

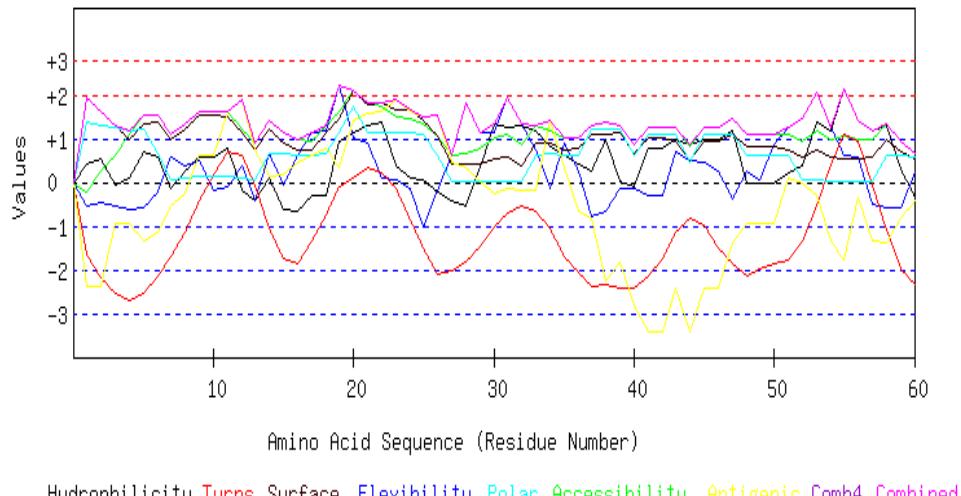
The server displays 1.[GRAPHICAL RESULT](#) 2.[TABULAR RESULT](#) 3.[Overlap Display](#)

seqname=  
Seq=VEEMALAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVNGAGIDDAAVVTCRPDSLADAQQMVEALGRYGRLDGVVASGSNHVAPITEMAVEDFDAVM DANVRGAWLVCRAAGRVLLE QQQGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTAAEWGGHIGRVNALAPTVFRSAVTEWMFTDDPK GRATREAMLARIPLRRFAEPEDFVGALIYLLSDASSFYTGQVMYLDGGYTAC

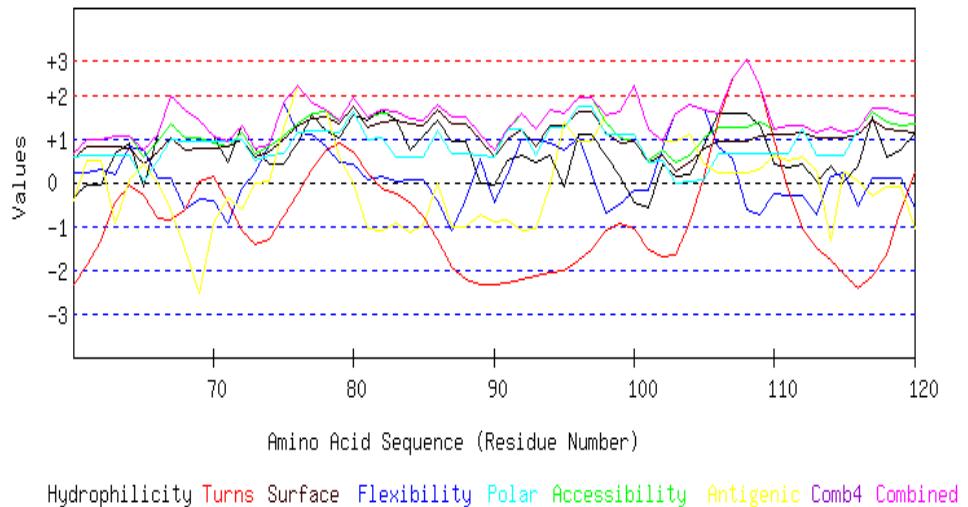
Length=270

#### GRAPHICAL RESULT

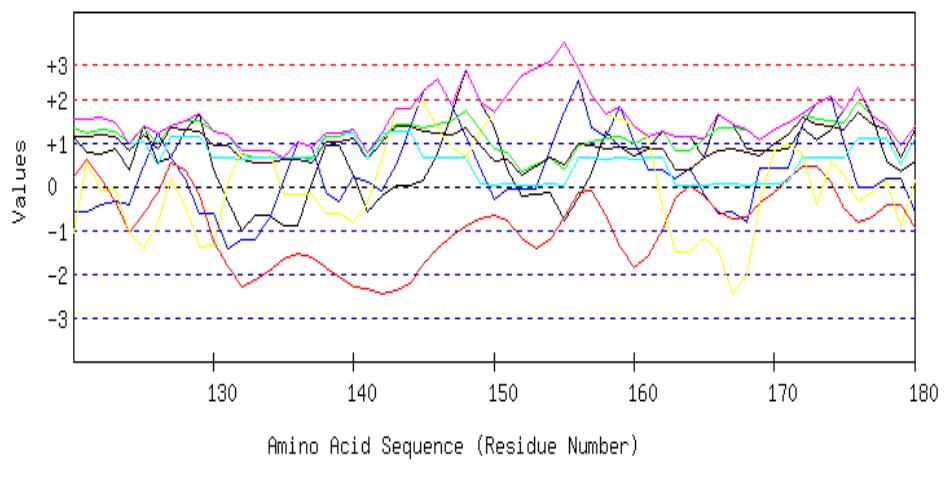
GRAPHICAL RESULT :: SEQ 1 to 60



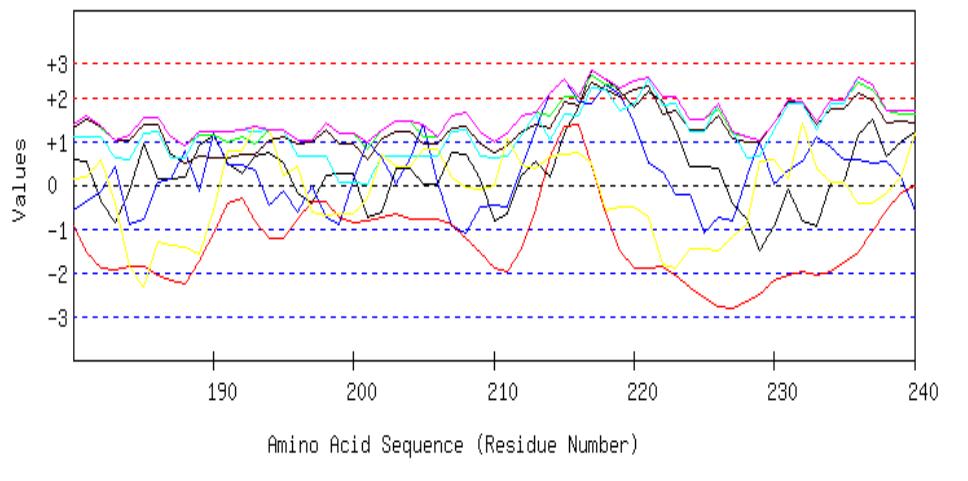
GRAPHICAL RESULT :: SEQ 61 to 120



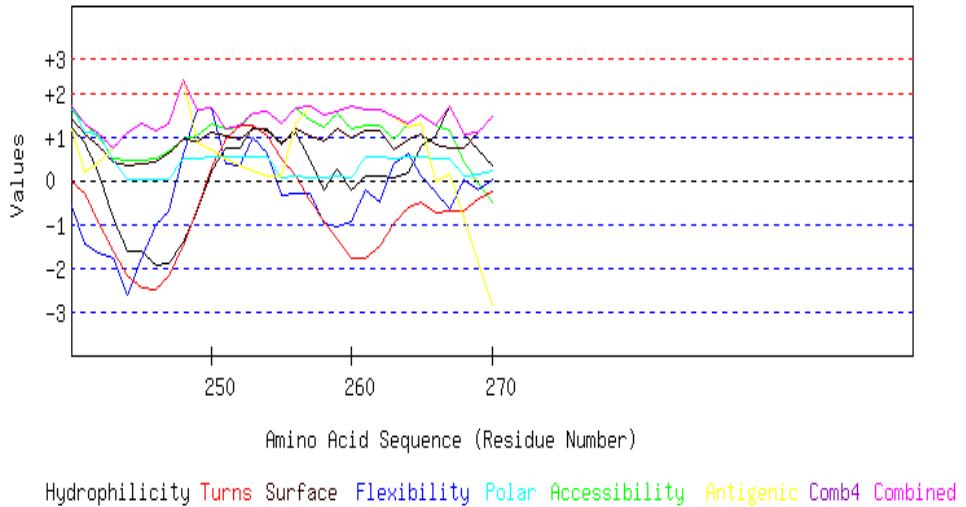
GRAPHICAL RESULT :: SEQ 121 to 180



GRAPHICAL RESULT :: SEQ 181 to 240



GRAPHICAL RESULT :: SEQ 241 to 300



## TOP

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## 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

VEEMALAAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLA  
ELVNGAGIDDAAVVTCRPDSLADAQQMVEAALGRYGRLDGVLVASGSNHVAPITEMAVED  
FDAVM DANVRGAWLVCRAAGRVLLEQQQGGSVLVSSVRGGLGNAAGYSAYCPSKAGTDL  
LAKTLAAEWGGHGIRVNALAPTVFRSAVTEWMFTDDPKGRATREAMLARIPLRRFAEPED  
FVGALIYLLSDASSFYTGQVMYLDGGYTAC

Length=270

A.A.	Parameter						Combined		
	Hydro	Flexi	Access	Turns	Surface	Polar	AntiPro	MAX	MIN

1	V	0.408	-0.538	-0.205	-1.675	1.923	1.368	-2.368	1.923	-2.368	-0.155
2	E	0.541	-0.442	0.253	-2.184	1.604	1.308	-2.368	1.604	-2.368	-0.184
3	E	-0.041	-0.526	0.627	-2.557	1.294	1.253	-0.923	1.294	-2.557	-0.125
4	M	0.092	-0.611	1.085	-2.701	0.975	1.193	-0.923	1.193	-2.701	-0.127
5	A	0.705	-0.593	1.533	-2.534	1.358	1.234	-1.348	1.533	-2.534	0.051
6	L	0.591	-0.234	1.533	-2.133	1.367	0.677	-1.158	1.533	-2.133	0.092
7	A	-0.136	0.580	1.085	-1.689	0.993	0.079	-0.543	1.085	-1.689	0.053
8	Q	0.263	0.375	1.337	-1.141	1.221	0.081	-0.268	1.337	-1.141	0.267
9	Q	0.572	0.511	1.636	-0.430	1.531	0.121	0.629	1.636	-0.430	0.653
10	V	0.572	-0.184	1.636	0.171	1.531	0.121	0.629	1.636	-0.184	0.639
11	P	0.800	-0.088	1.627	0.718	1.485	0.121	1.639	1.639	-0.088	0.900
12	N	-0.161	0.367	1.216	0.621	1.121	0.084	<b>1.911</b>	1.911	-0.161	0.737
13	L	-0.408	-0.416	0.889	-0.054	0.747	0.042	0.738	0.889	-0.416	0.220
14	G	0.092	0.644	1.440	-1.069	1.230	0.665	0.148	1.440	-1.069	0.450
15	L	-0.623	-0.080	1.132	-1.736	0.911	0.650	0.179	1.132	-1.736	0.062
16	A	-0.654	0.616	0.982	-1.868	0.756	0.630	0.452	0.982	-1.868	0.130
17	R	-0.307	1.155	0.945	-1.355	0.738	0.626	0.606	1.155	-1.355	0.344
18	F	-0.288	1.173	1.281	-0.723	1.157	0.668	0.769	1.281	-0.723	0.577
19	S	0.926	<b>2.203</b>	1.636	-0.107	1.467	1.152	0.325	2.203	-0.107	1.086
20	V	1.154	1.022	<b>2.085</b>	0.102	2.105	1.746	1.400	2.105	0.102	1.373
21	Q	1.299	0.914	1.804	0.357	1.786	1.142	1.561	1.804	0.357	1.266
22	D	1.375	0.099	1.730	0.214	1.813	1.139	1.631	1.813	0.099	1.143
23	K	0.383	0.051	1.496	-0.139	1.668	1.125	<b>1.906</b>	1.906	-0.139	0.927
24	S	0.111	-0.154	1.477	-0.845	1.658	1.125	1.639	1.658	-0.845	0.716
25	I	0.060	-1.009	1.346	-1.502	1.440	1.102	1.517	1.517	-1.502	0.422
26	L	-0.212	-0.194	1.066	-2.108	1.075	0.613	1.526	1.526	-2.108	0.253
27	I	-0.439	0.638	0.618	-2.038	0.437	0.019	0.451	0.638	-2.038	-0.045
28	T	-0.521	1.818	0.664	-1.837	0.437	0.019	0.332	1.818	-1.837	0.130
29	G	0.345	1.123	<b>0.795</b>	-1.441	0.410	0.017	0.010	1.123	-1.441	0.180
30	A	1.337	1.123	1.029	-1.022	0.556	0.032	-0.265	1.337	-1.022	0.398
31	T	1.261	1.936	1.085	-0.719	0.583	0.035	-0.151	1.936	-0.719	0.576
32	G	1.293	1.349	0.879	-0.526	0.382	0.015	-0.192	1.349	-0.526	0.457
33	S	1.198	0.722	1.318	-0.676	0.902	0.640	-0.193	1.318	-0.676	0.559
34	L	0.832	-0.134	1.197	-1.066	0.893	0.641	1.406	1.406	-1.066	0.538
35	G	0.636	0.884	1.001	-1.700	0.738	0.622	0.355	1.001	-1.700	0.362
36	R	0.408	0.257	1.010	-2.059	0.784	0.622	-0.655	1.010	-2.059	0.052
37	V	0.263	-0.761	1.290	-2.362	1.103	1.226	-0.816	1.290	-2.362	-0.008
38	A	0.977	-0.665	1.375	-2.354	1.093	1.221	-2.261	1.375	-2.354	-0.088
39	A	0.035	-0.126	1.300	-2.422	1.148	1.226	-1.827	1.300	-2.422	-0.095
40	R	-0.098	-0.126	<b>0.870</b>	-2.431	0.674	0.601	-2.836	0.870	-2.836	-0.478
41	A	0.768	-0.312	1.262	-2.140	1.002	1.089	-3.434	1.262	-3.434	-0.252
42	L	0.768	-0.312	1.262	-1.721	1.002	1.089	-3.434	1.262	-3.434	-0.192
43	A	0.996	0.706	1.253	-1.136	0.957	1.089	-2.423	1.253	-2.423	0.206
44	D	0.863	0.501	0.823	-0.832	0.483	0.464	-3.432	0.863	-3.432	-0.161
45	A	0.996	0.453	1.253	-0.977	0.957	1.089	-2.423	1.253	-2.423	0.193
46	G	0.996	0.249	1.253	-1.496	0.957	1.089	-2.423	1.253	-2.423	0.089
47	A	1.192	-0.378	1.449	-1.847	1.112	1.109	-1.372	1.449	-1.847	0.181
48	R	-0.022	0.249	1.094	-2.135	0.802	0.625	-0.928	1.094	-2.135	-0.045
49	L	-0.022	0.063	1.094	-1.981	0.802	0.625	-0.928	1.094	-1.981	-0.050
50	T	-0.022	0.876	1.094	-1.858	0.802	0.625	-0.928	1.094	-1.858	0.084
51	L	0.206	1.241	1.085	-1.791	0.756	0.625	0.082	1.241	-1.791	0.315
52	A	0.383	1.445	0.954	-1.355	0.592	0.041	-0.030	1.445	-1.355	0.290
53	G	1.375	<b>2.072</b>	1.188	-0.548	0.738	0.056	-0.305	2.072	-0.548	0.654
54	G	1.179	1.241	0.991	0.423	0.583	0.036	-1.356	1.241	-1.356	0.443
55	N	<b>2.121</b>	0.614	1.066	1.120	0.528	0.031	-1.791	2.121	-1.791	0.527
56	S	1.407	0.580	0.982	0.920	0.537	0.036	-0.346	1.407	-0.346	0.588
57	A	1.179	-0.480	0.991	-0.043	0.583	0.036	-1.356	1.179	-1.356	0.130
58	G	1.312	-0.576	1.328	-1.102	0.993	0.635	-1.383	1.328	-1.383	0.173
59	L	0.288	-0.595	0.945	-1.970	0.692	0.600	-0.835	0.945	-1.970	-0.125

60 A	-0.357	0.237	0.674	-2.328	0.528	0.582	-0.406	0.674	-2.328	-0.153
61 E	-0.047	0.237	0.973	-1.907	0.838	0.622	0.491	0.973	-1.907	0.172
62 L	-0.047	0.289	0.973	-1.323	0.838	0.622	0.491	0.973	-1.323	0.263
63 V	0.667	0.169	1.057	-0.469	0.829	0.617	-0.954	1.057	-0.954	0.274
64 N	0.895	0.804	1.047	-0.043	0.784	0.617	0.057	1.047	-0.043	0.594
65 G	-0.104	0.734	0.580	-0.309	0.401	0.019	0.405	0.734	-0.309	0.247
66 A	1.110	0.107	0.935	-0.806	0.711	0.503	-0.039	1.110	-0.806	0.360
67 G	1.976	0.107	1.328	-0.875	1.039	0.990	-0.637	1.976	-0.875	0.561
68 I	1.666	-0.617	1.029	-0.606	0.729	0.949	-1.534	1.666	-1.534	0.231
69 D	1.438	-0.388	1.038	0.036	0.774	0.949	-2.544	1.438	-2.544	0.186
70 D	1.072	-0.436	0.917	0.140	0.765	0.951	-0.946	1.072	-0.946	0.352
71 A	0.477	-0.939	0.804	-0.459	0.802	0.953	-0.357	0.953	-0.939	0.183
72 A	1.312	-0.126	1.141	-1.097	0.975	0.971	-0.638	1.312	-1.097	0.363
73 V	0.768	0.233	0.655	-1.412	0.565	0.500	-0.007	0.768	-1.412	0.186
74 V	0.402	0.868	0.814	-1.304	0.720	0.635	0.001	0.868	-1.304	0.305
75 T	0.402	1.820	1.057	-0.767	0.993	0.654	1.231	1.820	-0.767	0.770
76 C	0.901	1.125	1.328	-0.266	1.312	1.143	2.232	2.232	-0.266	1.111
77 R	1.546	1.089	1.599	0.298	1.476	1.162	1.803	1.803	0.298	1.282
78 P	1.198	0.814	1.636	0.731	1.494	1.166	1.650	1.650	0.731	1.241
79 D	1.002	0.455	1.440	0.894	1.339	1.146	0.599	1.440	0.455	0.982
80 S	1.546	0.407	1.926	0.697	1.750	1.617	-0.032	1.926	-0.032	1.130
81 L	1.413	0.043	1.496	0.230	1.276	0.992	-1.042	1.496	-1.042	0.630
82 A	1.660	0.133	1.580	-0.123	1.376	1.016	-1.098	1.660	-1.098	0.649
83 D	1.407	0.037	1.636	-0.273	1.431	0.569	-0.926	1.636	-0.926	0.554
84 A	0.730	0.073	1.477	-0.458	1.321	0.566	-1.141	1.477	-1.141	0.367
85 Q	1.078	0.073	1.440	-0.775	1.303	0.562	-0.987	1.440	-0.987	0.385
86 Q	1.438	-0.418	1.767	-1.294	1.668	1.162	-0.003	1.767	-1.294	0.617
87 M	0.939	-1.113	1.496	-1.920	1.349	0.673	-1.005	1.496	-1.920	0.060
88 V	0.939	-0.372	1.496	-2.212	1.349	0.673	-1.005	1.496	-2.212	0.124
89 E	-0.022	0.538	1.085	-2.358	0.984	0.636	-0.733	1.085	-2.358	0.019
90 A	-0.041	-0.444	0.748	-2.355	0.565	0.594	-0.896	0.748	-2.355	-0.261
91 A	0.490	0.183	1.188	-2.290	0.993	1.201	-0.842	1.201	-2.290	0.132
92 L	0.604	0.996	1.561	-2.223	1.239	1.219	-1.098	1.561	-2.223	0.328
93 G	0.471	0.996	1.225	-2.137	0.829	0.619	-1.072	1.225	-2.137	0.133
94 R	0.604	0.908	1.655	-2.045	1.303	1.244	-0.062	1.655	-2.045	0.515
95 Y	-0.111	0.722	1.571	-2.028	1.312	1.249	1.383	1.571	-2.028	0.586
96 G	1.103	1.032	1.926	-1.786	1.622	1.733	0.939	1.926	-1.786	0.939
97 R	1.103	0.201	1.926	-1.556	1.622	1.733	0.939	1.926	-1.556	0.852
98 L	0.604	-0.709	1.375	-1.109	1.139	1.110	1.528	1.528	-1.109	0.563
99 D	0.142	-0.504	1.038	-0.922	0.911	1.096	1.631	1.631	-0.922	0.485
100G	-0.452	-0.188	0.926	-1.064	0.948	1.097	2.220	2.220	-1.064	0.498
101V	-0.585	-0.188	0.496	-1.543	0.474	0.473	1.210	1.210	-1.543	0.048
102L	0.408	0.764	0.730	-1.685	0.619	0.487	0.935	0.935	-1.685	0.323
103V	0.136	1.577	0.449	-1.657	0.255	-0.002	0.945	1.577	-1.657	0.243
104A	0.187	1.764	0.608	-0.913	0.455	0.019	1.104	1.764	-0.913	0.461
105S	0.863	1.668	1.029	0.127	0.774	0.058	0.402	1.668	0.058	0.703
106G	1.578	0.812	1.272	1.424	0.938	0.672	0.235	1.578	0.235	0.990
107S	1.578	0.544	1.272	2.394	0.938	0.672	0.235	2.394	0.235	1.090
108N	1.578	-0.637	1.272	2.811	0.938	0.672	0.235	2.811	-0.637	0.981
109H	1.299	-0.755	1.365	2.224	1.057	0.671	0.295	2.224	-0.755	0.879
110V	0.433	-0.270	1.234	0.992	1.084	0.673	0.616	1.234	-0.270	0.680
111A	0.351	-0.288	1.281	-0.194	1.084	0.673	0.497	1.281	-0.288	0.486
112P	0.402	-0.288	1.309	-1.052	1.139	1.232	0.584	1.309	-1.052	0.475
113I	0.003	-0.743	1.141	-1.516	1.011	0.629	0.261	1.141	-1.516	0.112
114T	0.370	0.157	1.262	-1.789	1.020	0.627	-1.338	1.262	-1.789	0.044
115E	0.003	0.205	1.141	-2.085	1.011	0.629	0.261	1.141	-2.085	0.166
116M	0.364	-0.544	1.225	-2.429	1.103	1.209	0.015	1.225	-2.429	0.135
117A	1.502	0.109	1.636	-2.143	1.440	1.697	-0.316	1.697	-2.143	0.561
118V	0.591	0.109	1.375	-1.663	1.239	1.681	-0.105	1.681	-1.663	0.461

119E	0.730	0.109	1.318	-0.618	1.194	1.570	-0.088	1.570	-0.618	0.602
120D	1.129	-0.581	1.328	0.223	1.148	1.553	-1.043	1.553	-1.043	0.537
121F	0.762	-0.581	1.206	0.624	1.139	1.555	0.556	1.555	-0.581	0.752
122D	0.730	-0.406	1.318	0.180	1.194	1.570	-0.088	1.570	-0.406	0.643
123A	0.869	-0.336	1.262	-0.350	1.148	1.460	-0.070	1.460	-0.350	0.569
124V	0.370	-0.432	0.991	-1.058	0.829	0.971	-1.072	0.991	-1.072	0.086
125M	1.394	0.477	1.356	-0.635	1.185	1.007	-1.436	1.394	-1.436	0.478
126D	0.528	1.219	0.963	-0.138	0.856	0.520	-0.838	1.219	-0.838	0.444
127A	0.661	0.680	1.393	0.525	1.330	1.144	0.171	1.393	0.171	0.844
128N	1.255	0.177	1.505	0.400	1.294	1.143	-0.417	1.505	-0.417	0.765
129V	1.653	-0.637	1.515	-0.198	1.248	1.126	-1.372	1.653	-1.372	0.476
130R	0.389	-0.637	1.262	-1.251	0.948	0.662	-1.341	1.262	-1.341	0.005
131G	-0.325	-1.414	1.178	-1.817	0.957	0.667	0.104	1.178	-1.817	-0.093
132A	-1.002	-1.228	0.758	-2.302	0.638	0.628	0.806	0.806	-2.302	-0.243
133W	-0.680	-1.228	0.664	-2.152	0.556	0.644	0.839	0.839	-2.152	-0.194
134L	-0.680	-0.725	0.664	-1.956	0.556	0.644	0.839	0.839	-1.956	-0.094
135V	-0.907	0.107	0.674	-1.669	0.601	0.644	-0.171	0.674	-1.669	-0.103
136C	-0.907	1.016	0.674	-1.548	0.601	0.644	-0.171	1.016	-1.548	0.044
137R	0.085	0.884	0.646	-1.606	0.537	0.619	-0.193	0.884	-1.606	0.139
138A	0.933	-0.134	1.160	-1.879	1.002	1.239	-0.629	1.239	-1.879	0.242
139A	0.933	-0.338	1.160	-2.054	1.002	1.239	-0.629	1.239	-2.054	0.187
140G	0.263	0.237	1.290	-2.286	1.103	1.226	-0.816	1.290	-2.286	0.145
141R	-0.585	0.101	0.776	-2.350	0.638	0.607	-0.380	0.776	-2.350	-0.170
142V	-0.224	-0.086	1.103	-2.464	1.002	1.206	0.603	1.206	-2.464	0.163
143L	0.022	0.501	1.431	-2.392	1.376	1.249	1.777	1.777	-2.392	0.566
144L	0.022	1.333	1.431	-2.222	1.376	1.249	1.777	1.777	-2.222	0.709
145E	0.136	<b>2.164</b>	1.328	-1.795	1.276	0.667	<b>1.941</b>	2.164	-1.795	0.817
146Q	0.730	<b>2.445</b>	1.440	-1.426	1.239	0.665	1.353	2.445	-1.426	0.921
147G	1.672	1.858	1.515	-1.145	1.185	0.660	0.918	1.858	-1.145	0.952
148Q	<b>2.665</b>	1.135	1.748	-0.880	1.330	0.674	0.643	2.665	-0.880	1.045
149G	<b>1.938</b>	0.439	1.300	-0.735	0.957	0.076	1.258	1.938	-0.735	0.748
150G	1.325	-0.284	0.851	-0.649	0.574	0.035	1.683	1.683	-0.649	0.505
151S	0.383	-0.056	0.776	-0.761	0.629	0.041	<b>2.118</b>	2.118	-0.761	0.447
152V	-0.231	-0.056	0.328	-1.163	0.246	0.000	<b>2.543</b>	2.543	-1.163	0.238
153V	-0.180	-0.056	0.487	-1.437	0.446	0.020	<b>2.703</b>	2.703	-1.437	0.283
154L	-0.129	0.854	0.646	-1.216	0.647	0.040	<b>2.862</b>	2.862	-1.216	0.529
155V	-0.774	1.686	0.375	-0.718	0.483	0.022	<b>3.291</b>	3.291	-0.774	0.623
156S	-0.275	<b>2.409</b>	0.926	-0.130	0.966	0.645	<b>2.702</b>	2.702	-0.275	1.035
157S	0.319	1.349	1.038	-0.116	0.929	0.643	<b>2.113</b>	2.113	-0.116	0.897
158V	1.261	1.121	1.113	-0.673	0.875	0.638	1.679	1.679	-0.673	0.859
159R	0.914	1.826	1.150	-1.375	0.893	0.642	1.525	1.826	-1.375	0.796
160G	0.863	1.012	0.991	-1.859	0.692	0.622	1.365	1.365	-1.859	0.527
161G	0.895	0.385	1.141	-1.584	0.847	0.642	1.093	1.141	-1.584	0.488
162L	1.261	0.385	1.262	-0.997	0.856	0.641	-0.506	1.262	-0.997	0.415
163G	1.129	0.183	0.832	-0.252	0.382	0.016	-1.516	1.129	-1.516	0.111
164N	1.129	0.411	0.832	0.017	0.382	0.016	-1.516	1.129	-1.516	0.182
165A	0.648	-0.198	1.094	-0.215	0.665	0.035	-1.184	1.094	-1.184	0.121
166A	1.641	-0.605	1.328	-0.589	0.811	0.050	-1.459	1.641	-1.459	0.168
167G	1.413	-0.568	1.337	-0.748	0.856	0.050	-2.469	1.413	-2.469	-0.018
168Y	0.850	-0.837	1.290	-0.712	0.784	0.029	-2.024	1.290	-2.024	-0.089
169S	0.806	0.425	1.075	-0.362	0.692	0.046	-0.392	1.075	-0.392	0.327
170A	0.806	0.401	1.318	-0.137	0.966	0.065	0.838	1.318	-0.137	0.608
171Y	0.857	0.401	1.477	0.172	1.166	0.085	0.998	1.477	0.085	0.737
172C	1.337	1.435	1.674	0.446	1.567	0.661	0.731	1.674	0.446	1.122
173P	1.059	1.890	1.524	0.451	1.412	0.641	-0.439	1.890	-0.439	0.934
174S	1.287	<b>2.070</b>	1.515	0.083	1.367	0.641	0.571	2.070	0.083	1.076
175K	1.736	1.010	1.459	-0.517	1.285	0.641	0.280	1.736	-0.517	0.842
176A	<b>2.279</b>	-0.026	<b>1.945</b>	-0.826	1.695	1.112	-0.351	2.279	-0.826	0.833
177G	1.565	-0.026	1.617	-0.702	1.431	1.099	-0.136	1.617	-0.702	0.693

178T	0.572	0.179	1.384	-0.439	1.285	1.084	0.139	1.384	-0.439	0.601
179D	0.345	0.179	0.935	-0.426	0.647	0.490	-0.936	0.935	-0.936	0.176
180L	0.572	-0.564	1.384	-0.948	1.285	1.084	0.139	1.384	-0.948	0.422
181L	0.541	-0.360	1.589	-1.530	1.485	1.104	0.179	1.589	-1.530	0.430
182A	-0.370	-0.156	1.309	-1.889	1.339	1.090	0.573	1.339	-1.889	0.271
183K	-0.869	0.419	1.038	-1.958	1.020	0.601	-0.428	1.038	-1.958	-0.025
184T	-0.155	-0.915	1.122	-1.843	1.011	0.595	-1.873	1.122	-1.873	-0.294
185L	0.920	-0.779	1.533	-1.876	1.367	1.189	-2.334	1.533	-2.334	0.003
186A	0.155	0.053	1.552	-2.073	1.385	1.215	-1.302	1.552	-2.073	0.141
187A	0.155	0.143	1.094	-2.176	0.701	0.620	-1.366	1.094	-2.176	-0.118
188E	0.187	0.770	0.889	-2.252	0.501	0.600	-1.407	0.889	-2.252	-0.102
189W	0.901	-0.130	1.132	-1.726	0.665	1.215	-1.575	1.215	-1.726	0.069
190G	1.129	1.187	1.122	-1.090	0.619	1.215	-0.564	1.215	-1.090	0.517
191G	0.490	0.463	0.982	-0.425	0.601	1.216	0.768	1.216	-0.425	0.585
192H	0.263	0.445	1.085	-0.306	0.711	1.241	0.793	1.241	-0.306	0.605
193G	0.661	0.355	0.945	-0.804	0.683	1.218	1.360	1.360	-0.804	0.631
194I	0.743	-0.476	1.253	-1.214	1.039	1.258	1.246	1.258	-1.214	0.550
195R	0.515	-0.152	1.262	-1.207	1.084	1.258	0.236	1.262	-1.207	0.428
196V	-0.199	-0.607	1.019	-0.764	0.920	0.644	0.403	1.019	-0.764	0.202
197N	-0.427	-0.019	1.029	-0.363	0.966	0.644	-0.607	1.029	-0.607	0.175
198A	0.212	-0.725	1.412	-0.361	1.257	0.661	-0.709	1.412	-0.725	0.250
199L	0.275	-0.899	1.178	-0.735	0.938	0.056	-0.667	1.178	-0.899	0.021
200A	0.275	0.119	1.178	-0.840	0.938	0.056	-0.667	1.178	-0.840	0.151
201P	-0.749	0.974	0.814	-0.833	0.583	0.020	-0.303	0.974	-0.833	0.072
202T	-0.616	0.616	1.244	-0.733	1.057	0.645	0.706	1.244	-0.733	0.417
203V	0.376	0.029	1.477	-0.667	1.203	0.659	0.431	1.477	-0.667	0.501
204F	0.376	0.616	1.477	-0.769	1.203	0.659	0.431	1.477	-0.769	0.570
205R	0.010	1.365	1.113	-0.779	0.920	0.642	0.800	1.365	-0.779	0.581
206S	0.010	0.049	1.113	-0.789	0.920	0.642	0.800	1.113	-0.789	0.392
207A	0.737	-0.921	1.561	-0.882	1.294	1.240	0.185	1.561	-0.921	0.459
208V	0.686	-1.095	1.646	-1.205	1.358	1.261	-0.044	1.646	-1.205	0.372
209T	0.155	-0.508	1.206	-1.532	0.929	0.653	-0.098	1.206	-1.532	0.115
210E	-0.838	-0.460	0.991	-1.893	0.729	0.638	-0.007	0.991	-1.893	-0.120
211W	-0.642	-0.496	1.188	-1.961	0.884	0.657	1.044	1.188	-1.961	0.096
212M	0.225	0.365	1.580	-1.473	1.212	1.145	0.446	1.580	-1.473	0.500
213F	0.528	1.311	1.655	-0.550	1.376	1.614	0.397	1.655	-0.550	0.904
214T	0.168	2.112	1.571	0.661	1.285	1.033	0.643	2.112	0.168	1.068
215D	1.160	2.435	2.001	1.360	1.905	1.603	0.686	2.435	0.686	1.593
216D	1.786	1.896	2.001	1.364	1.813	1.585	0.741	2.001	0.741	1.598
217P	2.633	1.848	2.496	0.464	2.333	2.206	0.489	2.633	0.464	1.781
218K	2.437	2.303	2.300	-0.648	2.178	2.186	-0.562	2.437	-0.648	1.456
219G	2.134	2.046	2.225	-1.501	2.014	1.717	-0.512	2.225	-1.501	1.160
220R	1.767	1.419	2.384	-1.896	2.169	1.853	-0.504	2.384	-1.896	1.027
221A	2.128	0.491	2.468	-1.915	2.260	2.433	-0.750	2.468	-1.915	1.016
222T	1.900	0.287	2.019	-1.858	1.622	1.839	-1.825	2.019	-1.858	0.569
223R	1.274	-0.204	2.019	-2.051	1.713	1.856	-1.881	2.019	-2.051	0.390
224E	0.427	-0.204	1.505	-2.330	1.248	1.237	-1.445	1.505	-2.330	0.063
225A	0.427	-1.103	1.505	-2.577	1.248	1.237	-1.445	1.505	-2.577	-0.101
226M	0.364	-0.745	1.739	-2.769	1.567	1.841	-1.487	1.841	-2.769	0.073
227L	-0.408	-0.835	1.169	-2.815	1.075	1.218	-1.164	1.218	-2.815	-0.251
228A	-0.768	0.183	1.085	-2.650	0.984	0.638	-0.918	1.085	-2.650	-0.207
229R	-1.482	0.996	1.001	-2.489	0.993	0.643	0.527	1.001	-2.489	0.027
230I	-0.951	0.009	1.440	-2.191	1.422	1.251	0.582	1.440	-2.191	0.223
231P	-0.104	0.333	1.954	-2.071	1.886	1.870	0.146	1.954	-2.071	0.573
232L	-0.819	0.550	1.889	-1.997	1.841	1.874	1.407	1.889	-1.997	0.678
233R	-0.951	1.113	1.459	-2.065	1.367	1.250	0.398	1.459	-2.065	0.367
234R	0.048	0.874	1.926	-1.967	1.750	1.847	0.050	1.926	-1.967	0.647
235F	0.048	0.600	1.926	-1.785	1.750	1.847	0.050	1.926	-1.785	0.633
236A	1.122	0.600	2.337	-1.535	2.105	2.441	-0.412	2.441	-1.535	0.951

237E	1.489	0.503	<b>2.178</b>	-1.071	1.950	<b>2.306</b>	-0.420	2.306	-1.071	0.991
238P	0.642	0.556	1.683	-0.595	1.431	1.685	-0.168	1.685	-0.595	0.748
239E	0.990	0.197	1.627	-0.181	1.467	1.683	0.170	1.683	-0.181	0.850
240D	1.217	-0.583	1.617	-0.037	1.422	1.683	1.180	1.683	-0.583	0.929
241F	0.857	-1.446	1.290	-0.290	1.057	1.083	0.197	1.290	-1.446	0.392
242V	0.142	-1.678	0.963	-0.989	0.793	1.070	0.412	1.070	-1.678	0.102
243G	-0.857	-1.787	0.496	-1.634	0.410	0.472	0.760	0.760	-1.787	-0.306
244A	-1.609	-2.618	0.477	-2.183	0.328	0.002	1.101	1.101	-2.618	-0.643
245L	-1.609	-1.763	0.459	-2.461	0.382	0.003	1.284	1.284	-2.461	-0.529
246I	-1.957	-1.019	0.496	-2.489	0.401	0.007	1.131	1.131	-2.489	-0.490
247Y	-1.906	-0.695	0.655	-2.193	0.601	0.027	1.290	1.290	-2.193	-0.317
248L	-1.407	0.568	0.926	-1.490	0.920	0.516	<b>2.291</b>	2.291	-1.490	0.332
249L	-0.692	1.627	1.010	-0.668	0.911	0.511	0.846	1.627	-0.692	0.507
250S	0.225	1.658	1.300	0.315	1.084	0.529	0.684	1.658	0.225	0.828
251D	0.756	0.395	1.197	0.932	1.002	0.530	0.512	1.197	0.395	0.761
252A	0.756	0.347	1.216	1.247	0.948	0.529	0.328	1.247	0.328	0.767
253S	1.217	0.974	1.552	1.261	1.175	0.543	0.225	1.552	0.225	0.993
254S	1.135	0.610	1.599	1.034	1.175	0.542	0.106	1.599	0.106	0.886
255F	0.863	-0.342	1.318	0.511	0.811	0.053	0.116	1.318	-0.342	0.476
256Y	1.110	-0.282	1.646	0.095	1.185	0.096	1.289	1.646	-0.282	0.734
257T	0.465	-0.282	1.375	-0.460	1.020	0.077	1.718	1.718	-0.460	0.559
258G	-0.212	-0.977	1.216	-0.942	0.911	0.074	1.503	1.503	-0.977	0.225
259Q	0.250	-1.065	1.533	-1.346	1.194	0.090	1.584	1.584	-1.346	0.320
260V	-0.212	-0.929	1.197	-1.773	0.966	0.076	1.687	1.687	-1.773	0.144
261M	0.092	-0.206	1.272	-1.799	1.130	0.545	1.637	1.637	-1.799	0.382
262Y	0.092	-0.498	1.272	-1.517	1.130	0.545	1.637	1.637	-1.517	0.380
263L	0.073	0.399	0.935	-0.989	0.711	0.502	1.474	1.474	-0.989	0.444
264D	0.187	0.604	1.309	-0.637	0.957	0.520	1.217	1.309	-0.637	0.594
265G	0.781	0.101	1.515	-0.493	1.066	0.523	1.313	1.515	-0.493	0.687
266G	1.034	-0.272	1.262	-0.751	0.829	0.503	-0.029	1.262	-0.751	0.368
267Y	1.704	-0.645	1.132	-0.705	0.729	0.516	0.158	1.704	-0.705	0.413
268T	1.072	0.017	0.403	-0.712	0.729	0.087	-0.843	1.072	-0.843	0.107
269A	0.711	-0.220	-0.046	-0.419	1.093	0.147	-1.853	1.093	-1.853	-0.084
270C	0.351	0.035	-0.495	-0.240	1.458	0.207	-2.864	1.458	-2.864	-0.221

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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	<sup>1</sup> VEEMALAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVN GAGIDDAAVVTCRPSLADAQQMVEAALGRYGRLDGVLVASGSNHVAPITEMAVEVEDFDAM VRGAWLVCRAAGRVLLEQQQGGSVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG GHGIRVNALAPTVFRSAVTEWMFTDDPKGRATREAMLARIPLRRFAEPEPDVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>
Hydrophilicity	<sup>1</sup> VEEMALAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVN <u>GAGIDDAAVVTCRPSLADAQQMVEAALGRYGRLDGVLVASGSNHVAPITEMAVEVEDFDAM</u> VRGAWLVCRAAGRVLLE <u>EQQQGGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG</u> GHGIRVNALAPTVFRSAVTEWMFTDDPKGRAT <u>TDDPKGRATREA</u> MARIPLRRFAEPEPDVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>
Flexibility	<sup>1</sup> VEEMALAQQVPNLGL <u>ARFSVQDK</u> SILITGATGSLGRVAARALADAGARL <u>TLAGGNS</u> AGLAELVN GAGIDDAAVVTCRPSLADAQQMVEAALGRYGRLDGVLVASGSNHVAPITEMAVEVEDFDAM VRGAWLVCRAAGRVLLE <u>EQQQGGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG</u> GHGIRVNALAPTVFRSAVTEWMFTDDPKGRAT <u>REAMLARIPLRRFAEPEPD</u> FVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>
Accessibility	<sup>1</sup> VEEMALAQQVPNLGLA <u>RFSVQDK</u> SILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVN GAGIDDAAVVTCRPSLADAQQMVEAAL <u>GRYGRLDG</u> VVASGSNHVAPITEMAVEVEDFDAM VRGAWLVCRAAGRVLLE <u>EQQQGGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG</u> GHGIRVNALAPTVFRSAVTEWMFTDDPKGRAT <u>REAMLARIPLRRFAEPEPD</u> FVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>
Turns	<sup>1</sup> VEEMALAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVN GAGIDDAAVVTCRPSLADAQQMVEAALGRYGRLDGVLV <u>SGSNHVA</u> PITEMAVEVEDFDAM VRGAWLVCRAAGRVLLE <u>EQQQGGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG</u> GHGIRVNALAPTVFRSAVTEWMFTDDPKGRATREAMLARIPLRRFAEPEPDVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>
Exposed Surface	<sup>1</sup> VEEMALAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVN GAGIDDAAVVTCRPSLADAQQMVEAALGRYGRLDGVLVASGSNHVAPITEMAVEVEDFDAM VRGAWLVCRAAGRVLLE <u>EQQQGGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG</u> GHGIRVNALAPTVFRSAVTEWMFTDDPKGRAT <u>REAMLARIPLRRFAEPEPD</u> FVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>
Polarity	<sup>1</sup> VEEMALAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVN GAGIDDAAVVTCRPSLADAQQMVEAALGRYGRLDGVLVASGSNHVAPITEMAVEVEDFDAM VRGAWLVCRAAGRVLLE <u>EQQQGGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG</u> GHGIRVNALAPTVFRSAVTEWMFTDDPKGRAT <u>REAMLARIPLRRFAEPEPD</u> FVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>
Antigenic Propensity	<sup>1</sup> VEEMALAQQVPNLGLARFSVQDKSILITGATGSLGRVAARALADAGARLTLAGGNSAGLAELVN GAGIDDAAVVTCRPSLADAQQMVEAALGRYGRLDGVLV <u>ASGSNHVAPITEMAVEVEDFDAM</u> VRGAWLVCRAAGRVLLE <u>EQQQGGGSVVLVSSVRGGLGNAAGYSAYCPSKAGTDLLAKTLAAEWG</u> GHGIRVNALAPTVFRSAVTEWMFTDDPKGRATREAMLARIPLRRFAEPEPDVGALIYLLSDASSF YTGQVMYLDGGYTAC <sup>270</sup>

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