# **Supporting Information**

## pDobz/pDobb Protected Diaminodiacid as Novel Building Block for Peptide Disulfide-bond Mimics Synthesis

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### 1. Experimental Section

#### 1.1 Synthesis of contryphan-Vn

To investigate reduction stability and protease stability, we synthesized the native model peptide contryphan-Vn shown in Scheme S1.



**Scheme S1.** Route of synthesis of contryphan-Vn; reagents and conditions: a) TFA/EDT/TIPs/water (95:2:2:1, v/v/v/v), rt, 2 h; b) Guanidine hydrochloride, sodium dihydrogen phosphate, PBS buffer (containing 10% DMSO), rt, 24 h; The resinbound peptides were protected on side chains at asterisk sites. The following protecting groups for amino acid side chains were used: tert-butyl (tBu; for Asp), trityl (Trt; for Cys) and tert-butyloxycarbonyl (Boc; for Lys and Trp).

## 1.2 Characterization of contryphan-Vn



**Figure S1.** Contryphan-Vn was obtained as a white powder (46.8 mg, 43% yield according to initial resin load, 97.2% purity). a) HPLC trace of purified contryphan-Vn. Gradient: 90-0% of buffer B in 25 min with C18 column (5  $\mu$ m, 4.6×250 mm). b) HR-MS spectrum of contryphan-Vn (calc. for C<sub>50</sub>H<sub>65</sub>N<sub>13</sub>O<sub>11</sub>S<sub>2</sub> 1087.4368, found [M+H]<sup>+</sup> 1088.4383).

#### 2. HRMS data of small molecular compounds



e S2. HR-MS spectrum of compound 6 (calc. for  $C_8H_{14}N_4O_4$  230.1015, found [M-H]<sup>-</sup> 229.1149).



re S3. HR-MS spectrum of compound 7 (calc. for  $C_{30}H_{40}B_2N_4O_8$  606.3032, found  $[M+H]^+$  607.3137).



**re S4.** HR-MS spectrum of compound 9 (calc. for  $C_{24}H_{25}NO_4$  391.1784; found  $[M+H]^+$  392.1682).



**ure S5.** HR-MS spectrum of compound 10 (calc. for  $C_{54}H_{65}B_2N_5O_{12}$  997.4816; found [M+H]<sup>+</sup> 998.4977).



gure S6. HR-MS spectrum of compound 4 (calc. for  $C_{50}H_{57}B_2N_5O_{12}$  941.4190; found [M-H]<sup>-</sup> 940.4025).

## 3. <sup>1</sup>H and <sup>13</sup>C NMR spectra of compounds





180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 -10 f1 (ppm)









