Target Thresholds >= 25 <= 20 <= 5 <= 8.5 Ignore Ignore >= 75

Design	Stable-Stable	Unstable-Unstable	RLSASA_paratope	RLSASA_epitope	Unstable_	With_Stal	Stablizied_Percentage
insulin_epitope1_design440_0	20.83	33.33	11.6	2.79	8	6	75
insulin_epitope1_design444_1	3.7	44.44	8.62	3.55	8	8	100
insulin_epitope1_design490_1	18.75	25	11.51	4.8	5	4	80
insulin_epitope1_design632_0	0	38.1	12.51	3.87	6	5	83.33
insulin_epitope1_design642_0	3.33	50	13.17	2.64	6	5	83.33
insulin_epitope1_design821_0	14.29	50	14.98	3.21	11	10	90.91
insulin_epitope2_design12_0	29.41	41.18	3.77	2.14	2	2	100
insulin_epitope2_design16_0	36.11	16.67	2.34	6.6	1	1	100
insulin_epitope2_design16_1	21.62	18.92	4.15	4.85	9	7	77.78
insulin_epitope2_design180_1	20.83	33.33	11.76	3.8	6	4	66.67
insulin_epitope2_design207_0	20	35	3.45	2.93	5	5	100
insulin_epitope2_design207_1	13.04	26.09	7.1	6.08	5	5	100
insulin_epitope2_design223_0	18.52	37.04	6.92	4.96	5	4	80
insulin_epitope2_design230_0	15.79	36.84	7.82	4.15	10	9	90
insulin_epitope2_design230_1	20	23.33	8.06	7.23	8	7	87.5
insulin_epitope2_design277_0	44.74	18.42	7.58	5.94	6	6	100
insulin_epitope2_design289_0	31.03	20.69	7.21	1.87	4	3	75
insulin_epitope2_design345_1	28	24	9.74	4.19	7	6	85.71
insulin_epitope2_design350_1	25	25	11.1	4.54	8	6	75
insulin_epitope2_design36_1	30	40	7.12	4.01	9	7	77.78
insulin_epitope2_design376_0	28.12	28.12	5.61	6.34	8	5	62.5
insulin_epitope2_design376_1	21.21	39.39	3.88	2.94	6	6	100
insulin_epitope2_design422_0	29.03	29.03	5.22	3.82	4	2	50
insulin_epitope2_design432_0	8.7	43.48	11.05	4.23	8	6	75
insulin_epitope2_design480_0	25.64	28.21	5.61	6.5	7	4	57.14
insulin_epitope2_design480_1	32.26	22.58	2.77	4.36	8	7	87.5
insulin_epitope2_design481_0	20	20	4.33	5.96	9	6	66.67
insulin_epitope2_design494_1	23.08	30.77	5.77	5.56	6	3	50
insulin_epitope2_design49_1	18.52	29.63	7.13	6.59	8	8	100
insulin_epitope2_design538_1	34.38	15.62	4.11	5.22	1	0	0

insulin_epitope2_design567_1	7.5	47.5	9.58	3.89	13	11	84.62
insulin_epitope2_design56_1	21.88	34.38	10.31	4.83	9	7	77.78
insulin_epitope2_design580_1	15.15	39.39	12.86	6	10	8	80
insulin_epitope2_design586_0	42.86	14.29	12.38	3.3	3	3	100
insulin_epitope2_design586_1	33.33	18.52	11.78	4.42	6	5	83.33
insulin_epitope2_design590_1	20	32	2.49	2.98	6	6	100
insulin_epitope2_design592_1	19.35	32.26	5.65	4.38	5	5	100
insulin_epitope2_design642_1	29.63	14.81	9.65	3.3	7	7	100
insulin_epitope2_design651_0	13.33	40	2.91	3.83	9	6	66.67
insulin_epitope2_design661_0	26.09	17.39	3.75	2.89	3	3	100
insulin_epitope2_design6_0	13.04	34.78	9.61	5.05	9	8	88.89
insulin_epitope2_design700_1	12.12	30.3	5.97	6.98	11	9	81.82
insulin_epitope2_design724_1	40.54	24.32	3.74	4.03	6	6	100
insulin_epitope2_design727_1	31.25	28.12	9.58	6.54	7	5	71.43
insulin_epitope2_design751_1	7.69	53.85	9.76	2.47	9	7	77.78
insulin_epitope2_design766_0	8.82	44.12	5.35	7.46	10	9	90
insulin_epitope2_design783_1	16.67	36.67	10.39	2.4	9	9	100
insulin_epitope2_design790_1	37.5	29.17	3.06	2.79	4	2	50
insulin_epitope2_design791_1	21.88	31.25	14.79	3.93	4	3	75
insulin_epitope2_design819_0	35.48	16.13	7.17	3.29	5	5	100
insulin_epitope2_design861_0	9.38	37.5	4.4	5.17	10	9	90
insulin_epitope2_design861_1	15.62	31.25	5.14	5.47	7	6	85.71
insulin_epitope2_design89_1	25.81	32.26	4.17	2.34	6	2	33.33
insulin_epitope2_design952_0	26.67	30	5.63	2.1	7	5	71.43
insulin_epitope3_design229_0	3.45	51.72	10.66	4.46	7	6	85.71
insulin_epitope3_design626_1	29.03	25.81	6.68	4.87	4	4	100
insulin_epitope3_design805_1	0	29.17	12.2	5.34	4	2	50
insulin_epitope3_design843_0	7.69	42.31	9.66	5.68	5	4	80

<= 10 ·	<= -2.5	>= 0.65	Primary Reason
PAE (dG/BSA	SC	
8	-3.323	0.744	Need an epitope 1 design and these two appear equivalent so I've listed both
7	-3.665	0.727	
8	-3.271	0.748	Need an epitope 1 design and these two appear equivalent so I've listed both
5	-3.265	0.736	
8	-3.658	0.72	
5	-3.718	0.708	
6	-3.368	0.717	
5	-3.592	0.719	Meets all stability criteria
5	-3.576	0.71	Good SC and generally good stability criteria
5	-3.347	0.738	
7	-3.409	0.738	
8	-3.272	0.735	
6	-3.487	0.734	
6	-3.367	0.707	
7	-3.388	0.715	
8	-3.596	0.751	Generally strong stability criteria
8	-3.635	0.726	Generally strong stability criteria
6	-3.643	0.761	
7	-3.639	0.763	
6	-3.682	0.728	
5	-3.722	0.717	
6	-3.852	0.735	
7	-3.225	0.712	
5	-3.369	0.702	
7	-3.34	0.702	
8	-3.895	0.774	Meets almost all criteria
4	-3.34	0.704	
8	-3.488	0.743	
6	-3.14		
6	-3.572	0.709	Good stability criteria - the 0% is unimportant because it was 0 / 1

6	-3.367	0.727
8	-3.55	0.732
6	-3.542	0.748
5	-3.83	0.782
5	-3.27	0.735
8	-3.647	0.722
7	-3.658	0.722
6	-3.301	0.707
6	-3.641	0.721
5	-3.414	0.713
6	-3.381	0.741
6	-3.62	0.731
6	-3.772	0.74
5	-3.597	0.727
7	-3.304	0.74
6	-3.293	0.728
		0.725
6		0.732
6		0.759
7		0.718
6 7	-3.061 -3.349	0.732 0.727
6	-3.859	0.727
5		0.732
7	-3.19	0.718
7		0.776
7	-3.913	0.764
8	-3.375	0.7