



ATW0092
APTAMER INFORMATION
D5A4Z

Description:

- *Identifiers:* D5A4Z
- *Number of DNA nucleotides:* 32 bases
- *Target for selection:* p202 ERK1 (aa 190-208) CIADPEHDHTGFL(t)EYVATR

Aptamer was selected at Base Pair Biotechnologies from a randomized 32-mer DNA library against the target peptide. Proprietary methods were then used to select the aptamer.

Aptamer folding instruction before use:

Once the aptamer is in its working concentration, it needs to be heated to 85-95°C for 5 minutes, and then cooled to room temperature (~ 15 minutes) before use.

Validation data by Microscale Thermophoresis (MST) analysis:

- *Buffer used for validation:* 1X PBS, 1mM MgCl₂, 0.02% Tween, + 20% Human Serum
- *Average K_d:* 12.9 ± 4.3 nM

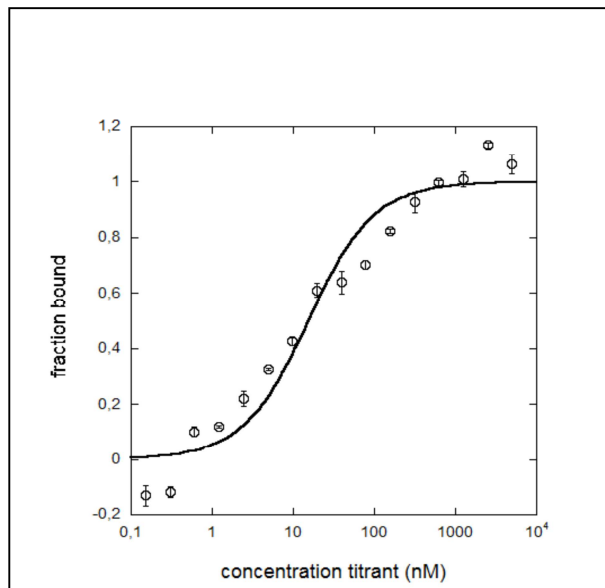


Figure 1. D5A4Z - p202 ERK1 (aa 190-208) peptide binding. The normalized fluorescence of the aptamer is plotted versus the titrated target concentration.

Target for MST analysis: ERK1 (aa 190-208) CIADPEHDHTGFLTEYVATR

- *Buffer used for validation:* 1X PBS, 1mM MgCl₂, 0.02% Tween, + 20% Human Serum
- *Average K_d:* No Binding indicated

