**Proteins** 

## **UNC-2170**

Cat. No.: HY-115531 CAS No.: 1648707-58-7 Molecular Formula:  $C_{14}H_{21}BrN_{2}O$ Molecular Weight: 313.23

Others Target: Pathway: Others

Storage: 4°C, protect from light

\* In solvent: -80°C, 6 months; -20°C, 1 month (protect from light)

**Product** Data Sheet

## **SOLVENT & SOLUBILITY**

In Vitro

Methanol: 125 mg/mL (399.07 mM; Need ultrasonic) DMSO: 100 mg/mL (319.25 mM; Need ultrasonic)

Preparing Stock Solutions	Solvent Mass Concentration	1 mg	5 mg	10 mg
	1 mM	3.1925 mL	15.9627 mL	31.9254 mL
	5 mM	0.6385 mL	3.1925 mL	6.3851 mL
	10 mM	0.3193 mL	1.5963 mL	3.1925 mL

Please refer to the solubility information to select the appropriate solvent.

In Vivo

- 1. Add each solvent one by one: 10% DMSO >> 40% PEG300 >> 5% Tween-80 >> 45% saline Solubility: ≥ 2.5 mg/mL (7.98 mM); Clear solution
- 2. Add each solvent one by one: 10% DMSO >> 90% (20% SBE-β-CD in saline) Solubility: ≥ 2.5 mg/mL (7.98 mM); Clear solution
- 3. Add each solvent one by one: 10% DMSO >> 90% corn oil Solubility: ≥ 2.5 mg/mL (7.98 mM); Clear solution

## **BIOLOGICAL ACTIVITY**

Description

UNC-2170 is a functionally active, fragment-like ligand for 53BP1 (IC  $_{50}$ =29  $\mu$ M; K $_{d}$ =22  $\mu$ M). UNC-2170 shows at least 17-fold selectivity for 53BP1 as compared to nine other methyl-lysine (Kme) reader proteins. 53BP1 is a Kme binding protein that plays a central role in DNA Damage Repair (DDR) pathways and is recruited to sites of double-strand breaks (DSB)<sup>[1]</sup>.

In Vitro

 $UNC-2170~(500~\mu\text{M})~results~in~a~significant~increase~in~soluble~53BP1~as~compared~to~untreated~lysates~or~lysates~treated~with~as~compared~to~untreated~lysates~treated~with~lysates~treated~lysates~treated~lysates~treated~lysates~treated~lysates~treated~lysates~treated~lysates~treated~lysates~treated~lysates~lysat$ UNC2892, the negative control compound. UNC-2170 (30-100 μM; naive splenocytes cultured with LPS and IL-4 for 3.5 days) clearly phenocopies the reduction in CSR seen in 53BP1 mutant B cells<sup>[1]</sup>.

MCE has not independently confirmed the accuracy of these methods. They are for reference only.

REFERENCES				
[1]. Perfetti MT, et al. Identi	fication of a fragment-like small	molecule ligand for the methyl-	lysine binding protein, 53BP1. ACS Chem Biol. 2015	;10(4):1072-1081.
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	Tel: 609-228-6898	Fax: 609-228-5909	E-mail: tech@MedChemExpress.com nouth Junction, NJ 08852, USA	
	Address: .	i Deer Park Dr, Suite Q, Monin	louth Junction, NJ 08852, USA	

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