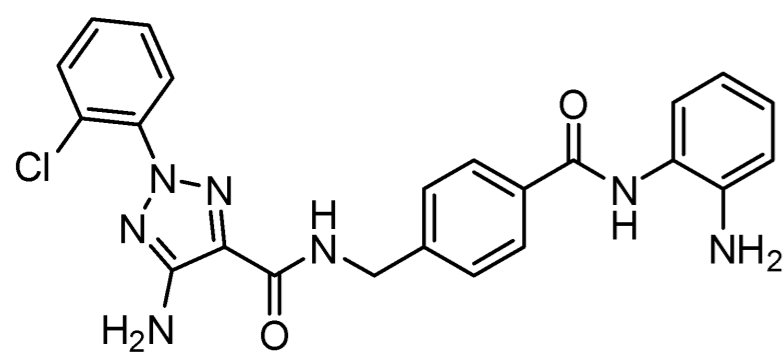


# Small Molecule Highlights

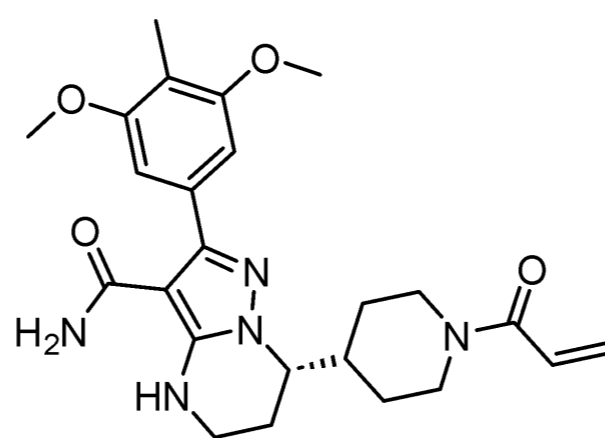
Snapshots from Recent Literature in Target-oriented Drug Design



**Compound 19h** **HDAC** **Oncology**

Triazole-based histone deacetylase (HDAC) inhibitor (Class I selective)  
Inhibitor Design: Structural screen of various surface cap groups  
HDAC1, HDAC2, HDAC3 IC<sub>50</sub> = 47, 125, 450 nM, HDAC4-11 IC<sub>50</sub> > 10 μM  
Anti-proliferative activity: MC38, HCT116, A549, PC-9, ES-2, DoHH2  
MOA: HDACi results in cell cycle arrest, and induces apoptosis (FACS)  
*In vivo* efficacy in both MC38, and HCT116 Xenograft models (TGI > 80%)

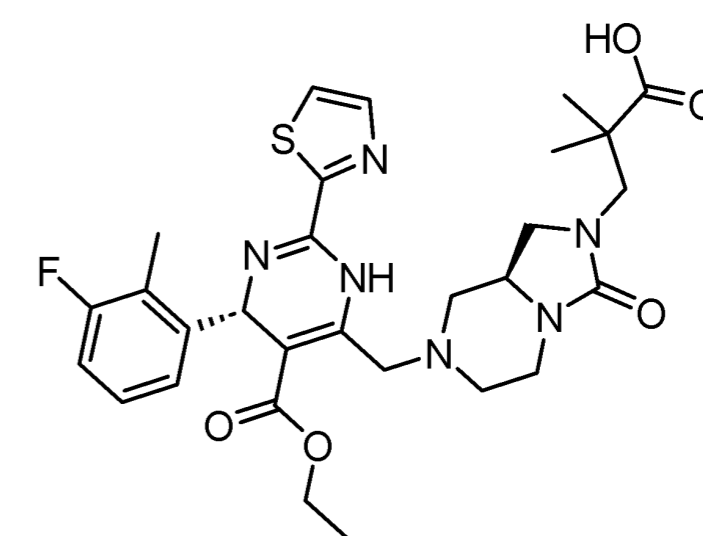
*J. Med. Chem.*  
SIAIS/XMU, China



**BGB-8035** **BTK** **Oncology**

Covalent inhibitor of Bruton's Tyrosine Kinase (BTK)  
Structural optimization of Zanubrutinib (Selectivity, Kinase DFG-out)  
Covalent mechanism: Standard acrylamide electrophile (Cys481, BTK)  
BTK, EGFR, TEC IC<sub>50</sub> = 1.1, 621, 99 nM (Off-targets: TXK, NLK, BMX)  
Cellular p223-BTK IC<sub>50</sub> = 14 nM, pEGFR IC<sub>50</sub> > 10 μM, pTEC IC<sub>50</sub> = 223 nM  
*In vivo* efficacy in REC-1 Xenografts (TGI = 79%) and a CIA rat model

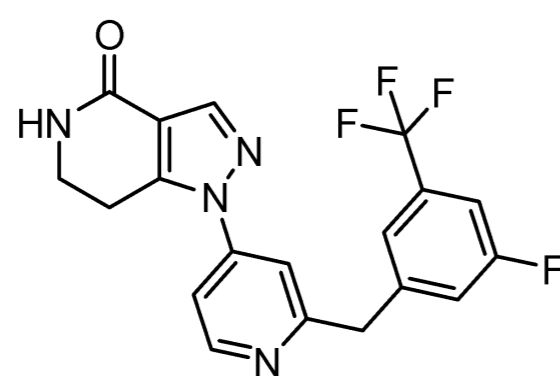
*J. Med. Chem.*  
BeiGene, China



**RG7907** **HBV** **Antiviral**

Hepatitis B Virus (HBV) core protein allosteric modulator (anti-viral)  
Structural core: Hetero-aryldihydropyrimidine (Derivatization of HAP\_R01)  
HBV DNA (HepG.2.2.15) EC<sub>50</sub> = 6 nM, HepDE19 cells CC<sub>50</sub> > 100 μM, LogD = 1.75  
Co-crystal structural determination of Cp149 Y132A: PDB 8I71 (Trp125, H<sub>2</sub>O)  
*In vivo* PK characterization: Rat, monkey, minipig (oral bioavailability observed)  
*In vivo* Efficacy in adeno-associated virus-HBV mouse model (20 mg/kg, q.d.)

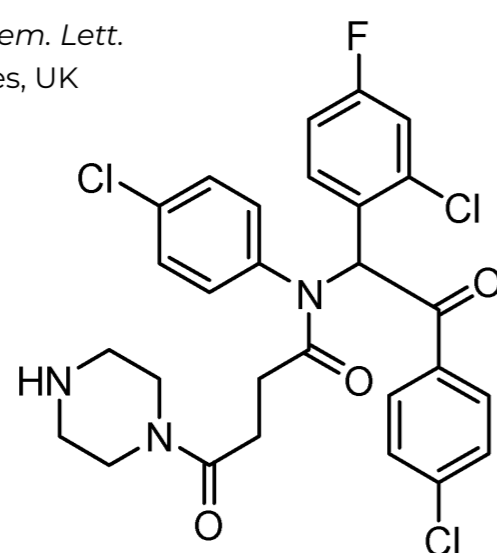
*J. Med. Chem.*  
CICoR, Roche, China



**HTL0041178** **GPR52** **Neurology**

GPR52 agonist (Lactam-pyrrolo core), identified via FBDD  
GPR52: orphan G-as G protein-coupled receptor, GPCR (D1/D2)  
cLogP, tPSA, GPR52 pEC<sub>50</sub>, Chrom. LLE = 3.1, 60 Å<sup>2</sup>, 7.5, 3.6  
Molecular docking studies: Superposition with c17 control analog  
*In vivo* PK (3 mg/kg, p.o.): C<sub>max</sub>, F, F<sub>abs</sub> % = 486 ng/mL, 95, 80%  
Efficacy in *d*-amphetamine-induced hyperactivity/locomotion in rats

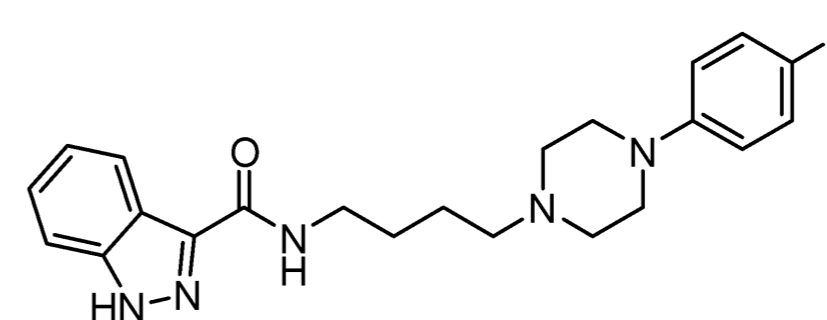
*ACS. Med. Chem. Lett.*  
Sosei Heptares, UK



**Compound C16** **p53-MDM2/MDMX** **Oncology**

α-aminoketone 4-ring dual inhibitor of p53-MDM2/MDMX  
p53-MDM2/MDMX: Regulates p53 transcriptional response  
Structural optimization: MDM2/MDMX K<sub>i</sub> = 0.23/2.45 μM  
Antiproliferative activity: HCT116/SH-SY5Y IC<sub>50</sub> = 0.68/0.54 μM  
MM/GBSA calculations of binding hot-spot residues of MDM2/MDMX  
Inhibition of HCT116 migration/invasion + reactivation of p53 activity

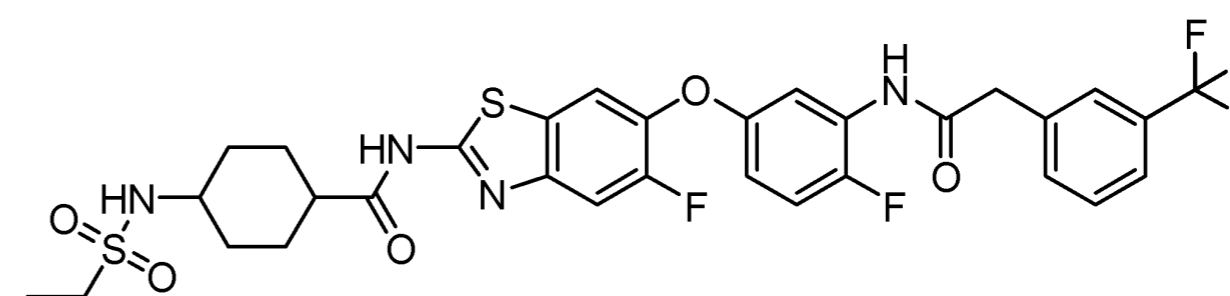
*Eur. J. Med. Chem.*  
NUCM/SUCM, China



**Compound 11** **DA/5-HT** **Neurology**

Multi-target arylpiperazine-based binder of aminergic GPCRs (anti-psychotic)  
Polypharmacology: Antagonist of D2/5-HT2A, Agonist of 5-HT1A receptor  
Structural optimization of D2AAK3 analog (Linker, aryl group derivatization)  
D2, 5-HT1A, 5-HT2A K<sub>i</sub> (Competitive-radio ligand binding) = 596, 56.6, 66.7 nM  
5-HT2C, H1 K<sub>i</sub> (Competitive-radio ligand binding) = 552, 1141 nM  
5-HT1A Agonism (cAMP) EC<sub>50</sub> = 160 nM, 5-HT2A Antagonism (IP) K<sub>b</sub> = 96.4 nM

*Eur. J. Med. Chem.*  
Medical University of Lublin, Poland



**SZM-1209** **RIPK1** **Respiratory**

Benzothiazole Receptor-Interacting Protein Kinase 1 (RIPK1) inhibitor  
RIPK1: Target for Acute lung injury/acute respiratory distress syndrome (ALI/ARDS)  
Inhibitor Design: Ligand based scaffold hopping (related necroptosis inhibitors)  
Antinecroptosis EC<sub>50</sub> = 22 nM, Cytotoxicity CC<sub>50</sub> > 100 μM (Improve. TAK-632)  
RIPK1 K<sub>d</sub> = 85 nM, RIPK3 K<sub>d</sub> > 10 μM (Recombinant protein) (>100x FS for RIPK1)  
*In vivo* Anti-inflammatory effects of SZM-1209 in an NNK-induced ALI model

*J. Med. Chem.*  
SMMU, China

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