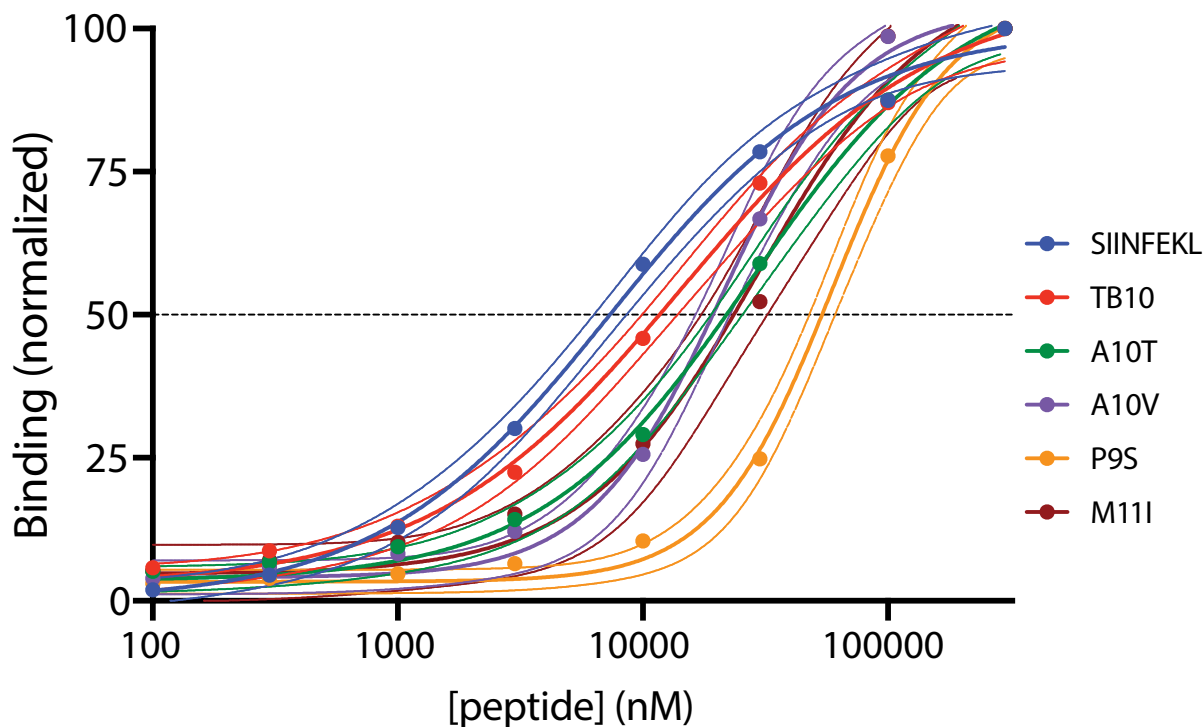


Peptide binding to H-2 K^b



	SIINFEKL	TB10	A10T	A10V	P9S	M11I	IMANAPAM
[Agonist] vs. response - Find ECanything							Ambiguous
Best-fit values							
ECF	7318	14363	28078	21278	59349	29705	~ 115602
F	= 50.00	= 50.00	= 50.00	= 50.00	= 50.00	= 50.00	= 50.00
HillSlope	0.9188	0.8498	1.008	1.593	1.807	1.222	~ 8.816
Bottom	-0.1011	2.707	3.466	4.088	3.341	4.703	15.19
Top	99.97	106.3	109.7	103.6	105.6	110.3	100
logECF	3.864	4.157	4.448	4.328	4.773	4.473	~ 5.063
Span	100.1	103.6	106.2	99.56	102.3	105.6	84.83
Goodness of Fit							
Degrees of Freedom	7	7	7	7	7	7	7
R squared	0.9979	0.9974	0.9973	0.9956	0.9967	0.987	0.8943
Sum of Squares	32.71	36.56	35.27	67.72	39.22	185.8	775.6
Sy.x	2.162	2.285	2.245	3.11	2.367	5.152	10.53
Constraints							
F	F = 50	F = 50	F = 50	F = 50	F = 50	F = 50	F = 50
Number of points							
# of X values	11	11	11	11	11	11	11
# Y values analyzed	11	11	11	11	11	11	11