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Erol C. Vatanserver^a, Jeffrey Kang^b, Alfred Tuley^a, E. Sally Ward^{b,c,d,*} and Wenshe Ray Liu^{a,e,f,*}

^a The Texas A&M Drug Discovery Laboratory, Department of Chemistry, Texas A&M University, College Station, Texas 77843, United States

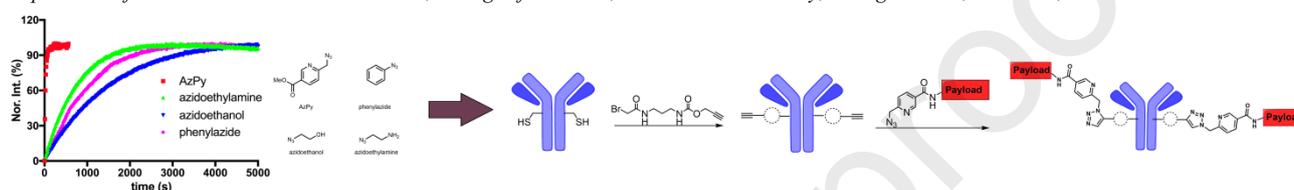
^b Department of Molecular and Cellular Medicine, Texas A&M University Health Science Center, College Station, TX 77843, USA

^c Department of Microbial Pathogenesis and Immunology, Texas A&M University Health Science Center, Bryan, TX 77807, USA

^d Centre for Cancer Immunology, Cancer Sciences Unit, Faculty of Medicine, University of Southampton, Southampton SO16 4YD, UK

^e Department of Biochemistry and Biophysics, Texas A&M University, College Station, TX 77843, USA

^f Department of Molecular and Cellular Medicine, College of Medicine, Texas A&M University, College Station, TX 77843, USA



TRANSLATING SMALL MOLECULE KINETICS INTO OPTIMAL ADC SYNTHESIS



An Optimal “Click” Formulation Strategy for Antibody-Drug Conjugate Synthesis

Erol C. Vatanserver^a, Jeffrey Kang^b, Alfred Tuley^a, E. Sally Ward^{b,c,d,*} and Wenshe Ray Liu^{a,e,f,*}

^a The Texas A&M Drug Discovery Laboratory, Department of Chemistry, Texas A&M University, College Station, Texas 77843, United States

^b Department of Molecular and Cellular Medicine, Texas A&M University Health Science Center, College Station, TX 77843, USA

^c Department of Microbial Pathogenesis and Immunology, Texas A&M University Health Science Center, Bryan, TX 77807, USA

^d Centre for Cancer Immunology, Cancer Sciences Unit, Faculty of Medicine, University of Southampton, Southampton SO16 4YD, UK

^e Department of Biochemistry and Biophysics, Texas A&M University, College Station, TX 77843, USA

^f Department of Molecular and Cellular Medicine, College of Medicine, Texas A&M University, College Station, TX 77843, USA

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ABSTRACT

As a versatile reaction for bioconjugation, Cu(I)-catalyzed alkyne-azide cycloaddition (CuAAC) has enormous potential in the synthesis of antibody-drug conjugates (ADCs). In order to optimize CuAAC-based ADC synthesis, we characterized kinetically different formulation processes by mimicking ADC synthesis using small molecules and subsequently revealed unique kinetic behaviors of different combinations of alkyne and azide conditions. Our results indicate that under ADC synthesis conditions, for an alkyne-containing drug, its concentration has minimal impact on the reaction rate when an antibody has a non-metal-chelating azide but is proportional to concentration when an antibody contains a metal-chelating azide; however, for an alkyne-containing antibody, the ADC synthesis rate is proportional to the concentration of a drug with a non-metal-chelating azide but displays almost no dependence on drug concentration with a metal-chelating azide. Based on our results, we designed and tested an optimal “click” formulation strategy that allowed rapid and cost-effective synthesis of a new ADC.

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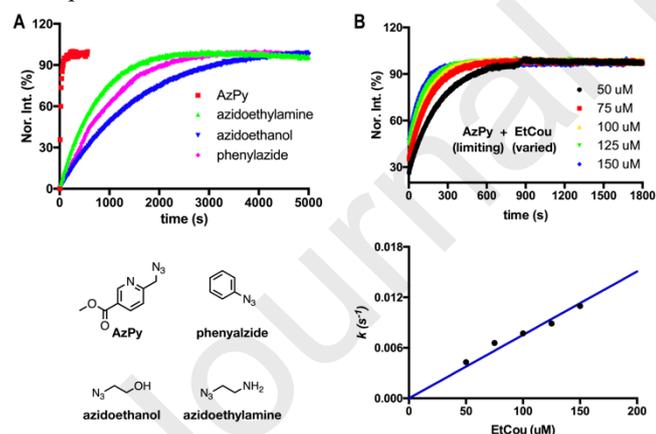
1. Introduction

Antibody-drug conjugates (ADCs) consist of cytotoxic drugs that are covalently conjugated to antibodies. The antibody component of an ADC directs the binding to antigens expressed on tumor cells and enables the specific delivery of their conjugated drugs. Consequently, ADCs combine the anti-tumor efficacy of chemotherapeutic drugs, whilst minimizing their significant systemic cytotoxicity.¹ With 9 ADCs approved by FDA and more than 150 in clinical trials,² the development of ADCs has revolutionized cancer treatment. This rapidly expanding use of ADCs has largely overshadowed their manufacturing challenges. How exactly antibodies and drugs are conjugated plays a significant role in defining both the cost and therapeutic index of an ADC.^{3,4} To date, the most broadly used reaction for antibody-drug conjugation is the covalent addition of a cysteine thiolate in an antibody to a maleimide in a drug (Figure 1A).⁵ In comparison to random conjugation to lysine residues of an antibody that are more abundant than cysteine residues, this reaction typically leads to more homogenous ADCs. The fast reaction kinetics ($734 \text{ M}^{-1}\text{s}^{-1}$)⁶ also allows the use of a low concentration of a drug (in the μM range) to achieve efficient coupling to the antibody, which significantly simplifies the manufacturing process of ADCs. Since most drugs used during ADC synthesis are complicated natural products and therefore expensive to produce, the use of a low drug

concentration also substantially reduces the manufacturing cost of ADCs. However, although efficacious, the cysteine-maleimide reaction has pitfalls. Maleimide reacts with lysine at a low level to form side products.⁷ The reaction itself is also slowly reversible, which leads to the release of the conjugated drug that can contribute to systemic cytotoxicity.⁸ For these reasons, substantial efforts to either improve the cysteine-maleimide chemistry^{9, 10} or develop novel conjugation strategies have been made.¹¹⁻¹⁹ As the founding click reaction for bioconjugation, Cu(I)-catalyzed alkyne-azide cycloaddition (CuAAC) has been explored for ADC synthesis.²⁰ In this study, we show how this versatile reaction can be favorably optimized for this purpose.

tested 2-propyn-1-ol concentrations, the collected data of fluorescence intensity versus time fit perfectly to the one-phase exponential production formation model, indicating that AzCou followed first-order kinetics to convert to product under all tested conditions (Figure 2A). However, unexpectedly the reaction trajectories for all 2-propyn-1-ol concentrations were essentially the same and all of the determined apparent reaction rate constants were almost equal. We originally tested pseudo first order conditions²⁵ by varying the 2-propyn-1-ol concentration with the aim of determining a second order rate constant between an azide-alkyne bimolecular reaction under a fixed Cu(I)-ligand condition. Apparently, the reaction did not follow a simple bimolecular reaction model. The involvement of the Cu(I) catalyst complicated the process. The independence of determined apparent rate constants from 2-propyn-1-ol concentrations might have been due to the formation of an alkyne-Cu(I) resting state before the involvement of an azide, or two parallel alkyne-Cu(I) and azide-Cu(I) chelating processes in which the alkyne-Cu(I) chelation was significantly faster than the other process.

AzCou is a non-metal-chelating azide. The independence of the rate of its CuAAC reaction from the concentration of excessive 2-propyn-1-ol might represent a general feature of CuAAC reactions. To explore this possibility, we tested AzCou reactions with five other alkynes in a setup similar to that for 2-propyn-1-ol. We chose five phenylacetylenes due to their significantly different electrochemical features of *p*-substitutes that substantially influence their host compounds' alkyne group.²⁶ Should a simple bimolecular model apply, they would behave very differently in reaction kinetics with AzCou. However, for all five phenylacetylenes, their reaction trajectories at different concentrations were very similar (Figure 2B-F). Although determined apparent k values at different alkyne concentrations showed some variations, they were very alike. They deviated far from predictions from a real bimolecular reaction in which we



expected a linear dependence of a determined apparent k value on the concentration of an excessively used reactant. The average k values for all tested alkynes were also very similar, indicating that the reaction kinetics were minimally dependent on both the identity and concentration of excessive alkynes when they reacted with limiting non-metal-chelating AzCou.

Since both the identity and concentration of an excessive alkyne have minimal impacts on the CuAAC reaction kinetics of non-metal-chelating AzCou, we reasoned that in a similar setup the azide identity might significantly influence the reaction kinetics. In order to compare reactivities of different azides under the condition of a limiting azide and an excess alkyne concentration, we used EtCou. When we reacted 100 μM EtCou in excess with 20 μM concentrations of several different azides,

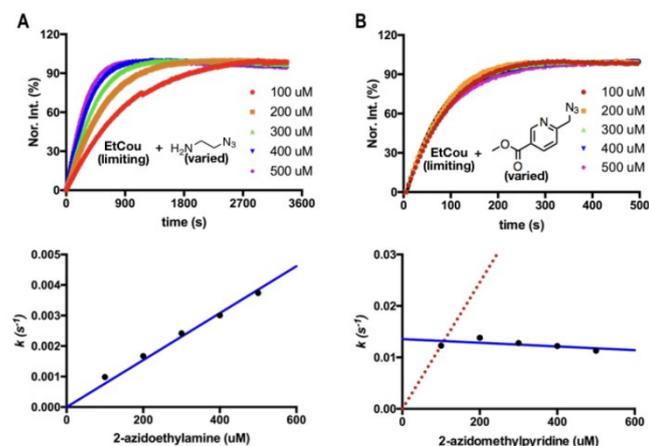
comparison, a metal-chelating azide, AzPy²⁷ was also included in the analysis. Overall and consistent with earlier studies, AzPy reacted more rapidly than all other non-metal-chelating azides. For non-metal-chelating azides, azidoethylamine reacted faster than phenylazide and phenylazide reacted faster than azidoethanol. Azidoethylamine reacted more quickly than the other two compounds, probably due to the chelating propensity of its amine toward Cu(I). To confirm that the minimal dependence of the reaction kinetics on the concentration of an excessive alkyne when it reacts with a limiting non-metal-chelating azide is a general phenomenon, we carried out analyses of reactions between EtCou and different non-metal-chelating azides by fixing the azide concentration and varying the EtCou concentration. For azidoethanol, reactions were too slow to determine k but had similar trajectories (SI, Figure S3). For azidoethylamine and phenylazide, reactions displayed similar kinetics at different EtCou concentrations (SI, Figures S4-5).

Figure 3. (A) CuAAC reaction trajectories between limiting concentrations of different azides (20 μM) and excess concentrations of EtCou (100 μM). (B) The one-phase exponential increase of product fluorescence when a limiting concentration (5 μM) of AzPy was reacted with excess concentrations of EtCou (50-150 μM).

Based on all reactions we have analyzed between limiting non-metal-chelating azides and excess alkyne concentrations, we conclude that the minimal dependence of the reaction kinetics on the concentration of an alkyne is a general observation.

Figure 4. The one-phase exponential increase of product fluorescence when a limiting concentration (20 μM) of EtCou reacted with excess concentrations (100-500 μM) of azidoethylamine

alkyne-containing drug may react with metal-chelating AzPy conjugated to an antibody. We reacted limiting AzPy with different concentrations of excessive EtCou and observed that the EtCou concentration significantly influenced the reaction rate



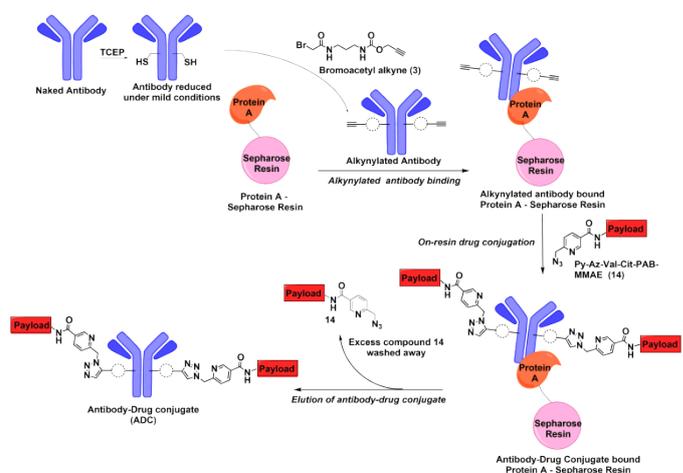
(Figure 3B). The determined k values were linearly proportional to EtCou trajectories. It is obvious that metal-chelating and non-metal-chelating azides involve very different CuAAC reaction kinetics. The AzPy-Cu(I) chelation is apparently able to outcompete the rapid alkyne-Cu(I) formation, leading to the observed dependence of the alkyne concentration. Based on all the data related to limiting azides reacting with excessive alkynes, we can deduce that for the reaction between an azide-containing antibody and an alkyne-containing drug to generate an ADC, the most kinetically favorable option is to react a metal-chelating antibody azide with a drug alkyne and improve the kinetics by increasing the drug concentration. When a non-metal-chelating antibody azide is used, changing the drug concentration has little impact on the reaction kinetics and more favorable reaction kinetics can be achieved by changing the azide identity in the antibody for example, from a standard alkyl azide to an aromatic azide.

containing antibody to an azide-containing drug that are present as limiting and excess reactant concentrations, respectively, we continued testing the reaction kinetics of limiting EtCou concentration with excess azidoethylamine concentration under pseudo first order conditions. By fixing the EtCou concentration and varying the azidoethylamine concentration, we observed that the azidoethylamine concentration significantly influenced the reaction kinetics and the determined apparent rate constants were linearly proportional to the azidoethylamine concentrations (Figure 4A). Similarly, we tested the reaction of limiting EtCou concentration with excess concentration of metal-chelating AzPy. Intriguingly, reactions at all AzPy concentrations displayed similar reaction trajectories and all determined k values were very similar (Figure 4B), indicating minimal influence of the azide concentration on the reaction kinetics under these conditions. Although we used substantially higher concentrations of Cu(I) in the EtCou-azidoethylamine reactions than in the EtCou-AzPy reactions, the reaction at the highest azidoethylamine concentration was still much slower than an EtCou-AzPy reaction, indicating that under these conditions a metal-chelating drug azide remains an optimal choice for favorable kinetics. We

Figure 5. Catch-and-release strategy to synthesize ADCs

might also use the alkyne-Cu(I) resting state theory or two parallel alkyne-Cu(I) and azide-Cu(I) chelating processes to explain the observations in Figure 4. When a non-metal-chelating azide is used, the alkyne-Cu(I) chelation is so fast that the kinetic dependence on the azide is observable. However, when a metal-chelating azide is used, the azide-Cu(I) chelation outcompetes the alkyne-Cu(I) chelation so that the reaction kinetics is solely determined by the alkyne-Cu(I) chelation. The implication of the data shown in Figure 4 for ADC generation is obvious. For the reaction between an alkyne-containing antibody and an azide-containing drug to generate ADCs, the most kinetically favorable option is to react an antibody alkyne with a metal-chelating drug azide although changing the drug concentration does not influence the overall kinetics. When a non-metal-chelating drug azide is used, improving the drug concentration will significantly increase the reaction kinetics and one may also achieve more favorable reaction kinetics by changing the azide identity in the drug from a standard alkyl azide to an aromatic azide.

Next, we wanted to verify that the conditions used in small molecule kinetics can be translated to generate protein conjugates



as well. In order to demonstrate this, we built a model system that allows us to monitor protein conjugation kinetics by FRET analysis. We site-specifically incorporated L-propargyl-lysine (Prock)²⁸ into superfolder green fluorescent protein (sfGFP) to generate our model protein (sfGFP134Prock). We synthesized compound (16) which is a metal-chelating azide with coumarin

dye can form a FRET pair upon conjugation (SI, Figure S12). After our first conjugation attempt with 10 μM CuSO_4 and 20 μM BTAA, we did not detect a FRET signal. This might be due to the presence of 6xHistag on the sfGFP134ProcK or another metal binding motif. Next, we tried the reaction with 40 μM CuSO_4 and 80 μM BTAA and the reaction appeared to reach a plateau in 10 minutes (SI, Figure S13). This result showed us that, from a kinetics point of view, it is reasonable to keep the concentration of CuSO_4 around 50 μM when carrying out click reactions with proteins.

Table 1. Relationships of different alkyne-azide combinations for the ADC synthesis

Antibody	Drug	Reaction	Effect of drug concentration on the reaction rate
Non-metal-chelating azide	Alkyne	Slow	Minimal
Metal-chelating azide	Alkyne	Fast	Linear
Alkyne	Non-metal-chelating azide	Slow	Linear
Alkyne	Metal-chelating azide	Fast	Minimal

Based on all of the kinetic observations, we can summarize kinetic features of the CuAAC-based ADC synthesis as shown in Table 1. Given the nature of antibodies, an ADC conjugation procedure that requires a short reaction time and facile purification is optimal. For achieving a short reaction time, a metal-chelating azide either on an antibody or a drug is an apparent choice. When a metal-chelating antibody azide is used, one may also improve the reaction kinetics by increasing the drug alkyne concentration. However, loading a metal-chelating azide onto an antibody either through genetic incorporation or conjugation to antibody thiolates will require conditions that may reduce the relatively labile azide functionality, leading to heterogeneity in the conjugated ADC. On the contrary, an antibody alkyne is far more stable. A cheaply produced alkyne such as bromoacetyl alkyne (compound **3** in Figure 5) can be rapidly coupled to an antibody by reaction with cysteine thiols. Using a catch-and-release method, exceedingly excessive bromoacetyl alkyne (**3**) can be added to achieve rapid synthesis of an alkyne-containing antibody and the residual alkyne compound can be quickly washed away. The coupled antibody alkyne can then react with a metal-chelating drug azide for the efficient synthesis of the ADC. Although the independence of the reaction kinetics during these conditions on the drug concentration does not allow the improvement of kinetics by increasing the drug concentration, it does provide a manufacturing benefit. This unique kinetic behavior means that only a minimally excessive metal-chelating azide is required for efficient ADC conjugation. Therefore, substantial manufacturing costs can be reduced with very low amounts of cytotoxic waste. To demonstrate all of these advantages related to reacting an antibody alkyne with a metal-chelating drug azide, we designed a catch-and-release synthetic route as shown in Figure 5.

In our study, we used the HER2 specific antibody, pertuzumab²⁹ as a model antibody. Pertuzumab is a monoclonal antibody used for the treatment of metastatic HER2-positive breast cancer. Wild-type pertuzumab has two disulfide bonds bridging the two heavy chains in the hinge region, and two disulfide bonds bridging each heavy chain hinge region with the two antibody light chains. These disulphides are reducible under mild conditions. To reduce the drug to antibody ratio, we used a

one reducible disulfide bond in the hinge region.³⁰ We first reduced pertuzumab-2C under mild conditions with TCEP then, reacted it with 4 mM bromoacetyl alkyne (**3**) for 30 mins to load the alkyne group. Next, we incubated the antibody with protein A-Sepharose and for 2 hrs to capture the alkynylated antibody on the resin. We subsequently washed out residual alkyne compound and provided 1.2 eq of AzPy-Payload (Py-Az-Val-Cit-PAB-MMAE, **14**) compound per alkynylated cysteine (there are 2 alkynylated cysteines for each antibody molecule) in the antibody (see Experimental Methods section for details). The provided AzPy-payload concentration was deemed to be in excess. Following 2 hrs reaction, we washed out the residual AzPy-payload and eluted the conjugated pertuzumab. The overall preparation time of the ADC is 9 hours. We analyzed the eluted ADC using electrospray ionization mass spectrometry (SI, Figure S17) that showed close to quantitative conversion, demonstrating the high efficiency of this newly designed ADC synthetic route. The addition of the drug molecule to heavy chain was identified by the change in mass and the shift in retention time of the heavy chain. The conditions and chromatogram for the LC-MS analysis can be found in the SI.

3. Conclusions

In conclusion, we have kinetically characterized CuAAC-based ADC conjugation processes using small molecule mimics. Our data reveal unique kinetic behaviors when different azide-alkyne combinations are used. During an ADC conjugation setup in which a limiting antibody concentration reacts with an excess drug concentration, our data indicates that a drug alkyne reacts slowly with a non-metal-chelating antibody azide and its concentration has minimal influence on the overall reaction kinetics, however it reacts rapidly with a metal-chelating antibody azide and its concentration is linearly proportional to the reaction rate constant. When a limiting antibody alkyne concentration is reacted with an excess concentration of azide-containing drug, our data implies that it reacts slowly when the drug azide is non-metal-chelating but the drug concentration is linearly proportional to the reaction rate constant. However, its reaction with a metal-chelating drug azide is exceedingly fast but not kinetically related to the drug concentration. Based on these observations, we designed an efficient and cost effective CuAAC-based ADC conjugation method and demonstrated that an ADC can be rapidly synthesized. The CuAAC-based linker is more stable than a traditional cysteine-maleimide linker leading to much lower non-targeted drug release.²⁰ Given the broad applications³¹ of CuAAC in many basic research and technological areas, we believe our kinetic observations will also assist widely in optimizing conditions for these applications.

4. Experimental Methods

Organic synthesis procedures, on-protein kinetics, LC-MS data for characterization of ADCs can be found in the Supplementary Information.

4.1 General method for kinetics measurement

Stock solutions for all azide and alkyne compounds were prepared in 100x working concentrations in DMSO. BTAA stock solutions were 250x the working concentration in DMSO. CuSO_4 stock solutions were 100x the working concentration. 250 mM sodium ascorbate stock solution in water was used in all analysis.

0.15M phosphate buffer, pH 7.0, 25% DMSO is added to the reaction well (or cuvette). Next, BTAA, CuSO_4 were added successively followed by azide derivative. After that, ascorbate

was

alkyne was added to initiate the click reaction. After addition of each reagent, the reaction mixture was gently mixed by pipetting up and down. Unless otherwise stated, a plate reader was used for all measurements. For AzCou excitation at 404 nM and emission at 480 nM was used. For EtCou excitation at 320 nM and emission at 400 nM was used. Data were fit to the equation: $I_t = I_0 + I_c \times (1 - \exp(-k \times t))$ where I_0 is the background fluorescent intensity, I_t the fluorescent intensity at a specified time, I_c the fluorescent intensity change when limiting reagent is fully converted to the product, and k the observed apparent rate constant.

Table 2. AzCou with excess alkyne concentrations (except 2-propyn-1-ol where 50 μM CuSO_4 , 300 μM BTAA and 50 μM AzCou, 2.5 mM sodium ascorbate used with 0.25, 0.5, 1, 2, 4 mM 2-propyn-1-ol)

Reagent	Final Concentration
CuSO_4	50 μM
BTAA	100 μM
AzCou	10 μM
Sodium ascorbate	2.5 mM
Alkyne Derivative	100, 250, 500, 750, 1000 μM

Table 3. CuAAC reaction trajectories between different limiting azide concentrations and excess EtCou concentrations

Reagent	Final Concentration
CuSO_4	50 μM
BTAA	100 μM
Azide derivative	20 μM
Sodium ascorbate	2.5 mM
7-Ethynylcoumarin	100 μM

Table 4. Limiting AzPy concentration reacted with excess EtCou concentrations (fluorometer was used)

Reagent	Final Concentration
CuSO_4	10 μM
BTAA	20 μM
AzPy	5 μM
Sodium ascorbate	2.5 mM
7-Ethynylcoumarin	50, 75, 100, 125, 150 μM

Table 5. EtCou reacted with excess AzPy concentrations

Reagent	Final Concentration
CuSO_4	10 μM
BTAA	20 μM
AzPy	100, 200, 300, 400, 500 μM
Sodium ascorbate	2.5 mM
7-Ethynylcoumarin	20 μM

Table 6. EtCou reacted with excess azidoethylamine concentrations

Reagent	Final Concentration
CuSO_4	10 μM
BTAA	20 μM
2-azidoethylamine	100, 200, 300, 400, 500 μM
Sodium ascorbate	2.5 mM
7-Ethynylcoumarin	20 μM

4.2 Catch-and-release method

reducible cysteines in its hinge region.³⁰

First, 100 μL 27 μM of Per-2C in PBS buffer was reduced by 200 μM TCEP at 37°C for 2 hrs. Then, bromoacetyl alkyne compound (3) (400 mM stock solution in DMSO) was added to the solution at a final concentration of 4 mM. The reaction was incubated at room temperature for 30 minutes. Next, 100 μL Protein A-Sepharose resin slurry in PBS was added to this reaction mixture. The mixture was incubated on a shaker at room temperature for 2 hours.

After that, an empty spin column was used for the washing step. After initial centrifugation, 300 μL PBS were added to the slurry and the resin washed 5 times with 300 μL PBS. All the centrifugations were carried out at 1000g for 1 min at room temperature.

After the final centrifugation step, the total slurry volume was brought to 100 μL .

Table 7. The 2x click reaction mixture (200 μL reaction mix)

2 x Click Reaction mixture
172 μL PBS
16.7 μL DMSO
1.6 μL BTAA (25 mM stock solution in DMSO)
4 μL Cu(II)SO_4 (5mM stock solution in sterile water)
1.3 μL azide (10 mM stock solution in DMSO)
4 μL sodium ascorbate (250 mM stock solution in sterile water)

The above mixture (2x click reaction mixture) was prepared in the given order and incubated for 1 min after sodium ascorbate addition. Next, 100 μL of the above solution was added to 100 μL resin suspension. The tube was covered in aluminum foil and placed on a shaker at room temperature for 2 hours.

Next, the slurry was added to a spin column for the final washing. After initial centrifugation, the resin was washed with 3 x 300 μL PBS containing 1mM EDTA. For the fourth and fifth washes, 1 mM phosphate buffer, pH 7.4, with 150 mM NaCl used reduce the trace amounts of phosphate in the slurry. ADC was eluted from protein A-Sepharose using acidic 0.1M glycine, 150 mM NaCl, pH 2.8 buffer for ~30 seconds with gentle shaking.

Three sequential elutions (90 μL each) were carried out to elute the ADCs. Each 90 μL eluent was neutralized immediately by adding 15 μL 2M Tris, 150 mM NaCl, pH 8.0.

The three eluates were combined and EDTA added to a final concentration of 1 mM, followed by dialysis against PBS. The ADCs were characterized by LC-MS (SI, Figure S16, S17).

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Supplementary material is available.

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Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

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