

Supplementary Information for

Elucidation of inhibitor-binding pocket of D-amino acid oxidase using docking simulation and *N*-sulfanylethylanilide-based labeling technology

Taiki Kohiki,^{a†} Yusuke Kato,^{b†} Yusuke Nishikawa,^a Kazuko Yorita,^b Ikuko Sagawa,^c Masaya Denda,^a Tsubasa Inokuma,^a Akira Shigenaga^{*a,d} Kiyoshi Fukui,^{*b} and Akira Otaka^{*a}

^a Institute of Biomedical Sciences and Graduate School of Pharmaceutical Sciences, Tokushima University, Shomachi, Tokushima 770-8505, Japan.

^b Division of Enzyme Pathophysiology, The Institute for Enzyme Research (KOSOKEN), Tokushima University, 3-18-15 Kuramoto, Tokushima 770-8503, Japan.

^c Support Center for Advanced Medical Sciences, Institute of Biomedical Sciences, Tokushima University Graduate School, Shomachi, Tokushima 770-8505, Japan

^d PRESTO, Japan Science and Technology Agency (JST), 4-1-8 Honcho, Kawaguchi, Saitama 332-0012, Japan.

[†]TK and YK contributed equally to this work.

E-mail:

shigenaga.akira@tokushima-u.ac.jp (A. Shigenaga)

kiyo.fukui@tokushima-u.ac.jp (K. Fukui)

aotaka@tokushima-u.ac.jp (A. Otaka)

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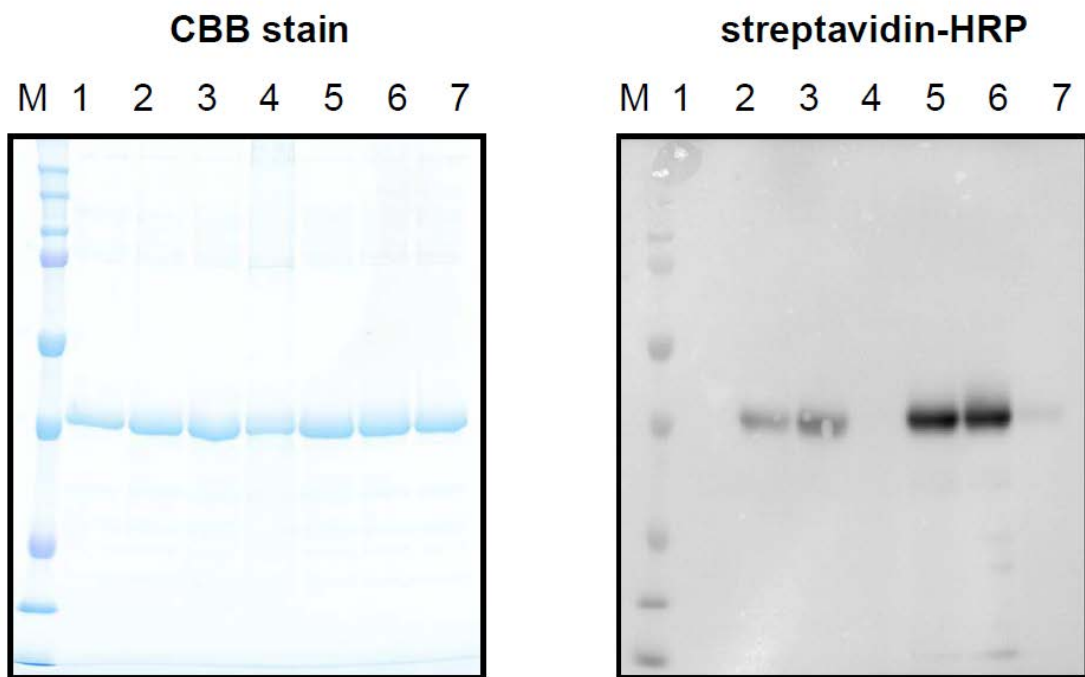


Fig. S1 An original picture of Fig. 4C. Lane 4 and 7 in Fig. S1 correspond to lane 1 and 2 in Fig. 4C, respectively.

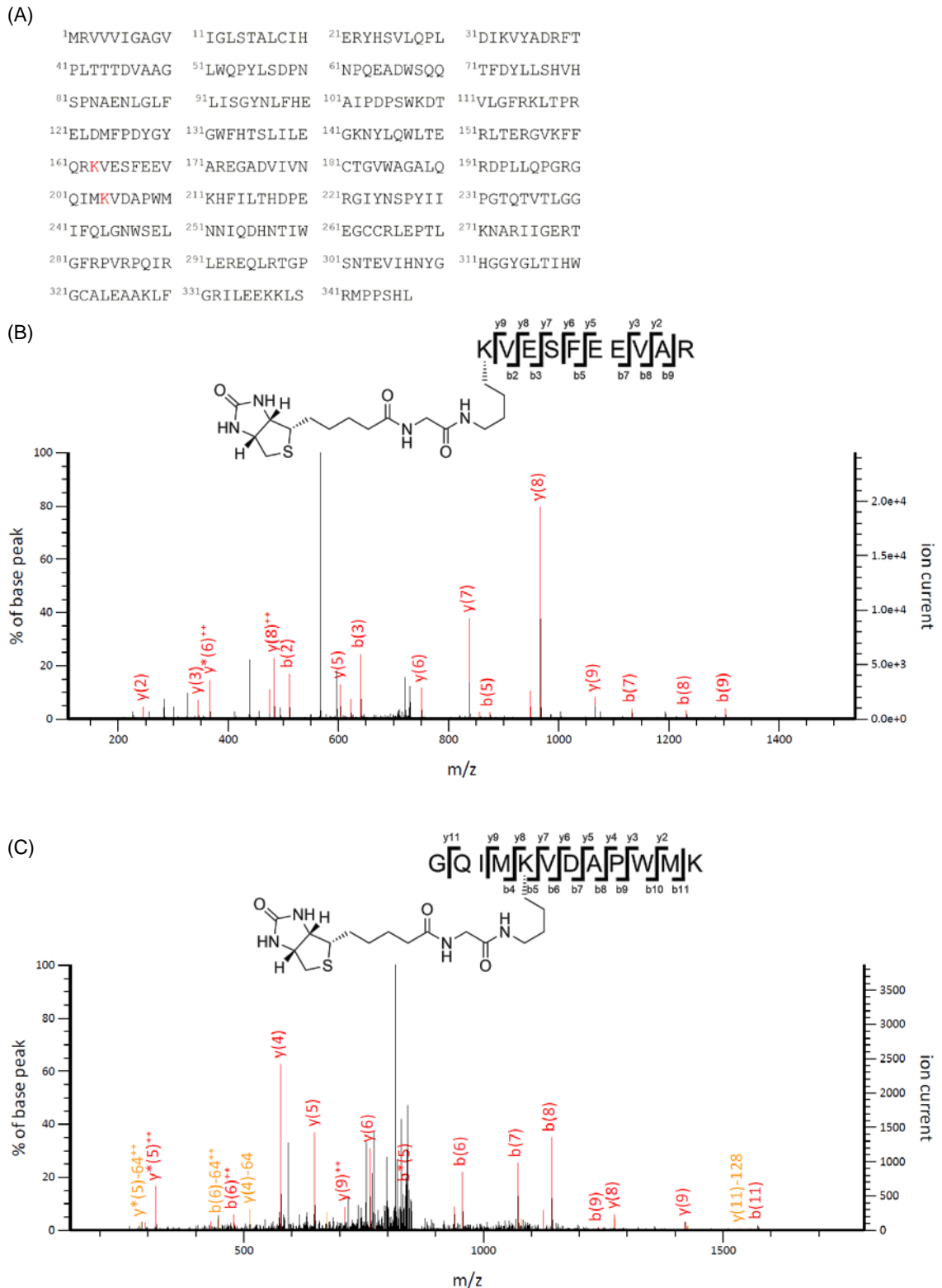
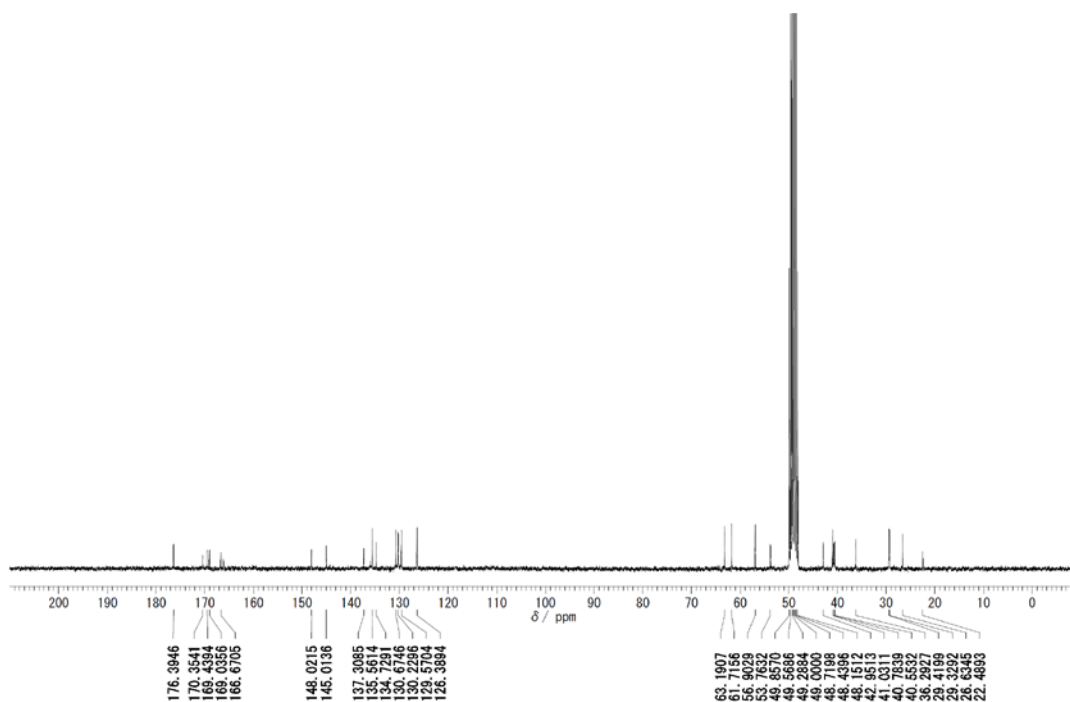
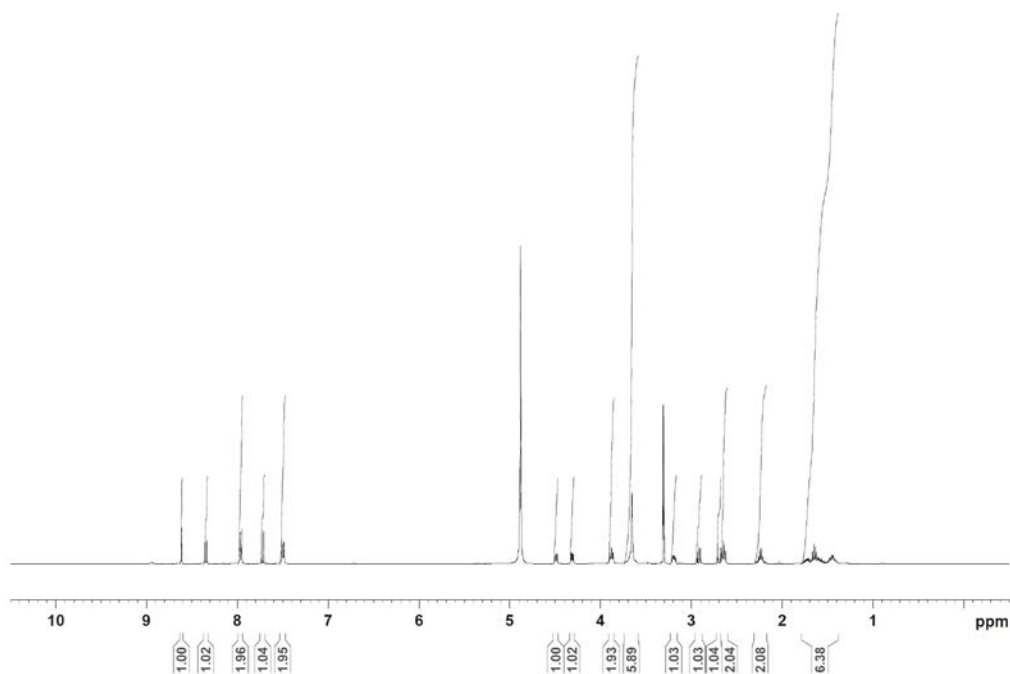
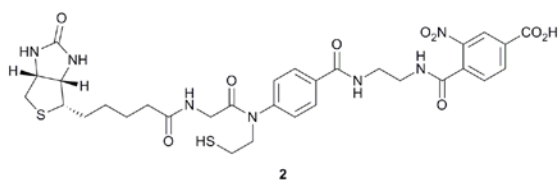
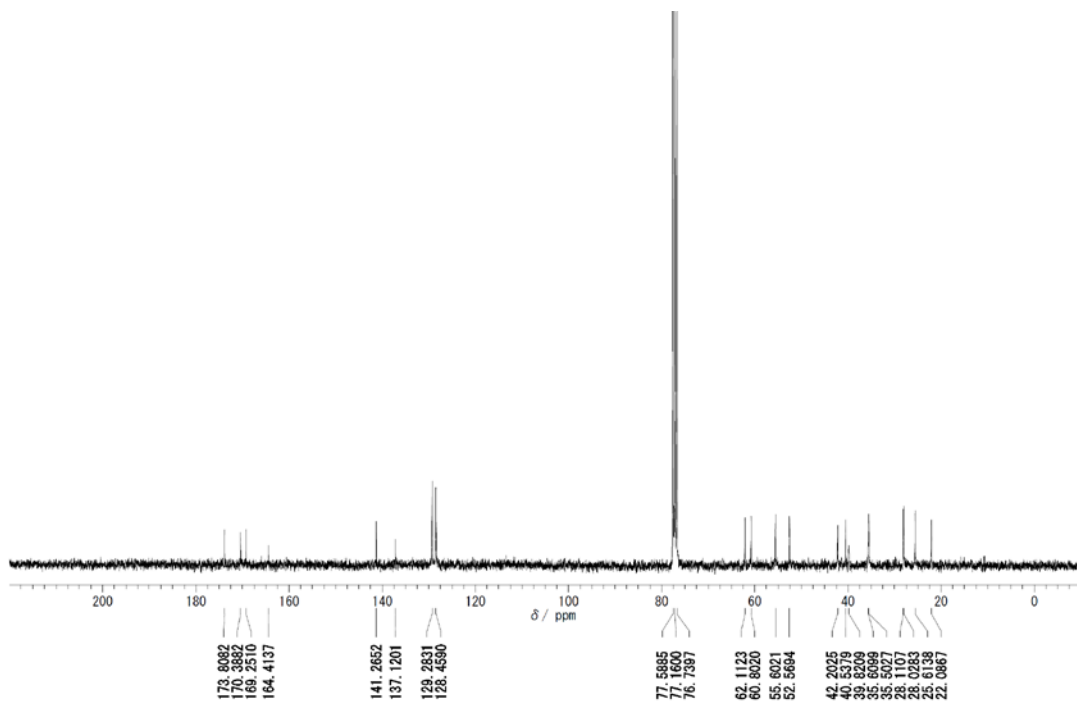
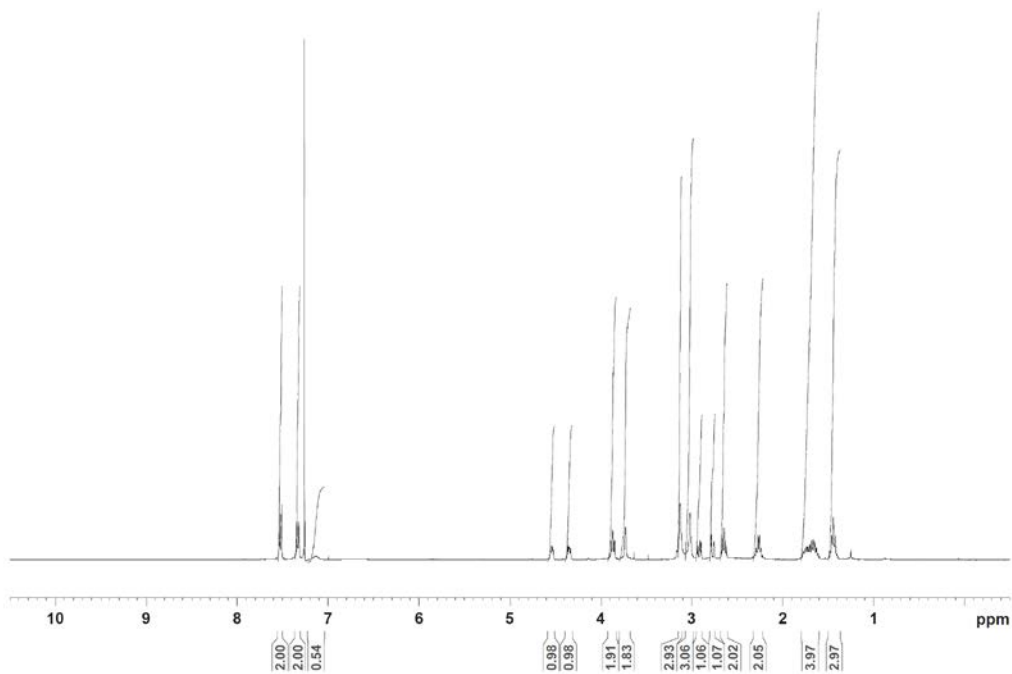
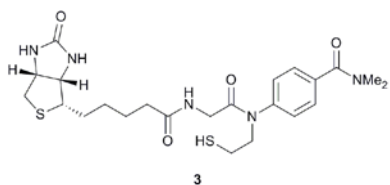
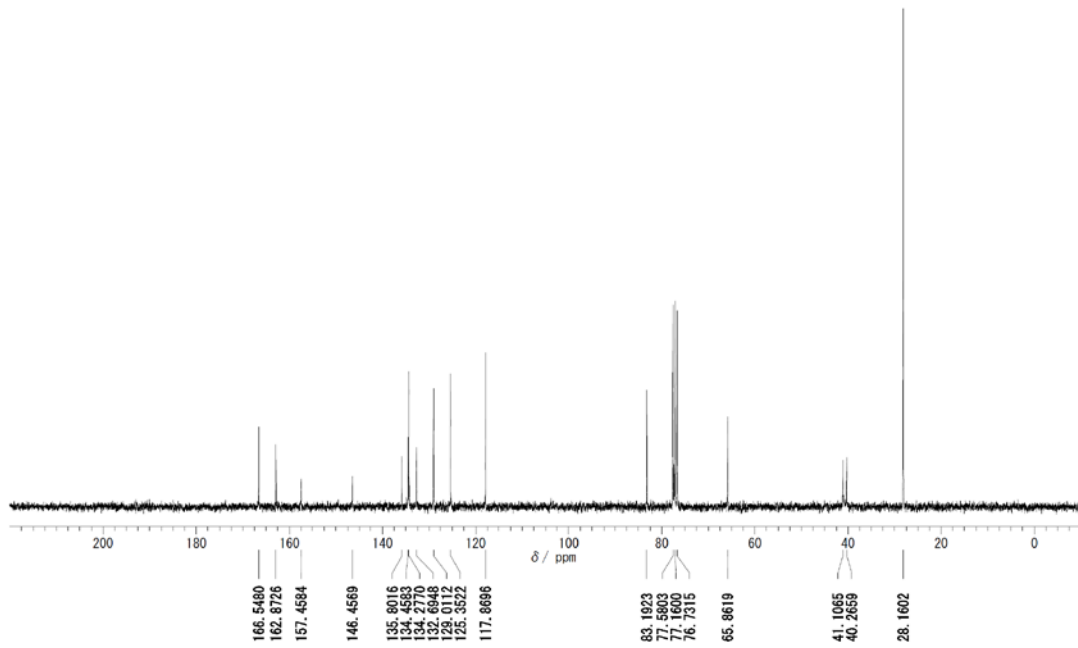
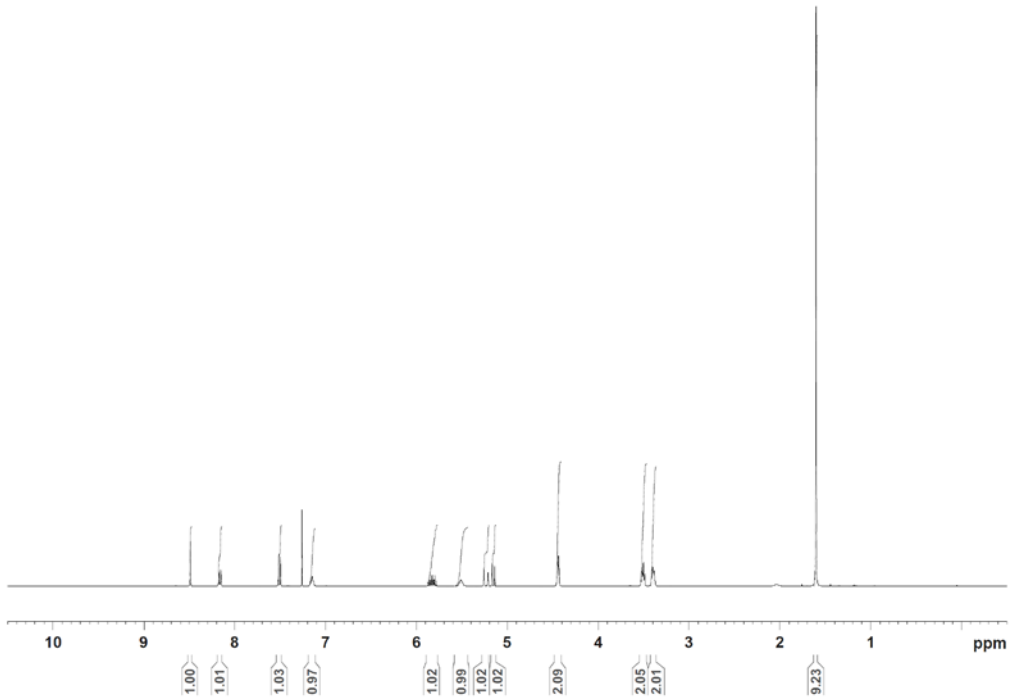
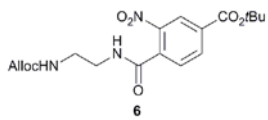
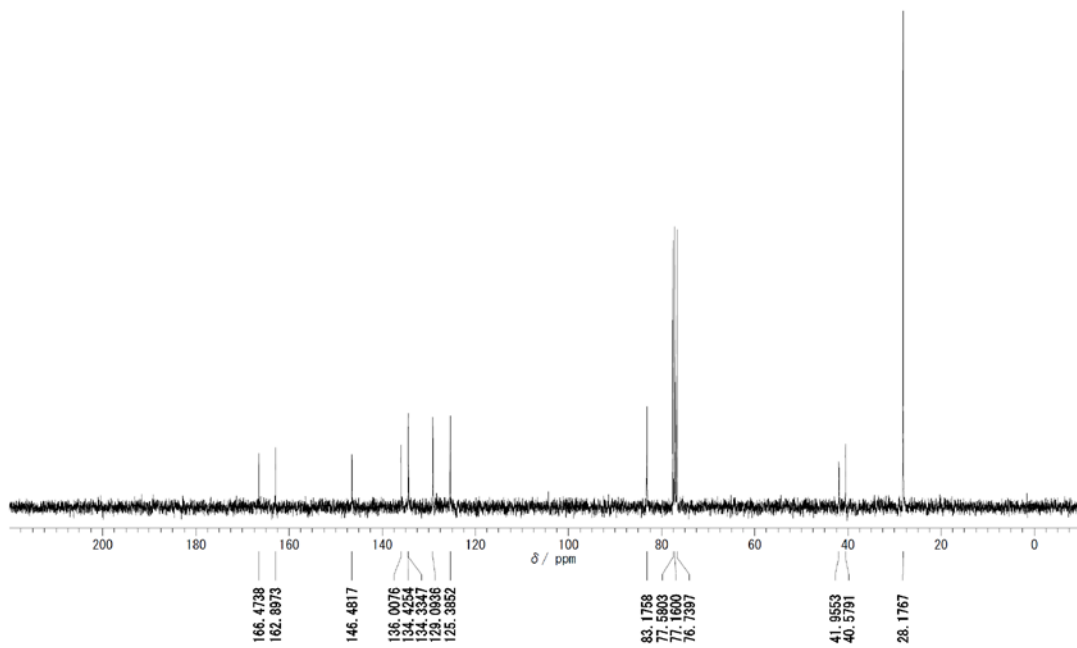
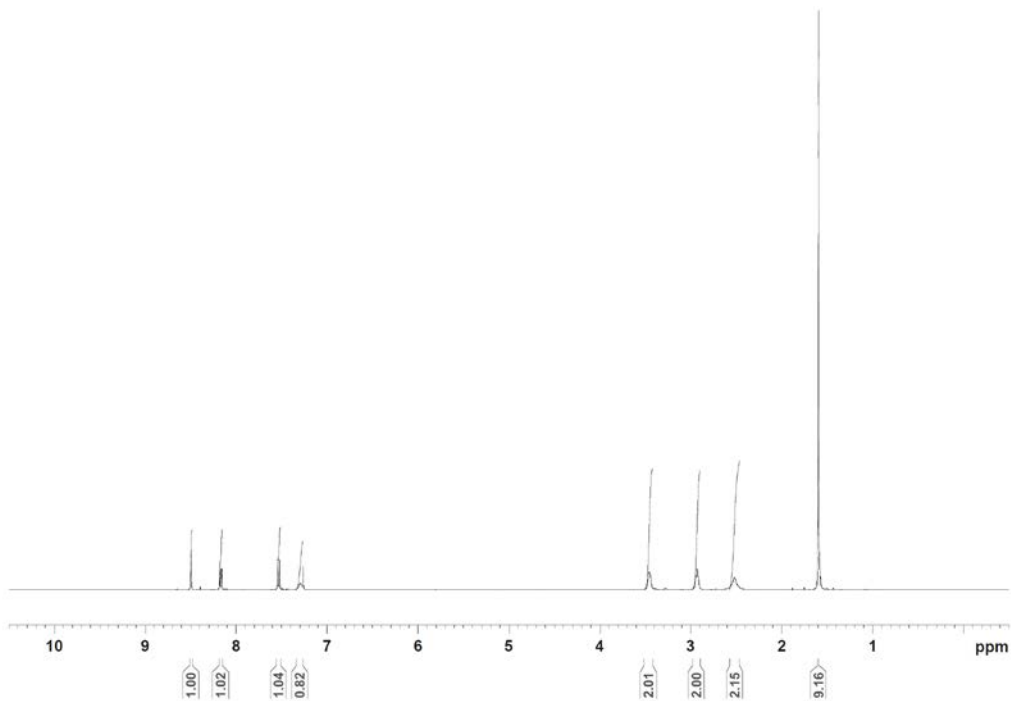
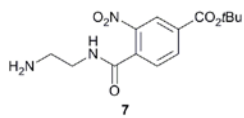


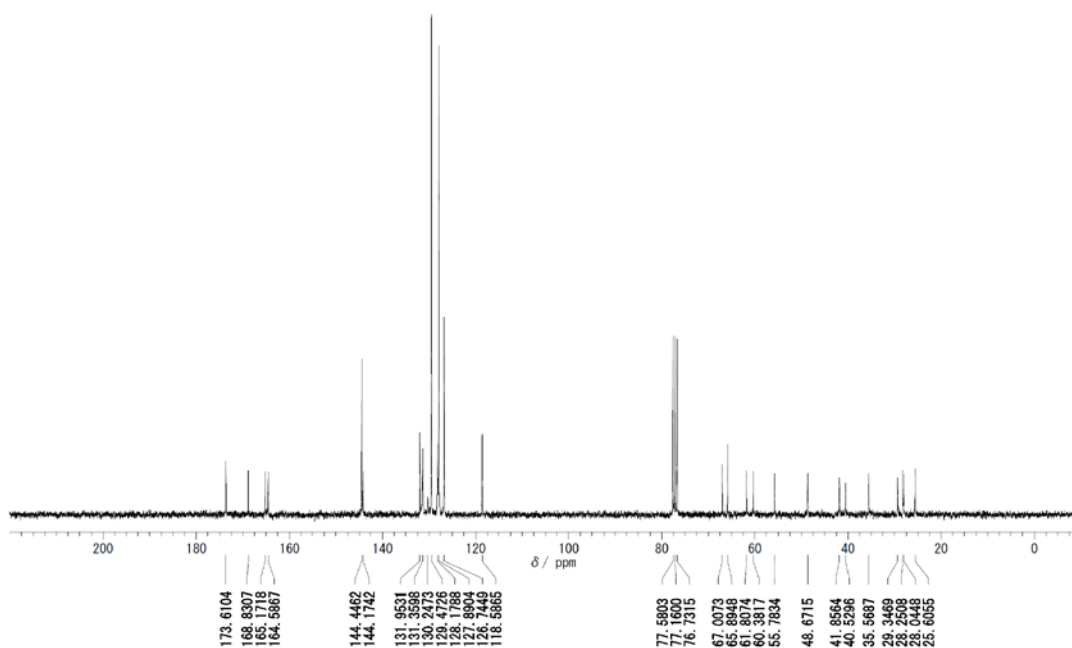
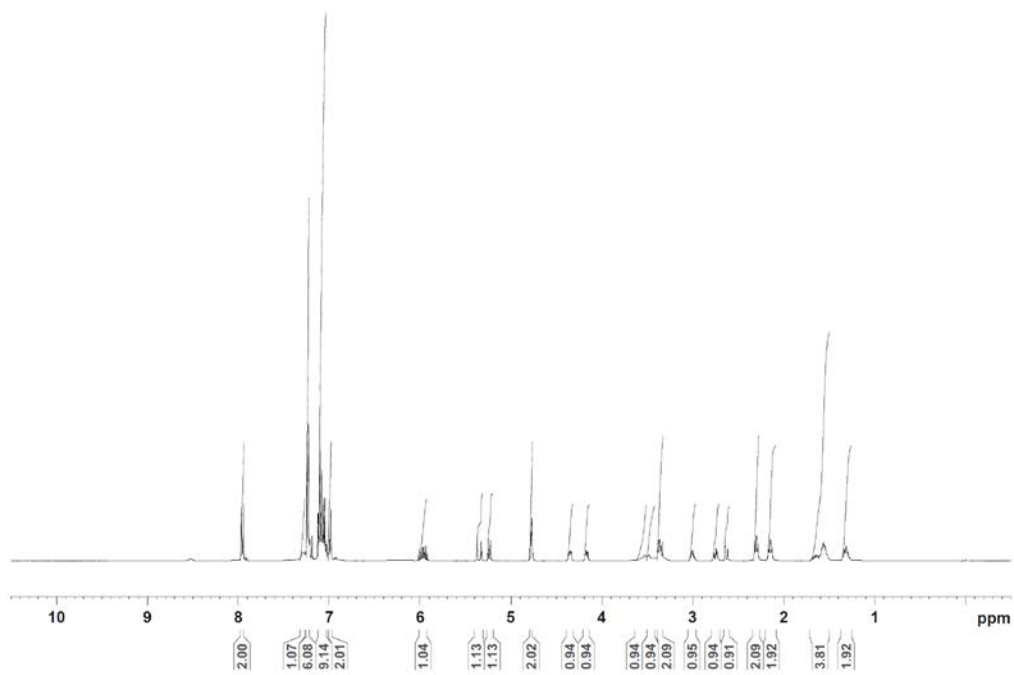
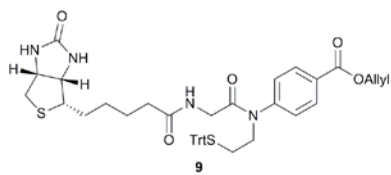
Fig. S2 Mass spectral analysis of the labeled residues of DAO using SEAL reagent 2. (A) A primary sequence of DAO. (B and C) nanoLC-MS/MS analysis of the fragments labeled at Lys163 (B) and 204 (C).

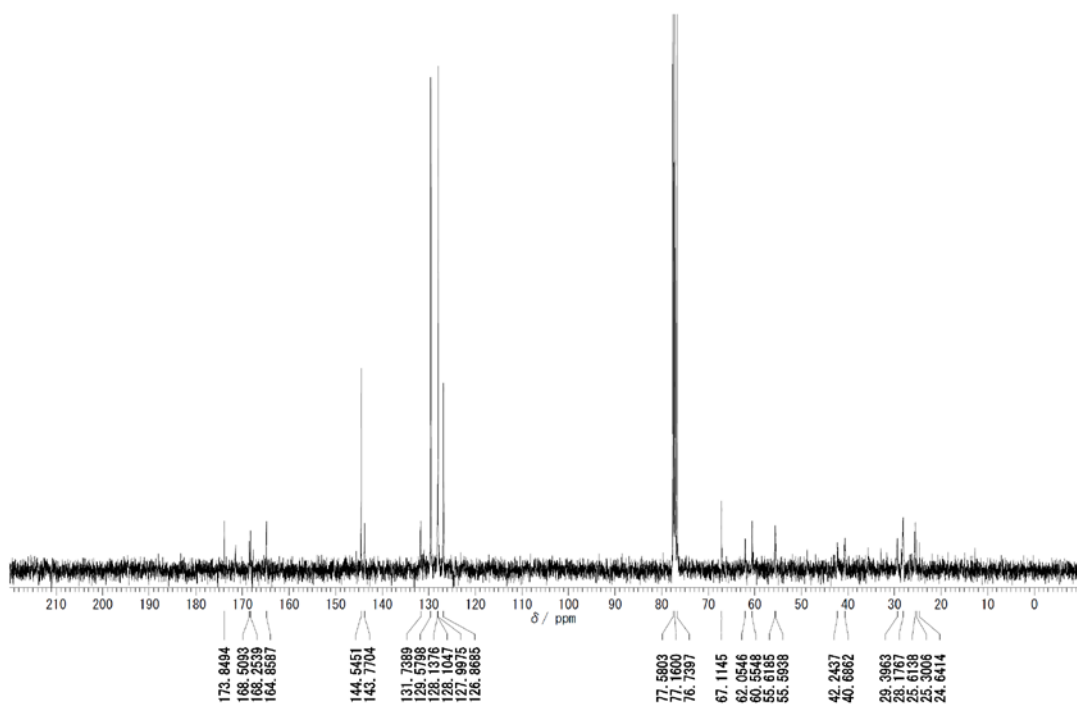
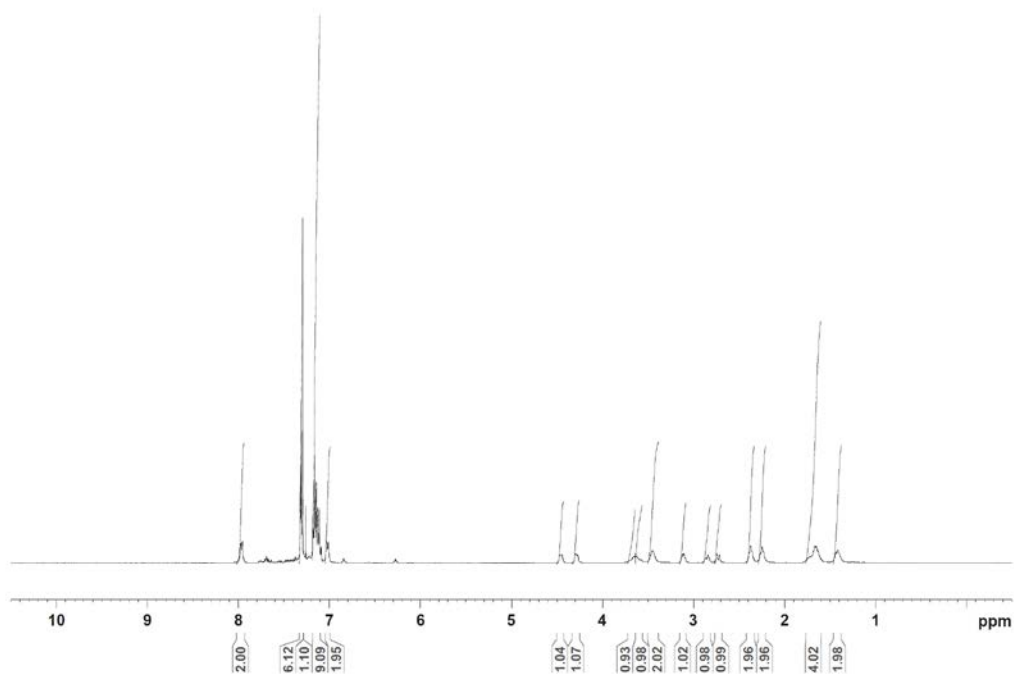
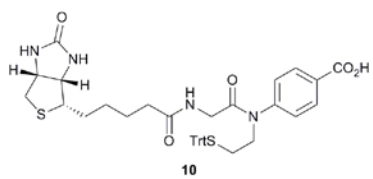


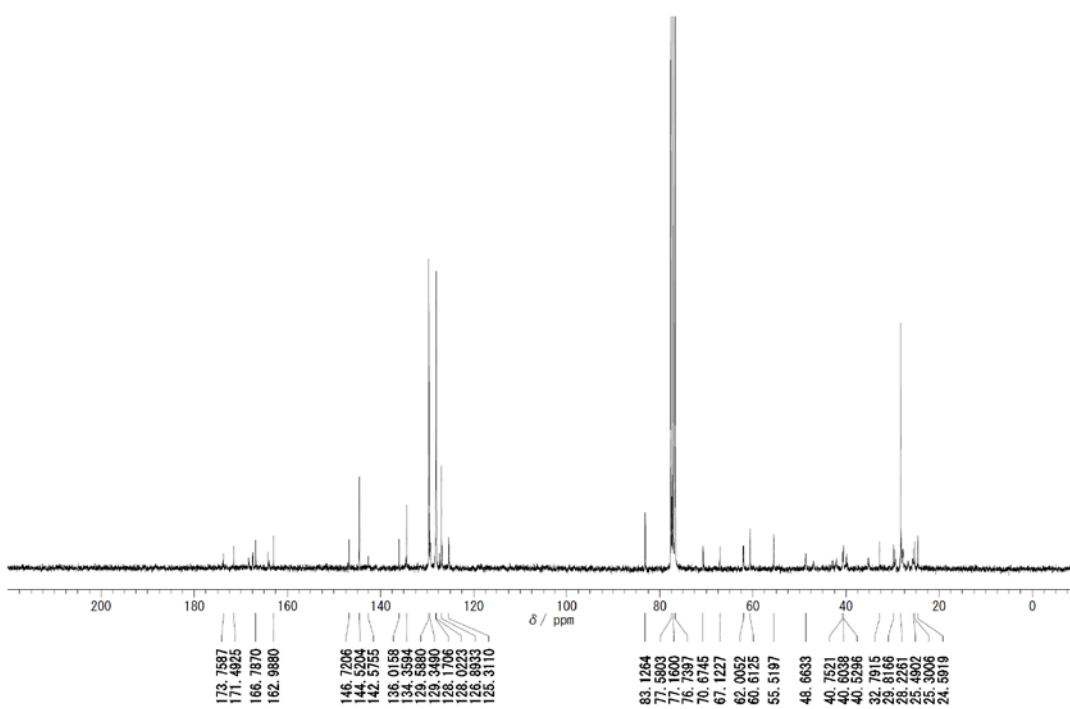
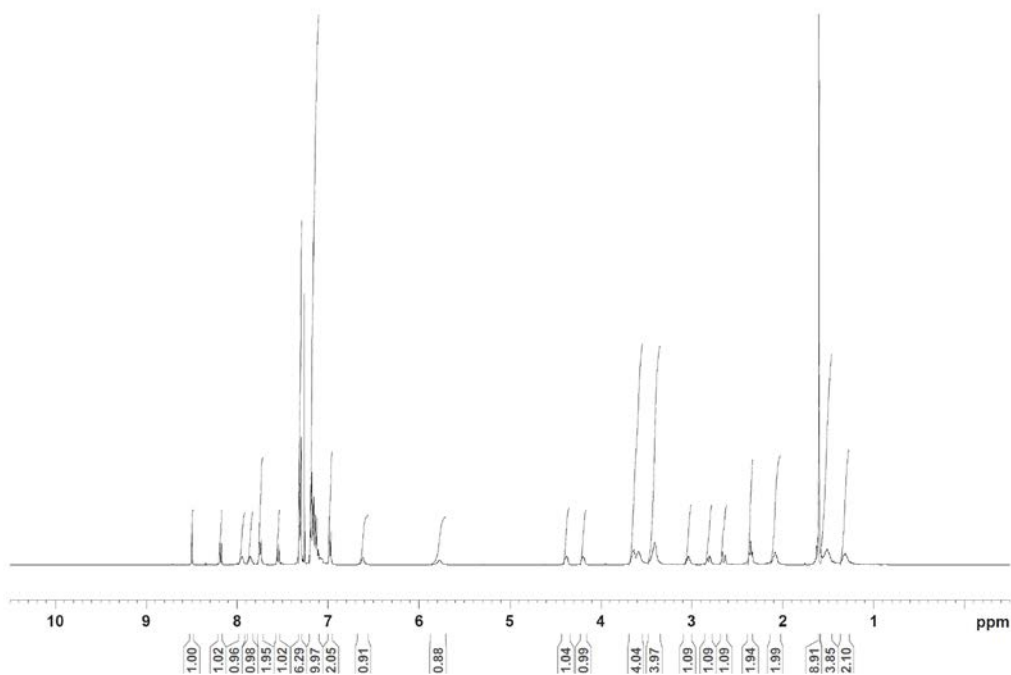
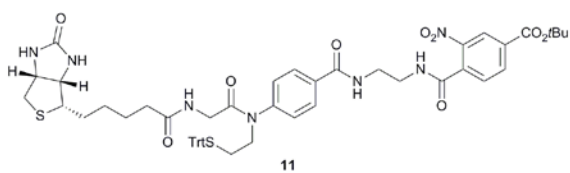


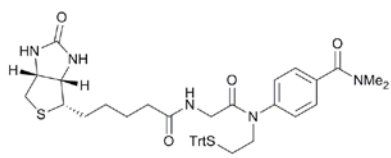












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