

Sensorgrams (upper charts) and steady-state fit (lower charts) obtained for each replicate of peptide 14 / Phd_{STm}⁵²⁻⁷³ L67A (analyte) binding to Doc_{STm} (ligand). Six concentrations of the analyte were flown in multiple individual cycles, ranging from 500 nM to 2.06 nM (3-fold dilutions). The association and dissociation times were, respectively, 100 s and 300 s. For replicate 3 the association and dissociation times were, respectively, 80 s and 200 s

Summary of the fitted parameters obtained from each SPR measurement of peptide 14.

Replicate	K _D (nM)	RMAX (RU)	Offset (RU)	Chi ² (RU ²)
1	24.96 ± 0.50	26.8 ± 0.1	2.10 ± 0.11	0.009
2	23.93 ± 0.94	24.5 ± 0.2	2.10 ± 0.20	0.028
3	15.4 ± 2.13	15.8 ± 0.5	0.90 ± 0.54	0.131