Mad1 (6-21)

Cat. No.:	HY-P3242			
CAS No.:	880150-82-3			
Molecular Formula:	$C_{84}H_{140}N_{24}O_{26}S_{2}$			
Molecular Weight:	1966.29			
Sequence Shortening:	RMNIQMLLEAADYLER			
Target:	Others			
Pathway:	Others			
Storage:	Sealed storage, away from moisture and light, under nitrogen			
	Powder -8	80°C	2 years	
	-1	20°C	1 year	
	* In solvent : -80°C, 6 months; -20°C, 1 month (sealed storage, away from moisture			
	and light, under nitrogen)			

BIOLOGICAL ACTIVITY

Description	Mad1 (6-21) is the 6-21 fragment of Mad1 protein. Mad1 (6-21) binds to mammalian Sin3A PAH2 with a K_d of ~29 nM ^[1] .
In Vitro	The PAH2 domain of mSin3A adopts a left-handed, up-and-down, four-helix bundle structure with residues in all four helices as well as in the turn regions defining a compact structural domain with an extensive hydrophobic core. Helices $\alpha 1$ and $\alpha 2$ form a deep hydrophobic pocket, which constitutes the primary interaction surface for the Mad1 (6-21) peptide. The Mad1 (6-21) forms an amphipathic α helix in the complex and interacts with PAH2 mainly through the apolar surface of the helix ^[1] . MCE has not independently confirmed the accuracy of these methods. They are for reference only.

REFERENCES

[1]. K Brubaker, et al. Solution structure of the interacting domains of the Mad-Sin3 complex: implications for recruitment of a chromatin-modifying complex. Cell. 2000 Nov 10;103(4):655-65.

Caution: Product has not been fully validated for medical applications. For research use only.

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Product Data Sheet

