

BcePred Prediction Server

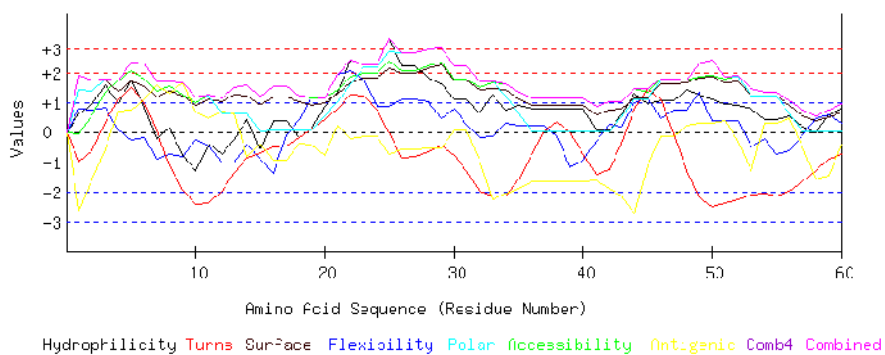
The server displays 1. [GRAPHICAL RESULT](#) 2. [TABULAR RESULT](#) 3. [Overlap Display](#)

seqname=
Seq= MTEHTDFELLELATPYALNAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSA
ATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTV
AEQVLTAPDVRTVSRPLGAGTATVVFSDRNTGLLVMMNVAPP SRGTVYQMWLLGGAKGP
RSAGTMGTAAVTPSTTATLTDLGASTALFTVEPGTGSPQPTGTILAEPLG

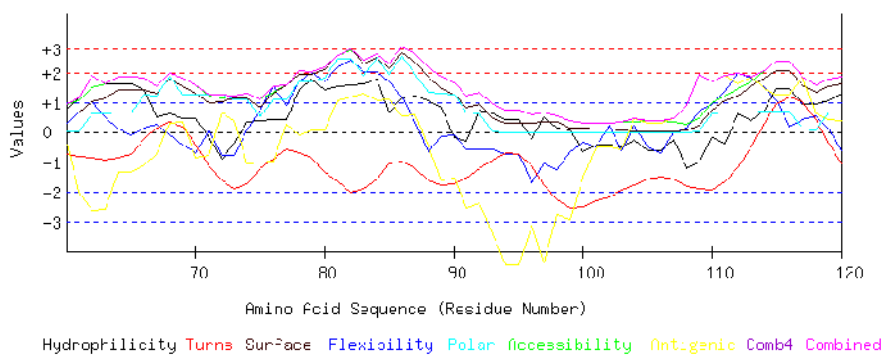
Length=232

GRAPHICAL RESULT

GRAPHICAL RESULT :: SEQ 1 to 60



GRAPHICAL RESULT :: SEQ 61 to 120



5	T	1.748	-0.252	2.057	1.495	1.686	2.322	0.720	2.322	-0.252	1.397
6	D	0.838	-0.168	1.776	0.967	1.540	2.307	1.114	2.307	-0.168	1.196
7	F	-0.237	-0.911	1.365	0.079	1.185	1.713	1.576	1.713	-0.911	0.681
8	E	0.123	-0.737	1.533	-1.153	1.376	1.693	1.282	1.693	-1.153	0.588
9	L	-0.787	-0.821	1.253	-1.937	1.230	1.678	1.676	1.678	-1.937	0.328
10	L	-1.286	-0.258	0.982	-2.418	0.911	1.189	0.675	1.189	-2.418	-0.029
11	E	-0.376	-0.460	1.244	-2.382	1.112	1.205	0.465	1.244	-2.382	0.115
12	L	-0.736	-1.035	1.160	-1.963	1.020	0.625	0.711	1.160	-1.963	-0.031
13	A	-0.275	-1.035	1.496	-1.377	1.248	0.638	0.608	1.496	-1.377	0.186
14	T	0.440	-0.426	1.580	-0.850	1.239	0.633	-0.837	1.580	-0.850	0.254
15	P	-0.635	-0.917	1.169	-0.701	0.884	0.039	-0.376	1.169	-0.917	-0.077
16	Y	0.389	-1.372	1.552	-0.517	1.185	0.074	-0.924	1.552	-1.372	0.055
17	A	0.389	-0.110	1.552	-0.476	1.185	0.074	-0.924	1.552	-0.924	0.242
18	L	-0.174	0.429	1.234	-0.189	1.020	0.056	-0.376	1.234	-0.376	0.286
19	N	0.104	1.173	1.141	0.116	0.902	0.057	-0.436	1.173	-0.436	0.437
20	A	0.857	1.139	1.160	0.428	0.984	0.526	-0.777	1.160	-0.777	0.617
21	V	1.356	1.952	1.431	0.796	1.303	1.015	0.224	1.952	0.224	1.154
22	S	2.431	2.048	1.842	1.253	1.658	1.609	-0.237	2.431	-0.237	1.515
23	D	2.254	1.732	1.973	1.236	1.823	2.194	-0.125	2.254	-0.125	1.584
24	D	2.254	0.868	1.973	0.740	1.823	2.194	-0.125	2.254	-0.125	1.389
25	E	3.120	0.868	2.365	-0.018	2.151	2.681	-0.723	3.120	-0.723	1.492
26	R	2.203	1.107	2.075	-0.877	1.977	2.663	-0.561	2.663	-0.877	1.227
27	A	2.203	1.107	2.075	-0.814	1.977	2.663	-0.561	2.663	-0.814	1.236
28	D	1.837	1.010	2.234	-0.672	2.132	2.798	-0.553	2.798	-0.672	1.255
29	I	1.609	0.471	2.337	-0.451	2.242	2.824	-0.527	2.824	-0.527	1.215
30	D	1.110	0.796	1.786	-0.782	1.759	2.200	0.062	2.200	-0.782	0.990
31	R	1.110	0.257	1.786	-1.284	1.759	2.200	0.062	2.200	-1.284	0.841
32	R	0.610	-0.198	1.515	-1.972	1.440	1.711	-0.939	1.711	-1.972	0.310
33	V	1.249	-0.156	1.655	-2.152	1.458	1.710	-2.270	1.710	-2.270	0.213
34	A	0.749	0.299	1.627	-2.154	1.412	1.240	-2.042	1.627	-2.154	0.162
35	A	0.895	0.203	1.346	-1.573	1.093	0.635	-1.881	1.346	-1.881	0.103
36	A	0.762	0.203	1.160	-0.742	0.893	0.029	-1.660	1.160	-1.660	0.092
37	P	0.762	0.203	1.160	0.004	0.893	0.029	-1.660	1.160	-1.660	0.199
38	S	0.762	-0.156	1.160	0.337	0.893	0.029	-1.660	1.160	-1.660	0.195
39	P	0.762	-1.186	1.160	-0.007	0.893	0.029	-1.660	1.160	-1.660	-0.001
40	V	0.762	-0.935	1.160	-0.755	0.893	0.029	-1.660	1.160	-1.660	-0.072
41	A	0.048	-0.300	0.851	-1.414	0.574	0.015	-1.629	0.851	-1.629	-0.265
42	A	0.079	0.275	1.001	-1.270	0.729	0.035	-1.902	1.001	-1.902	-0.150
43	A	0.579	0.179	1.029	-0.381	0.774	0.505	-2.131	1.029	-2.131	0.079
44	F	1.306	0.992	1.477	0.800	1.148	1.103	-2.746	1.477	-2.746	0.583
45	N	0.939	1.167	1.356	1.474	1.139	1.105	-1.147	1.474	-1.147	0.862
46	D	1.072	0.461	1.786	1.107	1.613	1.729	-0.138	1.786	-0.138	1.090
47	E	1.072	0.736	1.786	-0.042	1.613	1.729	-0.138	1.786	-0.138	0.965
48	V	1.420	0.736	1.730	-1.281	1.649	1.727	0.200	1.730	-1.281	0.883
49	R	1.242	1.323	1.860	-2.165	1.813	2.311	0.312	2.311	-2.165	0.957
50	A	1.103	0.395	1.917	-2.492	1.859	2.421	0.295	2.421	-2.492	0.785
51	V	0.939	0.395	1.786	-2.360	1.649	1.842	0.362	1.842	-2.360	0.659
52	R	0.907	0.395	1.898	-2.277	1.704	1.857	-0.282	1.898	-2.277	0.600
53	E	0.775	-0.514	1.468	-2.099	1.230	1.233	-1.291	1.468	-2.099	0.114
54	T	0.408	-0.234	1.346	-2.076	1.221	1.234	0.308	1.346	-2.076	0.315
55	M	0.408	-0.725	1.346	-2.134	1.221	1.234	0.308	1.346	-2.134	0.237
56	A	0.553	-0.611	1.066	-1.980	0.902	0.630	0.468	1.066	-1.980	0.147
57	V	0.193	-0.120	0.739	-1.679	0.537	0.030	-0.516	0.739	-1.679	-0.116
58	V	-0.003	0.467	0.543	-1.286	0.382	0.010	-1.566	0.543	-1.566	-0.207
59	S	0.591	0.564	0.748	-0.901	0.492	0.013	-1.470	0.748	-1.470	0.005
60	A	0.787	0.283	0.945	-0.758	0.647	0.033	-0.420	0.945	-0.758	0.217
61	A	1.154	0.642	1.066	-0.803	0.656	0.031	-2.018	1.154	-2.018	0.104
62	T	1.881	1.000	1.515	-0.841	1.030	0.629	-2.634	1.881	-2.634	0.369
63	T	1.603	0.509	1.608	-0.946	1.148	0.628	-2.573	1.608	-2.573	0.283
64	A	1.603	0.109	1.851	-0.868	1.422	0.647	-1.343	1.851	-1.343	0.489
65	E	1.603	-0.096	1.851	-0.749	1.422	0.647	-1.343	1.851	-1.343	0.476
66	P	1.407	0.143	1.814	-0.222	1.440	1.247	-1.117	1.814	-1.117	0.673
67	P	0.496	0.275	1.533	0.143	1.294	1.233	-0.723	1.533	-0.723	0.607
68	A	0.629	-0.084	1.963	0.360	1.768	1.857	0.287	1.963	-0.084	0.969
69	H	0.465	-0.408	1.832	0.164	1.558	1.278	0.354	1.832	-0.408	0.749
70	L	0.465	-0.703	1.589	-0.405	1.285	1.259	-0.876	1.589	-0.876	0.373
71	R	-0.174	0.041	1.206	-1.139	0.993	1.242	-0.775	1.242	-1.139	0.199
72	T	-0.888	-0.773	1.122	-1.626	1.002	1.247	0.671	1.247	-1.626	0.108
73	A	-0.389	-0.773	1.234	-1.894	1.148	1.116	0.394	1.234	-1.894	0.120
74	I	0.326	0.059	1.318	-1.760	1.139	1.111	-1.051	1.318	-1.760	0.163
75	L	0.389	0.742	1.085	-1.218	0.820	0.506	-1.009	1.085	-1.218	0.188
76	D	0.421	1.521	1.337	-0.881	1.303	1.081	-0.985	1.521	-0.985	0.542
77	A	0.421	0.886	1.580	-0.561	1.576	1.100	0.245	1.580	-0.561	0.750
78	T	1.420	1.700	2.047	-0.698	1.959	1.697	-0.103	2.047	-0.698	1.146
79	K	1.767	2.022	2.010	-0.871	1.941	1.693	0.051	2.022	-0.871	1.231
80	P	1.401	1.682	2.169	-1.332	2.096	1.829	0.059	2.169	-1.332	1.129
81	E	1.533	2.178	2.599	-1.658	2.570	2.454	1.068	2.599	-1.658	1.535
82	V	1.584	2.417	2.730	-2.039	2.789	2.476	1.191	2.789	-2.039	1.592
83	R	1.634	2.010	2.431	-1.890	2.306	1.902	1.286	2.431	-1.890	1.383
84	R	1.767	2.010	2.617	-1.589	2.506	2.507	1.065	2.617	-1.589	1.555
85	Q	0.642	1.688	2.309	-1.055	2.160	1.933	1.113	2.309	-1.055	1.256
86	S	1.141	1.197	2.860	-0.977	2.643	2.556	0.524	2.860	-0.977	1.421
87	R	1.205	0.341	2.627	-1.200	2.324	1.951	0.565	2.627	-1.200	1.116
88	W	1.072	-0.647	2.197	-1.613	1.850	1.327	-0.444	2.197	-1.613	0.534
89	R	0.825	-0.144	1.870	-1.776	1.476	1.284	-1.617	1.870	-1.776	0.274
90	T	-0.167	-0.102	1.655	-1.729	1.276	1.269	-1.526	1.655	-1.729	0.096
91	A	-0.300	-0.593	1.225	-1.524	0.802	0.644	-2.535	1.225	-2.535	-0.326
92	A	0.743	-0.593	1.356	-1.204	0.938	0.639	-2.398	1.356	-2.398	-0.074
93	F	0.610	-0.593	0.926	-0.928	0.465	0.014	-3.407	0.926	-3.407	-0.416
94	A	0.414	-0.743	0.730	-0.709	0.310	-0.006	-4.458	0.730	-4.458	-0.637
95	S	0.414	-0.743	0.730	-0.744	0.310	-0.006	-4.458	0.730	-4.458	-0.642
96	A	-0.224	-1.695	0.589	-1.140	0.291	-0.004	-3.126	0.589	-3.126	-0.758
97	A	0.490	-1.067	0.655	-1.795	0.337	-0.008	-4.387	0.655	-4.387	-0.825
98	A	0.123	-1.272	0.533	-2.262	0.328	-0.007	-2.788	0.533	-2.788	-0.763
99	I	0.073	-0.645	0.375	-2.555	0.127	-0.027	-2.948	0.375	-2.948	-0.800
100A		-0.642	-0.320	0.290	-2.495	0.136	-0.021	-1.503	0.290	-2.495	-0.651
101V		-0.414	-0.494	0.281	-2.304	0.091	-0.021	-0.492	0.281	-2.304	-0.479
102G		-0.414	0.229	0.281	-2.178	0.091	-0.021	-0.492	0.281	-2.178	-0.358
103L		-0.490	-0.603	0.356	-1.945	0.063	-0.019	-0.563	0.356	-1.945	-0.457
104G		-0.262	0.229	0.346	-1.779	0.018	-0.019	0.447	0.346	-1.779	-0.146
105A		-0.610	-0.494	0.384	-1.584	0.036	-0.015	0.294	0.384	-1.584	-0.284
106F		-0.610	-0.699	0.384	-1.511	0.036	-0.015	0.294	0.384	-1.511	-0.303
107G		-0.262	-0.034	0.346	-1.585	0.018	-0.019	0.447	0.346	-1.585	-0.155
108L		-1.204	0.153	0.272	-1.825	0.073	-0.014	0.882	0.272	-1.825	-0.238
109G		-1.008	0.716	0.468	-1.891	0.228	0.006	1.933	1.933	-1.891	0.064

110V	-0.161	0.944	0.963	-1.955	0.747	0.627	1.681	1.681	-1.955	0.407
111L	-0.389	1.399	1.216	-1.679	1.066	0.646	1.901	1.901	-1.679	0.594
112T	0.604	1.962	1.449	-1.178	1.212	0.660	1.625	1.962	-1.178	0.905
113R	0.376	1.830	1.702	-0.486	1.531	0.679	1.845	1.845	-0.486	1.068
114P	0.743	1.507	2.066	0.287	1.813	0.697	1.476	2.066	0.287	1.227
115S	1.457	1.052	2.393	0.954	2.078	0.710	1.261	2.393	0.710	1.415
116P	1.457	0.197	2.393	1.168	2.078	0.710	1.261	2.393	0.197	1.324
117F	0.958	0.413	1.842	1.014	1.595	0.087	1.851	1.851	0.087	1.109
118P	0.958	0.546	1.599	0.439	1.321	0.068	0.621	1.599	0.068	0.793
119T	1.040	0.091	1.776	-0.375	1.531	0.648	0.435	1.776	-0.375	0.735
120V	1.287	-0.605	1.860	-1.040	1.631	0.671	0.378	1.860	-1.040	0.598
121A	0.920	-0.017	1.496	-1.535	1.349	0.654	0.747	1.496	-1.535	0.516
122E	0.206	-0.017	1.169	-1.726	1.084	0.640	0.962	1.169	-1.726	0.331
123Q	0.206	-0.234	1.169	-1.682	1.084	0.640	0.962	1.169	-1.682	0.306
124V	0.572	-0.186	1.290	-1.607	1.093	0.639	-0.637	1.290	-1.607	0.166
125L	0.572	-0.186	1.533	-1.392	1.367	0.658	0.593	1.533	-1.392	0.449
126T	0.711	0.832	1.477	-0.893	1.321	0.547	0.610	1.477	-0.893	0.658
127A	0.098	0.832	1.029	-0.393	0.938	0.506	1.036	1.036	-0.393	0.578
128P	0.598	0.736	1.580	-0.025	1.422	1.129	0.446	1.580	-0.025	0.841
129D	1.508	1.233	1.860	-0.008	1.567	1.144	0.052	1.860	-0.008	1.051
130V	0.945	1.507	1.543	-0.525	1.403	1.125	0.600	1.543	-0.525	0.943
131R	1.224	1.962	1.692	-0.776	1.558	1.146	1.770	1.962	-0.776	1.225
132T	1.356	0.944	1.879	-0.939	1.759	1.751	1.549	1.879	-0.939	1.186
133V	0.857	1.080	1.851	-0.699	1.713	1.281	1.778	1.851	-0.699	1.123
134S	0.509	1.177	1.889	-0.575	1.731	1.285	1.624	1.889	-0.575	1.091
135R	0.604	0.948	1.449	-0.585	1.212	0.660	1.625	1.625	-0.585	0.845
136P	0.408	0.626	1.253	-1.009	1.057	0.641	0.575	1.253	-1.009	0.507
137L	1.002	0.267	1.365	-1.297	1.020	0.639	-0.014	1.365	-1.297	0.426
138G	0.920	0.962	1.412	-1.592	1.020	0.639	-0.133	1.412	-1.592	0.461
139A	0.787	0.239	0.982	-1.531	0.547	0.014	-1.142	0.982	-1.531	-0.015
140G	0.983	0.143	0.935	-1.364	0.428	0.015	-1.321	0.983	-1.364	-0.026
141T	1.331	-0.659	0.898	-1.159	0.410	0.011	-1.168	1.331	-1.168	-0.048
142A	0.737	-0.294	0.786	-1.115	0.446	0.013	-0.579	0.786	-1.115	-0.001
143T	0.022	0.519	0.720	-1.173	0.401	0.017	0.682	0.720	-1.173	0.170
144V	0.073	0.568	0.879	-1.040	0.601	0.037	0.842	0.879	-1.040	0.280
145V	0.010	1.477	1.113	-0.916	0.920	0.642	0.800	1.477	-0.916	0.578
146F	0.509	2.182	1.384	-0.318	1.239	1.131	1.801	2.182	-0.318	1.133
147S	0.446	2.848	1.617	-0.102	1.558	1.735	1.760	2.848	-0.102	1.409
148R	1.122	2.619	2.038	0.432	1.877	1.775	1.058	2.619	0.432	1.560
149D	1.685	1.601	2.356	0.702	2.041	1.793	0.510	2.356	0.510	1.527
150R	2.627	0.858	2.412	1.068	2.041	1.789	0.259	2.627	0.259	1.579
151N	1.634	-0.052	2.178	0.901	1.895	1.774	0.534	2.178	-0.052	1.266
152T	0.787	-0.775	1.664	0.438	1.431	1.155	0.970	1.664	-0.775	0.810
153G	-0.079	-0.657	1.272	-0.705	1.103	0.667	1.568	1.568	-0.705	0.453
154L	-0.610	-0.675	0.832	-1.605	0.674	0.060	1.513	1.513	-1.605	0.027
155L	-0.610	-0.566	0.832	-1.921	0.674	0.060	1.513	1.513	-1.921	-0.002
156V	-0.496	-0.362	0.935	-1.134	0.829	0.081	1.360	1.360	-1.134	0.173
157M	-1.090	0.093	0.823	0.176	0.866	0.082	1.948	1.948	-1.090	0.414
158N	-0.376	0.566	0.907	1.322	0.856	0.077	0.503	1.322	-0.376	0.551
159N	0.338	0.812	1.234	1.660	1.121	0.090	0.288	1.660	0.090	0.792
160V	0.705	1.016	1.599	1.119	1.403	0.108	-0.081	1.599	-0.081	0.838
161A	1.382	1.740	1.758	0.651	1.513	0.111	0.134	1.758	0.111	1.041
162P	1.205	2.231	1.889	0.288	1.677	0.695	0.246	2.231	0.246	1.176
163P	1.122	1.776	1.580	0.225	1.321	0.654	0.360	1.776	0.225	1.005
164S	1.685	1.010	1.898	0.096	1.485	0.672	-0.188	1.898	-0.188	0.951
165R	1.318	0.646	1.776	-0.385	1.476	0.674	1.410	1.776	-0.385	0.988
166G	1.065	-0.282	1.786	-0.886	1.440	0.674	1.523	1.786	-0.886	0.760
167T	1.312	-1.412	1.870	-1.070	1.540	0.698	1.466	1.870	-1.412	0.629
168V	0.636	-2.107	1.711	-1.263	1.431	0.695	1.251	1.711	-2.107	0.336
169Y	-0.262	-2.215	1.300	-1.370	0.975	0.096	1.274	1.300	-2.215	-0.029
170Q	-1.204	-1.182	1.225	-1.711	1.030	0.101	1.709	1.709	-1.711	-0.005
171M	-2.115	-1.045	0.945	-2.150	0.884	0.086	2.103	2.103	-2.150	-0.185
172W	-1.520	-0.931	1.057	-2.406	0.847	0.085	1.514	1.514	-2.406	-0.194
173L	-1.040	0.403	0.795	-2.447	0.565	0.065	1.183	1.183	-2.447	-0.068
174L	-1.286	1.235	0.468	-2.338	0.191	0.023	0.009	1.235	-2.338	-0.243
175G	-0.661	1.798	0.926	-2.135	0.784	0.601	0.130	1.798	-2.135	0.206
176G	0.332	1.984	0.898	-2.049	0.720	0.575	0.108	1.984	-2.049	0.367
177A	1.046	2.213	1.225	-1.895	0.984	0.589	-0.107	2.213	-1.895	0.579
178K	1.894	2.213	1.739	-1.755	1.449	1.208	-0.543	2.213	-1.755	0.886
179G	1.944	2.008	1.898	-1.286	1.649	1.228	-0.384	2.008	-1.286	1.008
180P	1.717	1.872	1.907	-0.928	1.695	1.228	-1.394	1.907	-1.394	0.871
181R	1.944	1.399	1.898	-0.616	1.649	1.228	-0.384	1.944	-0.616	1.017
182S	1.913	1.213	1.646	-0.552	1.166	0.654	-0.408	1.913	-0.552	0.804
183A	1.287	0.848	1.646	-0.763	1.257	0.671	-0.464	1.646	-0.763	0.640
184G	1.514	0.848	1.393	-1.153	0.938	0.652	-0.683	1.514	-1.153	0.501
185T	1.578	0.221	1.160	-1.332	0.619	0.047	-0.642	1.578	-1.332	0.236
186M	1.299	-0.366	1.010	-1.568	0.465	0.027	-1.812	1.299	-1.812	-0.135
187G	1.299	0.239	1.010	-1.536	0.465	0.027	-1.812	1.299	-1.812	-0.044
188T	0.705	-0.030	0.898	-1.534	0.501	0.028	-1.223	0.898	-1.534	-0.094
189A	0.705	0.335	0.898	-1.567	0.501	0.028	-1.223	0.898	-1.567	-0.046
190A	1.103	0.826	1.150	-1.400	0.729	0.030	-0.948	1.150	-1.400	0.213
191V	1.154	1.317	1.309	-0.901	0.929	0.050	-0.788	1.317	-0.901	0.439
192T	1.154	1.413	1.309	-0.215	0.929	0.050	-0.788	1.413	-0.788	0.550
193P	1.350	1.413	1.505	0.535	1.084	0.070	0.262	1.505	0.070	0.889
194S	1.350	0.850	1.505	0.751	1.084	0.070	0.262	1.505	0.070	0.839
195T	1.913	0.485	1.823	0.575	1.248	0.089	-0.286	1.913	-0.286	0.835
196T	1.002	0.534	1.543	-0.063	1.103	0.074	0.109	1.543	-0.063	0.614
197A	1.198	-0.162	1.496	-0.538	0.984	0.075	-0.070	1.496	-0.538	0.426
198T	1.420	0.465	1.617	-0.674	1.148	0.544	-0.239	1.617	-0.674	0.612
199L	0.509	-0.026	1.337	-0.511	1.002	0.529	0.155	1.337	-0.511	0.428
200T	0.541	1.034	1.132	-0.274	0.802	0.509	0.114	1.132	-0.274	0.551
201D	0.541	1.034	1.132	-0.294	0.802	0.509	0.114	1.132	-0.294	0.548
202L	0.623	0.495	1.085	-0.490	0.802	0.510	0.234	1.085	-0.490	0.465
203G	1.533	0.495	1.365	-0.624	0.948	0.524	-0.161	1.533	-0.624	0.583
204A	1.337	-0.132	1.169	-0.515	0.793	0.504	-1.212	1.337	-1.212	0.278
205S	0.123	-0.306	0.814	-0.491	0.483	0.021	-0.768	0.814	-0.768	-0.018
206T	0.838	-0.671	0.898	-0.613	0.474	0.015	-2.213	0.898	-2.213	-0.182
207A	-0.104	-1.258	0.842	-1.069	0.474	0.019	-1.962	0.842	-1.962	-0.437
208L	0.092	-0.683	1.038	-1.337	0.629	0.039	-0.911	1.038	-1.337	-0.162
209A	-0.553	-0.120	0.767	-1.396	0.465	0.021	-0.482	0.767	-1.396	-0.185
210F	-0.389	0.507	0.898	-1.221	0.674	0.601	-0.549	0.898	-1.221	0.074
211T	-0.389	1.173	1.141	-1.028	0.948	0.620	0.681	1.173	-1.028	0.449
212V	0.553	1.309	1.216	-1.057	0.893	0.614	0.246	1.309	-1.057	0.539
213E	0.749	2.261	1.412	-0.966	1.048	0.634	1.297	2.261	-0.966	0.919
214P	1.691	2.044	1.468	-0.885	1.048	0.630	1.046	2.044	-0.885	1.006

215G	1.774	<u>2.176</u>	1.421	-0.563	1.048	0.630	1.165	2.176	-0.563	1.093
216T	<u>2.140</u>	1.908	1.786	-0.220	1.330	0.647	0.797	2.140	-0.220	1.198
217G	<u>2.026</u>	1.908	1.786	0.222	1.339	0.090	0.986	2.026	0.090	1.194
218S	<u>2.026</u>	1.908	1.786	0.534	1.339	0.090	0.986	2.026	0.090	1.239
219P	<u>1.995</u>	1.543	<u>1.991</u>	0.770	1.540	0.110	1.027	1.995	0.110	1.282
220Q	<u>2.026</u>	0.860	1.786	0.612	1.339	0.090	0.986	2.026	0.090	1.100
221P	<u>1.995</u>	0.165	<u>1.991</u>	0.399	1.540	0.110	1.027	1.995	0.110	1.032
222T	1.078	-0.194	1.702	-0.171	1.367	0.092	1.188	1.702	-0.194	0.723
223G	0.364	-0.110	1.375	-0.754	1.103	0.078	1.403	1.403	-0.754	0.494
224T	0.117	-0.941	1.047	-1.409	0.729	0.036	0.230	1.047	-1.409	-0.027
225I	0.477	-1.073	1.132	-1.987	0.820	0.617	-0.016	1.132	-1.987	-0.004
226L	-0.433	-0.953	0.851	-2.374	0.674	0.602	0.378	0.851	-2.374	-0.179
227A	-0.661	-0.122	1.103	-2.280	0.993	0.621	0.598	1.103	-2.280	0.036
228E	-1.571	0.133	0.823	-2.125	0.847	0.606	0.992	0.992	-2.125	-0.042
229L	-0.705	-0.188	0.954	-1.724	0.820	0.605	0.671	0.954	-1.724	0.062
230P	-0.123	0.271	0.580	-1.457	1.130	0.659	-0.775	1.130	-1.457	0.041
231L	-0.256	0.167	0.122	-1.252	1.449	0.719	-0.775	1.449	-1.252	0.025
232G	-0.749	0.626	-0.663	-1.081	1.403	0.180	-1.758	1.403	-1.758	-0.292

[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	<u>¹MTEHTDFELLELATPYALNAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTVAEQVLTAPDVRTV</u>
Hydrophilicity	¹ MTEHTDFELLELATPYAL <u>NAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTVAEQVLTAPDVRTV</u>
Flexibility	¹ MTEHTDFELLELATPYAL <u>NAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTVAEQVLTAPDVRTV</u>
Accessibility	¹ <u>MTEHTDFELLELATPYALNAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTVAEQVLTAPDVRTV</u>
Turns	¹ MTEHTDFELLELATPYALNAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTVAEQVLTAPDVRTV
Exposed Surface	¹ MTEHTDFELLELATPYALNAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDAT <u>KPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTVAEQVLTAPDVRTV</u>
Polarity	¹ <u>MTEHTDFELLELATPYALNAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGAFGLGVLTRPSPPTVAEQVLTAPDVRTV</u>
Antigenic Propensity	¹ MTEHTDFELLELATPYALNAVSDDERADIDRRVAAAPSPVAAAFNDEVRAVRETMVVSAATTAEPHAHLRTAILDATKPEVRRQSRWRTAAFASAAAIAVGLGA <u>FGLGVLTRPSPPTVAEQVLTAPDVRTV</u>

[TOP](#)

[Home](#)