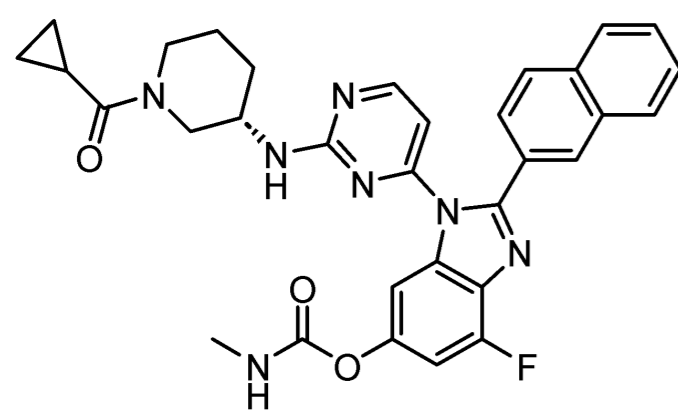


# Small Molecule Highlights

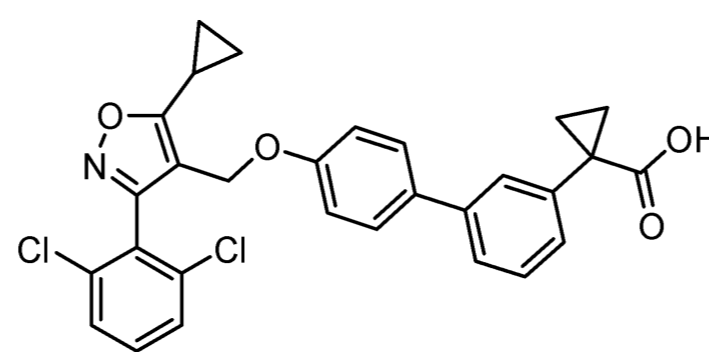
Snapshots from Recent Literature in Target-oriented Drug Design



**Compound 3h** **JNK3** **Neurology**

*N*-monomethyl carbamate *c*-Jun N-terminal kinase 3 inhibitor (JNK3) JNK3/Alzheimer's (AD) via APP, Aβ<sub>42</sub>, AMPK, Tau = Neuronal cell death  
Inhibitor design: Pharmacodynamic-focused structure optimizations  
*In vitro* kinase activity JNK1, JNK2, JNK3 IC<sub>50</sub> = >10,000, 2202, 21 nM  
Induced fit docking (H-bonds): Lys68, Met149, Gln155, H<sub>2</sub>O (PDB 4KKH)  
Neuroprotection in cells, Efficacy in 3xTg mice (restore cognitive function)

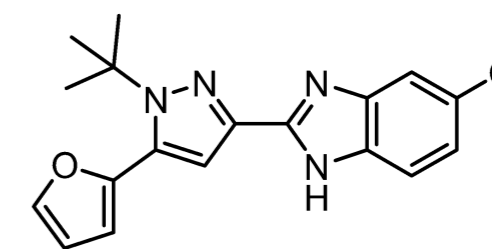
*J. Med. Chem.*  
CPHU/Samjin Pharma, South Korea



**ZLY28** **FXR/FABP1** **Metabolism**

First-in-Class intestinal restricted FXR/FABP1 dual modulator  
FXR/FABP1: Drug targets for nonalcoholic steatohepatitis (NASH)  
Inhibitor design: Comprehensive multi-parameter optimizations  
FXR FRET IC<sub>50</sub> = 143 nM, Cell-based IC<sub>50</sub> = 491 nM, FABP1 IC<sub>50</sub> = 2.7 μM  
LM % rem. mouse, rat, beagle, human (at 1 hr) = 64.3, 68.6, 57.4, 71.6 %  
Efficacy in histological, and *in vivo* NASH models/liver lipid metabolism

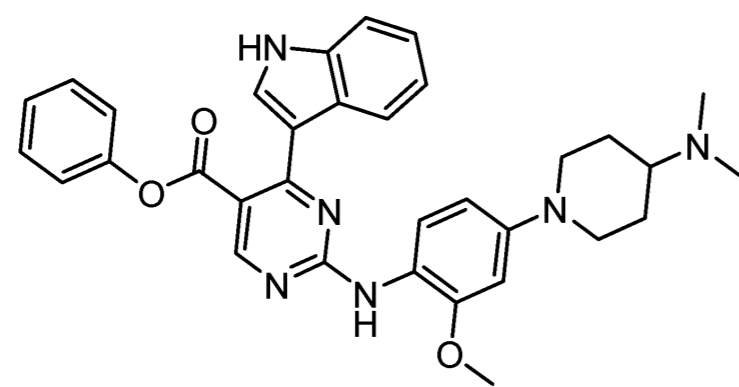
*J. Med. Chem.*  
GPU, China



**Compound 50** **P2Y<sub>6</sub>R** **Inflammation**

Pyrazole benzimidazole-based selective P2Y<sub>6</sub>R antagonist  
P2Y<sub>6</sub>R: Inflammatory disease (Ulcerative colitis/Acute lung injury)  
Inhibitor design: MD, virtual screen (50K), SBDD towards HitID  
P2Y<sub>6</sub>R % Inh. (1 μM) = 132.71%, IC<sub>50</sub> = 5.91 nM (5x Improv. MRS2578)  
Docking Interactions: Asp179, Leu180, Pro183, Tyr262, Leu263, Tyr283  
*In vivo* efficacy in DSS-induced ulcerative colitis/LPS-induced ALI models

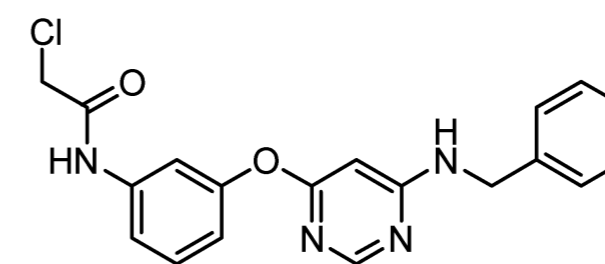
*J. Med. Chem.*  
SU/CPU, China



**Compound 13** **EGFR** **Oncology**

Reversible aminopyrimidine as double-mutant EGFR<sup>L858R/C797S</sup> inhibitor  
Rationale: Address Osimertinib resistance due to EGFR<sup>L858R/C797S</sup> mutations  
Inhibitor design: Structure-based strategy (targeting hydrophobic back pocket)  
HTRF EGFR<sup>WT</sup>, EGFR<sup>L858R/C797S</sup>, EGFR<sup>L858R/C797S/T790M</sup> IC<sub>50</sub> = 0.008, 0.020, 0.64 nM  
Cell activity EGFR<sup>WT</sup> A431 IC<sub>50</sub> = 1837 nM, EGFR<sup>L858R/C797S</sup> Ba/F3 IC<sub>50</sub> = 7.1 nM  
Co-crystal EGFR<sup>T790M/V948R</sup> PDB 7ZYQ: Gln791, Met793, hydrophobic interactions

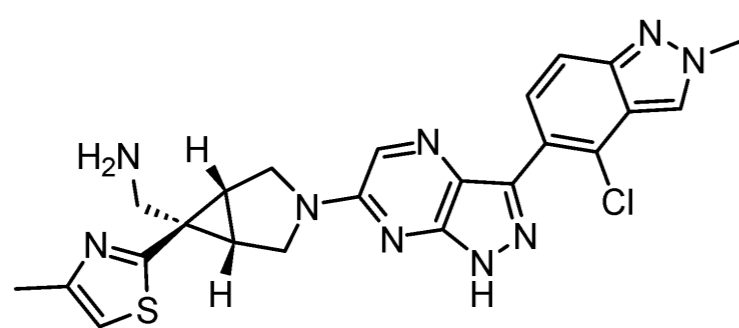
*J. Med. Chem.*  
DDHD/ZIW, Germany



**DK-2403** **MAP2K7** **Oncology**

Covalent, selective MAP2K7 inhibitor (chloroacetamide E+)  
MAP2K7: Pediatric T-cell acute lymphoblastic leukemia (T-ALL)  
Inhibitor design: Structure-based, One-pot synthesis, Cys218  
MAP2K7 ADP Glo-assay IC<sub>50</sub> = 10 nM (no pre-incubation, IC<sub>50</sub> = 93 nM)  
Selectivity across 90 kinases (wild-type), 1 off-target: EGFR (at 1 μM)  
Cell activity KOPT-K1, ALL-SIL IC<sub>50</sub> = 2.8, 1.1 μM (persisted post wash-out)

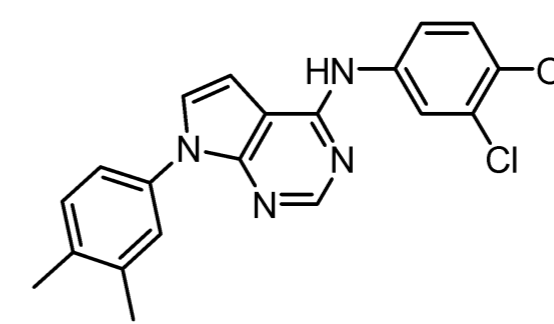
*ACS Med. Chem. Lett.*  
Northwestern University, USA



**Compound 25** **SHP2** **Oncology**

Allosteric SH2-protein tyrosine phosphatase inhibitor (bicyclo[3.1.0]hexane)  
SHP2: Nonreceptor PTPase, roles in Ras/MAPK, JAK/STAT, PI3K/AKT (oncology)  
Inhibitor design: Ligand-based strategies (pyrazopyrazines conformational analysis)  
*In vitro* SHP2 IC<sub>50</sub> = 5 nM, pERK SHP2 IC<sub>50</sub> = 25 nM, hERG (Tox.) IC<sub>50</sub> >30 μM  
*In vivo* PK (SD rats, 1 mg/kg, p.o.): T<sub>1/2</sub>, F%, C<sub>Max</sub> = 1.9 hrs, 1.2%, 0.01 μM  
P-gp substrate (in rats), likely responsible for poor oral bioavailability

*ACS Med. Chem. Lett.*  
IRBM/Promidis, Italy



**ARUK2002821** **PI5P4Kα** **Oncology**

Pyrrolopyrimidine phosphatidylinositol 5-phosphate 4-kinase inhibitor  
PI5P4K: Roles in cancer, neurodegeneration, immunology (+ PIP/PIP<sub>2</sub>)  
Inhibitor design: Virtual screen/Scaffold-hopping strategy (known PI5P4Ki)  
ADP-glo assay PI5P4Kα pIC<sub>50</sub> = 8.0, PI5P4Ky+ pIC<sub>50</sub> <4.7 (XlogP = 5.8)  
InCELL Pulse (Cell Target Eng.) PI5P4Kα pIC<sub>50</sub> = 6.6, MLM T<sub>1/2</sub> = 11 mins  
Co-crystal structure with PI5P4Kα (PDB 8C8C) + conformational analysis

*RSC Med. Chem.*  
University of Cambridge, UK

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