Exo-Linker: Positional Reconfiguration Driving Significant Advances in ADC Stability and Efficacy

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Abstract

Antibody-drug conjugates (ADCs) have transformed targeted cancer therapy by combining the specificity of monoclonal antibodies with the cytotoxic potency of small-molecule drugs. However, payload instability, hydrophobicity, and premature cleavage limit their efficacy and safety. This study presents Exo-Linker technology as a novel solution to these issues. By repositioning cleavable peptide linkers like Glu-Val-Cit and Glu-Glu-Val-Cit at the exo-position of the p-aminobenzyl carbamate moiety, Exo-Linkers improve stability, hydrophilicity, and resistance to enzymatic degradation.

Key findings highlight the superior pharmacokinetics, tumor-suppressive efficacy, and enzymatic stability of Exo-Linker ADCs in preclinical models, significantly outperforming traditional Val-Cit-based linkers. Integration with the second-generation AJICAP platform broadens therapeutic windows, ensures precise drug-to-antibody ratio control, and minimizes aggregation. Exo-Linkers establish a new standard for ADC development, overcoming critical limitations of traditional linkers while enabling safer and more effective cancer treatments. This innovative approach redefines the therapeutic landscape, enhances patient outcomes, and broadens the scope of ADC applications.

Keywords: antibody-drug conjugate, stable linker, site-specific conjugation, enzymatic cleavable linker, AJICAP

1. Introduction

The rapid advancement of Antibody-drug conjugates (ADCs) has revolutionized targeted cancer therapies, providing a powerful strategy to enhance therapeutic precision and efficacy^{1 2}. ADCs utilize the specificity of monoclonal antibodies (mAbs) to deliver highly cytotoxic small-molecule payloads directly to tumor cells through specialized chemical linkers, minimizing systemic toxicity. Over 13 ADCs have received U.S. Food and Drug Administration (FDA) approval, with more than 100 candidates in various stages of clinical development ³. Despite this progress, significant challenges persist, particularly regarding conventional linker technologies' stability, safety, and efficacy limitations (Figure 1) ^{4 5}. This study presents the Exo-Linker platform, a novel approach that overcomes these issues by incorporating advanced linker chemistry, conjugation methods, and ADC optimization to enhance therapeutic outcomes ^{6 7}.



Figure 1. Illustration of Exo-linker technology. a) traditional Val-Cit linker, b) Exo-cleavable linker

2. Current Challenges in Linker Technology

2.1. Role of Linkers in ADC Design

Linkers are essential to ADC functionality, directly impacting structural homogeneity, pharmacokinetics (PK), and therapeutic safety margins. They are broadly classified as cleavable or non-

cleavable. Linkers must remain stable in systemic circulation regardless of type to prevent premature payload release while ensuring efficient and selective drug delivery at the tumor site. Cleavable linkers, especially enzyme-cleavable designs, are preferred for clinical applications. The valine-citrulline (Val-Cit) linker is the most widely used, relying on cathepsin B-mediated cleavage. This mechanism enables precise payload release after ADC internalization and lysosomal degradation, making it a cornerstone in ADC design for targeting various cancers.

2.2. Limitations of Val-Cit Linkers

Despite extensive validation in primate and human plasma models, Val-Cit linkers have critical limitations that restrict their broader applicability in ADCs:

1. Hydrophobicity and aggregation:

The p-aminobenzyl carbamate (PAB) moiety in Val-Cit linkers adds significant hydrophobicity, limiting the drug-to-antibody ratio (DAR) to modest levels ⁸. Efforts to increase the DAR often cause ADC aggregation, reducing efficacy, pharmacokinetic stability, and safety.

2. Premature cleavage:

Val-Cit linkers are prone to enzymatic degradation by non-specific enzymes such as carboxylesterase Ces1C ⁹ and human neutrophil elastase (NE) ¹⁰. This degradation can cause premature payload release, off-target toxicity, and adverse clinical outcomes, including neutropenia, ultimately compromising the therapeutic potential of ADCs with Val-Cit linkers.

3. Payload compatibility:

The hydrophobicity of Val-Cit linkers limits their compatibility with hydrophobic payloads, often necessitating co-solvents or complex manufacturing processes to ensure payload solubility and stability. These requirements increase production costs and complicate ADC development pipelines.

These challenges highlight the need for novel linker technologies that address the limitations of Val-Cit linkers while preserving or enhancing therapeutic efficacy and safety ¹¹.

3. Exo-Linker Technology: Design and Advantages

Exo-Linker technology represents a paradigm shift in ADC development by addressing the limitations of traditional Val-Cit linkers (Figure 2). This innovative approach repositions cleavable peptide linkers, such as Glu-Val-Cit (EVC) —a linear variant resistant to CES1C ¹² —and Glu-Glu-Val-Cit (EEVC), to the exo-position of the PAB moiety. By leveraging the hydrophilicity of tetrapeptides and optimizing structural design, Exo-Linkers improve ADC stability, mitigate hydrophobicity-related challenges, and enhance therapeutic efficacy.

3.1. Molecular Design

The Exo-Linker incorporates glutamic acid residues that confer resistance to enzymatic degradation by non-cathepsin B enzymes, like carboxylesterase Ces1C and NE. This repositioning masks the payload's hydrophobicity and shortens the structural distance between the antibody and payload, improving plasma stability, reducing aggregation, and enhancing intracellular payload release.



The exo-linker undergoes cleavage by Cathepsin B, a process that is identical to that of the conventional Val-Cit linker.

Major advantage of exo-linkers

- 1. Increasing mouse plasma stability due to hydrophilic moiety
- 2. Shielding effect by hydrophilic moiety
- 3. Increasing shielding effect by antibody
- 4. CMC advantage due to small molecular size
- 5. Adjustable hydrophilic units depending on hydrophobicity of payload

Figure 2. Design and benefits of exo-linkers

3.2. Key Advantages

• Improved Stability:

Exo-Linkers demonstrate exceptional plasma stability. Studies in mouse plasma showed that free payload concentrations remained below 5% after four days of incubation, underscoring the linker's resistance to premature cleavage.

Improved Hydrophilicity:

Hydrophobic interaction chromatography (HIC) analyses revealed faster retention times for ADCs with Exo-Linkers than those with traditional linear Val-Cit and EVC linkers. This highlights the improved hydrophilic profile of Exo-Linkers, which reduces aggregation and enhances pharmacokinetics ¹³.

Resistance to NE-mediated cleavage:

Val-Cit linkers are susceptible to NE, which cleaves the bond between valine and citrulline, causing premature detachment of Cit-PAB payloads, ¹⁴ increasing potential off-target toxicity and side effects. In our study, Exo-Linkers remained intact under NE exposure, preventing premature payload release and enhancing stability. To confirm this, we tested ADCs with Val-Cit and Exo-Linkers in an NE-sensitive setup using HER2-low MCF-7 cells (Figure 3). Without NE, all ADCs except MMAE performed similarly ¹⁵. However, with NE, Val-Cit ADCs exhibited higher cytotoxicity, likely due to the hydrophobicity of Cit-PAB-MMAE, which enhances cell penetration. In contrast, Exo-Linker ADCs avoided this effect, maintaining stability and minimizing off-target risks. These results show that Exo-Linkers overcome the critical weaknesses of Val-Cit linkers, offering a safer and more reliable option for ADC development. This technology marks a significant advancement in enhancing ADC safety and therapeutic potential.



Figure 3. Evaluation of off-target toxicity of ADCs using an *in vitro* cytotoxicity assay: (A, B) Schematic representation of the assay under a) NE-depleted and b) NE-pre-treated conditions. Anti-HER2 ADCs treated or untreated with NE in the NE-reaction buffer and incubated with MCF-7 cells (C, D) Cell viability after incubation with (c) untreated and (d) NE-treated ADCs and MMAE.

• Simplified Manufacturing:

The small molecular size of Exo-Linkers streamlines production and quality control, offering a distinct advantage in manufacturing efficiency. Their intrinsic hydrophilicity eliminates the need for co-solvents during payload conjugation, enhancing compatibility with various payloads and enabling the design of ADCs with higher DARs.

4. Functional Insights into Exo-Linker Technology

4.1. Payload Compatibility and Hydrophilicity

The compatibility of Exo-Linkers with hydrophobic payloads was demonstrated using pyrene as a model compound. Comparative analyses showed that ADCs with a high DAR of 8, synthesized using Exo-Linkers, exhibited significantly reduced aggregation compared to those with Val-Cit or linear EVC linkers. LogP evaluations ¹⁶ confirmed the enhanced hydrophilic properties of Exo-Linkers, demonstrating their ability to mask payload hydrophobicity. This hydrophilic advantage contributes to improved pharmacokinetic stability and reduced off-target interactions, which are critical factors in ADC performance.

4.2. In Vivo Efficacy

Exo-Linker ADCs have demonstrated superior therapeutic efficacy in xenograft models. In studies with HER2-positive NCI-N87 cells, ADC consiting of APL-1091 or APL-1092 (Figure 4) showed better tumor suppression than traditional ADCs with linear EVC or Val-Cit linkers. Exo-Linker ADCs achieved these

results at lower doses, indicating improved therapeutic potency and a broader therapeutic index than conventional designs.



Figure 4. Chemical structure of Exo-linker payloads.

4.3. Pharmacokinetics

Rat PK studies validated the stability of Exo-Linker ADCs, showing minimal reductions in DAR over 21 days, in contrast to the significant payload detachment observed with traditional Val-Cit linker-based ADCs ¹⁷. This improved DAR retention demonstrates Exo-Linkers' ability to maintain payload integrity in vivo, resulting in more consistent therapeutic outcomes and extended efficacy over time.

4.4. Resistance to enzymatic interference

The robustness of Exo-Linkers against enzymatic degradation was demonstrated in cytotoxicity assays using NE. Traditional Val-Cit ADCs showed increased off-target toxicity due to premature payload release from NE cleavage. In contrast, Exo-Linker ADCs retained their payloads and resisted NEmediated interference. This enzymatic stability reduces side effects and ensures targeted payload delivery, improving safety and efficacy.

5. Integration with AJICAP Technology

Integrating Exo-Linker technology with the AJICAP Second Generation Platform marks a significant advancement in ADC development ¹⁸ ¹⁹. Developed by Ajinomoto, the AJICAP method enables site-specific conjugation by introducing thiol groups at predetermined antibody sites, such as Lys248, ensuring precise DAR control and reducing heterogeneity ²⁰. This site-specific approach enhances the advanced properties of Exo-Linkers, offering synergistic benefits for ADC development:

- Expanded therapeutic window
- Improved Solubility and Stability
- Superior Antitumor Efficacy



Figure 5. Comparison of therapeutic indices of three ADC designs: (1) a stochastic DAR = 4 ADC synthesized using the interchain break conjugation method with MC-Val-Cit-PAB-MMAE (left), (2) a site-specific DAR = 2 ADC generated through the AJICAP conjugation method with MC-Val-Cit-PAB-MMAE (middle), and (3) a site-specific DAR=2 ADC using the AJICAP conjugation method with exo-linker-MMAE (right). Each design illustrates the impact of conjugation strategy and linker technology on the safety and efficacy profile of the ADC.

Figure 5 illustrates a comparative analysis of therapeutic indices for three linker technologies and conjugation strategies used in ADCs. The therapeutic index, defined as the ratio of the maximum tolerated dose (MTD) to the minimum effective dose (MED), is crucial for assessing the safety and efficacy profiles of ADCs. The first category represents ADCs using traditional interchain break

conjugation methods with MC-Val-Cit-PAB MMAE under good laboratory practice²¹. These ADCs have a limited MTD, indicating an increased likelihood of dose-limiting toxicity at elevated concentrations ²². Additionally, a relatively high MED is needed to achieve therapeutic efficacy. The narrow therapeutic window highlights the challenges in balancing safety and efficacy for ADCs with traditional linker technologies. In contrast, the DAR = 2 site-specific ADC incorporating the AJICAP conjugation platform with MC-Val-Cit-PAB-MMAE linkers shows a remarkable improvement in MTD, approximately an 8-fold increase than traditional counterparts ¹⁷. This improvement significantly expands the therapeutic window, reflecting the superior pharmacokinetics and reduced systemic toxicity enabled by site-specific conjugation. The AJICAP system overcomes critical limitations of conventional ADCs by enhancing homogeneity and DAR control. Finally, site-specific ADCs synthesized with AJICAP and Exo-Linker technologies extend the therapeutic index by reducing the MED. Exo-Linker technology, with exo-positioned peptide linkers (e.g., Glu-Val-Cit), enhances ADC hydrophilicity, reduces aggregation, and improves plasma stability. It mitigates premature cleavage from enzymatic interference (e.g., carboxylesterase and neutrophil elastase) and simplifies conjugation. As a result, Exo-Linker ADCs show optimal pharmacokinetic stability and a broader therapeutic window, enabling lower doses and reduced toxicity, even at higher concentrations.

Experimental procedures

Reagents and Cells

Trastuzumab (Herceptin), a human monoclonal IgG1 antibody, was purchased from Midwinter. The AJICAP peptide reagent was prepared as previously reported ²³. MC-VC-PABC-MMAE and deruxtecan were purchased from NJ Biopharmaceuticals, LLC (USA). APL-1081, APL-1082, APL-1091, and APL-1092 were prepared as previously reported ⁶. All other chemical reagents were purchased from Sigma-Aldrich (USA).

The MCF-7 cell line was obtained from the Japanese Collection of Research Bioresources Cell Bank (Japan). The SKBR-3 cell line was obtained from the Memorial Sloan Kettering Cancer Center (USA).

• Experimental procedure for AJICAP site-specific DAR = 2 ADC

Trastuzumab was converted to DAR = 2 ADC as previously described ²³.

• Experimental procedure for stochastic DAR = 4 ADC

Trastuzumab was converted to stochastic DAR = 4 ADC as previously described ²⁴.

• Instruments and analytical methods

ADC concentration and recovery were measured using a Solo-VPE system with the Slope Spectroscopy method ²⁵.

RP-HPLC analysis of ADCs was performed using AdvanceBio RP-mAb Diphenyl, 2.1×100 mm, 3.5μ m column (Agilent), as previously reported ²⁶.

SEC-HPLC analysis of ADCs was performed using a Waters ACQUITY UPLC Protein BEH SEC column (200 Å, 4.6×300 mm, 1.7μ m), as previously reported ¹⁶.

In-vitro study

The in vitro cytotoxicity assay was performed using a previously reported method ²⁷.

In vivo xenograft study

Cells

NCI-N87 cells (Cat # CRL-5822) were purchased from ATCC. Studies were performed using a previously established procedure ¹⁷.

Conclusion

Exo-Linker technology significantly advances ADC development, offering enhanced stability, hydrophilicity, and resistance to enzymatic degradation. Its synergy with innovative conjugation methods like AJICAP highlights its transformative potential. By overcoming the limitations of conventional linkers, Exo-Linkers set a new standard for safer, more effective ADCs, shaping the future of targeted cancer therapy.

The exo-linker technology introduced in this study is a significant advancement in biopharmaceutical innovation, already employed in collaboration with over 40 pharmaceutical companies through strategic partnerships and licensing agreements. It exemplifies Ajinomoto Group's commitment to advancing biomanufacturing platforms and expanding its contract development and manufacturing organization (CDMO) business.

Ajinomoto's proprietary technologies include AJIPHASE, a highly efficient liquid-phase synthesis platform for large-scale peptide and nucleic acid production; CORYNEX, an advanced protein expression system; and innovative methodologies like FMR-based biomanufacturing ²⁸ and enzymatic protein modification ²⁹. These innovations provide a strong foundation for meeting modern drug discovery's diverse and complex demands.

The development of exo-linkers is poised to play a pivotal role in advancing next-generation therapeutic platforms. By integrating this technology into broader biopharmaceutical applications, Ajinomoto aims to enhance the precision, scalability, and efficiency of drug discovery and manufacturing, reinforcing its leadership in the evolving biopharmaceutical landscape.

References

1. Tarantino, P.; Ricciuti, B.; Pradhan, S. M.; Tolaney, S. M., Optimizing the safety of antibody-drug conjugates for patients with solid tumours. *Nat Rev Clin Oncol* **2023**.

2. Yamazaki, S.; Matsuda, Y., Tag-Free Enzymatic Modification for Antibody–Drug Conjugate Production. *ChemistrySelect* **2022**, *7* (48) e202203753.

3. Colombo, R.; Tarantino, P.; Rich, J. R.; LoRusso, P. M.; de Vries, E. G. E., The Journey of Antibody–Drug Conjugates: Lessons Learned from 40 Years of Development. *Cancer Discovery* **2024**, *14* (11), 2089-2108.

4. Balamkundu, S.; Liu, C. F., Lysosomal-Cleavable Peptide Linkers in Antibody-Drug Conjugates. *Biomedicines* **2023**, *11* (11).

5. Fujii, T.; Matsuda, Y., Novel formats of antibody conjugates: recent advances in payload diversity, conjugation, and linker chemistry. *Expert Opin Biol Ther* **2023**, *23* (11), 1053-1065.

6. Watanabe, T.; Arashida, N.; Fujii, T.; Shikida, N.; Ito, K.; Shimbo, K.; Seki, T.; Iwai, Y.; Hirama, R.; Hatada, N., Exo-Cleavable Linkers: Enhanced Stability and Therapeutic Efficacy in Antibody–Drug Conjugates. *J Med Chem* **2024**, *67* (20), 18124-18138..

7. Watanabe, T.; Fujii, T.; Matsuda, Y., Exo-Cleavable Linkers: Enhancing Stability and Therapeutic Efficacy in Antibody-Drug Conjugates. *Journal of Synthetic Organic Chemistry, Japan* **2024**, *82* (11), 1117-1124.

Lucas, A. T.; Price, L. S. L.; Schorzman, A. N.; Storrie, M.; Piscitelli, J. A.; Razo, J.;
 Zamboni, W. C., Factors Affecting the Pharmacology of Antibody-Drug Conjugates. *Antibodies (Basel)* 2018, 7 (1).

9. Dorywalska, M.; Dushin, R.; Moine, L.; Farias, S. E.; Zhou, D.; Navaratnam, T.; Lui, V.; Hasa-Moreno, A.; Casas, M. G.; Tran, T. T.; Delaria, K.; Liu, S. H.; Foletti, D.; O'Donnell, C. J.; Pons, J.; Shelton, D. L.; Rajpal, A.; Strop, P., Molecular Basis of Valine-Citrulline-PABC Linker Instability in Site-Specific ADCs and Its Mitigation by Linker Design. *Mol Cancer Ther* **2016**, *15* (5), 958-70.

10. Zhao, H.; Gulesserian, S.; Malinao, M. C.; Ganesan, S. K.; Song, J.; Chang, M. S.; Williams, M. M.; Zeng, Z.; Mattie, M.; Mendelsohn, B. A.; Stover, D. R.; Donate, F., A Potential Mechanism for ADC-Induced Neutropenia: Role of Neutrophils in Their Own Demise. *Mol Cancer Ther* **2017**, *16* (9), 1866-1876.

11. Bulger, P. G.; Conlon, D. A.; Cink, R. D.; Fernandez-Cerezo, L.; Zhang, Q.; Thirumalairajan, S.; Raglione, T.; Liang, R.; Zhou, J.; Chalgeri, A., Drug-Linkers in Antibody–Drug Conjugates: Perspective on Current Industry Practices. *Organic Process Research & Development* **2023**, *27* (7), 1248-1257.

12. Anami, Y.; Yamazaki, C. M.; Xiong, W.; Gui, X.; Zhang, N.; An, Z.; Tsuchikama, K., Glutamic acid-valine-citrulline linkers ensure stability and efficacy of antibody-drug conjugates in mice. *Nat Commun* **2018**, *9* (1), 2512.

Lyon, R. P.; Bovee, T. D.; Doronina, S. O.; Burke, P. J.; Hunter, J. H.; Neff-LaFord, H.
D.; Jonas, M.; Anderson, M. E.; Setter, J. R.; Senter, P. D., Reducing hydrophobicity of homogeneous antibody-drug conjugates improves pharmacokinetics and therapeutic index. *Nat Biotechnol* 2015, *33* (7), 733-5.

14. Miller, J. T.; Vitro, C. N.; Fang, S.; Benjamin, S. R.; Tumey, L. N., Enzyme-Agnostic Lysosomal Screen Identifies New Legumain-Cleavable ADC Linkers. *Bioconjug Chem* **2021**, *32* (4), 842-858.

Ha, S. Y. Y.; Anami, Y.; Yamazaki, C. M.; Xiong, W.; Haase, C. M.; Olson, S. D.;
Lee, J.; Ueno, N. T.; Zhang, N.; An, Z.; Tsuchikama, K., An Enzymatically Cleavable Tripeptide
Linker for Maximizing the Therapeutic Index of Antibody-Drug Conjugates. *Mol Cancer Ther* 2022, *21* (9), 1449-1461.

16. Fujii, T.; Reiling, C.; Quinn, C.; Kliman, M.; Mendelsohn, B. A.; Matsuda, Y., Physical characteristics comparison between maytansinoid-based and auristatin-based antibody-drug conjugates. *Explor Target Antitumor Ther* **2021**, *2* (6), 576-585.

17. Matsuda, Y.; Seki, T.; Yamada, K.; Ooba, Y.; Takahashi, K.; Fujii, T.; Kawaguchi, S.; Narita, T.; Nakayama, A.; Kitahara, Y.; Mendelsohn, B. A.; Okuzumi, T., Chemical Site-Specific Conjugation Platform to Improve the Pharmacokinetics and Therapeutic Index of Antibody-Drug Conjugates. *Mol Pharm* **2021**, *18* (11), 4058-4066.

18. Fujii, T.; Matsuda, Y.; Seki, T.; Shikida, N.; Iwai, Y.; Ooba, Y.; Takahashi, K.; Isokawa, M.; Kawaguchi, S.; Hatada, N.; Watanabe, T.; Takasugi, R.; Nakayama, A.; Shimbo, K.; Mendelsohn, B. A.; Okuzumi, T.; Yamada, K., AJICAP Second Generation: Improved Chemical Site-Specific Conjugation Technology for Antibody-Drug Conjugate Production. *Bioconjug Chem* **2023**, *34* (4), 728-38.

19. Fujii, T.; Ito, K.; Takahashi, K.; Aoki, T.; Takasugi, R.; Seki, T.; Iwai, Y.; Watanabe, T.; Hirama, R.; Tsumura, R.; Fuchigami, H.; Yasunaga, M.; Matsuda, Y., Bispecific Antibodies Produced via Chemical Site-Specific Conjugation Technology: AJICAP Second-Generation. *ACS Med Chem Lett* **2023**, *14* (12), 1767-1773.

20. Matsuda, Y.; Shikida, N.; Hatada, N.; Yamada, K.; Seki, T.; Nakahara, Y.; Endo, Y.; Shimbo, K.; Takahashi, K.; Nakayama, A.; Mendelsohn, B. A.; Fujii, T.; Okuzumi, T.; Hirasawa, S., AJICAP-M: Traceless Affinity Peptide Mediated Conjugation Technology for Site-Selective Antibody-Drug Conjugate Synthesis. *Org Lett* **2024**, *26* (27), 5597-5601.

21. Tawfiq, Z.; Matsuda, Y.; Alfonso, M. J.; Clancy, C.; Robles, V.; Leung, M.; Mendelsohn, B. A., Analytical Comparison of Antibody-drug Conjugates Based on Good Manufacturing Practice Strategies. *Anal Sci* **2020**, *36* (7), 871-875.

22. Tawfiq, Z.; Caiazza, N. C.; Kambourakis, S.; Matsuda, Y.; Griffin, B.; Lippmeier, J. C.; Mendelsohn, B. A., Synthesis and Biological Evaluation of Antibody Drug Conjugates Based on an Antibody Expression System: Conamax. *ACS Omega* **2020**, *5* (13), 7193-7200.

23. Watanabe, T.; Fujii, T.; Stofleth, J. T.; Takasugi, R.; Takahashi, K.; Matsuda, Y., Scale-Up Synthesis of Site-Specific Antibody–Drug Conjugates Using AJICAP Second-Generation Technology. *Org. Process Res. Dev.* **2023**, *27*(6), 1136–1143.

24. Matsuda, Y.; Tawfiq, Z.; Leung, M.; Mendelsohn, B. A., Insight into Temperature Dependency and Design of Experiments towards Process Development for Cysteine-Based Antibody-Drug Conjugates. *ChemistrySelect* **2020**, *5* (28), 8435-8439.

25. Matsuda, Y.; Chakrabarti, A.; Takahashi, K.; Yamada, K.; Nakata, K.; Okuzumi, T.; Mendelsohn, B. A., Chromatographic analysis of site-specific antibody-drug conjugates produced by AJICAP first-generation technology using a recombinant Fcgammallla receptor-ligand affinity column. *J Chromatogr B Analyt Technol Biomed Life Sci* **2021**, *1177*, 122753.

26. Matsuda, Y.; Leung, M.; Tawfiq, Z.; Fujii, T.; Mendelsohn, B. A., In-situ Reverse Phased HPLC Analysis of Intact Antibody-Drug Conjugates. *Anal Sci* **2021**, *37* (8), 1171-1176.

Yamazaki, S.; Ito, K.; Aoki, T.; Arashida, N.; Watanabe, T.; Fujii, T.; Matsuda, Y.,
Biological Evaluation of Antibody-Drug Conjugates Produced by Tag-Free Lipoate Ligase A
Modification. *Biochemistry* 2024, 63 (5), 644–650.

28. Nakahara, Y.; Kawaguchi, T.; Matsuda, Y.; Endo, Y.; Date, M.; Takahashi, K.; Kato, K.; Okasora, T.; Ejima, D.; Nagaki, A., Spatiotemporal Control of Protein Refolding through Flash-Change Reaction Conditions. *Langmuir* **2024**, *40* (16), 8483-8492.

29. Yamazaki, S.; Shikida, N.; Takahashi, K.; Matsuda, Y.; Inoue, K.; Shimbo, K.; Mihara, Y., Lipoate-acid ligase a modification of native antibody: Synthesis and conjugation site analysis. *Bioorg Med Chem Lett* **2021**, *51*, 128360.

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