

Application Note 27 Using Reichert Surface Plasmon Resonance (SPR) for screening of small molecule inhibitors

Surface Plasmon Resonance (SPR) can be used as an effective tool for screening many samples against a target or targets immobilized on a sensor chip.¹⁻³ Often researchers do initial research showing that a certain molecule with a particular molecular backbone can act as an inhibitor to a given protein. They then follow up with analogs to see if efficacy can be improved by making structural changes to the original molecule including addition of other functional groups. In the example highlighted here, researchers at the University of Pittsburgh used SPR to help characterize inhibitor analogs and determine structure-activity relationships which ultimately could aid in the development of a new class of anti-retroviral drugs.¹



Experimental

Background

Earlier research identified a small molecule inhibitor of HIV- 1 Nef which was shown to block several of its activities related to enhancement of viral infectivity and replication.¹ In the current research, 216 analogs of the original compound were synthesized and screened against recombinant Nef protein using SPR. The analogs all contained the same hydroxypyrazole scaffold from the original inhibitor (core molecule in the inset above) but with modifications at the A, B and C positions. From the SPR data obtained, compounds were compared based on affinity, extent of binding, and residence time (which is calculated from the reciprocal of the dissociation rate constant $(1/k_{off})$).¹

Studying the SPR results, the top scoring inhibitors were found to share certain features, including a slow off rate. None of the top analogs returned to baseline over the course of a 3 minute dissociation, suggesting that these inhibitors might induce a change in the conformation of Nef that enables tight binding. However, no covalent bond was formed, since the analogs dissociated in weak base. The Pitt Researchers developed an "Activity Score" which combined multiple SPR parameters with anti- retroviral activity into a single numerical value for each analog providing an unbiased ranking of the compounds which helped them rank the analogs for further testing.¹





Conditions

- Instrument: Reichert4SPR
- Sensor Chip: Carboxymethyl Dextran Temperature: 25°C
- Target: Recombinant, full-length HIV-1 Nef protein (NL4-3 allele)
- Analyte: Various
- Running Buffer: PBS + 1% DMSO
- Flow Rate: 50 µL/min
- Association Time: 1.5 minutes
- Dissociation Time: 3 minutes
- Regeneration: 5 mM NaOH for 30 seconds

Results

Forty-five of the analogs showed no binding. The remainder were ranked based on SPR kinetics, the size of the SPR response and their activity from a cell-based HIV-1 infectivity assay. Sensorgrams were fit to either a 1:1 model or a conformational change model. Affinities varied widely from low micromolar to picomolar. The top six analogs bound Nef with K_p values ranging from about 0.65 pM (FC-7943) to 10 nM (FC-8698).¹

Results for two of the top six compounds are shown here:





Figure 1⁴. The SPR sensorgrams obtained for compound 7902 are show fit to a 2-step model. Compound 7902 has the structure shown in the inset and shares, at the A-position, a phenyl group substituted at the para position by a hydrophobic trifluoromethyl group, with three of the other top compounds. The K_p is 93.6 pM and the residence time is 3.79e5 minutes, with 96.1% infectivity inhibition at 1 μ M. Each analog concentration was measured in triplicate (30 μ M, red; 10 μ M, orange; 3.3 μ M, green; 1.1 μ M, blue), with the fitted curves shown in black.



Figure 2⁴. The SPR sensorgrams obtained for compound 8698 are shown here, fit to a 1:1 model. Compound 8698 has the structure shown in the inset and shares a chloro- benzimidazole group at the C-position with one other of the top compounds. The K_p is 13 nM, the residence time is 780 minutes, with 100% infectivity inhibition at 1 μ M. Analog concentrations as per the legend to Figure 1



Summary

The screening example outlined here highlights how useful it is to do in vitro testing in real time using SPR. For this example, several hundred inhibitors were tested and ranked based on their kinetics and activity. By using SPR to help create an "Activity Score", researchers were able to determine the top compounds to take into further testing. Their structure–activity approach led these researchers to identify new analogs with high affinity for Nef that also show strong antiretroviral activity and maintain the ability to reverse down regulation by Nef of MHC-I at the cell surface.¹

Reference

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