## **Supporting Information**

## Development of Novel Bone Targeting Peptide-Drug Conjugate of 13aminomethyl-15-thiomatrine for Osteoporosis Therapy

Jia Su<sup>a†</sup>, Chao Liu<sup>b†</sup>, Haohao Bai<sup>b</sup>, Wei Cong<sup>b</sup>, Hua Tang<sup>b</sup>, Honggang Hu<sup>b</sup>, Li Su<sup>b\*</sup>, Shipeng He<sup>b\*</sup> and Yong Wang<sup>a\*</sup>

 <sup>a</sup> Department of Orthopaedics, Wenzhou Hospital of Integrated Traditional Chinese and Western Medicine, Zhejiang, China.
<sup>b</sup> Institute of Translational Medicine, Shanghai University, Shanghai, China.



Scheme S1. The synthesis route of BTM19-4. Reagents and conditions: a) TFE/DCM (1:3, v/v), rt, 4 h, 87%; b) 4-aminophenyl methanol, HOBt, DIC, DMF, rt, 2 h, 74%; c) i) Triphosgene, activated carbon, THF, rt, 12 h; ii) M19, Et<sub>3</sub>N, DMF, rt, 12 h, 76% in two steps; d) TFA/water/EDT/TIPs (95:2:2:1, v/v/v/v), rt, 2 h, 59%.



Figure S1. (A) Illustration of the chemical stability study of M19 vs. PDCs in water/MeCN solution at room temperature; Chromatograms of chemical stability study of M19 (B), BTM19-1 (C), BTM19-2 (D), BTM19-3 (E).



**Figure S2.** (A) Mechanism of the drug release from self-immolative spacer PABC and Cathepsin K substrate; Chromatograms of drug release study of **BTM19-1** (B), **BTM19-2** (C) and **BTM19-3** (D) at pH 5.5 (37 °C) with cathepsin K.



Figure S3. Chromatograms of proteolytic stability study of BTM19-1 (A), BTM19-2 (B), BTM19-3 (C) under a-chymotrypsin treatment.



Figure S4. Chromatograms of binding study of BTM19-1 (A), BTM19-2 (B), BTM19-3 (C) and BTM19-4 (D) to hydroxyapatite at pH=5.5 and 37 °C.



Figure S5. Quantification data of cytotoxic study after M19 (A), BTM19-1 (B), BTM19-2 (C) and BTM19-3 (D) treatment on RAW264.7 cell measured by CCK-8 assay. Data points are displayed as the mean value SEM of duplicate independent experiments. (\*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001).



Figure S6. Formation of TRAP-positive cells from RAW264.7 cells with RANKL, M-CSF and treated with M19 (A), BTM19-1 (B), BTM19-2 (C) and BTM19-3 (D).



**Figure S7. 6** as white powder, 1.43 g, 82% yield. **A**) Structure of **BTM19-1-2**; **B**) HPLC trace of purified **BTM19-1-2**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-1-2** (calcd. for C<sub>81</sub>H<sub>132</sub>N<sub>12</sub>O<sub>28</sub>S 1752.8995; found [M+2H]<sup>2+</sup> 877.4609).



**Figure S8.** 7 as white powder, 1.12 g, 79% yield. **A**) Structure of **BTM19-1-3**; **B**) HPLC trace of purified **BTM19-1-3**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-1-3** (calcd. for C<sub>88</sub>H<sub>139</sub>N<sub>13</sub>O<sub>28</sub>S 1857.9573; found [M+2H]<sup>2+</sup> 929.9841).



**Figure S9. 8** as white powder, 1.1 g, 78% yield. **A**) Structure of **BTM19-1-4**; **B**) HPLC trace of purified **BTM19-1-4**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-1-4** (calcd. for C<sub>105</sub>H<sub>164</sub>N<sub>16</sub>O<sub>29</sub>S<sub>2</sub> 2177.1292; found [M+2H]<sup>2+</sup> 1089.5528).



Figure S10. BTM19-1 as lyophilized white powder, 492 mg, 62% yield. A) Structure of BTM19-1; B) HPLC trace of purified BTM19-1. Gradient: 90-0% of buffer B in 15 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). C) HR-MS spectrum of BTM19-



1 (calcd. for  $C_{68}H_{100}N_{16}O_{26}S$  1588.6715; found [M+H]<sup>+</sup> 1589.6329).

**Figure S11. 9** as white powder, 1.5 g, 83% yield. **A**) Structure of **BTM19-2-2**; **B**) HPLC trace of purified **BTM19-2-2**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-2-2** (calcd. for C<sub>83</sub>H<sub>136</sub>N<sub>12</sub>O<sub>29</sub>S 1796.9257; found [M+2H]<sup>2+</sup> 899.4961).



**Figure S12. 10** as white powder, 1.18 g, 75% yield. **A**) Structure of **BTM19-2-3**; **B**) HPLC trace of purified **BTM19-2-3**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-2-3** (calcd. for C<sub>90</sub>H<sub>143</sub>N<sub>13</sub>O<sub>29</sub>S 1901.9835; found [M+2H]<sup>2+</sup> 951.9905).



**Figure S13. 11** as white powder, 1.07 g, 85% yield. **A**) Structure of **BTM19-2-4**; **B**) HPLC trace of purified **BTM19-2-4**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-2-4** (calcd. for C<sub>107</sub>H<sub>168</sub>N<sub>16</sub>O<sub>30</sub>S<sub>2</sub> 2221.1554; found [M+2H]<sup>2+</sup> 1111.5709).



**Figure S14. BTM19-2** as lyophilized white powder, 493 mg, 63% yield. A) Structure of **BTM19-2**; B) HPLC trace of purified **BTM19-2**. Gradient: 90-0% of buffer B in 15 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). C) HR-MS spectrum of **BTM19-2** (calcd. for C<sub>70</sub>H<sub>104</sub>N<sub>16</sub>O<sub>27</sub>S 1632.9678; found [M+2H]<sup>2+</sup> 817.4979).



**Figure S15. 12** as white powder, 1.47 g, 80% yield. **A**) Structure of **BTM19-3-2**; **B**) HPLC trace of purified BTM19-3-2. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-3-2** (calcd. for C<sub>85</sub>H<sub>140</sub>N<sub>12</sub>O<sub>30</sub>S 1840.9519; found [M+2H]<sup>2+</sup> 921.4898).



**Figure S16. 13** as white powder, 1.15 g, 74% yield. **A**) Structure of **BTM19-3-3**; B) HPLC trace of purified **BTM19-3-3**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). C) HR-MS spectrum of **BTM19-3-3** (calcd. for C<sub>92</sub>H<sub>147</sub>N<sub>13</sub>O<sub>30</sub>S 1946.0098; found [M+2H]<sup>2+</sup> 974.0215).



**Figure S17. 14** as white powder 1.09 g, 86% yield. **A**) Structure of **BTM19-3-4**; B) HPLC trace of purified **BTM19-3-4**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-3-4** (calcd. for C<sub>109</sub>H<sub>172</sub>N<sub>16</sub>O<sub>31</sub>S<sub>2</sub> 2265.1816; found [M+2H]<sup>2+</sup> 1133.5828).



**Figure S18. BTM19-3** as lyophilized white powder, 443 mg, 60% yield. A) Structure of **BTM19-3**; **B**) HPLC trace of purified **BTM19-3**. Gradient: 90-0% of buffer B in 15 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). C) HR-MS spectrum of **BTM19-3** (calcd. for C<sub>72</sub>H<sub>108</sub>N<sub>16</sub>O<sub>28</sub>S 1676.7240; found [M+2H]<sup>2+</sup> 839.3631).



**Figure S19. 16** as white powder, 1.6 g, 87% yield. **A**) Structure of **BTM19-4-2**; **B**) HPLC trace of purified BTM19-4-2. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-4-2** (calcd. for C<sub>90</sub>H<sub>150</sub>N<sub>14</sub>O<sub>24</sub>S 1843.0668; found [M+2H]<sup>2+</sup> 922.5041).



**Figure S20. 17** as white powder 1.25 g, 74% yield. **A**) Structure of **BTM19-4-3**; **B**) HPLC trace of purified **BTM19-4-3**. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-4-3** (calcd. for C<sub>97</sub>H<sub>157</sub>N<sub>15</sub>O<sub>24</sub>S 1948.1247; found [M+2H]<sup>2+</sup> 975.0709).



**Figure S21. 18** as white powder, 1.05 g, 76% yield. **A**) Structure of **BTM19-4-4**; **B**) HPLC trace of purified BTM19-4-4. Gradient: 90-0% of buffer B in 20 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-4-4** (calcd. for C<sub>114</sub>H<sub>182</sub>N<sub>18</sub>O<sub>25</sub>S<sub>2</sub> 2267.2965; found [M+3H]<sup>3+</sup> 756.7441).



**Figure S22. BTM19-4** as lyophilized white powder, 458 mg, 59% yield. **A**) Structure of BTM19-4; **B**) HPLC trace of purified **BTM19-4**. Gradient: 90-0% of buffer B in 15 min with C18 column (5  $\mu$ m, 2.5 mm×250 mm). **C**) HR-MS spectrum of **BTM19-4** (calcd. for C<sub>80</sub>H<sub>126</sub>N<sub>18</sub>O<sub>20</sub>S 1690.9116; found [M+2H]<sup>2+</sup> 846.4346).



Figure S23. ESI-MS spectrum of 2 (calculated for  $C_{15}H_{22}N_2S$  262.15; found [M+H]<sup>+</sup> 263.2).



Figure S24. <sup>1</sup>H-NMR data of 2.



Figure S25. <sup>13</sup>C-NMR data of 2.



Figure S26. ESI-MS spectrum of 3 (calculated for  $C_{16}H_{27}N_3S$  293.19; found [M+H]<sup>+</sup> 294.2).



Figure S27. <sup>1</sup>H-NMR data of 2.



Figure S28. <sup>13</sup>C-NMR data of 2.