



Combining peptide fragments for the high-throughput generation and screening of large libraries

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Established method to synthesize thiol peptides

Testing dimerization reaction









LCMS chromatograms of test dimerizations and analysis of results



Transfer of 2 nmol per peptide, final concentration of peptides 100 µM * Impurity on the column, ** overlapping product peak

Synthesis of dimer library and screening against thrombin

HS A

Reduced thiol peptides are stored in 1536 microwell plate

Ready to screen, no purification needed!

Workflow for synthesizing dimer libraries



Acoustic droplet ejection to transfer nanoliter volumes of peptides to dimerization plate

To facilitate screening of a large library, dimers are screened in mixtures. Peptides are organized in groups of 6. In each screening well, two groups are pooled resulting in a mixture of 12 peptides forming 72 dimers. In total a library of 853 471 dimers was screened.

Mixtures that decreased the proteolytic activity of thrombin in the first step are deconvoluted by resynthesizing and screening all 72 dimers of the mixture in parallel.

After identification of the active dimer in the mixture, the compound is resynthesized and purified for characterization and IC₅₀ determination.



218 pools of 6 peptides per pool

Large screen: Transfer of 10 pmol per peptide \rightarrow screening concentration for each individual dimer ~ 100 nM Deconvolution: Transfer of 2.5 pmol per peptide \rightarrow screening concentration for each individual dimer ~ 170 nM

References

[1] Schüttel, M., et al. "Solid-phase peptide synthesis in 384-well plates." Journal of Peptide Science 30.4 (2024): e3555. [2] Bognar, Z., et al. "Solid-phase peptide synthesis on disulfide-linker resin followed by reductive release affords pure thiol-functionalized peptides." Organic & Biomolecular Chemistry 20.29 (2022): 5699-5703.

A library of 853 471 peptides was screened against the model target thrombin.

Nanomolar thrombin inhibitors were identified.