# Dimerization of a heat shock protein 90 inhibitor enhances inhibitory activity

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<sup>13</sup> C NMR	<u>3</u>
LCMS-ESI/UV	<u>4</u>
HRMS (ESI-TOF)	<u>5</u>
SM122-PEG-9 DIMER:	
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<sup>13</sup> C NMR	7
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## <sup>1</sup>H NMR SM122-PEG-5 DIMER



## <sup>13</sup>C NMR SM122-PEG-5 DIMER



## LCMS-ESI/UV SM122-PEG-5 DIMER





#### HRMS (ESI-TOF) SM122-PEG-5 DIMER

Exact mass=1921.0293, found 1921.0233 (M + Na<sup>+</sup> + H<sup>+</sup>)



## <sup>1</sup>H NMR SM122-PEG-9 DIMER



## <sup>13</sup>C NMR SM122-PEG-9 DIMER



#### LCMS-ESI/UV SM122-PEG-9 DIMER





Electronic Supplementary Material (ESI) for Organic & Biomolecular Chemistry This journal is The Royal Society of Chemistry 2013

Supplementary material

#### HRMS (ESI-TOF) SM122-PEG-9 DIMER

Exact mass=2097.1342, found 2097.1290  $(M + Na^{+} + H^{+})$ 





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## <sup>1</sup>H NMR SM122-PEG-4



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## <sup>13</sup>C NMR SM122-PEG-4



## LCMS-ESI/UV SM122-PEG-4 ==== Shimadzu LCMSsolution Analysis Report ====



#### HRMS (ESI-TOF) SM122-PEG-4

Exact mass=1038.5528, found 1038.5504  $(M + Na^{+})$ 





## <sup>1</sup>H NMR SM122-PEG-8



## <sup>13</sup>C NMR SM122-PEG-8





## LCMS-ESI/UV SM122-PEG-8 ==== Shimadzu LCMSsolution Analysis Report ====



## HRMS (ESI-TOF) SM122-PEG-8

Exact mass=1214.6577, found 1214.6552 (M + Na<sup>+</sup>)





Solubility Data Internal Standard Coumermycin A1 (10 μM; MW: 1110.08)



Co-injection of Coumermycin A1 (10 µM) and SM122 (MW: 782.44) solubility sample



Co-injection of Coumermycin A1 (10 µM) and SM122-PEG-5 DIMER solubility sample



Co-injection of Coumermycin A1 (10 µM) and SM122-PEG-9 DIMER solubility sample

# ==== Shimadzu LCMSsolution Analysis Report ====



Co-injection of Coumermycin A1 (10 µM) and SM122-PEG4 solubility sample

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Co-injection of Coumermycin A1 (100 µM) and SM122-PEG8 solubility sample



Test	CA-1	SM122-PEG-4	SM122-PEG-5	SM122-PEG-8	SM122-PEG-9	SM122
rest			DIMER		DIMER	
1	847455	514647				
2	718067	484622				
3	694257	433552				
1	698723		285549			
2	717285		240109			
3	651601		236551			
1	6995989			11897539		
2	6799004			11119149		
3	5775530			9614665		
1	869070				350779	
2	524656				263268	
3	526123				299311	
1	581530					41858
2	542472					66946
3	805189					45169

#### **Supplemental Table 1: Peak area values**

#### **Supplemental Table 2: Solubility values**

11	v	
Compound	(µg/mL)	(μM)
SM122	$0.66 \pm 0.28$	$0.84 \pm 0.35$
SM122-PEG-4	$6.45 \pm 0.36$	$6.36 \pm 0.35$
SM122-PEG-5 DIMER	$7.00 \pm 0.71$	$3.69 \pm 0.37$
SM122-PEG-8	$198.64 \pm 3.89$	$166.69 \pm 3.27$
SM122-PEG-9 DIMER	$10.18 \pm 1.72$	$4.91\pm0.83$

#### **Calculation method:**

Solubility ( $\mu$ M) = (peak area of compound/ peak area of CA-1) × concentration of CA-1<sup>\*</sup> Solubility ( $\mu$ g/mL) = Solubility ( $\mu$ M) × MW of compound (g/mol) × 10<sup>-3</sup> (L/mL) × 10<sup>6</sup> ( $\mu$ g/g) <sup>\*</sup> For compound SM122, SM122-PEG-5 DIMER, SM122-PEG-9 DIMER, and SM122-PEG-

4, concentration of CA-1 is 10  $\mu$ M; for SM122-PEG-8, concentration of CA-1 is 100  $\mu$ M.

**Solubility Method:** ~5 mg of SM122-PEG conjugates or DIMERs was stirred in 0.3 mL of Milli-Q water for over 24 hours. The filtrate was filtered through a 45 micron filter to remove all insoluble particulates. The concentration of each sample was measured by LCMS utilizing Coumermycin A1 (CA-1) with known concentrations as the internal standard. The solubility value in water was then calculated (Table 1).

#### **Biological data**





**Supplemental Figure 1a-c:** Luciferase activity or % inhibition of luciferase activity in rabbit reticulocyte lysate (RRL) refolding system treated with indicated concentrations of 17AAG, SM122, PEG conjugates or DIMERs. RRL was pre-incubated with compounds for 5 hours. Error bars indicate the standard deviation.

Compound	Concentration (µM)	% Inhibition of luciferase activity
17AAG	1	43.34 ± 3.15
SM122	1	$27.27 \pm 4.82$
SM122-PEG-4	1	$2.94 \pm 1.72$
SM122-PEG-5 DIMER	1	$52.56 \pm 1.22$
SM122-PEG-8	1	$3.63 \pm 1.32$
SM122-PEG-9 DIMER	1	$53.30 \pm 2.61$

## Supplemental Table 3: Inhibition of Luciferase activity.



**Supplemental Figure 2a-b:** The inhibitory effects of 17AAG, SM122, SM122-PEG-8 and SM122-PEG-9 DIMER on the binding between Hsp90 and its co-chaperone HOP and FKBP52.