# **Screening Libraries**

# **NQTrp**

Cat. No.: HY-19738 CAS No.: 185351-19-3 Molecular Formula:  $C_{21}H_{16}N_2O_4$ Molecular Weight: 360.36 Target: Tau Protein

Pathway: **Neuronal Signaling** -20°C, protect from light Storage:

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

**Product** Data Sheet

# **SOLVENT & SOLUBILITY**

In Vitro

DMSO: 125 mg/mL (346.88 mM; ultrasonic and warming and heat to 60°C)

	Solvent Mass Concentration	1 mg	5 mg	10 mg
Preparing Stock Solutions	1 mM	2.7750 mL	13.8750 mL	27.7500 mL
ototic ookations	5 mM	0.5550 mL	2.7750 mL	5.5500 mL
	10 mM	0.2775 mL	1.3875 mL	2.7750 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: ≥ 2.08 mg/mL (5.77 mM); Clear solution
- 2. Add each solvent one by one: 10% DMSO >> 90% (20% SBE-β-CD in saline) Solubility: ≥ 2.08 mg/mL (5.77 mM); Clear solution

# **BIOLOGICAL ACTIVITY**

D	esc	rip	tio	n

NQTrp, an aromatic naphthoquinone-tryptophan hybrid molecule, an inhibitor of the aggregation of the tau protein with generic anti-amyloidogenic effects. NQTrp inhibits the in vitro aggregation of hexapeptide (41GCWMLY46 within the Nterminus of  $\gamma D$ -crystallin) as well as full-length  $\gamma D$ -crystallin<sup>[1]</sup>.

# In Vitro

In TEM analysis, GDC6 peptide fibrils (50 μM) in the presence of NQTrp 5:1, 1:1, 1:5 ratio (GDC6: NQTrp). NQTrp effectively inhibits the fibrillation of the GDC6 peptide (50 μM) in a dose-dependent manner and prevents the formation of elongated amyloid fibrils<sup>[1]</sup>.

NQTrp has no toxic effect toward retinal cell culture and reduces the cytotoxicity induced by aggregates of the hexapeptide( $IC_{50}=70 \mu M$ ) in ARPE-19 cells<sup>[1]</sup>.

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

FERENCES					
	An amyloidogenic hexapeptide hybrids. Int J Biol Macromol. 20	from the cataract-associated γE 020 Aug 15;157:424-433.	-crystallin is a model for the	e full-length protein and i	s inhibited by
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