

G9a/GLP selective methyltransferase chemical probe, UNC0638

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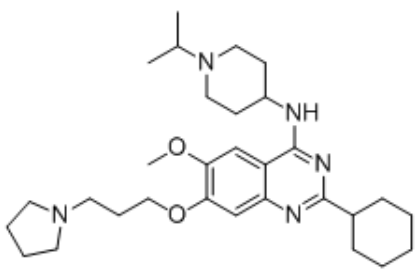
UNC0638	Target Activity			
	Target	K _D / nM (ITC)	IC ₅₀ / nM (Activity)	T _m shift / °C ¹
	G9a (EHMT2)		<15 (Hill Slope 1.3)	4
	GLP (EHMT1)		19±1 (Hill Slope 0.8)	8
2-cyclohexyl-N-(1-isopropyl piperidin-4-yl)-6-methoxy-7-(3-(pyrrolidin-1-yl)propoxy)quinazolin-4-amine	SETD7		>10,000	nt
	SETD8		>10,000	nd
	PRMT3		>10,000	nd
	SUV39H2		>10,000	nt
	DOT1L		nt	nd
	PRDM1		nt	nd
	PRDM10		nt	nd
	PRDM12		nt	nd
	SMYD3		nt	nd
	JMJD2E		4660 (Alphascreen)	nt
	HTATIP		nt	nd
	DNMT1		1287**	nt
	MLL		>10,000**	nt
	EZH2		>10,000**	nt
	PRMT1		>10,000**	nt
	SUV39H1		>10,000**	nt
	SUV39H2		>10,000**	nt
G9a		91**	nt	
nt=not tested, nd=not detected, ¹ singlicate @ 100 μM **Tested at BPS Bioscience using different assay format				
Selectivity within target family	See above, additional screening ongoing.			
Selectivity beyond target family	<ul style="list-style-type: none"> Ricerca Hit Profiling Screen of 29 targets, measuring inhibition of radioligand binding at 1 μM. <30% Inhibition for 26 receptors. >30% Inhibition for muscarinic M₂ (64%), adrenergic alpha_{1A} (90%) and alpha_{1B} (69%) receptors. 			
Physicochemical properties	MW = 509.7		cLogP 5.78	
	Soluble in DMSO at least up to 10mM.			
Storage	Stable as solid in the dark at -20°C.			
Cellular activity	<ul style="list-style-type: none"> Significant reduction in H3K9 dimethylation at 100nM in MDA-MB231 cells as measured by fluorescence immunostaining (Fig. 1 below). Good separation of functional and toxic effects (Fig. 2 below) 			
Co-crystal structures	Structure solved PDB 3RJW			
Primary reference	Vedadi et al, Nat Chem Biol. 7:566-74 (2011).			
Material availability	Available through Sigma-Aldrich (Item U4885).			
Notes	An inactive structurally-related compound will be available shortly for use as a negative control.			
Further information	See probe website: http://www.thesgc.org/chemical_probes/UNC0638			
Funding	The work at SGC is supported by Ontario Research Fund Grant RE-03-003, and the work at The University of North Carolina CICBDD is supported by NIH grant Number RC1GM090732.			

Fig 1 (a) H3K9m2 immunostaining in MDA-MB231 cells with G9a and EHMT1 knockdown and cells treated with UNC638 and BIX-01294 inhibitors. BIX-01294 was identified as EHMT1 and 2 inhibitor, effective in mES cells at about 1uM (Kubicek et al, 2007 Molecular Cell 25, 473-481) **(b)** UNC638 and BIX-01294 dose-response on H3K9 dimethylation levels in MDA-MB231 cells at 48h.

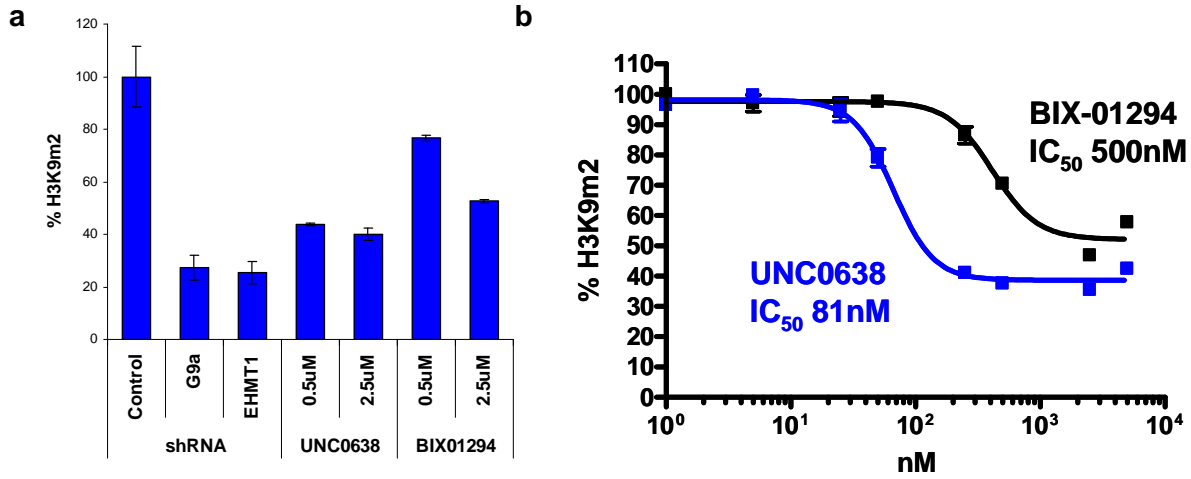
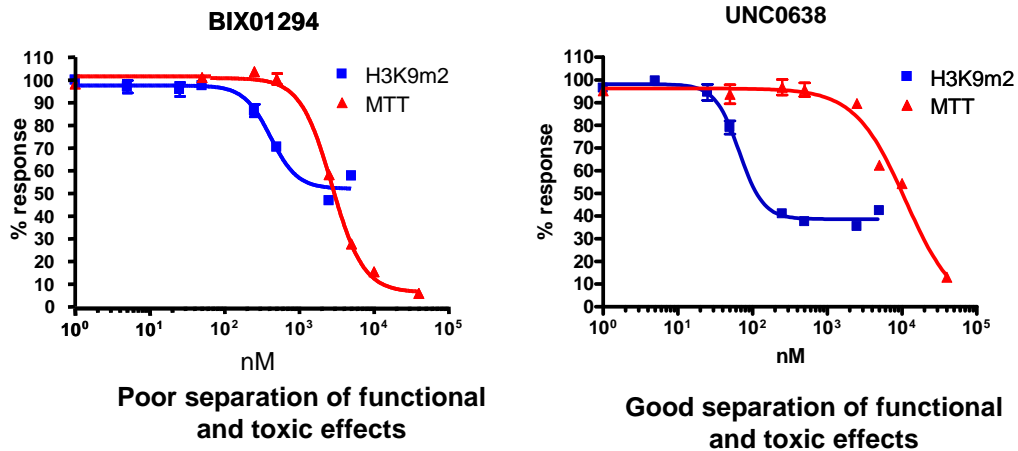


Fig 2 Effect of UNC638 and BIX-01294 on H3K9 dimethylation levels and cell toxicity measured by MTT assay at 48h in MDA-MB231 cells.



	In Vitro G9a IC ₅₀ (nM)	H3K9Me2 48h IC ₅₀ (nM)	Cell Tox 48h EC ₅₀ (nM)	Tox/Func Ratio
BIX-01294	133 ± 15	500 ± 43	2805	5.6
UNC638	<15	81 ± 9	11190	138