5-Carboxylate Substituted Piperazine-2-ones as Potent and Selective MMP-2 Inhibitors

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Matrix metalloproteinases (MMP) are metalloproteases. which contain zinc at their activation site.¹ These enzymes decompose and reconstruct structural proteins in the extracellular matrix such as membrane collagen, aggrecan. fibronectin, and laminin. As a result, MMPs play an important role in the development of chronic diseases such as multiple sclerosis, arthritis. fibrosis and other inflammation, and the growth and metastasis of malignant tumors.^{2,3} For this reasons, MMPs are attractive targets for drug candidates which can inhibit and potentially treat such diseases.⁴

The majority of MMPs are divided into four groups that include collagenase (MMP-1, -8, -13). gelatinase (MMP-2, -9). stromelysins (MMP-1, -10, -3) and membrane type MMPs (MMP-14, -15, 16, 17).⁵ Of these groups, the gelatinase MMPs are an attractive target because the inhibitors of MMP-2 are useful for anti-cancer therapy. Many MMP-2 inhibitors were developed and underwent clinical tests. However, owing to the inhibitors bearing low selectivity to MMP-2, the majority of these compounds were unsuccessful in the clinical tests.⁶ We, therefore, have concentrated our efforts on the development of selective MMP-2 inhibitors while sparing other MMPs. Herein we report the use of piperazinones bearing zinc binding groups (ZBG) for selective inhibition of MMP-2.

MMPs inhibitors based upon piperazinone have some advantages over substrate analog inhibitors. Substrate analog inhibitors suffer from hydrolysis by peptidase *in vivo*, whereas conformationally restricted peptidomimetics such as piperazinone are not hydrolyzed by peptidase and retain their activity *in vivo*.⁷ Moreover, piperazinone based inhibitors can be easily modified with diverse P₁ and P₂ substituents that can improve inhibitory potency and selectivity

(Figure 1).8

Firstly, we synthesized piperazinone derivatives containing a hydroxamic acid as the zinc binding group because the hydroxamic acid has the high affinity for the catalytic zinc ion in the active site of MMPs and can bind to Glu-202 and Ala-165 in active site of MMPs via hydrogen bond.⁹

The synthetic route for creating piperazinone derivatives containing a hydroxamic acid group was outlined in Scheme 1.10

Piperazinone derivatives were evaluated for inhibition of MMP-2, MMP-9 and MMP-1, Table 1.¹¹ Although these inhibitors strongly inhibited MMP-2, they suffered from just 10-fold selectivity over MMP-1. To improve selective inhibition of the MMP-2 over other MMPs, we examined piperazinone derivatives containing a carboxylic acid group as a selective MMP-2 inhibitor. By replacing the strong ZBG hydroxamic acid with the relatively less stronger ZBG carboxylic acid, we expected that the selectivity of the

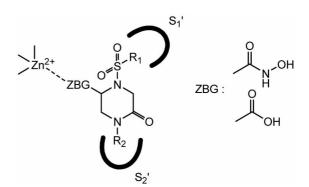
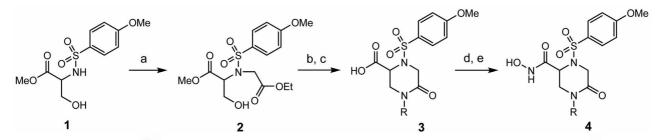


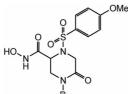
Figure 1. Schematic representation for binding mode of piperazinone based MMP-2 inhibitors.



Scheme 1. Reagents and conditions: (a) Ethyl bromoacetate, K_2CO_3 , DMF; (b) MsCl, Et₃N, CH_2Cl_2 ; R-NH₂, CH_2Cl_2 ; (c) LiOH, THF; (d) O-Benzyl-hydroxylamine, DCC; (e) Pd/C, H₂, EtOH

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Table 1. IC_{50} values for hydroxamic acid based inhibitors against MMP-1, MMP-2, and MMP-9



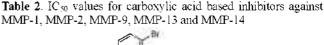
Compds	R -	$IC_{50}(\mu M)$		
		MMP-1	MMP-2	MMP-9
-4a	Н	0.089	0.051	0.036
4b	CH3	0.351	0.014	0.013
4c	CH ₃ (CH ₂) ₂	0.485	0.040	0.052
4d	Allyl	0.292	0.016	0.009
4e	Ph-CH ₂	0.013	0.005	0.015
4f	o-Pyridine-CH2 HCl	0.461	0.170	0146
4g	o-Pyridine-C ₂ H ₄ HCl	0.353	0.084	0.084
4h	Piperidine-C ₂ H ₄ HCl	0.176	0.030	0.027
-li	Morphorine-C ₂ H ₄ HCl	0.200	0.038	0.114

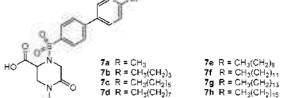
inhibitors against MMPs is strongly dependent on the substituents on the piperazinone derivatives. From X-ray crystallographic analysis, nuclear magnetic resonance (NMR) analysis and homology modeling, the S₁' pockets of MMP-2 are a relatively deeper than these of MMP-1 and, consequently, incorporation of an extended P₁' group (*e.g.*, bromo-biphenyl) leads to selective inhibition.^{12,13}

Carboxylic acid based inhibitors with bromo-biphenyl moiety were synthesized based upon Scheme I and evaluated for the inhibition of MMPs including MMP-1, MMP-2. MMP-9, MMP-13, and MMP-14, Table 2. As expected, compound 7a containing the bromo-biphenyl group as the P₁' group was found to have the high potency (IC₅₀ = 0.7 nM) for MMP-2 with the high selectivity over MMP-1 and MMP-14. However, compound 7a exhibited low selectivity over MMP-13 which also has a relatively deep hydrophobic S₁' pocket. We explored the structure activity relationship (SAR) of compound 7a with the intent of improving its selectivity while maintaining the inhibitory potency by modifying the P₂' substituent with versatile alkyl chains.

Modification of the P_2 ' substituent with different alkyl chains induced the strong effect on the inhibitory potency and selectivity. Especially, compound 7b was an active inhibitor (IC₅₀ = 1 nM) of MMP-2 with > 1000-fold selectivity over MMP-1, MMP-13, and MMP-14 by SAR. Moreover, compound 7b had the high inhibitory potency against MMP-2 with greater selectivity than the hydroxamic acid based inhibitor (Marimastat) and the carboxylic acid based inhibitor (BAY 12-9566).^{4,14}

In conclusion, carboxylic acid derivatives of piperazin-2one were synthesized and evaluated as the inhibitors of MMP-2. The derivatives were highly potent inhibitors of MMP-2 with > 1000 fold selectivity over MMP-1, MMP-13, MMP-14 and may thus be potentially useful scaffold in the Communications to the Editor





Compds	IC 50(µM)					
compas	MMP-1	MMP-2	MMP-9	MMP-13	MMP-14	
	3.857	0.0007	0.0016	0.008	>10	
7b	>25	0.001	0.152	1.130	>10	
7c	>25	0.001	0.076	0.0082	>10	
7d	>25	0.004	0.089	0.917	>10	
7e	>25	0.074	2.235	2.460	>10	
7f	>25	1.195	15.12	2.350	>10	
7g	>25	1.821	>25	>10	nd^a	
7h	>25	0.981	>25	>10	nda	
Marimastat	0.0008^{b}	0.0004 '	0.0008	0.0012°	nda	
BAY 12-9566	$>5^{b}$	0.011°	0.301*	1.47°	nd ^o	

^aIC₅₀ was not determined. ^bIC₅₀ Literature values (ref. 5, 14)

design of inhibitors that are effective for other zinc proteases of medicinal interest.

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