	1.001	1.595
886L	-0.477	1.595	1.412	-1.250	1.130	0.691	1.001	1.595	
-1.250	0.586	0.484	1.595	1.589	-0.048	1.267	0.686	1.020	1.595
887G	0.484	1.595	1.589	-0.048	1.267	0.686	1.020	1.595	
-0.048	0.942	1.476	0.968	1.823	0.973	1.412	0.700	0.745	1.823
888N	1.476	0.968	1.823	0.973	1.412	0.700	0.745	1.823	
0.700	1.157	0.629	0.449	1.309	1.299	0.948	0.081	1.180	1.309
889T	0.629	0.449	1.309	1.299	0.948	0.081	1.180	1.309	
0.081	0.842	1.344	0.586	1.393	0.928	0.938	0.076	-0.265	1.393
890T	0.714	0.714	0.586	1.393	0.928	0.938	0.076	-0.265	1.393
891S	1.116	0.453	1.561	0.573	1.157	0.696	0.002	1.561	
0.002	0.794	1.034	-0.576	1.253	0.087	0.802	0.655	0.116	1.253
892L	0.481	0.838	-0.013	1.300	0.268	0.920	0.654	0.295	1.300
893A	0.838	0.838	-0.013	1.300	0.268	0.920	0.654	0.295	1.300
-0.013	0.609	-0.073	0.818	1.038	0.494	0.720	0.638	0.505	1.038
894H	-0.073	0.592	0.818	1.038	0.494	0.720	0.638	0.505	1.038
-0.073	0.592	-0.351	0.728	1.132	0.639	0.838	0.637	0.565	1.132
895G	-0.351	0.591	0.640	1.664	0.463	1.467	1.227	0.195	1.664
-0.351	0.598	0.591	0.640	1.664	0.463	1.467	1.227	0.195	1.664
896P	0.591	0.591	0.281	1.664	0.126	1.467	1.227	0.195	1.664
0.195	0.892	1.091	0.455	1.776	-0.327	1.613	1.096	-0.081	1.776
897F	0.793	0.803	-0.310	1.786	-0.528	1.658	1.096	-1.091	1.786
0.126	0.591	0.863	-0.310	1.786	-0.528	1.658	1.096	-1.091	1.786
898P	1.091	0.863	-1.346	1.543	-0.633	1.385	1.077	-2.321	1.543
-0.327	0.081	1.325	-1.442	1.860	-0.672	1.668	1.092	-2.240	1.860
900A	0.863	0.863	-1.442	1.543	-0.633	1.385	1.077	-2.321	1.543
-2.321	0.081	0.227	1.325	-1.442	1.860	-0.672	1.668	1.092	1.860
901D	0.610	0.610	-1.442	1.533	-0.944	1.403	1.078	-2.025	1.533
-2.240	0.031	0.515	-0.867	0.963	-1.332	0.756	0.485	-1.502	0.963
902A	0.016	-0.211	-0.376	1.234	-1.401	1.075	0.974	-0.501	1.234
-1.502	-0.211	0.515	-0.376	1.234	-1.401	1.075	0.974	-0.501	1.234
904Y	0.217	0.376	-0.066	1.290	-1.414	1.121	1.085	-0.518	1.290
-1.401	0.268	0.572	0.714	1.487	-0.901	1.276	1.105	0.533	1.487
905L	0.572	0.684	0.403	1.365	-0.619	1.267	1.106	2.132	2.132
-0.901	0.206	0.837	0.403	1.365	-0.619	1.267	1.106	2.132	2.132

908E	0.819	0.223	1.440	-0.643	1.394	1.686	1.773	1.773
-0.643	0.956							
909T	1.280	-0.448	1.776	-0.971	1.622	1.700	1.671	1.776
-0.971	0.947							
910V	1.647	-0.448	2.141	-1.159	1.905	1.718	1.302	2.141
-1.159	1.015							
911E	0.781	-0.448	1.748	-1.353	1.576	1.230	1.899	1.899
-1.353	0.776							
912Y	0.667	-0.414	1.748	-1.050	1.586	0.673	2.089	2.089
-1.050	0.757							
913P	0.104	0.620	1.431	-0.959	1.422	0.655	2.637	2.637
-0.959	0.844							
914V	0.781	1.093	1.851	-0.501	1.741	0.694	1.935	1.935
-0.501	1.085							
915Q	0.648	1.093	1.515	-0.187	1.330	0.094	1.962	1.962
-0.187	0.922							
916V	1.129	1.415	1.711	0.230	1.731	0.670	1.695	1.731
0.230	1.226							
917N	0.762	2.138	1.346	0.168	1.449	0.652	2.064	2.138
0.168	1.226							
918G	1.261	2.343	1.898	-0.322	1.932	1.275	1.474	2.343
-0.322	1.409							
919K	1.242	1.619	1.561	-1.249	1.513	1.233	1.311	1.619
-1.249	1.033							
920V	1.742	0.692	2.113	-1.919	1.996	1.856	0.721	2.113
-1.919	1.029							
921R	1.065	0.692	1.692	-2.406	1.677	1.817	1.423	1.817
-2.406	0.851							
922G	0.471	-0.122	1.580	-2.425	1.713	1.819	2.012	2.012
-2.425	0.721							
923R	-0.123	-0.749	1.010	-2.378	1.066	1.225	2.535	2.535
-2.378	0.370							
924V	0.244	-1.023	1.132	-2.358	1.075	1.224	0.936	1.224
-2.358	0.176							
925V	0.111	-0.436	0.702	-2.289	0.601	0.599	-0.073	0.702
-2.289	-0.112							
926V	0.383	0.199	0.982	-1.958	0.966	1.088	-0.082	1.088
-1.958	0.225							
927A	0.446	0.870	0.748	-1.404	0.647	0.483	-0.040	0.870
-1.404	0.250							
928A	1.312	1.445	1.141	-0.441	0.975	0.971	-0.638	1.445
-0.638	0.681							
929D	2.039	1.936	1.589	0.311	1.349	1.569	-1.253	2.039
-1.253	1.077							
930T	2.766	1.193	2.038	0.666	1.722	2.167	-1.868	2.766
-1.868	1.241							
931D	2.962	1.533	2.234	0.373	1.877	2.187	-0.817	2.962
-0.817	1.479							
932E	2.248	0.994	2.150	-0.285	1.886	2.192	0.628	2.248
-0.285	1.402							
933E	1.976	0.419	2.328	-1.086	2.205	2.298	0.702	2.328
-1.086	1.263							
934T	1.780	-0.252	2.132	-1.551	2.050	2.278	-0.349	2.278
-1.551	0.870							
935L	1.280	-0.947	1.860	-1.960	1.731	1.789	-1.350	1.860
-1.960	0.343							
936K	0.553	-0.252	1.412	-2.147	1.358	1.191	-0.735	1.412
-2.147	0.197							
937A	-0.521	-0.544	1.001	-2.261	1.002	0.597	-0.274	1.002

967L	-1.773	-1.135	-0.065	-0.652	0.811	0.086	1.428	1.428
-1.773	-0.186							
968V	-1.192	-0.677	-0.439	-1.288	1.121	0.141	-0.017	1.121
-1.288	-0.336							
969I	-0.958	-0.326	-0.775	-1.484	1.449	0.199	-1.616	1.449
-1.616	-0.501							

TOP

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 VTEPTAGPGVPRADDADSDVPRYRYTAELAARLERTWQENWARLGTNFVPNPVGSLAPPDGAAVPDDKLFVQDMFPYPSGEGLHVGHPLGYIATDVYARYFRMVGRNVLHALGFDAFGLPAEQYAVQTGTHPRTRTEANVVNFRRQLGRLGFHDSRRSFSTTDVDFYRWTQWIFLQIYNNAWFDTTANKARPISELVAEFGARCLDGGRDWAKLTAGERADVIDEYRLVYRADSLVNWCPLGTVLANEEVTADGRSDRGNFPVFRKRLRQWMMRITAYADRLLDDLDVLDWPEQVKTMQRNWIGRSTGAVALFSARAASDDGFEVDIEVFTTRPDTLFGATYLVLAPEHDLVDELVAASWPAGVNPLWTYGGGTPGEAIAAYRRAIAAKSDLERQESREKTGVFLGSYAINPANGEPVPIFIADYVLAGYGTGAIMAVPGHDQRDWDFARAFGLPIVEVIAGGNISESAYTGDGILVNSDYLNGMSVPAAKRAIVDRLESAGRGRARIEFKLRDWLFARQRYWGEFPPIVYDSDGRPHALDEAALPVELPDVPDYSPLVFDPPDDADSEPSPLAKATEWVHVVDLGLKPYSRDTNVMPQWAGSSWYE LRYTDPHNSERFCAKEANEAYWMGPRPAEHGPDDPGGVDLVGGAEHAVLHLLYSRFWHKVLYDLGHVSSREPYRRLVNQGYIQAYAYTDARGSYVPAEQVIERGDRFVYPGPDGEVEVFQEFKGIGKSLKNSVSPDEICDAYGADTLRVYEMSMGPLEASRPWATKD VVGAYRFLQRVWRLVVDEHTGETRVADGVELIDTLRALHRTIVGVSEDFAALRNTATAKLIEYTNHLLTKHRDAVPRAAVEPLVQMLAPLAPHIAEELWLRLGNTTSLAHG PFPKADAAYLVDETVEYPVQVNGKVRGRVVVAADTDEETLKAAVLTDEKVQAFLAGATPRKVIVVAGRLVNLVI ⁹⁶⁹
Hydrophilicity	¹ V <u>T</u> ESPTAG <u>P</u> GGV <u>PR</u> ADDAD <u>S</u> DV <u>P</u> RYRYTAELAARLERTWQENWARLGTNFVPNPVGSLAPPDGAAVPDDKLFVQDMFPYPSGEGLHVGHPLGYIATDVYARYFRMVGRNVLHALGFDAFGLPAEQYAVQTGTH <u>P</u> RTRTEANVVNFRRQLGRLGF <u>G</u> HDSRRSFSTTDVDFYRWTQWIFLQIYNNAWF <u>DT</u> TANKAR <u>P</u> ISELVAEFGARCLDGGRDWAKL <u>T</u> AGER

	<p><u>ADVIDEYRLVYRADSLVNWCPLGLTVLA</u><u>NEEV TADGRSDRGN</u><u>FPVFRKRLRQWM</u> MRITAYADRLLDDLDVLDWPEQVKTMQRNWIGRSTGAVALF<u>SARAASDDGFE</u>VDIE VFTTRPDTLFGATYLVLAPEHDLVDELVAASWPAGVNPLWTY<u>GGGTPGEA</u>IAAYRR AIAAK<u>SDLERQESREKTGV</u>FLGSYAI<u>NPANGE</u>VPIFIADYVLAGYGTGAIMAV<u>PGHD</u> <u>QRDWDFARAFGLPIVEVIAGGNI</u><u>SESAYTGDGILVNSDYLNGMSVPAAKRAIVDRLE</u> <u>SAGRGRARIEFKL RDWL FARQRYWGEFPPIV</u><u>YDSDGRPHAL</u>LDEAALPVELPDVPDY SPVL<u>FDPDDADSEPSPP</u>LA KATEWVHV DLDLG DGL<u>KPYRSRTDN</u>VM P QWAGSSWYE LRY<u>TDPHN SERFC</u>AKENEAYWMGPR<u>PAEHGPDDPGGV</u>DLYVGGAEHAVLHLLYSR FWHKVLYDLGHVSSREP YRRLVNQGYI QAYAYTDARGSYVPAEQVIERGDRFV<u>YPG</u> <u>PDGEVE</u>V F Q EFG KIG KSL<u>KNSVSPDE</u>I CDAYGADTLRVYEMSMGPLEASRPWATKD VVGAYRFLQRVWRLV<u>VDEHTGETRVADG</u>VEL DIDTLRALHRTIVGVSEDFAAL<u>RNN</u> <u>TATAK</u>LIEYT NHLT<u>KKHRDA</u>V P RAAVEPLVQMLAPLAPHIAEELWLRLGNTTSLAHG PFPKADAAYLVDETVEY PVQVNGKVRGRVV<u>VAADTDEETLK</u>AAVL<u>TDEKVQA</u>FLA GATPRKVIVVAGRLVNLVI⁹⁶⁹</p>
Flexibility	<p>¹ VTESPTAGPGGVPRADDADSDVPRYRYTAELAARLERTWQENWARL GTFNVPNPV GSLAPPDGAAVPDDKL FVQDMFPYPSGEGLHV GHPLGYI ATDVYARYFRM VGRNV L HALGFDAFGLPAEQYAVQT<u>GTHPRTRTEANV</u><u>VNFRRQLGR</u><u>LGFGHDSRRSF</u>TTDV DFYRWTQWIFLQIYNAWFDTTANKARPIS E LVAEFESGARCL DGG RDWA KL TAGE R ADVIDEYRLVYRADSLVNWCPLGLTVLAN<u>EEVTADGRSDRGNF</u><u>PVFRKRL</u>RQWM MRITAYADRLLDDLDVLDWPEQVKTMQR<u>NWIGRST</u>GAVALFSARAASDDGFEVDIE VFTTRPDTLFGATYLVLAPEHDLVDELVAASWPAGVNPLWTY<u>GGGTP</u>GEAIAAYRR AIA<u>AKSDLERQESREK</u>TGVFLGSYAINPANGEVPIFIADYVLAGYGTGAIMAV<u>PGHD</u> QRDWDFARAFGLPIVEVIAGGNISESAYTGDGILVNSDYLNGMSVPAAKRAIV<u>DRLE</u> <u>SAGRGRARIEFKL RDWL FARQRYWGEFPPIV</u><u>IVYDSDGR</u>PHALDEAALPVELPDVPDY SPVL<u>FDPDDADSEPSPP</u>LA KATEWVHV DLDLG DGL<u>KPYRSRTDN</u>VM P QWAGSSWYE LRY<u>TDPHN SERFC</u>AKENEAYWMGPR<u>PAEHGPDDPGGV</u>DLYVGGAEHAVLHLLYSR FWHKVLYD<u>LGHVSSREP YRRLVNQGYI QAYAYTDARGSY</u>GSYVPAEQVIERGDRFV<u>YPG</u> PDGEVEVF<u>QEFGKIGKSLKNSVSPDE</u>I CDAYGADTLRVYEMSMGPLEASRPWATKD VVGAYRFLQRVWRLV<u>VDEHTGETRVADG</u>VEL DIDTLRALHRTIVGVSEDFAALRNN TATAK LIEYT<u>NHLTKKHRDA</u>V P RAAVEPLVQMLAPLAPHIAEELWLRLGNTTSLAHG PFPKADAAYLVDETVEY PVQVNGKVRGRVV<u>VAADTDEETLK</u>AAVL<u>TDEKVQA</u>FLA GATPRKVIVVAGRLVNLVI⁹⁶⁹</p>
Accessibility	<p>¹ VTESPTAGPGGV<u>PRADDADSDVPRYRYTAELAARLERTWQENWARL</u> GTFNVPNPV GSLAPPDGAAVPDDKL FV<u>QDMFPYPSGEGLHV GHPLGYI</u>ATDVYARYFRM VGRNV L HALGFDAFGLPA<u>EQYAVQTGTHPRTRTEANV</u><u>VNFRRQLGR</u><u>LGFGHDSRRSF</u>TTDV DFYRWTQWIFLQIYNAWFDTTANKARPIS E LVAEFESGARCL DGG RDWA KL TAGE R ADVIDEYRLVYRADSLVNWCPLGLTVLA<u>NEEV TADGRSDRGNF</u><u>PVFRKRL</u>RQWM MRITAYADRLLDDLDV<u>DWPEQVKTMQR</u>WIGRSTGAVALFSARAASDDGFEVDIE VFTTRPDTLFGATYLVLAPEHDLVDELVAASWPAGVNPLWTY<u>GGGTP</u>GEAIAAYRR AIA<u>AKSDLERQESREKTGV</u>FLGSYAI<u>NPANGE</u>VPIFIADYVLAGYGTGAIMAV<u>PGHD</u> <u>QRDWDFAR</u>AFGLPIVEVIAGGNISESAYTGDGILVNSDYLNGMSVPAA<u>KRAIVDRLE</u> <u>SAGRGRARIEFKL RDWL FARQRYWGEFPPIV</u><u>YDSDGRPHAL</u>DEAALPVEL<u>PDV</u>PDY SPVL<u>FDPDDADSEPSPP</u>LA KATEWVHV DLDLG<u>DGLKPYRSRTDN</u>VM P QWAGSSWYE LRY<u>TDPHN SERFC</u>AKENEAYWMGPR<u>PAEHGPDDPGGV</u>DLYVGGAEHAVLHLLYSR FWHKVLYDLG<u>HVSSREP YRRLVNQGYI QAYAYTDARGSY</u>VPAEQVIERGDRFV<u>YPG</u> PDGEVEVF<u>QEFGKIGKSLKNSVSPDE</u>I CDAYGADTLRVYEMSMGPLEASRPWATKD</p>

	VVG <u>AYRFLQR</u> VWRLVV <u>DEHTGETR</u> VADGVELDIDLALHRTIVGVSEDFAA <u>LRNN</u> <u>TATAKLIETNHLTKKHRAVPR</u> AAVEPLVQMLAPLAPHIAEELWLRLGNTTSLAHG PFPKADAAYLV <u>DETVEYPVQVNNGKVRGRVVVAADTDEETLKAAVL</u> <u>LTDEKVQAFLA</u> <u>GATPRKVIVVAGRLVNLVI</u> ⁹⁶⁹
Turns	¹ VTESPTAGPGGVPRADDADSDVPRYRYTAELAARLERTWQENWARLGTNFVPNPV GSLAPPDGAAVPDDKLFWQDMFPYPSGEGLHVGHPLGYIATDVYARYFRMVGRNVL HALGFDAFGLPAEQYAVQTGTHPRTRTEANVNFRRQLGRLGFHDSRRSFSTTDV DFYRWTQWIFLQIYNAWFDTTANKARPISELVAEFESGARCLDGGRDWAKLTAGER ADVIDEYRLVYRADSLVNWCPLGLTVLANEEVTADGRSDRGNFPVFRKRLRQWM MRITAYADRLLDDLDVLDWPEQVKTMQRNWIGRSTGAVALFSARAASDDGFEVDIE VFTTRPDTLFGATYLVLAPEHDLVDELVAASWPAGVNPLWTYGGGTPGEAIAAYRR AIAAKSDLERQESREKTGVFLGSYAINPANGEPVPIFIADYVLAGYGTGAIMAVPGHD QRDWDFARAFGLPIVEVIAGGNISESAYTGDGILVNSDYLNGMSVPAAKRAIVDRLE SAGRGRARIEFKLRDWLFARQRYWGEFPPIVYDSDGRPHALDEAALPVELPDVPDY SPVLFDPPDDADSEPSPLAKATEWVHVLDLGDKPYSRDTNVMPQWAGSSWYE <u>LRYTDPHN</u> SERFCAKE <u>NEAYWMGPRPAEHGPDDPGGV</u> DLYVGGAEHAVLHLLYSR FWHKVLYDLGHVSSREP <u>YRRLVNQGYI</u> QAYAYTDARGSYVPAEQVIERGDRFVYPG PDGEVEVFQEFGKIGKSLKNSVSPDEICDAYGADTLRVYEMSMGPLEASRPWATKD VVGAYRFLQRVWRLVVDEHTGETRVADGVELDIDLALHRTIVGVSEDFAALRNN TATAKLIETNHLTKKHRAVPR <u>A</u> AAVEPLVQMLAPLAPHIAEELWLRLGNTTSLAHG PFPKADAAYLV <u>DETVEYPVQVNNGKVRGRVVVAADTDEETLKAAVL</u> <u>TDEKVQAFLA</u> <u>GATPRKVIVVAGRLVNLVI</u> ⁹⁶⁹
Exposed Surface	¹ VTESPTAGPGGVPRADDADSDVPRYRYTAELAARLERTWQENWARLGTNFVPNPV GSLAPPDGAAVPDDKLFWQDMFPYPSGEGLHVGHPLGYIATDVYARYFRMVGRNVL HALGFDAFGLPAEQYAVQTGTHPRTRTEANVNFRRQLGRLGFHDSRRSFSTTDV DFYRWTQWIFLQIYNAWFDTTANKARPISELVAEFESGARCLDGGRDWAKLTAGER ADVIDEYRLVYRADSLVNWCPLGLTVLANEEVTADGRSDRGNFPV <u>FRKRLRQWM</u> MRITAYADRLLDDLDVLDWPEQVKTMQRNWIGRSTGAVALFSARAASDDGFEVDIE VFTTRPDTLFGATYLVLAPEHDLVDELVAASWPAGVNPLWTYGGGTPGEAIAAYRR AIAA <u>KSDLERQESREKTGV</u> FLGSYAINPANGEPVPIFIADYVLAGYGTGAIMAVPGHD QRDWDFARAFGLPIVEVIAGGNISESAYTGDGILVNSDYLNGMSVPAAKRAIVDRLE SAGRGRARIEFKLRDWLFARQRYWGEFPPIVYDSDGRPHALDEAALPVELPDVPDY SPVLFDPPDDADSEPSPLAKATEWVHVLDLGDKPYSRDTNVMPQWAGSSWYE <u>LRYTDPHN</u> SERFCAKE <u>NEAYWMGPRPAEHGPDDPGGV</u> DLYVGGAEHAVLHLLYSR FWHKVLYDLGHV <u>SSREP</u> <u>YRRLVNQGYI</u> QAYAYTDARGSYVPAEQVIERGDRFVYPG PDGEVEVFQEFGKIGKSLKNSVSPDEICDAYGADTLRVYEMSMGPLEASRPWATKD VVGAYRFLQRVWRLVVDEHTGETRVADGVELDIDLALHRTIVGVSEDFAALRNN TATAKLIETN <u>HLTKKHRAV</u> PRAAAVEPLVQMLAPLAPHIAEELWLRLGNTTSLAHG PFPKADAAYLV <u>DETVEYPVQVNNGKVRGRVVVAADTDEETLKAAVL</u> <u>TDEKVQAFLA</u> <u>GATPRKVIVVAGRLVNLVI</u> ⁹⁶⁹
Polarity	¹ VTESPTAGPGGV <u>PRADDADSDV</u> <u>PRYRYTAELAARLERTWQEN</u> WARLGTNFVPNPV GSLAPPDGAAVPDDKLFWQDMFPYPSGEGLHVGHPLGYIATDVYARYFRMVGRNVL HALGFDAFGLPAEQYAVQT <u>GTHPRTRTEAN</u> VNFRRQLGRLGFHDSRRSFSTTDV DFYRWTQWIFLQIYNAWFDTTAN <u>KARPISELVA</u> EFESGARCLDGGRDWAKLTAGER <u>ADVIDEYRLVYR</u> ADSLVNWCPLGLTVLANEEVT <u>ADGRSDRGNFPVFRKRLRQWM</u>

	MRITAYADRLLDDLDVLDWPEQVKTMQRNWIGRSTGAVALFSARAAS <u>DDGFEVDIE</u> <u>VFTTRPDTLFGATYLVLAPEHDLVDEL</u> VAASWPAGVNPLWTYGGGTPGEAIAAYRR <u>AIAAKSDLERQESREKTGVFLGSYAINPANGE</u> PVPIFIADYVLAGYGTGAIMAV <u>PGHD</u> <u>QRDWDFAR</u> AFGLPIVEVIAGGNISESAYTGDGILVNSDYLNGMSVPAA <u>KRAIVDRLE</u> <u>SAGRGRARIEFKLRDWLFARQRYWGE</u> PFPIVY <u>DSDGRPHALDEAALPVELPDV</u> PDY SPVL <u>FDPDDADSEP</u> SPPLA <u>KATEWVH</u> VDDLGDGLKPYSRDTNVMPQWAGSSWYE LRYTDPHNSERFCAKEANEAYWM <u>GPRPAEHGPDDP</u> GGVDLYVGG <u>AEHAVLHLLYSR</u> <u>FWHKVLYDLGHVSSREPYRRLV</u> NQGYIQAYAYTDARGSYVP <u>AEQVIERGDRFV</u> YPG PDG <u>EVEVFQE</u> FGKIGKSLKNSVSPDEICDAYGADTLRVYEMSMGPLEASRPWATKD VVGAYRFLQRVVW <u>RLVVDEHTGETRV</u> ADGVELDID <u>TLRALHRTIVGV</u> SEDFAALRNN TATAKLIET <u>TNHLTKKHRDAVPR</u> AAVEPLVQMLAPL <u>APHIAEELWLRL</u> GNTTSLAHG PFPKADAAYLVDETVEYPVQV <u>NGKVRGRVV</u> VAAADTDEETLKAAVLTDEKVQAFLA GATPRKVIVVAGRLVNLVI ⁹⁶⁹
Antigenic Propensity	¹ VTESPTAGPGGVPRADDADSDVPRYRYTAELAARLERTWQENWARLGTNFNPV GSLAPPDGAAVPDDKLVQDMFPYPSGE <u>GLHVGHPLGY</u> IATDVYARYFRM <u>VGRNVL</u> <u>HALGFDAFGLPAEQYAVQTGTHPRTRTEANVVNFRRQLGRLGFHDSRRSFSTTDV</u> DFYRWTQ <u>WIFLQIY</u> NAWFDTTANKARPISELVAEFESGARCLDGGRDWAKLTAGER ADVIDEYRLVYRAD <u>SLVNWCPLGLTVL</u> ANEEVTADGRSDRGNFPVFRKRLRQWM MRITAYADR <u>LLDDLDVLD</u> WPEQVKTMQRNWIGRSTGAVALFSARAASDDGFE <u>VDIE</u> <u>VFT</u> TRPDTLFGATYLVLAPEH <u>DLVDEL</u> VAASWPAGVNPLWTYGGGTPGEAIAAYRR AIAAKSDLERQESREKT <u>GVFLGSY</u> AINPANG <u>EPVPIFI</u> ADYVLAGYGTGAIMAVPGHD QRDWDFARA <u>FGLPIVEI</u> AGGNISESAYTGDGIL <u>LVNSDYL</u> NGMSVPAAKRAIVDRLE SAGRGRARIEFKLRDWLFARQRYWG <u>EPFPIVYDS</u> DSDGRPHALDEAAL <u>LPVELPDV</u> PDY <u>SPVLFD</u> DDADSEPSPLAKAT <u>EWVHVLDL</u> GDGLKPYSRDTNVMPQWAGSSWYE LRYTDPHNSERFCAKEANEAYWM <u>GPRPAEHGPDDP</u> GGVDLYVGG <u>AEHAVLHLLYSR</u> <u>FWHKVLYDLGHVS</u> SREPYRR <u>LVNQGYI</u> QAYAYTDARGSYVPAAEQVIERGDRFVYPG PDGE <u>VEVFQE</u> FGKIGKSLKNS <u>VSPDEICDAY</u> GADTLRVYEMSMGPLEASRPWATKD VVGAYRFLQRVVW <u>RLVVDEHTGETRV</u> ADGVELDIDTLRAL <u>HRTIVGV</u> SEDFAALRNN TATAKLIET <u>TNHLTKKHRDAVPR</u> AAVEPLVQMLAPLAPHIAEELWLRLGNTTSLAHG PFPKADAAYLVDETVEYPVQV <u>NGKVRGRVV</u> VAAADTDEETLKAAVLTDEKVQAFLA GA <u>TPRKVIVVAGRLVNLVI</u> ⁹⁶⁹

[TOP](#)

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