

Systematic SAR Studies of BMS-182,193, A Novel Inhibitor of HIV Protease

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Following the identification of the lead compound BMS-182,193, a novel aminoalcohol inhibitor of HIV protease, our effort was focused on the improvement of its intrinsic potency and other *in vitro* properties including therapeutic index. The above goals have been achieved by the modification of the P₁' and P₂' side chains of BMS-182,193 through the introduction of various polar groups. In general, substitution at the P₁' side chain of BMS-182,193 resulted in improvements in the therapeutic index as well as in intrinsic potency. Certain modifications at the P₂' side chain lead to significant increases in intrinsic potency which could be explained by the binding mode to HIV protease developed from molecular modeling studies. This poster will disclose our SAR results in those respects.

Potent Inhibitors of HIV Protease: P₂ and P₃ Extended Analogs of BMS-182,193

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As part of a program to design potent inhibitors of HIV protease, we have synthesized and evaluated a number of analogs of an aminoalcohol-derived lead compound BMS-182,193 (HIV-Pr IC₅₀ = 150 nM; HIV-1 ED₅₀ = 80 nM). Extension of the core of BMS-182,193 via an amide linkage into the S₂ and S₃ binding sites of HIV protease resulted in compounds with up to 50-fold enhancement in potency against the enzyme, as well as substantial improvement in the antiviral activity in cell culture. A proposed binding mode for the various P₂ and P₃ extended analogs based on structure-activity relationships and molecular modeling will be presented.