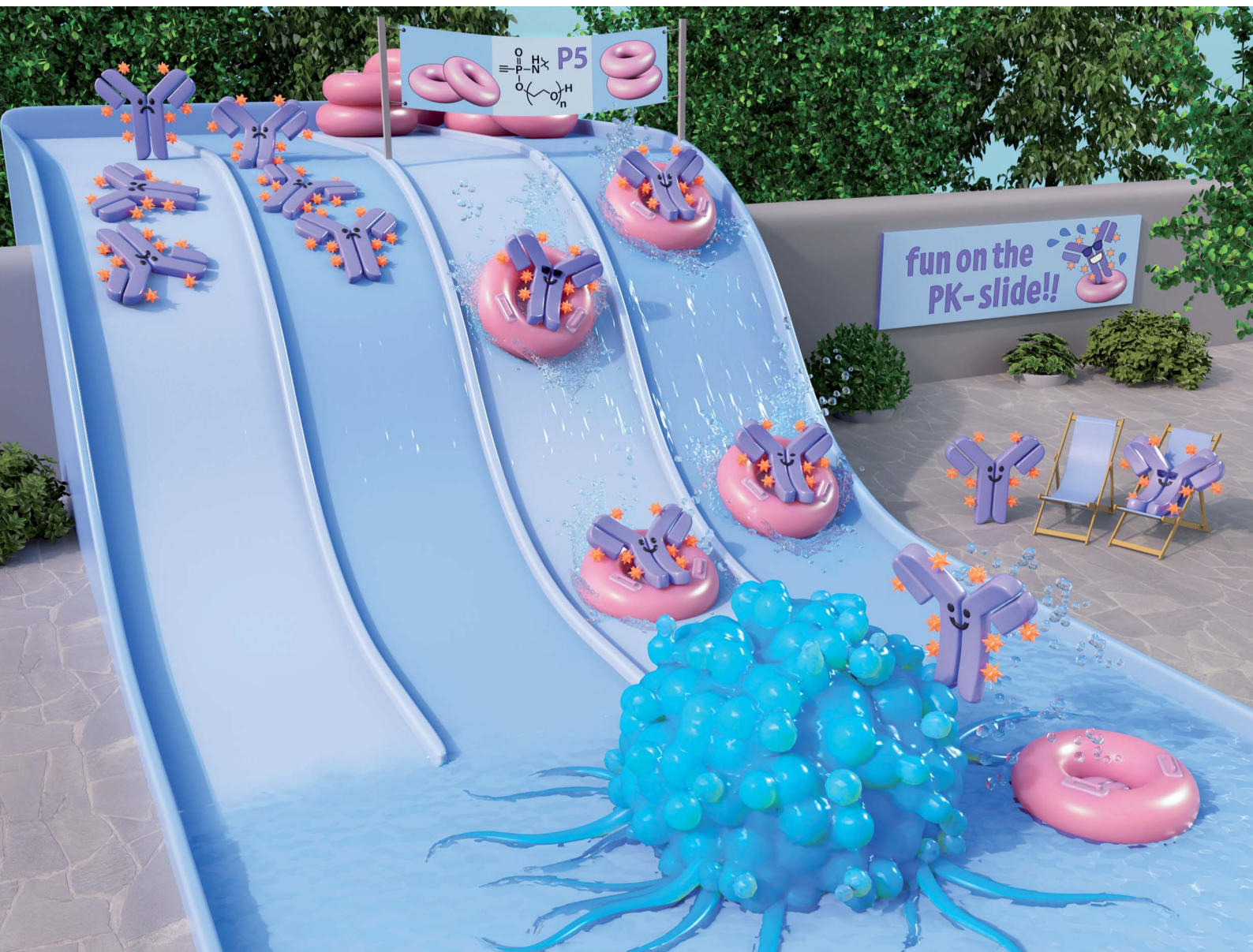


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Compact hydrophilic electrophiles enable highly efficacious high DAR ADCs with excellent *in vivo* PK profile†

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The recent success of antibody–drug conjugates (ADC), exemplified by seven new FDA-approvals within three years, has led to increased attention for antibody based targeted therapeutics and fueled efforts to develop new drug-linker technologies for improved next generation ADCs. We present a highly efficient phosphoramidate-based conjugation handle that combines a discrete hydrophilic PEG-substituent, an established linker-payload and a cysteine-selective electrophile in one compact building block. This reactive entity provides homogeneous ADCs with a high drug-to-antibody ratio (DAR) of 8 in a one-pot reduction and alkylation protocol from non-engineered antibodies. The compact branched PEG-architecture introduces hydrophilicity without increasing the distance between antibody and payload, allowing the generation of the first homogeneous DAR 8 ADC from VC-PAB-MMAE without increased *in vivo* clearance rates. This high DAR ADC exhibits excellent *in vivo* stability and increased antitumor activity in tumour xenograft models relative to the established FDA approved VC-PAB-MMAE ADC Adcetris, clearly showing the benefit of the phosphoramidate based building-blocks as a general tool for the efficient and stable antibody-based delivery of highly hydrophobic linker-payload systems.

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Introduction

The site-specific and chemoselective modification of proteins with unnatural functionalities like fluorescent dyes or cytotoxic drugs has become an indispensable strategy to advance the molecular life sciences and to develop biological therapeutics.¹ In particular antibody–drug conjugates (ADCs) receive increasing attention for the targeted treatment of diverse malignant diseases, as exemplified by seven new FDA approvals within the past three years, accumulating to eleven marketed ADCs in total.^{2–4} Most of these ADCs are produced *via* cysteine-selective bioconjugation strategies that exploit the unique nucleophilicity of thiols, liberated by reduction of interchain disulfides of monoclonal antibodies, to yield chemically defined conjugates.^{5–7} Strikingly, 9 out of 11 marketed ADCs are manufactured using maleimide functionalized linkers despite their known instability towards retro-Michael additions and thiol-

exchange reactions in the presence of external thiols.^{8–10} In particular the transfer of cytotoxic payload to endogenous serum proteins such as albumin was identified as the main pathway of these undesired decompositions.¹¹ The resulting drug loss in early days of circulation increases off target toxicity and reduces the ADC efficacy. To address these stability issues in the process of developing safer and more efficacious next generation ADCs, several strategies have been pursued to improve maleimide-based bioconjugates by promoting hydrolysis of the thiosuccinimide linkage to yield stable thioethers.^{12,13} Beyond that, a set of new cysteine-selective conjugation platforms including α,β -unsaturated acrylamides,^{14–16} unsaturated bissulfones,^{17,18} sulfonamides,¹⁹ vinyl²⁰ and divinylpyrimidines,²¹ divinyltriazines,²² as well as disubstituted maleimides^{23–25} and pyridazinediones^{26,27} have been developed to produce stable bioconjugates. However, most of these conjugation platforms lack ultimate evaluation in an *in vivo* setting, thus falling short their ambition to enable next generation therapeutics.

Recently, our laboratory contributed to this growing toolbox for cysteine-selective conjugation methods by introducing electrophilic vinyl- and ethynylphosphoramidates for the modification of peptides and proteins (Fig. 1a).^{28,29} Those unsaturated phosphoramidates exhibit excellent thiol selectivity and outstanding linkage stability against thiol exchange and retro addition. Along with these favored characteristics,

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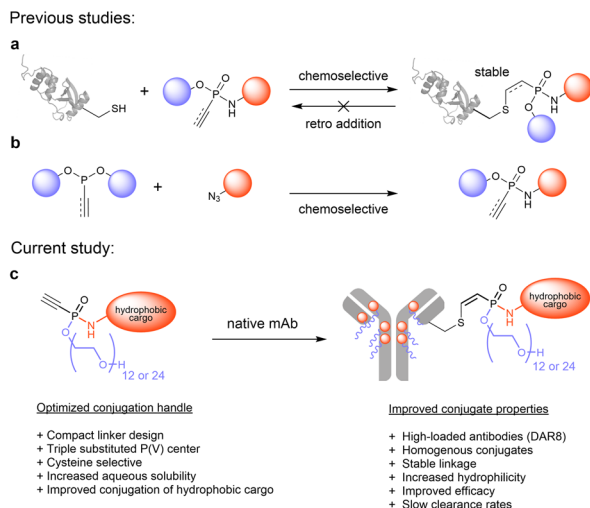


Fig. 1 Cysteine-selective phosphoramidate electrophiles. (a): Thiol-selective bioconjugation reaction of ethynylphosphoramidates with cysteine containing proteins yield highly stable conjugates. (b): Synthesis of ethynylphosphoramidates by chemoselective Staudinger-phosphonite reactions from ethynylphosphonites and azides. (c): Compact hydrophilic phosphoramidate building blocks that allow convenient ADC generation from native antibodies.

ethynylphosphoramidates are conveniently accessible from various azides and diverse phosphonites through chemoselective Staudinger phosphonite reactions (SPHR) (Fig. 1b). Previously, we applied ethynyl phosphoramidates for the synthesis of ADCs employing commonly used payloads such as the tubulin inhibitors MMAF and MMAE to generate efficacious stably linked DAR 4 ADCs.³⁰ In a head-to-head comparison to the analogous maleimide linked Adcetris, we showed that a phosphoramidate-based ADC displays a drastically improved serum stability which translated into a beneficial *in vivo* anti-tumor activity.³⁰

Aside from improving linkage stability, another strategy to enhance the *in vivo* efficacy of ADCs is to increase the drug loading to a maximum of eight drugs per antibody (DAR 8). Although such high-loaded ADCs usually exhibit greater potency *in vitro* compared to their lower modified analogues, they are often inferior *in vivo* owing to their increased overall hydrophobicity that triggers accumulation and aggregation in the liver.^{31,32} These processes ultimately lead to fast clearance from blood circulation and decreased payload delivery to the tumor. Therefore, the four marketed ADCs Adcetris, Polivy, Padcev and Tivdak, all conjugated with the hydrophobic linker-payload maleido-caproyl-valine-citrulline-PAB-MMAE (mc-VC-PAB-MMAE), are limited to an average DAR of 3.5 to 4. To overcome this limitation, efforts towards more hydrophilic linker that enable the generation of high DAR MMAE conjugates with acceptable pharmacokinetic profile are ongoing. It has been demonstrated that the replacement of the hydrophobic Val-Cit-PAB cleavage site with a hydrophilic glucuronide linker that contains branched PEG polymers or polysarcosine had a significant effect on clearance from circulation *in vivo* and

improved the pharmacokinetic profile of DAR 8 MMAE ADCs.^{33–35}

In our present study, we introduce a new phosphoramidate based hydrophilic conjugation handle that allows the straightforward modification of monoclonal antibodies with challenging hydrophobic payloads to yield highly modified antibody conjugates without impairing their favourable pharmacokinetic profiles (Fig. 1c). To that end, we exploited the intrinsic compact structure of ethynylphosphoramidates, and designed cysteine-selective conjugation handles with hydrophilic *O*-substituents that are directly incorporated into the phosphorous core of the electrophile. The resulting compact branched structures improve the aqueous solubility and conjugation efficiency of highly hydrophobic linker-payloads as the established Val-Cit-PAB-MMAE that is part of five marketed ADCs (Fig. 1c).

In the course of this study, we identified PEG₁₂ and PEG₂₄ as most suitable hydrophilic substituents that enable the generation of the first ethynylphosphoramidate-based ADCs with eight MMAE payloads per antibody. Most importantly, our strategy succeeds without changing the VC-PAB cleavage site to the much more polar glucuronide linker. These promising bioconjugates display improved *in vitro* and *in vivo* efficacy while maintaining exquisite PK profiles.

Results and discussion

We started our investigation by evaluating the impact of different water-soluble polar phosphoramidate-*O*-substituents to mediate solubility of a hydrophobic cargo. For this purpose, we chose the fluorescent dye cyanine 5 (Cy5) as non-toxic hydrophobic model and equipped the Cy5-phosphoramidate with *O*-substituents such as diethylene glycol (PEG₂), dodecethylene glycol (PEG₁₂) or choline to obtain hydrophilic conjugation handles that can compensate for the hydrophobicity of the fluorophore. To that end, bis(diisopropylamino)-chlorophosphine **1** was alkylated with ethynylmagnesium bromide, followed by 1*H*-tetrazole mediated substitution of the diisopropylamino groups with the respective alcohol to prepare phosphonites **3a–c** (Fig. 2a). The resulting highly hydrophilic phosphonites, were directly used in a subsequent SPHR with Cy5 azide **4** (for synthesis refer to ESI†) to give phosphoramidates **5a–c**. To initially evaluate the solubility enhancing properties of the different *O*-substituents we set up a shake-flask solubility assay and determined the maximum solubility of phosphoramidates **5a–c** in the aqueous system that is usually used for phosphoramidate bioconjugation reactions. For comparison we also included the previously reported Cy5 phosphoramidate **5d** with ethyl- *O*-substituent to our evaluation.²⁹

As expected, ethyl phosphoramidate **5d** displayed low solubility under the tested conditions with a saturation concentration of 36 μM (Fig. 2c). Changing the substituent to the more polar diethylene glycol significantly increased the aqueous solubility of the Cy5 conjugate **5a** to 175 μM . Interestingly, the cationic choline substituent did not significantly improve the solubility of the Cy5 phosphoramidate **5c**, which is displayed by the lowest measured solubility of 31 μM . In contrast, the PEG₁₂



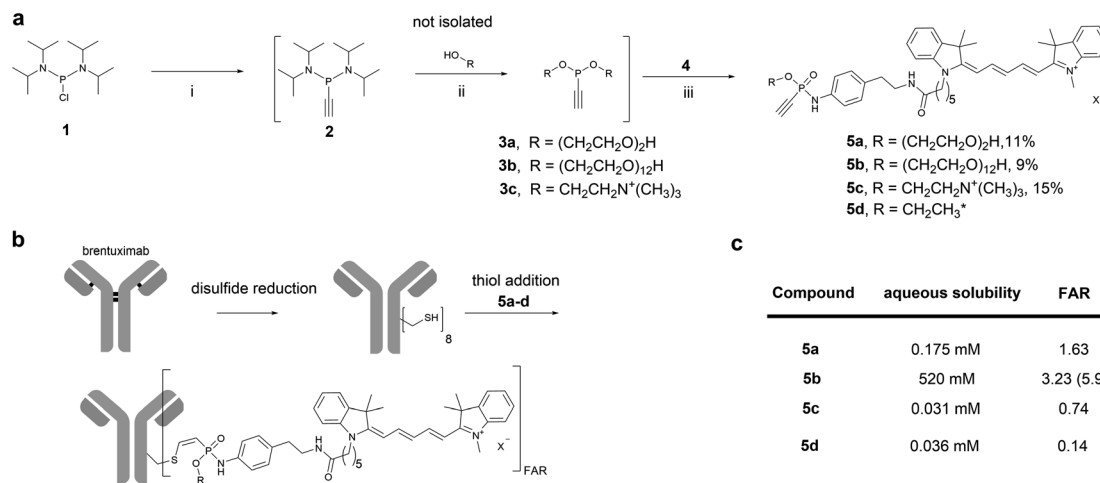


Fig. 2 a): Sequential one-pot synthesis of differently *O*-substituted Cy5-phosphonamidates. (i): Ethynyl magnesium bromide, THF, 0°C-rt, 30 min.; (ii): R-OH, 1*H* tetrazole, MeCN, 0°C-rt, 2-3 h; (iii): Cy5-azide 4, DMF, rt, overnight. *Synthesized according to previous report.²⁹ (b): Scheme for the sequential reduction and alkylation protocol for the modification of brentuximab with Cy5-phosphonamidates **5a-d**. Reduction: DTT, 50 mM sodium borate in PBS (pH 8.0), 37 °C, 40 min. Thiol addition: 50 mM Tris-HCl, 1 mM EDTA, 100 mM NaCl, pH 8.5, 5% DMSO, 14 °C, 16 h. (c): Table summarizing properties of phosphonamidates **5a-d**; FAR = fluorophore to antibody ratio.

substituent drastically increased the solubility of phosphonamidate **5b** to 520 mM. This immense increase in solubility qualified longer polyethylene glycol substituted phosphonamidates as most promising candidates for improving the conjugation of hydrophobic cargo to antibodies.

Next, we evaluated the set of fluorescent phosphonamidates in conjugation experiments with the therapeutically relevant monoclonal antibody (mAb) brentuximab (Fig. 2b). After reduction of intramolecular disulfides, we incubated the mAb with 10 equivalents excess of Cy5 **5a-d** for 16 h at 14 °C, using only a limited amount of DMSO cosolvent. After purification we obtained brentuximab-Cy5 conjugates with fluorophore to antibody ratios (FAR) ranging from 0.1 to 2.8 (Fig. 2c). Under the tested conjugation conditions, we observed the expected trend of increasing FAR for differently substituted phosphonamidates from low soluble ethyl **5d** and choline **5c** over moderately soluble PEG₂ **5a** to highly soluble PEG₁₂ **5b**, confirming the expected correlation between aqueous solubility and conjugation efficiency. The low FAR of ethyl- (FAR 0.2) and choline (FAR 0.3) substituted conjugates **5d** and **5c** is a result of their low solubility (30 μM) in the reaction buffer, which reduces the effective concentration to only one equivalent labeling reagent per antibody. This unfavored stoichiometry is improved when the more polar PEG₂- and PEG₁₂ phosphonamidates **5a** and **5b** are used. Especially the PEG₁₂ *O*-substituent increases aqueous solubility of the Cy5-phosphonamidate in buffer which allows the use of 100 equivalent excess during conjugation reactions yielding in an improved FAR of 5.9.

Encouraged by the beneficial solubility and conjugation efficiency of the PEG₁₂ substituted phosphonamidate, we decided to employ this structural motif to synthesize homogeneously modified ADCs from the challenging hydrophobic linker-payload Val-Cit-PAB-MMAE. In the linker-payload design of compound **7**, depicted in Fig. 3a, the PEG-substituent is

directly incorporated into the electrophilic phosphonamidate unit for the cysteine addition, which delivers a branched conjugation handle that introduces hydrophilicity to the hydrophobic linker-payload without increasing the distance between antibody and payload. This compact structure limits exposure of the hydrophobic linker-payload and should reduce unfavored accumulation and aggregation of highly conjugated ADCs. To synthesize the respective MMAE-phosphonamidate we used phosphonite **3b** in a SPHR with azido benzoic acid succinimide ester and reacted the resulting NHS-phosphonamidate building block **6** with VC-PAB-MMAE (Fig. 3a). With the resulting compact drug-linker conjugate **7** in hand, we set up an initial screening experiment to identify optimal conditions for a one-pot reduction and alkylation protocol that yields a DAR 8 ADC from native antibodies (Fig. 3b). To that end, we co-incubated brentuximab at 10 mg mL⁻¹ with 10 equivalents of **7** and increasing equivalents of TCEP. In this set of experiments, we observed that a maximum degree of modification with a DAR of 6.9 is reached with 8 equivalents of the reductant (Fig. 3c). Those 8 equivalents were carried over to a second experiment in which we increased the stoichiometry of **7** to 12 and 15 equivalents relative to the antibody. We identified that a slight excess of 12 equivalents of **7** is sufficient to generate a fully modified DAR 8 ADC in a one-pot reduction and alkylation protocol. Remarkably, this translates to only 1.5 equivalents **7** per cysteine, which allows a reagent efficient ADC production.

Next, we started the evaluation of the DAR 8 brentuximab-7 by determining its *in vitro* potency in a cell-based viability assay with three CD30-overexpressing cell lines and compared the cell killing activity to a DAR 4 brentuximab-7 and Adcetris (Fig. 4a-c). Across all tested antigen positive cell lines, we observed nearly identical cell-killing behavior for Adcetris and the DAR 4 brentuximab-7. These results indicate that the modification of



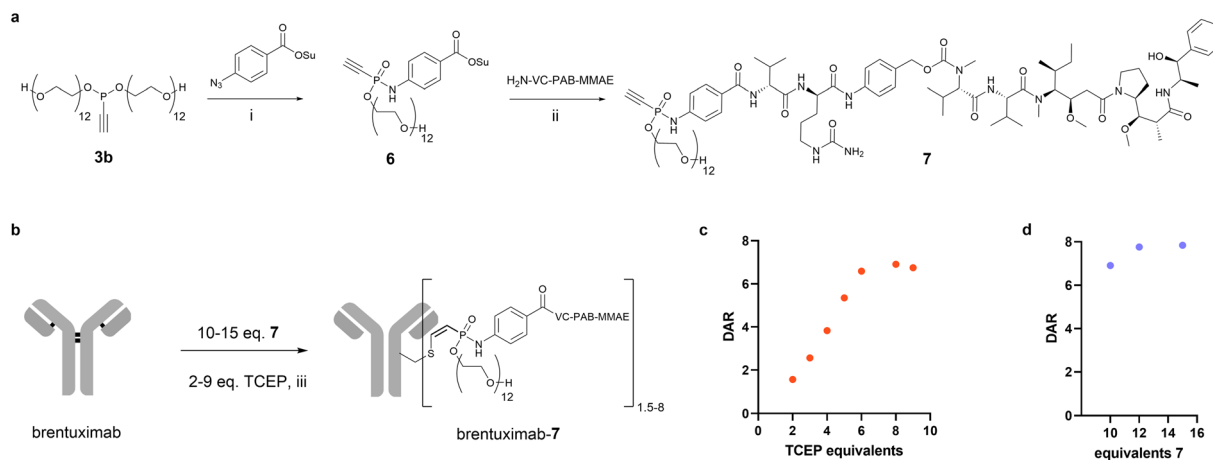


Fig. 3 Synthesis of ethynylphosphonamidate 7 and optimization of conjugation conditions to brentuximab. (a): Synthesis scheme for the preparation of ethynylphosphonamidate 7. (i): 4-Azidobenzoic acid-*N*-hydroxysuccinimide ester, DMF, rt, overnight, 68%; (ii): VC-PAB-MMAE, DIPEA, DMSO, overnight, 35%. (b): Equivalent screen of TCEP and 7 for one-pot reduction and alkylation protocol of brentuximab. (iii): 50 mM Tris-HCl, 1 mM EDTA, 100 mM NaCl, pH 8.5, 5% DMSO, 14 °C, 16 h. (c): DAR of brentuximab-7 depending on equivalents of TCEP with 10 equivalents 7; (d): DAR of brentuximab-7 depending on equivalents of 7 with 8 equivalents TCEP.

brentuximab with PEG₁₂ phosphonamidate 7 has negligible effect on the recognition, binding, and subsequent internalization. Furthermore, both ADCs had no effect on the HL60 control cell line (Fig. 4d). When we further compared the IC₅₀ values for the DAR 4 ADCs with the high loaded DAR 8 ADC, we observed a clear increase in potency, which is in-line with the higher payload loading of the DAR 8 brentuximab-7. To further evaluate cathepsin B cleavage and the traceless release of MMAE from the VC-PAB-linker after ADC internalization, we set up an indirect bystander assay for which we incubated CD30 positive L-540 cells with Adcetris, brentuximab-7 DAR4 or DAR8. In this setting, the ADC is internalized and exposed to intracellular cathepsin B activity. If the VC-peptide is recognized and cleaved by cathepsin B, free MMAE can passively diffuse across the cell membrane into the cell culture medium while uncleaved cysteine adducts would be trapped intracellularly. The supernatant of L-540 cells was then transferred to CD30 negative HL-60 cells and viability was assayed. As shown in Fig. 4e, we observed the same trend of potency for the three ADCs, indicating that PEG₁₂ substituted phosphonamidate VC-PAB-MMAE does not inhibit the cathepsin B mediated payload release in an *in vitro* potency setting.

To investigate, if the improved *in vitro* cell killing of DAR 8 brentuximab-7 is also translated to more efficient tumor growth inhibition, we assessed the *in vivo* efficacy of brentuximab-7 in a Karpas 299 derived tumor xenograft model. To that end, we adapted a previously reported study design, which revealed high antitumor activity of a DAR 4 phosphonamidate linked MMAE brentuximab. In the latter study a dosage of two times 1 milligram per kilogram of body weight (mg kg⁻¹) lead to complete tumor remission, while a reduced dosing of two times 0.5 mg kg⁻¹ displayed decreased activity with only partial tumor growth regression.³⁰ We assumed that the decreased antitumor activity, observed for the low dosing scheme, would substantially improve once the DAR is increased from 4 to 8, as long as the

high DAR does not trigger accelerated ADC clearance. To challenge this assumption, we treated four groups of 10 mice with either brentuximab-7 (DAR 8 and DAR 4), Adcetris (DAR 4) or PBS as vehicle control. All groups received four doses of 0.5 mg kg⁻¹ ADC every third day and tumor growth was monitored over an extended period of 74 days (Fig. 4g). Under these experimental settings we observed only temporary and incomplete tumor regression for Adcetris with 30% remission rate, which is in line with earlier reports.³¹ Remarkably, our DAR 4 brentuximab-7 displays improved efficacy compared to Adcetris with higher remission rate of 60%. This is also in line with our previous *in vivo* efficacy experiments and again can be explained with the high stability of phosphonamidate-linked conjugates that leads to prolonged payload delivery to the tumor when compared to less stable maleimide conjugates. The improved activity of brentuximab-7 is even more pronounced when the DAR is increased to 8 drugs per antibody. The DAR 8 treated group showed an excellent response to the treatment and was in tumor remission already a few days after ADC treatment. Only one mouse exhibited a tumor relapse and had to be sacrificed at day 25, while no relapse was observed for the remaining group over the whole observation period of 74 days, resulting in an excellent remission rate of 90%. This favored *in vivo* activity at a low dosing of 0.5 mg kg⁻¹ of the high loaded brentuximab-7 indicates that its plasma clearance is not affected by the payload increase and that the here presented compact branched PEG₁₂ phosphonamidate moderates the additional hydrophobicity to yield a sufficiently hydrophilic ADC. These results are in line with similar studies that investigated the effect of alternative hydrophilic linker on *in vivo* activity of high DAR ADCs.³³⁻³⁵ Notably, in these studies the highly validated but hydrophobic Val-Cit-PAB cleavage site had to be replaced with a glucuronide-cleavage site to achieve similar results. Thus, brentuximab-7 is the first example of a stable conjugated Val-Cit-PAB-MMAE based DAR 8 ADC with improved *in vivo* efficacy.



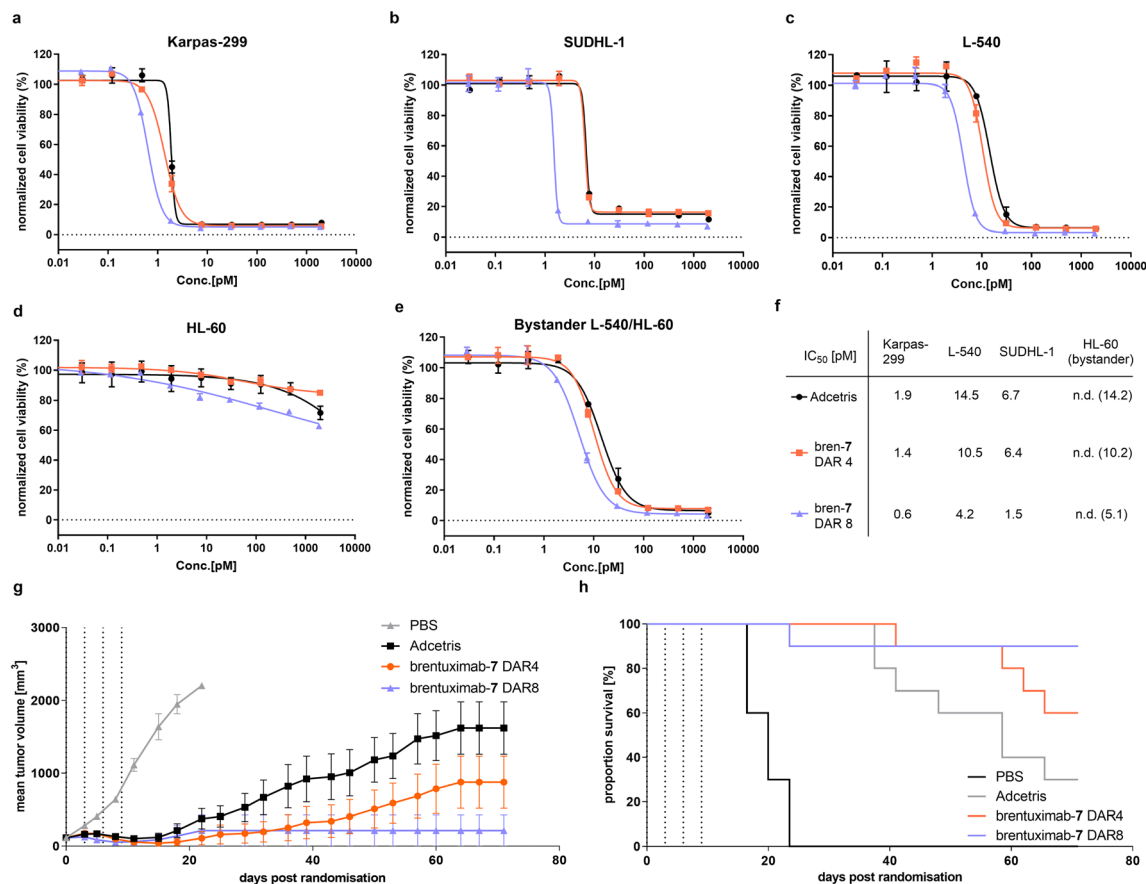


Fig. 4 *In vitro* and *in vivo* efficacy evaluation of brentuximab-7. (a–c): *In vitro* cell viability of different CD30 overexpressing cell lines in response to increasing concentration of ADC brentuximab-7 DAR 4 (orange) and DAR 8 (violet) in comparison to Adcetris (black). (d): *In vitro* cell viability of CD30 negative control cell line HL-60 in response to increasing concentration of ADC brentuximab-7 DAR 4 (orange) and DAR 8 (violet) in comparison to Adcetris (black). (e): *In vitro* cell viability of CD30 negative cell line HL-60 in response to supernatants of L-540 cells pre-incubated with increasing concentration of ADCs brentuximab-7 DAR 4 (orange) and DAR 8 (violet) in comparison to Adcetris (black). (f): Table summarizing IC₅₀ for cell viability of the tested ADCs. (g): Anti-tumor activity of ADC brentuximab-7 DAR 4 (orange) and DAR 8 (violet) in comparison to Adcetris (black) and vehicle control (PBS, grey) in Karpas 299 tumor xenograft model in SCID mice. Treatment of 0.5 mg kg⁻¹ was administered 4 times on day 0, 3, 6 and 9 post randomizations. Data are mean of *n* = 10 animals per group. (h): Kaplan–Meier survival plot of anti-tumor activity study shown in (g).

In the final part of our study, we focused on the pharmacokinetic behaviour of Val-Cit-PAB-MMAE based ADCs and set out to investigate if the here presented compact hydrophilic conjugation handle can compensate the additional hydrophobicity of a DAR 8 ADC to sufficiently ensure ADC circulation without enhanced clearance in comparison to the established DAR 4 Adcetris. As one can anticipate that the relative clearance of an ADC can be predicted from its overall hydrophobicity, we analyzed brentuximab-7 by hydrophobic interaction chromatography (HIC) and compared it to Adcetris (Fig. 5b). In contrast to the heterogeneously modified Adcetris, that is mainly comprised of zero-, two-, four- and six-times modified mAb, brentuximab-7 is a highly homogeneous ADC, eluting in a single narrow peak at a similar retention time as the DAR 6 Adcetris and thus significantly earlier than the Adcetris DAR 8. This reflects a reduced hydrophobicity of high loaded brentuximab-MMAE conjugates when the PEG₁₂ phosphonamidate 7 is used for ADC preparation. To estimate if these beneficial properties of PEG₁₂ phosphonamidates can be

further optimized, we prepared another brentuximab-MMAE conjugate with a PEG₂₄-substituted phosphonamidate 8. The HIC analysis of brentuximab-8 displays a similar narrow peak shape as observed for brentuximab-7 with a shift to even lower retention times, corresponding to a homogeneous and significantly more hydrophilic species in the range of a DAR 5 Adcetris. These results indicated that a brentuximab-8 might exhibit a similar PK profile as the marketed DAR 4 Adcetris.

Finally, we wanted to investigate if the decreased hydrophobicity is sufficient to translate into an advantageous PK profile with slow plasma clearance. To that end, we focused on the most hydrophilic DAR 8 ADC discovered in this study and treated female Sprague-Dawley rats with 5 mg kg⁻¹ of brentuximab-8 or Adcetris. The levels of total antibody and intact ADC were determined by ELISA at different time points. As displayed in Fig. 5c, both curves for total antibody show a slow decrease of blood concentration during the first 7 days of the study, pointing towards slow clearance and acceptable blood circulation. Remarkably, the high loaded brentuximab-8 shows very



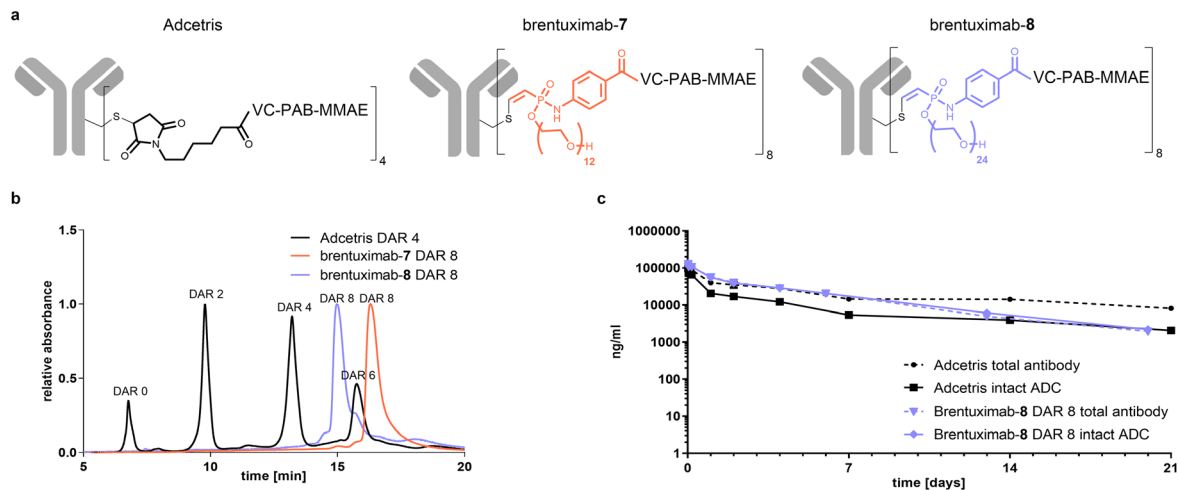


Fig. 5 a): Chemical structures of the different ADCs and their linker architectures compared in this study. (b): UV-traces of analytical hydrophobic interaction chromatography (HIC) of homogenous DAR 8 ADCs brentuximab-7 (orange) and brentuximab-8 (violet) in comparison to the heterogeneously modified DAR4 ADC Adcetris (grey). (c): Plasma pharmacokinetics of brentuximab-8 in female Sprague-Dawley rats. Data are mean of $n = 3$ animals per group and time point.

similar clearance compared to the total antibody of Adcetris. This slow clearance behavior indicates that the compact PEG₂₄-phosphonamidate compensates for the additional hydrophobicity introduced by increased DAR and that our here presented conjugation handle can improve the pharmacokinetic profile of high loaded ADCs, even when challenging hydrophobic linker-payloads, such as Val-Cit-PAB-MMAE are used. Looking closer at the intact ADC fraction of both conjugates, we see a clear drop for Adcetris during the first 7 days of circulation, reflecting a loss of payload, while the curve for intact brentuximab-8 perfectly correlates with its total antibody curve. This observation again confirms that the phosphonamidate linkage is highly stable *in vivo*, leading to higher concentration of intact ADC for longer periods of time while maleimide linked conjugates suffer from early drug loss and lower concentration of intact ADC. From day 7 onwards until the end of our observation period, both curves for total antibody and intact ADC of brentuximab-7 keep decreasing slowly to the level of intact Adcetris. This is a remarkable PK profile for a stable conjugated Val-Cit-PAB-MMAE based DAR 8 ADC, especially when keeping in mind that at this time of circulation the remaining Adcetris has a DAR far less than 4 due to linker payload loss caused by retro Michael addition. Moreover, those results confirm the excellent stability of the phosphonamidate linkage that has been measured in *ex vivo* settings, previously.²⁹

Conclusions

With this study, we demonstrate how the modular synthesis of ethynylphosphonamidates *via* chemoselective Staudinger-phosphonite reactions allows the straightforward preparation of hydrophilic labeling reagents that are suitable for the conjugation of challenging hydrophobic cargo molecules to non-engineered antibodies in a one-pot reduction and alkylation protocol. We identified ethynylphosphonamidates

substituted with hydrophilic PEG₁₂ as ideal conjugation handle that increases the aqueous solubility and conjugation efficiency of hydrophobic Cy5 dyes and employed the compact PEG₁₂ substituted ethynylphosphonamidate for the streamlined synthesis of homogeneously modified ADCs with eight drugs per antibody. Here, we chose the established linker-payload VC-PAB-MMAE as challenging hydrophobic model drug and demonstrated that the branched PEG₁₂ substituted ethynylphosphonamidate allows the generation of a DAR 8 brentuximab with only slight excess of 1.5 equivalents labelling reagent per cysteine. The resulting DAR 8 brentuximab-7 displays increased *in vitro* potency when compared to DAR 4 ADCs and most importantly maintains this advantage in an *in vivo* xenograft tumor model.

Here, the DAR 8 brentuximab-7 outperformed its DAR 4 analogue and the DAR 4 Adcetris, displaying an excellent *in vivo* efficacy, which indicates that the introduced hydrophilicity compensates for the overall hydrophobicity of 8 VC-PAB-MMAE units and sufficiently inhibits accumulation and aggregation of the ADC during circulation. Finally, we showed that the hydrophilicity of phosphonamidate linked ADCs can be further increased when longer PEG is incorporated into the conjugation handle and showed that a PEG₂₄ substituted VC-PAB-MMAE ethynylphosphonamidate delivers a DAR 8 brentuximab-8 that displays a remarkable pharmacokinetic profile with slow clearance from blood, which is in the range of lower modified Adcetris. In conclusion, the here presented PEG substituted compact phosphonamidate conjugation handle combines outstanding linkage stability and a branched hydrophilic architecture that enables the generation of highly efficacious DAR 8 ADCs from challenging hydrophobic payloads and native antibodies without impairing the favorable pharmacokinetic profile of monoclonal antibody conjugates. To the best of our knowledge, we here present the first example of a Val-Cit-PAB-based MMAE linker-payload that is applicable to generate



DAR 8 ADCs without enhanced clearance *in vivo*, emphasizing the potential of branched PEG phosphoramidates to conjugate even more hydrophobic and therefore demanding payloads to antibodies in high DAR. Consequently, we highlight how adequate linker design opens synthetic pathways to well defined protein-conjugates with improved properties that may help to develop future biological therapeutics with beneficial therapeutic indices.

Data availability

The datasets supporting this article have been uploaded as part of the ESI material.†

Author contributions

P. O., M.-A. K. and C. P. R. H. conceptualization; P. O., J. J., P. M., I. M., M.-A. K. and C. P. R. H. investigation; P. O., J. J., M.-A. K. formal analysis; P. O., M.-A. K. and C. P. R. H. writing-original draft; P. O., M.-A. K. and C. P. R. H. writing-review and editing; P. O., D. S., J. H., M.-A. K. and C. P. R. H. supervision and funding acquisition.

Conflicts of interest

The technology described in the manuscript is part of a pending patent application by P. O., J. J., P. M., I. M., D. S., H. L., J. H., and M.-A. K, C. P. R. H. P. O., P. M., I. M., D. S., J. H., and M.-A. K are employers of Tubulis.

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Notes and references

- D. Schumacher and C. P. R. Hackenberger, *Curr. Opin. Chem. Biol.*, 2014, **22**, 62–69.
- S. J. Walsh, J. D. Bargh, F. M. Dannheim, A. R. Hanby, H. Seki, A. J. Counsell, X. Ou, E. Fowler, N. Ashman, Y. Takada, A. Isidro-Llobet, J. S. Parker, J. S. Carroll and D. R. Spring, *Chem. Soc. Rev.*, 2021, **50**, 1305–1353.
- P. Khongorzul, C. J. Ling, F. U. Khan, A. U. Ihsan and J. Zhang, *Mol. Cancer Res.*, 2020, **18**, 3–19.
- J. Z. Drago, S. Modi and S. Chandarlapaty, *Nat. Rev. Clin. Oncol.*, 2021, **18**, 327–344.
- J. M. Chalker, G. J. L. Bernardes, Y. A. Lin and B. G. Davis, *Chem. – Asian J.*, 2009, **4**, 630–640.
- S. B. Gunnoo and A. Madder, *ChemBioChem*, 2016, **17**, 529–553.
- P. Ochtrop and C. P. R. Hackenberger, *Curr. Opin. Chem. Biol.*, 2020, **58**, 28–36.
- M. R. Lewis and J. E. Shively, *Bioconjugate Chem.*, 1998, **9**, 72–86.
- S. C. Alley, D. R. Benjamin, S. C. Jeffrey, N. M. Okeley, D. L. Meyer, R. J. Sanderson and P. D. Senter, *Bioconjugate Chem.*, 2008, **19**, 759–765.
- S. D. Fontaine, R. Reid, L. Robinson, G. W. Ashley and D. V. Santi, *Bioconjugate Chem.*, 2015, **26**, 145–152.
- C. Wei, G. Zhang, T. Clark, F. Barletta, L. N. Tumey, B. Rago, S. Hansel and X. Han, *Anal. Chem.*, 2016, **88**, 4979–4986.
- R. P. Lyon, J. R. Setter, T. D. Bovee, S. O. Doronina, J. H. Hunter, M. E. Anderson, C. L. Balasubramanian, S. M. Duniho, C. I. Leiske, F. Li and P. D. Senter, *Nat. Biotechnol.*, 2014, **32**, 1059–1062.
- J. M. J. M. Ravasco, H. Faustino, A. Trindade and P. M. P. Gois, *Chem. – Eur. J.*, 2019, **25**, 43–59.
- S. Ariyasu, H. Hayashi, B. Xing and S. Chiba, *Bioconjugate Chem.*, 2017, **28**, 897–902.
- B. Bernardim, P. M. S. D. Cal, M. J. Matos, B. L. Oliveira, N. Martínez-Sáez, I. S. Albuquerque, E. Perkins, F. Corzana, A. C. B. Burtoloso, G. Jiménez-Osés and G. J. L. Bernardes, *Nat. Commun.*, 2016, **7**, 13128.
- N. J. Smith, K. Rohlfing, L. A. Sawicki, P. M. Kharkar, S. J. Boyd, A. M. Kloxin and J. M. Fox, *Org. Biomol. Chem.*, 2018, **16**, 2164–2169.
- S. Shaunak, A. Godwin, J. W. Choi, S. Balan, E. Pedone, D. Vijayarangam, S. Heidelberger, I. Teo, M. Zloh and S. Brocchini, *Nat. Chem. Biol.*, 2006, **2**, 312–313.
- H. Khalili, A. Godwin, J. W. Choi, R. Lever and S. Brocchini, *Bioconjugate Chem.*, 2012, **23**, 2262–2277.
- R. Huang, Z. Li, Y. Sheng, J. Yu, Y. Wu, Y. Zhan, H. Chen and B. Jiang, *Org. Lett.*, 2018, **20**, 6526–6529.
- M. J. Matos, C. D. Navo, T. Hakala, X. Ferhati, A. Guerreiro, D. Hartmann, B. Bernardim, K. L. Saar, I. Compañón, F. Corzana, T. P. J. Knowles, G. Jiménez-Osés and G. J. L. Bernardes, *Angew. Chem., Int. Ed.*, 2019, **58**, 6640–6644.
- S. J. Walsh, S. Omarjee, W. R. J. D. Galloway, T. T. L. Kwan, H. F. Sore, J. S. Parker, M. Hyvönen, J. S. Carroll and D. R. Spring, *Chem. Sci.*, 2019, **10**, 694–700.
- A. J. Counsell, S. J. Walsh, N. S. Robertson, H. F. Sore and D. R. Spring, *Org. Biomol. Chem.*, 2020, **18**, 4739–4743.
- F. F. Schumacher, J. P. M. Nunes, A. Maruani, V. Chudasama, M. E. B. Smith, K. A. Chester, J. R. Baker and S. Caddick, *Org. Biomol. Chem.*, 2014, **12**, 7261–7269.
- J. P. M. Nunes, M. Morais, V. Vassileva, E. Robinson, V. S. Rajkumar, M. E. B. Smith, R. B. Pedley, S. Caddick, J. R. Baker and V. Chudasama, *Chem. Commun.*, 2015, **51**, 10624–10627.



- 25 M. Morais, J. P. M. Nunes, K. Karu, N. Forte, I. Benni, M. E. B. Smith, S. Caddick, V. Chudasama and J. R. Baker, *Org. Biomol. Chem.*, 2017, **15**, 2947–2952.
- 26 A. Maruani, M. E. B. Smith, E. Miranda, K. A. Chester, V. Chudasama and S. Caddick, *Nat. Commun.*, 2015, **6**, 2–10.
- 27 C. Bahou, D. A. Richards, A. Maruani, E. A. Love, F. Javaid, S. Caddick, J. R. Baker and V. Chudasama, *Org. Biomol. Chem.*, 2018, **16**, 1359–1366.
- 28 M.-A. Kasper, M. Glanz, A. Oder, P. Schmieder, J. P. von Kries and C. P. R. Hackenberger, *Chem. Sci.*, 2019, **10**, 6322–6329.
- 29 M. A. Kasper, M. Glanz, A. Stengl, M. Penkert, S. Klenk, T. Sauer, D. Schumacher, J. Helma, E. Krause, M. C. Cardoso, H. Leonhardt and C. P. R. Hackenberger, *Angew. Chem., Int. Ed.*, 2019, **58**, 11625–11630.
- 30 M. A. Kasper, A. Stengl, P. Ochtrup, M. Gerlach, T. Stoschek, D. Schumacher, J. Helma, M. Penkert, E. Krause, H. Leonhardt and C. P. R. Hackenberger, *Angew. Chem., Int. Ed.*, 2019, **58**, 11631–11636.
- 31 K. J. Hamblett, P. D. Senter, D. F. Chace, M. M. C. Sun, J. Lenox, C. G. Cervený, K. M. Kissler, S. X. Bernhardt, A. K. Kopcha, R. F. Zabinski, D. L. Meyer and J. A. Francisco, *Clin. Cancer Res.*, 2004, **10**, 7063–7070.
- 32 P. Malik, C. Phipps, A. Edginton and J. Blay, *Pharm. Res.*, 2017, **34**, 2579–2595.
- 33 R. P. Lyon, T. D. Bovee, S. O. Doronina, P. J. Burke, J. H. Hunter, H. D. Neff-Laford, M. Jonas, M. E. Anderson, J. R. Setter and P. D. Senter, *Nat. Biotechnol.*, 2015, **33**, 733–735.
- 34 W. Viricel, G. Fournet, S. Beaumel, E. Perrial, S. Papot, C. Dumontet and B. Joseph, *Chem. Sci.*, 2019, **10**, 4048–4053.
- 35 P. J. Burke, J. Z. Hamilton, S. C. Jeffrey, J. H. Hunter, S. O. Doronina, N. M. Okeley, J. B. Miyamoto, M. E. Anderson, I. J. Stone, M. L. Ulrich, J. K. Simmons, E. E. McKinney, P. D. Senter and R. P. Lyon, *Mol. Cancer Ther.*, 2017, **16**, 116–123.

