Additional file 2: Figure S2. Sequence alignment of sdAb A3 and variants.

	1	10	20	30	40	50	60	70	80	90	100	110	120	129
	I	+	+	+	+	+	+	+	+	+	+	+	+	1
83	EYOLYE	ESGGGLYO <mark>r</mark> e	DSLRLSCT	asgrtfs <mark>r</mark> aym(	HFROAPGKE	REFYAAISAA	PGTAYYAFYA	DSYRGRESIS	adsakntyyl	OMNSLKPEDT	AVYYCAADL	<hover sec<="" second="" th="" the=""><th>RSYDYHGOGT</th><th>OVTYSS</th></hover>	RSYDYHGOGT	OVTYSS
83+						C		C.						
A3+neg				D										
no no b				•••••		•••••	•••••	••••••	•••••					•••••

**Figure S2.** Sequence alignment of the SEB binding sdAb A3, A3+, and A3+neg using MultAlin [32]. The initial two amino acids (MA) and the amino acids added due to the restriction sites and the His-tag are not show above (AAALEHHHHH).