

A

UbV	Randomised Residues	Count	Max Values	
			Normalized Absorbance (BSA)	Normalized Absorbance (GST)
1	KFKLTSNRTRIAGMLIKKTHILKSRLARR	9	34.2	15.7
2	QFKYPGNRPRVAGKLQPVTHVLTLRVKAA	8	33.5	14.8
3	QFKRIGQRPSIAGKLLKRTHIVRL LHGDR	7	38.0	18.0
4	QFKLTGHSTRIAGKMRFKHFLLRPSLFR	7	36.7	17.0
5	SFRFTRKRSSIAGQLTKTYVLRFGVKK	5	38.8	19.1
6	RFKAQGERSRIAGMQRKTTLSRVRVYDS	5	36.0	17.1
7	KFKRNGNRPRIAGNQKVTHVSRPPGIPK	5	36.2	16.2
8	QRKLTGTRTSIAGNQLELTHLLRLRGAGS	4	28.6	11.4
9	LFKISGTRANITGKQKKVTHLRRIRGETV	3	38.2	18.2
10	NFKLSKRRTRIAGKQKKTHLLRLSAVER	3	35.4	16.2
11	PFKLTGKRSTIAGNMLKVTHLLRHSKVTA	3	35.6	13.0
12	QFKLRGKTTRIAGLKQKKHYLLRLRVRSY	3	28.9	12.2
13	QKKYPGTTTRIGGKQLKLTHVLIVRAPNI	2	34.7	17.2
14	QFKISGNRARITGKSQKKTHLFRRVGTLA	2	30.7	15.8
15	ERKFTGKKTSIAGKHLKRTHVFRLRDKLE	2	30.9	15.7
16	TFKLPGTRIRIAGYQLKKTHLGSLRDRIA	1	32.1	16.8
17	HFKPTWNRTRIAGMKQKKTFLVLTTRRRYE	1	31.9	16.8
18	QFRLAGRRTTRITGMHITKTHLWRIEAQPN	1	30.4	16.0
19	QFKFSRTRTRIAGGEQKKTHLLIPRAVA	1	26.6	14.0
20	MFKLPGKRRTRIAGKQAKVTHVLRRLRSVPL	1	26.3	13.8
21	QYKLRQPRTRIAGQLLIKTHVLTLRGSHA	1	25.6	13.4
22	PFKRAKKRTTIAGKHVKKTYLLRHRRTSG	1	25.5	13.4
23	PHKLTRPITNIAGKKQAKTHLFRLLRAHS	1	24.2	12.7
24	QFKPTRQRPRITGIQHKKTHLLRLLA AVR	1	23.4	12.3
25	KFKLTGSRSRVGGKKLMKTHLMRLRGHAS	1	22.5	11.8
26	QKKLMGERPSIAGKQRKHTHILRLRGPER	1	19.4	10.2
27	AFKFTRETSRIAGYQKTKHHLLRLRSTLS	1	15.5	8.2

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