

18.12.14 Parkville

A17 ₁₋₈ wt vs mutants	M1A
	S2A
	Y6A
CPMGs	Y7A

141218_D13_A17-peptides cpmg 16ms

- | | | | |
|----|---|------------------------------|--------------------------------|
| 1 | → | A17 ¹⁸ wt 40 μM | } binds like in previous exp!! |
| 2 | → | " + D13 10 μM | |
| 3 | → | A17 ₁₋₈ M1A 40 μM | } seems to bind OK |
| 4 | → | " + D13 10 μM | |
| 5 | → | A17 ₁₋₈ S2A 40 μM | } " |
| 6 | → | " + D13 10 μM | |
| 7 | → | A17 ₁₋₈ Y6A 40 μM | } reduced binding |
| 8 | → | " + D13 10 μM | |
| 9 | → | A17 ₁₋₈ Y7A 40 μM | } a bit less than wt |
| 10 | → | " + D13 10 μM | |

Scale will fall more fine numbers.
 > png files of the spectra pairs saved.

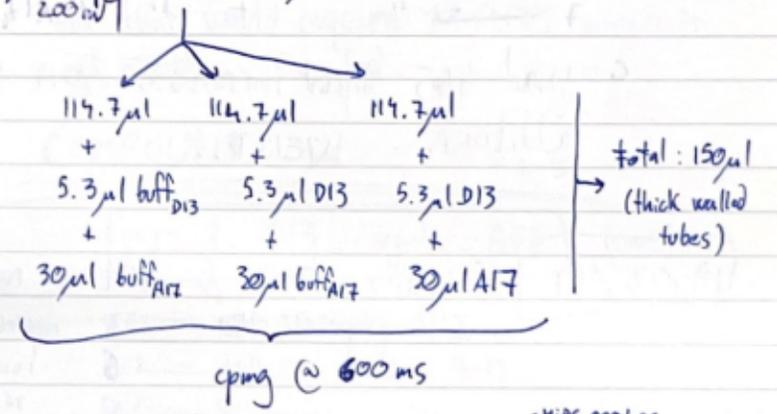
Also, test some of the D13 binders (fragm. screening) for competition w/ A17₁₋₈

→ deuterated (P) buff pH 7.5 (20mM NaH₂PO₄/Na₂HPO₄) + 250mM NaCl

↳ 40ml of 216mg NaH₂PO₄
 88mg Na₂HPO₄
 584mg NaCl
 D₂O

→ Prepare samples w/ 200 μM compound (stock: 200mM)
 ± 3 μM D13 (stock: 85 μM)
 ± 100 μM A17 (stock: 500 μM)

• 1 μl compound + 763.7 μl (P) buff



• For the 1st round, we'll run compounds

- MIPS 000 400
- MIPS 000 455
- MIPS 000 568
- MIPS 000 901