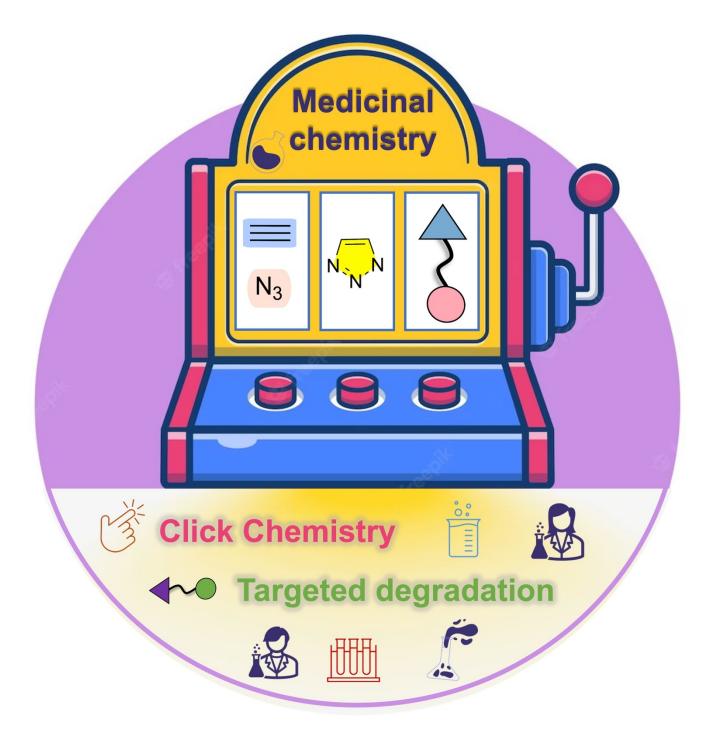




Click Chemistry and Targeted Degradation: A Winning Combination for Medicinal Chemists?

Anna Pasieka,^[a] Eleonora Diamanti,^[a] Elisa Uliassi,^[a] and Maria Laura Bolognesi^{*[a]}





Click chemistry is universally recognized as a powerful strategy for the fast and precise assembly of diverse building blocks. Targeted Protein Degradation (TPD) is a new therapeutic modality based on heterobifunctional small-molecule degraders that provides new opportunities to medicinal chemists dealing with undruggable targets and incurable diseases. Here, we highlight how very recently the TPD field and that of click

1. Introduction

Once a year the chemistry community stops research waiting for the announcement of its most prestigious award: the Nobel Prize. Last year, the Royal Swedish Academy of Sciences awarded The Nobel Prize in Chemistry 2022 to three great scientists: Carolyn R. Bertozzi, Morten Meldal and K. Barry Sharpless "for the development of click chemistry and bioorthogonal chemistry". The choice was not surprising, when looking at recent publications' trends in the field, where these two strategies are flourishing.

Prof. Sharpless and prof. Meldal independently reported the high-yielded and regioselective copper-catalyzed Huisgen cy-

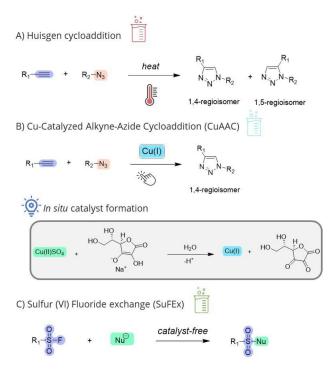


Figure 1. Click chemistry reactions: A) Huisgen cycloaddition reaction; B) Cucatalyzed azide-alkyne cycloaddition (CuAAC); C) sulfur(VI) fluoride exchange (SuFEx).

 [a] Dr. A. Pasieka, Dr. E. Diamanti, Dr. E. Uliassi, Prof. M. Laura Bolognesi Department of Pharmacy and Biotechnology Alma Mater Studiorum-University of Bologna Via Belmeloro 6, 40126 Bologna (Italy) E-mail: marialaura.bolognesi@unibo.it

© © 2023 The Authors. ChemMedChem published by Wiley-VCH GmbH. This is an open access article under the terms of the Creative Commons Attribution License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited. chemistry have merged, opening up the possibility for finetuning the properties of a degrader, chemically assembled through a "click" synthesis. By reviewing concrete examples, we want to provide the reader with the insight that the application of click and bioorthogonal chemistry in the TDP field may be a winning combination.

cloaddition reaction (Figure 1A), which now is recognized as the flagship of "click chemistry".^[1,2] Few years later, the idea that similarly simple and fast reactions could be performed in living organisms^[3] was introduced by prof. Bertozzi, who performed the first "bioorthogonal reaction" in cells, i.e., the Staudinger ligation of azides with triarylphosphines.^[4] Over the years, these two methodologies, have completely changed the field of material science, but also chemical biology, medicinal chemistry, and even drug discovery.^[5-7] In fact, these two new synthetic philosophies have become a perfect answer for the growing needs of pharma: in addition to the features mentioned above, click reactions satisfy the principles of green chemistry, such as prevention of hazardous waste and byproducts, minimized derivatization due to better selectivity and tolerance (insensitive to oxygen and water), atom economy, use of safer chemicals and solvents (environmentally benign), energy minimization (a lot of click reactions can occur with less or no heating conditions).

The term "click chemistry" was first coined in 2001^[8] and referred to modular, wide in scope, stereospecific reactions in simple conditions giving a single product in very high yields after an easy isolation. The used reagents and solvents were requested to be commonly available, benign and easily removed (e.g., water).^[8] Some reactions were found to meet these requirements and included in the list: e.g., cycloadditions of unsaturated species (especially 1,3-dipolar cycloaddition reactions), the Diels-Alder family of transformations, nucleophilic substitution chemistry, and carbonyl chemistry of the "non-aldol" type. However, the real breakthrough come from the discovery from both Sharpless and Meldal that copper(I) iodide substantially catalyzed the Huisgen 1,3-dipolar cycloaddition reaction between azides and terminal alkynes forming 1,4-regioisomers of 1,2,3-triazoles.

Thanks to this discovery, the current gold standard of click chemistry is the Cu-catalyzed azide-alkyne cycloaddition (CuAAC; Figure 1B).^[9] In most applications, copper(II) sulfate is used as a catalyst together with a reducing agent – sodium ascorbate – to produce Cu(I) *in situ*. In contrast to the small diversity of catalysts, the reaction can be set up with a great variety of solvents, depending on the solubility of substrates. The most common choice is a mixture of solvents: water and miscible (sometimes just partially) organic solvents e.g., *tert*-BuOH or different alcohols. Today, there are other popular examples of "click chemistry" reactions, e.g., thiol-ene reaction, [4+1] cycloadditions between isonitriles (isocyanides) and tetrazines, which have been discussed in other recent reviews.^[10-12]



The sulfur(VI) fluoride exchange (SuFEx) represents the most recent set of click chemistry transformations: a metal-free click chemistry reaction, discovered by Sharpless in 2014.^[13] Specifically, SuFEx exploited the elevated reactivity of sulfur(VI) fluorides with appropriate nucleophiles, including amines (Figure 1C).^[13,14] Similar to CuAAC, SuFEx reactions have been reported to be highly efficient (high conversion), water friendly and easy to set up, while significantly differing from the previous one for harnessing metal catalyst-free protocols. Moreover, the easy preparation and good properties of sulfonyl fluorides, such as the insensitivity to ambient oxygen and the hydrolytical stability, are other important advantages. Initially, sulfuryl fluoride (SO₂F₂) was used in form of gas, but, due to obvious drawbacks, over time it has been replaced with more stable, solid FSO₂-carrying reagents, such as fluorosulfuryl imidazolium salts.^[15] The most typical reactions revolve around the coupling of sulfonyl fluorides with silyl ethers or primary/ secondary amines resulting in the formation of S-O bond or sulfonamides, respectively.^[16,17]

To note, the rapid formation of sulfonamides has particularly attracted the interest of medicinal chemists.^[18] Another intriguing aspect is that SuFEx technology can be exploited not only for creating connections, but also as for preparing reagents in their own right.^[19,20]

The concept of "bioorthogonal chemistry" is very similar to "click chemistry", so that the two are often referred interchangeably. Yet, there is a crucial difference: for the first, the reagents are not toxic for living organisms and therefore, reactions can be performed *in vivo*.^[21,22] Three out of the most famous



Anna Pasieka obtained her PhD degree with distinction in Pharmaceutical Sciences under the supervision of Professor Barbara Malawska in October 2021 in Jagiellonian University in Kraków. In her PhD thesis she was focused on new multifunctional ligands with potential activity in Alzheimer's disease. During PhD studies, she was awarded a COST STSM within the "MuTaLig" Action spending three months at University of Barcelona. After graduation, she worked as medicinal chemist in the biopharmaceutical company in Warsaw. Since November 2022 Anna is a postdoc in the prof. Bolognesi's group and works in the research project "PROLEISH (PROTACs to treat leishmaniasis)".



Eleonora Diamanti was trained as medicinal chemist and did her Ph.D. with Prof. Piomelli (IIT, Genova) working one year with Prof. Aggarwal in Bristol on the total synthesis of a natural product to treat Mycobacterium tuberculosis. In 2016, she joined the Hirsch group as a postdoctoral fellow at the University of Groningen and she then moved to the Helmholtz institute (HIPS, Saarbrücken) focusing on the design and synthesis of antiinfective agents. Since 2023, she is Assistant Professor at the University of Bologna working on the development of PROTAC to combat the antimicrobial resistance problem. bioorthogonal transformations are (i) the Staudinger ligation^[4,23], (ii) the strain-promoted azide-alkyne cycloaddition (SPAAC), also called as "Cu-free click chemistry"^[21,24] and (iii) inverse electron demand Diels-Alder (IEDDA) cycloaddition (Figure 2).^[25-27] The Staudinger reaction is the first reported bioorthogonal reaction between an azide and a phosphine, resulting in an amide.[4,28] In turn, SPAAC reaction was developed by Bertozzi to eliminate the cytotoxic effect of copper in vivo. Instead of using the catalyst, the alkyne moiety is introduced in a strained difluorooctyne (DIFO), dibenzocyclooctyne (DBCO) or biarylazacyclooctynone (BARAC). These scaffolds allowed for the activation of the alkyne fragment and thus, the formation of the triazole ring.^[22,24,29] IEDDA is believed as the fastest bioorthogonal reaction (Figure 2). It is performed between tetrazine and trans-cyclooctene (TCO) derivatives and, since its description by Blackman, is successfully used for many biomedical applications.^[25-27,30]

The simplicity, speed, ease, and wide scope of click/ bioorthogonal chemistry have been so much appreciated that both strategies are widely used in the search of many drugs, e.g., anticancers, antivirals, peptidomimetics, antiparasitics or anti-inflammatories.^[31,32]

As highlighted by the Nobel Committee,^[33] thanks to its modular nature and robustness, the principle may be equated to an *"IKEA* 'flatpack' approach, in which all necessary components, the 'building blocks', along with a set of easy-to-use hardware, the 'reactions', were provided together with a reliable assembly instruction for almost anybody to follow". On this basis, it is intuitive that click chemistry is particularly



Elisa Uliassi received her M.Sc. degree in Pharmaceutical Chemistry and Technology in 2012, and her Ph.D. in Chemistry in 2016, from the University of Bologna. During her Ph.D. program, she was a visiting Ph.D. student in Prof. L. A. Romeiro's laboratory at Universidade Católica de Brasília, and then in Dr. K. A. Jacobson's laboratory at the National Institutes of Health (U.S.). She was a Visiting Researcher in Prof. M. J. Fuchter's laboratory at the Imperial College of London in 2021. She is currently an Assistant Professor at the Department of Pharmacy and Biotechnology of the University of Bologna.



Maria Laura Bolognesi is Professor of Medicinal Chemistry, Coordinator of the Biotechnological, Biocomputational, Pharmaceutical and Pharmacological Sciences PhD Program at the University of Bologna. Her research is concerned with the design, synthesis, and biological investigation of chimeric small molecules in the neurodegenerative and neglected tropical disease therapeutic areas. She has a track record of 200 publications in highranked scientific journals, including patents and patent applications, and numerous invited talks worldwide. Review doi.org/10.1002/cmdc.202300422



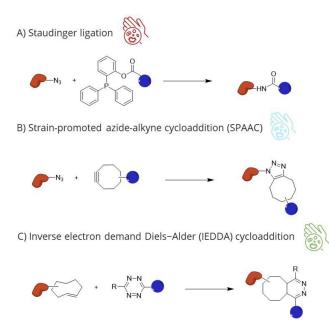


Figure 2. Bioorthogonal chemistry transformations: A) Staudinger ligation; B) strain-promoted azide-alkyne cycloaddition (SPAAC); C) inverse electron demand Diels-Alder (IEDDA) cycloaddition.

powerful in the synthesis of heterobifunctional molecules, i.e., molecules consisting of two protein-binding moieties joined *via* a linker of appropriate length.^[34] This particularly applies to degraders, an innovative class of heterobifunctional agents that have been recently brought to the forefront in medicinal chemistry.^[35-37]

There are different groups of degraders: PROTACs – proteolysis targeting chimeras, LYTACs – lysosome targeting chimeras,^[38] AUTACs – autophagy targeting chimeras,^[39,40] hydrophobic tagging (HyT) and RIBOTACs.^[41–46] They all consist of two *'building blocks'*: a ligand binding the target of interest and a moiety recruiting the target responsible of inducing degradation, connected *via* a suitable linker (Figure 3).

Specifically, PROTACs, featuring a ligand for the protein of interest (POI) joined with a spacer to an E3 ubiquitin ligase binder, promote POI ubiquitination through the formation of the ternary POI-PROTAC-E3 ligase complex, and the consequent selective degradation by the proteasome. It is clear that linker length, composition and rigidity are all crucial for creating a productive ternary complex and, consequently, for inducing the degradation process.^[47–49] As a result, the selectivity, ease, rapidity, and modularity of click ligations make them nearly ideally suited for the construction of degraders, a process that otherwise involves multi-step protocols and difficult purifications. These are very tedious tasks and performing successful linker chemistry, i.e., allowing the combination of the different recruiting elements in a biologically benign, fast, and selective fashion, remains a complex endeavor. Conversely, click chemistry may enable rapid structure-activity relationship (SAR) profiling, ensuring a fast exploration of the chemical space in terms of ligand assembly and probing different attachment points, as well as linker length.

However, it was not only the simplicity and speed of click and bioorthogonal chemistry that have attracted great interest for the development of new degraders. The introduction of the 1,2,3-triazole ring into a linker could change the physicochemical properties of the molecule in terms of topological polar surface area (TPSA) and lipophilicity (distribution coefficient, logD and cLogP) and hence, improve cell permeability, solubility or stability.^[50,51] Moreover, a triazole ring increases the rigidity of the linker, potentially allowing the proper proximity of POI and E3 ligase to form an effective ternary POI-PROTAC-E3 ligase complex.^[52,53] In turn, harnessing bioorthogonal chemistry may overcome problems related to PROTAC delivery into the cells (mainly due to their high molecular weight). This is achieved through the delivery of smaller POI- and E3 ligaseligand precursors, which can react directly in the cells providing the final degraders, termed CLIPTACs (i.e., in-cell click-formed proteolysis-targeting chimeras).[50,54]

Herein, it is our goal to highlight the most interesting, exciting, and useful points of intersection between click chemistry and degrader discovery, with a particular emphasis on PROTACs. Given the boom of this topic, the current review will focus on selected examples that aim to highlight the growing impact and potentialities offered by the combination of these two approaches.

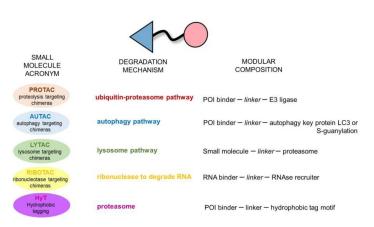


Figure 3. Groups of heterobifunctional degraders, together with their mechanism of degradation and modular composition.

2. Cu-Catalyzed Azide-Alkyne Cycloaddition (CuAAC)

2.1. "Click chemistry" and targeted degradation: the debut

In 2017, two back-to-back articles were published in a J. Med Chem Special issue dedicated to "Inducing Protein Degradation as a Therapeutic Strategy", which marked the debut of CuAAC in PROTAC synthesis.^[55] Jung's group synthetized triazole-based PROTACs targeting sirtuins (Sirt)^[56] – a family of histone deacetylases, involved in the pathogenesis of many diseases.^[57] The starting point was the Sirt2-selective and potent SirReal, selected as the POI ligand,^[58] and the well-known cereblon (CRBN) ligand thalidomide - chosen as E3 ligase binder. The final coupling of the alkynylated derivative of the POI ligand with a thalidomide-derived azide was performed in water: tert-BuOH (1:1) mixture by addition of copper sulphate and sodium ascorbate. The reaction was carried out at 60 °C for the first hour, and then at room temperature for 16 hours. The first-inclass triazole-based PROTAC (1) (Figure 4) was obtained in 33% yield. For the first time, 1- induced Sirt degradation demonstrated that the exploitation of a triazole linkage had the potential of accelerating the synthesis of novel degraders and inducing the degradation by creating the ternary complex.^[56]

Intriguingly, in the same special issue Wurz et al. proposed the first "click chemistry platform" to obtain rapidly and efficiently novel PROTACs by CuAAC.^[59] To validate this breakthrough idea, the authors focused on PROTACs directed to bromodomain BRD4. As the POI ligand, the well-known JQ-1 moiety binding to BRD4^[60] was converted from a *tert*-butyl ester to an azido derivative by reaction with 2-azidoethanamine. As E3 ligase binders, VHL and CRBN ligands functionalized with various PEG linkers terminating with alkyne functionalities were prepared. The typical CuAAC reaction conditions (CuSO₄, sodium ascorbate, 0.1 M in THF/water, room temperature, 3 hours) were used to obtain ten final compounds (2-11) in optimal yields (55-90%) (Figure 5). Proximity and protein degradation assays proved the ability of the triazole-based PROTACs 2-11 to create the ternary complex and thus, to induce the proteasome mediated degradation.^[59]

2.2. CuAAC reaction in the synthesis of new PROTACs

A similar synthetic procedure was used in the optimization of first-in-class cyclin-dependent kinase 2 (CDK2) selective degraders.^[61] As the POI ligand, CDK2 inhibitor J2 was selected (Figure 6). In turn, thalidomide was used as the E3 ligase binder. First, CDK2 binders with terminal alkyne group were prepared. Then, to optimize the linker length, various azide-terminated PEG-thalidomide were synthetized. Eventually, using CuAAC reaction (CuSO₄, sodium ascorbate, DMF/water, 70 °C, 4 hours) both fragments were rapidly combined to provide four final compounds with yields ranging from 30 to 60%. The most promising CDK2 degrader (12) showed not only the selective degradation of CDK2 protein but was also effective in acute myeloid leukemia (AML) primary cells. Therefore, triazole-based PROTAC 12 demonstrated a potential application in AML.^[61,62]

Fuchs and coworkers decided to examine the impact of triazole position within various linker on physicochemical and biological properties of novel CDK9 degraders.^[63] Pan-CDK9 inhibitor AT7519 was selected as the POI ligand^[64] and thalidomide as the E3 ligase binder. The series of ten PROTACs containing triazole-alkyl linkers of different length (13-22, Figure 7) were synthetized using CuAAC reaction (CuSO₄, sodium ascorbate, THF/water, RT, above 20 h) with yields ranging from 18 to 100%. Then, their degradation activity, kinetic solubility and lipophilicity were evaluated. The data confirmed that compounds with greater cell permeability have higher in vitro biological activity. Of note, while keeping constant the linker length between the POI and E3 ligase ligands, the proximity of the triazole ring to the piperidine or benzene ring significantly influenced the lipophilicity and aqueous solubility. It is important to highlight that the relationship between degradation activity and physicochemical properties is not always fully clear. PROTAC permeability seems to be an important factor, but not crucial to create a ternary complex and induce POI degradation. Nevertheless, this research has showed that slight differences in the structure and the use of click chemistry may help in improving drug-likeness of novel degraders, already at the early stage.^[63]

Thanks to its effectiveness, the Wurz's strategy^[59] was applied in the synthesis of PROTACs targeting glycogen synthase kinase 3 (GSK-3 β) for the treatment of Alzheimer's disease (AD). To explore the potential degraders, the well-

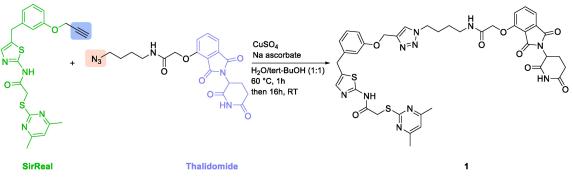


Figure 4. Synthesis of first-in-class triazole-based PROTAC (1) via CuAAC.

Review doi.org/10.1002/cmdc.202300422

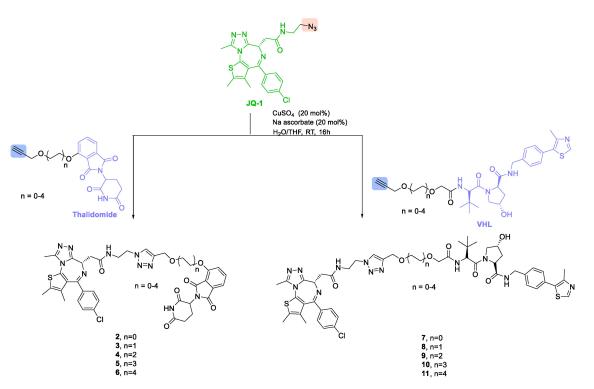


Figure 5. Click chemistry platform to obtain a library of novel PROTACs (2-11).

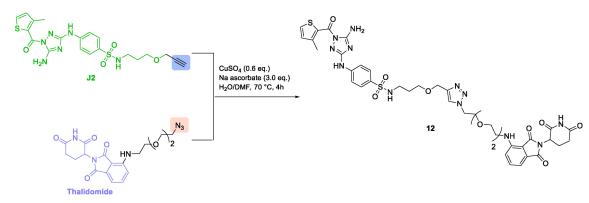


Figure 6. Synthesis of CDK2 selective PROTAC (12) via CuAAC.

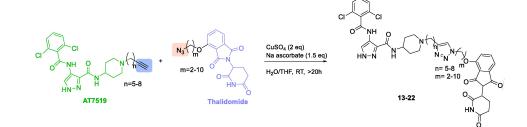


Figure 7. Synthesis of CDK9-directed PROTACs (13-22) via CuAAC.

known GSK-3 β inhibitor AZD2858 was used. In turn, as E3 ligase binders, a library of different CRBN (pomalidomide, lenalidomide) and von Hippel-Lindau (VHL) ligands were used. More than thirty triazole-based PROTACs were obtained in a click reaction (CuSO₄, sodium ascorbate, a mixture of water/tert-

BuOH/DCM/DMF, RT) with yields ranging from 20 to 80%. Among them, PT-65 (23) (Figure 8) was selected as the most promising agent with degradation effect in the nanomolar range and a positive outcome in an *in vivo* AD model.^[65] The SAR study showed that the linker featuring PEG and triazole

ChemMedChem 2023, 18, e202300422 (6 of 17)

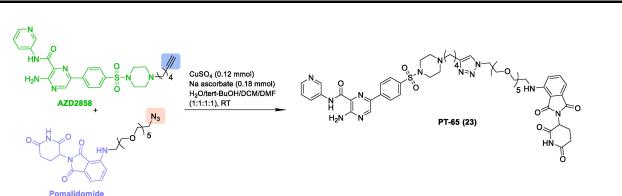


Figure 8. Synthesis of GSK-3 β -directed PROTAC 23 via CuAAC.

was the optimal for the biological activity. Moreover, with the increasing length of the linker, the activity was reduced. It was observed that the physicochemical properties of PROTACs played a key role in achieving the desired effect – three of the most active molecules showed similar logP, bioavailability and membrane permeability.^[65] This was not the only attempt to the search for new degraders in neurodegenerative diseases. Other series of GSK-3 β PROTACs based on the potent pyridinethiazole inhibitor was reported.^[66,67] Among these compounds, the triazole linker turned out to be a key spacer to form a productive ternary complex and thus, to induce the GSK-3 β degradation.^[66]

A slightly different condition of CuAAC reaction was used in the preparation of histone deacetylase 6 (HDAC6) degraders.^[68] HDAC6 is a protein responsible for deacetylation of some cytoplasmic proteins and its dysregulation is involved in cancer development.^[69] Yang and colleagues designed and synthetized four PROTACs consisting of the crebinostat structure (POI ligand)^[70] and thalidomide moiety connected *via* different triazole-PEG linkers. The click reactions were performed in a mixture of water/*tert*-BuOH with the addition of a complexing ligand - tris[(1-benzyl-1*H*-1,2,3-triazol-4-yl)methyl]amine (TBTA) in good yields (60-70%). The use of TBTA accelerates the CuAAC reaction by maintaining the Cu(I) oxidation state and avoiding the oxidation of catalytic Cu(I) by dissolved oxygen.^[71]

The aldehyde handle on the linker was then exploited for the condensation with the "HDAC biasing" acyl hydrazide reagent (Figure 9A). Compound 24 a with a 3-PEG linker displayed the highest degradation effect (Figure 9B).^[68] Further research resulted in the development of a second-generation HDAC6 degraders with improved potency and selectivity. The most potent compound (24 b) (Figure 9B) degraded HDAC6 at a nanomolar concentration and showed antiproliferative effect in multiple myeloma cells.^[72] Munoz et al. synthetized just one niclosamide based anti-cancer PROTAC via a CuAAC protocol. The PROTAC consists of a POI ligand, anthelmintic drug niclosamide, connected by a linker to the E3 ligase binder VH-032. Compared to the previously described conditions, the authors used different combination of reagents and solvents: copper iodide, hexamethylphosphoramide (HMPA) and N,Ndiisopropylethylamine (DIPEA), resulting in 38% yield. The addition of DIPEA accelerated copper(I)-acetylide formation, and thus, increased reaction rates.^[71] Then, an amide coupling with a VHL ligand was performed, leading to the final compound 25 (Figure 10). Despite the positive in vitro antiproliferative results, for 25 the degradation activity was not confirmed.^[73] This example, clearly demonstrates that the introduction of a triazole ring via click chemistry cannot be a one-size-fits-all-solution. On the other hand, CuAAC were profitably applied in the development of PROTACs targeting, among

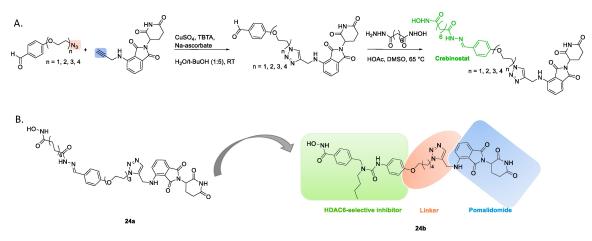
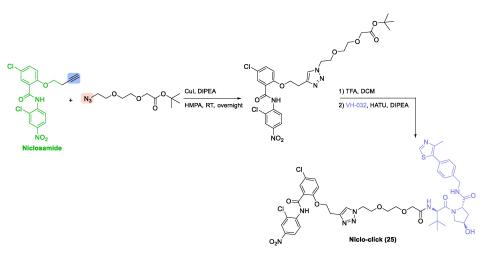


Figure 9. A) Synthesis of HDAC6-directed PROTACs via CuAAC; B) Design of second-generation HDAC6-directed PROTAC 24b from 24a.

ChemMedChem 2023, 18, e202300422 (7 of 17)

Review doi.org/10.1002/cmdc.202300422



cancer cells.^[77]

Figure 10. Synthesis of Niclo-click PROTAC 25 via CuAAC.

others, estrogen receptor alpha (ERa),^[74] Fms-like tyrosine receptor kinase 3 (FLT3),^[75] or cyclin-dependent kinase 6 (CDK6).^[76]

2.3. Dual PROTACs

The implementation of "click chemistry" has contributed to expand the research scope of PROTACs to new modalities, leading to the development of so-called *multitarget* PROTACs, i.e., single molecules able to recruit and degrade multiple target proteins.^[43] Based on this, dual triazole-based PROTACs targeting simultaneously the epidermal growth factor receptor (EGFR) and poly (ADP-ribose) polymerase (PARP), have been developed.^[77] The two POI warheads, namely gefitinib and olaparib, were combined *via* trifunctional natural amino acids (as star-type core linkers) with the respective E3 ligand, *via* a CuAAC protocol (Figure 11). This convergent synthetic strategy (CuSO₄, sodium ascorbate, water/THF, RT, maximum 3 h) resulted in the efficient development of eight dual PROTACs,

2.4. Hydrophobic tagging (HyT) via click chemistry

The CuAAC was also successfully exploited for the development of hydrophobic tag-based degraders. The hydrophobic tagging (HyT) builds on the idea that the introduction of fragments mimicking a partially denatured protein folding state into a targeted degrader, can induce the chaperone-mediated degradation by the proteasome.^[78] According to this strategy, Xie *et al.* proposed two series of selective androgen receptor degraders (SARDs) bearing a triazole-based linker.^[79] As the POI ligand, the AR agonist RU59063 was chosen,^[80] linked to an adamantyl moiety as a well-known hydrophobic tag (HyT39).^[81] These two fragments were connected *via* PEG, alkyl or aromatic

including four thalidomide-based and four VHL ligand-based

PROTACs, with yields ranging from 20 to 80%. PROTAC 26

turned out to be the most active of the series, allowing the simultaneous degradation of both EGFR and PARP targets in

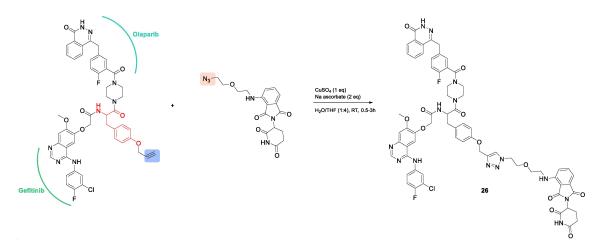


Figure 11. Synthesis of dual EGFR/PARP PROTAC 26 via CuAAC. The star-type core linker is highlighted in red.

ChemMedChem 2023, 18, e202300422 (8 of 17)



linkers of different lengths using CuAAC reaction (CuSO₄, sodium ascorbate, water/THF, RT, 0–2 h). Twenty-nine final compounds were obtained, with yields ranging from 46 to 86%. The most active compound **27** (Figure 12) displayed high AR binding affinity, good inhibitory activity against LNCaP prostate cancer cell line, and promising AR degradation effect.^[79] The positive degradation effect showed that triazole-based linkers can be used for the synthesis of different type of degraders.

2.5. Oligonucleotide-type click chemistry

Noteworthy, click chemistry is not only used in the synthesis of PROTACs featuring small molecules as POI ligands. It has also been successfully applied to assemble oligonucleotide, further confirming its versatility and wide scope. Naganuma et al. reported a series of oligonucleotide-type ER α degraders,^[82] synthesized via a CuAAC protocol. A decoy oligonucleotide able to bind to ERa receptor was first selected as POI ligand. Then, the alkyne-terminated decoy moiety (ER(dec), Figure 13) was connected via triazole linkers with different azide-terminated E3 ligase ligands: LCL161 - inhibitor of apoptosis protein ligand (IAP) (LCL-PEG3-N3), VH032 - ligand of von Hippel Lindau protein, and pomalidomide – CRBN ligand.^[83] The click reactions were performed in a mixture of common solvents (DMSO, methanol, water) with a catalyst in form of copper(I) salt (iodocopper(I) triethyl phosphite, Cul•P(OEt)₃) and triethylamine (TEA) addition at 40 °C. Implementing directly a copper(I) salts excludes the necessity of the use of reducing agents, but the copper(I) needs to be stabilized by bulky organic substituents, like triethyl phosphite.^[71] The addition of TEA increased the reaction rate.^[71] The most promising molecule in terms of highest ER α degradation effect was **LCL-ER(dec)** (28, Figure 13) consisting of the LCL161 moiety, a PEG3 linker and the ER α -binding decoy ER(dec).^[82]

Similarly, another class of ER α degraders,^[84] featuring the specific estrogen response element (ERE) directed to the DNAbinding motif of ER α was developed.^[85] The reaction between ERE and VHL-based E3 ligase was performed by CuAAC reaction and the obtained ERE-PROTAC displayed a satisfactory degradation effect in breast cancer cells.^[84] These two examples highlighted the versatility of click chemistry, expanding its use to the development of non-small molecule-based degraders.

2.6. DNA-encoded library screening

The use of click chemistry in the TPD field has been successfully employed in combination with emerging drug discovery technologies. Disch *et al.* proposed its use in the search of new bispecific ER α degraders,^[86] starting from DNA-encoded chemical library (DECL) screening against ER α .^[87] DECL is a technology that allows to synthetically generate millions of compounds in a rapid way and, to screen them against the selected biological target. In fact, the synthetized compounds are connected to DNA fragments which work like "bar codes", allowing a fast hit identification.^[87] In the PROTAC field, this is particularly

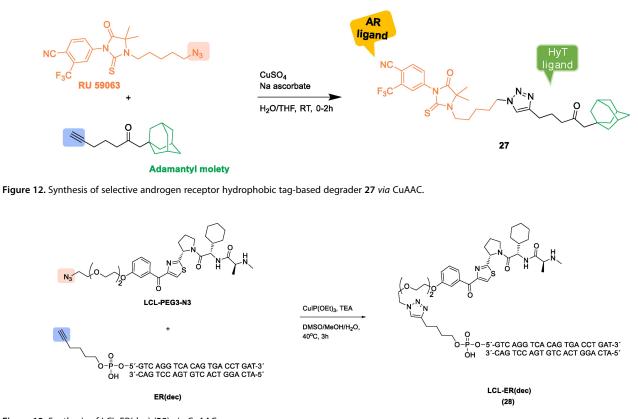


Figure 13. Synthesis of LCL-ER(dec) (28) via CuAAC.

© 2023 The Authors. ChemMedChem published by Wiley-VCH GmbH

ChemMedChem 2023, 18, e202300422 (9 of 17)

important, not only for the rapid identification of novel POI binders, but also because it allows to know the linker attachment point a priori, given the connected DNA fragment. Among the DECL, two promising molecules were selected as hits, leading to the identification of the $ER\alpha$ ligand, which was transformed into the corresponding azide derivative. Click chemistry synthesis between azide-derivatized $ER\alpha$ ligand and an alkyne-terminated E3 binder panel rapidly generated $ER\alpha$ degraders.^[86] The synthetic strategy involved the use of CuSO₄, sodium ascorbate, THPTA, water/DMSO, RT. To note, THPTA tris((1-hydroxy-propyl-1H-1,2,3-triazol-4-yl)methyl)amine, is a ligand exploited in CuAAC, which, similar to the previously mentioned TBTA, accelerates the cycloaddition by maintaining the Cu(I) oxidation state. The advantage of employing THPTA compared to TBTA is its water solubility.^[71] Thanks to the combination of DECL technology and click chemistry, Eratargeted PROTAC 29, with beneficial in vivo effects, was discovered (Figure 14).^[86] In addition, this work established that DECL information can be directly leveraged to efficiently produce PROTACs starting from an azide derivative of the POI ligand obtained by replacing the DNA tag, and then crossreacted with a series of E3 binder alkynes, utilizing click chemistry.

2.7. Solid-phase organic synthesis in combination with click chemistry

Another interesting idea proposed by Xu et al. is the combination of solid-phase organic synthesis (SPOS)[88] with click chemistry for the development of new PROTACs.^[89] Because of a difficult purification due to the highly polar character of the thalidomide scaffold, SPOS was proposed as a suitable solution.^[90] To perform CuAAC reaction, a supported azide reagent was prepared by linking an azide-terminated pomalidomide to the resin via an amine bond. In the next step, it was reacted with an alkyne-terminated POI ligand (TFC-007) under standard reaction conditions (Cul, DIPEA, DMF, RT), providing, after acidic resin cleavage, the triazole-based PROTAC 30 with 99% purity, but very poor yield (1%) (Figure 15). PROTAC 30 was found to have the highest degradation activity against hematopoietic prostaglandin D synthase (H-PGDS) with respect to the corresponding amide- and urea-based PROTACs. Although further optimization is required, once more, click chemistry combined with SPOS might facilitate the rapid development of PROTACs directed to different POIs.[89]

3. "Click" Reactions in cellulo

Despite of positive results of PROTACs in clinical trials, the development of degraders still shows several challenges. First at all, the physicochemical properties barely fulfill druglikeness

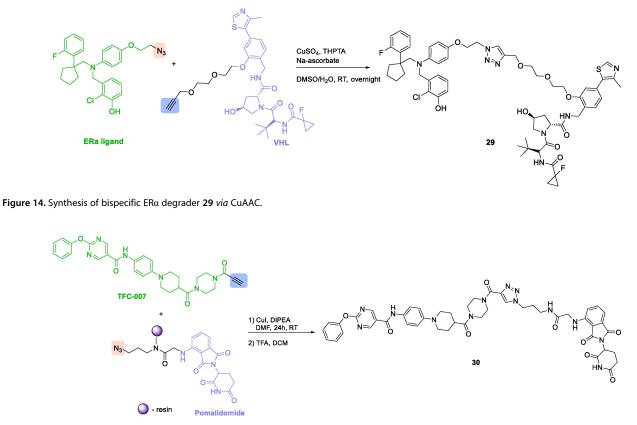


Figure 15. The synthesis of triazole-based PROTAC (30) via a solid-phase CuAAC

ChemMedChem 2023, 18, e202300422 (10 of 17)

8607187,20

5, 20, Downloaded from https://chemistry-europe.onlinelibary.wiley.com/doi/10.1002/cmdc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [0901/2024]. See the Terms and Conditions 0 https://onlinelibrary.wiley.com/temistry-europe.onlinelibary.or rules of use; OA articles are governed by the applicable Creative Commons.

rules, like the Lipinski Rule of Five (Ro5).^[91] Due to the high molecular weight and polar surface area (800–1000 Da and ~200 Å², respectively), PROTACs may encounter issues such as scarce cell permeability, low bioavailability, and poor water solubility.^[50,92,93] Therefore, strategies to overpass these problems and allowing a proper PROTAC delivery inside cells are being developed.^[51,54] In the following sections, we present selected biorthogonal chemistry examples of these novel approaches, where two click precursors undergo rapid and reversible covalent assembly *in situ*.

3.1. Inverse electron demand Diels-Alder (IEDDA) cycloaddition

One of the most promising approaches is *in-cell* click-formed proteolysis targeting chimeras (CLIPTACs). The idea of CLIPTACs was firstly proposed by Astex Pharmaceuticals^[94] and it was based on the inverse electron demand Diels-Alder (IEDDA) cycloaddition between tetrazine and TCO.^[27]

As the POI ligand, the well-known bromodomain-containing protein 4 (BRD4) ligand JQ1 was selected, [95] whereas thalidomide was used as the E3 ligase binder. Then, tetrazine-tagged thalidomide derivatives (Tz-thalidomide) and trans-cyclo-octene-tagged JQ1 (JQ1-TCO) (Figure 16) were designed and synthetized as smaller PROTAC precursors, able to cross cell membranes. Once inside the cells, these precursors promptly reacted and formed the desired JQ1-CLIPTAC (31, Figure 16). This approach was successfully validated in biological assays, where firstly the formation of 31 in solution was confirmed, and then the 31-induced degradation of BRD4 in cells was observed.^[94] The authors further corroborated this promising strategy by developing effective CLIPTACs directed to the extracellular-regulated protein kinase 1/2 (ERK1/2).^[94] It is worth noting that, in both cases, the preclicked CLIPTACs (prepared outside cells) did not show any cellular degradation effect, probably due to the lack of cell permeability.[94] This further confirmed that the observed degradation resulted from the self-assembled CLIPTAC following the precursors' entry into cells. In addition, the tagged E3 ligase precursors might be exploitable in click reactions with any suitably tagged POI ligand, making this approach largely appliable.

Indeed, the IEDDA cycloaddition approach has been used in the discovery of novel PROTACs targeting various kinases, including the platelet-derived growth factor receptor (PDGFR- β).^[96,97] Two POI ligands, i.e., linifanib,^[98] and S5^[99] were modified by introducing the norbornene group and in turn, E3 ligase binders by the tetrazine group (E3L-Tz). After the confirmation that the self-assembled PROTACs were formed inside the cells, PDGFR- β degradation was evaluated by Western blot. The observed effect was not fully satisfactory, probably due to low cell permeability of the E3L-Tz precursors.^[96] Collectively, in spite of significant advances, the development of cell-permeable PROTACs remains still a challenge.

3.2. Copper-free strain-promoted azide-alkyne cycloaddition (SPAAC)

Another bioorthogonal reaction, i.e., the strain-promoted azidealkyne cycloaddition (SPAAC)^[29] was exploited for the development of transcription factors (TF) PROTACs (TF-PROTACs).^[100] TF bind specific DNA fragments, and as a result, they control DNA transcription.^[101] TF are important biological targets involved in many diseases, especially cancers, but they are considered 'undruggable'.^[102] Therefore, PROTAC technology has successfully been combined with bioorthogonal chemistry, as a promising approach to address these undruggable targets. Liu *et al.* reported a click chemistry platform allowing the fast and easy synthesis of TF-PROTACs directed to the nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) *via* a copper-free SPAAC reaction.^[100] As E3 ligase binders, a library of eighteen bicyclooctyne (BCN)-terminated VHL ligands connected to various linkers was synthetized, exemplified by VHLL-

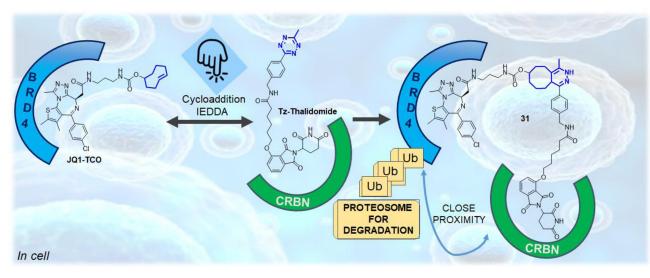


Figure 16. Synthesis of CLIPTAC 31 via IEDDA cycloaddition



BCN carrying a PEG linker (Figure 17). As the NF-κB binder, a commercially available azide-modified DNA oligomer (N_3 -NF-κB-ODN) was used. The copper-free SPAAC reactions between VHLL-BCN and N_3 -NF-κB-ODN were performed under physiological conditions (PBS, 37 °C), resulting in eighteen TF-PROTACs, named dNF-κBs, with 32 as the most promising one (Figure 17). Regarding the chemistry part, it is worth noting that, increasing the linker length resulted in lower reaction yields. As for the biological studies, the authors demonstrated that five out of eighteen dNF-κBs reduced p65 protein level. To validate the general utility of this methodology, another series of TF-PROTACs targeting E2F was successfully developed,^[100]

demonstrating that bioorthogonal chemistry reactions combined with the PROTAC technology hold great potential also for undruggable targets.

3.3. CuAAC catalyzed by intracellular copper

Si *et al.* proposed another interesting solution to increase the cell permeability and selectivity of anti-cancer PROTACs, based on a (non-bioorthogonal) CuAAC *in vivo* catalyzed by endoge-nous copper (Figure 18).^[103] Sorafenib^[104] – a multi-targeted

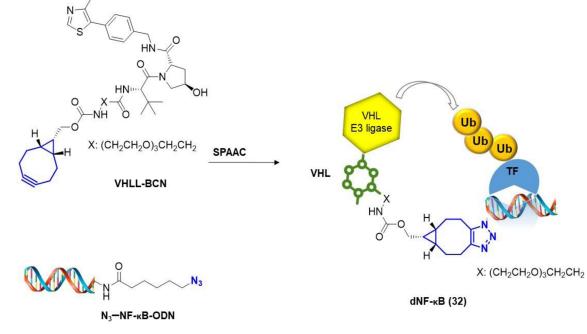


Figure 17. The synthesis of TF-PROTAC (32) via a SPAAC reaction.

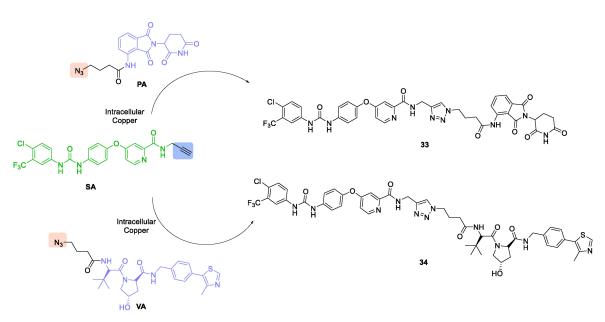


Figure 18. Synthesis anti-cancer PROTACs 33 and 34 via CuAAC reaction in vivo.

kinase inhibitor, was selected as the POI ligand and was converted into the alkyne terminated precursor (**SA**, Figure 18).

Azide-terminated VH032 and pomalidomide ligands (VA and PA, respectively in Figure 18) were synthesized as E3 ligase binders. The in situ self-assembly of the two clickable precursors into PROTACs 33 and 34 was catalyzed by the high level of endogenous copper in tumor cells.^[105] Once verified the formation of self-assembled PROTACs - predominately in tumor cells with respect of healthy ones, Western blot analysis showed that only cells treated with SA and VA remarkably degraded the vascular endothelial growth factor receptor 2 (VEGFR-2) and EPH receptor B4 (EphB4). The idea of generating tumor-specific self-assembled PROTACs by exploiting a click reaction between azides and alkynes together with a higher copper concentration in tumor, might be highly innovative. However, cell permeability and stability of click precursors need to be improved,^[103] especially in terms of alternative reactive groups, with more appropriate physicochemical properties.[54]

The bioorthogonal reactions are used not only to produce degraders directly in the cells, but also to enhance PROTACs' bioavailability in specific cells by the development of new delivery techniques, or to control the degradation process. In the following sections, we shortly describe other recent implementations of bioorthogonal reactions for the development of specific degraders.

3.4. Bioorthogonal polymeric PROTAC (POLY-PROTAC) nanoparticles

Gao *et al.* described a polymeric anti-BRD4 PROTAC nanoplatform to drive selective PROTAC accumulation in tumors.^[106] This strategy was based on the preparation of two different nanoparticles (NPs): azide-terminated NPs containing polymeric PROTAC (POLY-PROTAC) and DBCO-modified pretargeted NPs responsible for the tumor delivery. *In situ* bioorthogonal click reaction between these clickable groups enhanced intratumor accumulation and retention of azide-modified POLY-PROTAC NPs. Upon internalization into tumor cells, the POLY-PROTAC NPs released the PROTAC payload *via* a reductive/acidic cleavage. The released VHL-based PROTACs led to BRD4 degradation and anti-tumor effect in mice.^[106] This study might provide a versatile nanoplatform for enhancing tumor-specific PROTAC delivery.

3.5. Gold nanocluster PROTACs (GNCTACs)

Instead of NPs, Wang *et al.* proposed the use of gold nanoclusters (GNCs) to increase PROTAC bioavailability, especially for peptide-based PROTACs.^[107,108] A hybrid nanosized PROTAC (GNCTAC) was prepared by combining GNCs to a human epidermal growth factor receptor 2 (HER2) targeting peptide through gold–sulfur coordination. Then, GNCs were linked to a conjugation unit terminating with a DBCO moiety, which was combined to a CRBN ligand derivatized with an azide group *via* a SPAAC reaction. GNCTAC could efficiently accumulate at the tumor site and efficiently degrade HER2, as well as inhibit tumor growth *in vivo*.^[108] Collectively, GNCTACs could overcome the intrinsic limitations of peptide-based PROTACs (i.e., poor cell permeability and low stability), by efficiently delivering them into cells.

3.6. "Ligation to scavenging" approach

Jin *et al.* applied the biorthogonal reactions to the termination of the TPD process.^[109] This innovative strategy termed "ligation to scavenging", aimed to terminate the degradation thanks to the formation of the corresponding triazole-based inactive PROTACs and to eliminate rapidly them from living cells. The inactivation of PROTACs resulted from the IEDDA reaction between tetrazine-modified PROTACs (Tz-PROTACs) and TCO-modified polyamidoamine dendrimer^[110] (PAMAM-G5-TCO). This approach was successfully applied to terminate PARP degradation in living cells.^[109] Thus, this study provided a valuable approach to regulate PROTAC levels in living cells, which paves the way for a "controlled" TPD.

4. Sulfur(VI) Fluoride Exchange (SuFEx) Approach

4.1. SuFEx-enabled high-throughput medicinal chemistry

The easy and rapidity of SuFEx reaction was implemented in the development of a PROTAC targeting the transcriptional coactivator ENL,^[111] involved in the pathogenesis of acute myeloid leukemia (AML).^[112] Garnar-Wortzel et al. established an innovative high-throughput medicinal chemistry approach that leverages SuFEx reactivity to expedite chemical diversification of a starting ENL inhibitor hit.[111] Firstly, the high-throughput screening (HTS) of a big collection of small molecules led to the identification of an amido-imidazopyridine ENL inhibitor hit (HTS hit, Figure 19). Then, the high-throughput hit-to-lead process relied on parallel synthesis of hit analogs via SuFEx.^[113] The biocompatible character of SuFEx transformations allowed that the crude sulfonamide products could be tested directly in biological assays without further purification, thus expediting the medicinal chemistry process. Thanks to this approach, a potent and selective inhibitor SR-0813 was discovered (Figure 19). The last step was the synthesis of the first-in-class PROTAC targeting transcriptional coactivator ENL, SR-1114 (Figure 19), which consists of SR-0813 as the POI ligand, a PEG linker and a thalidomide moiety as the E3 ligase binder (Figure 19). The biological assays demonstrated its potential activity in AML and therefore, the success of the SuFEx-based approach to enable high-throughput medicinal chemistry, as well as its compatibility with cell-based drug discovery.^[111]

Review doi.org/10.1002/cmdc.202300422



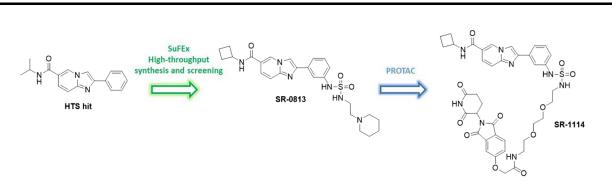


Figure 19. The identification of ENL PROTAC SR-1114 using SuFEx-enabled high-throughput synthesis and screening.

4.2. SuFEx platform for E3 ligase binders

Liu *et al.* applied the SuFEx chemistry for the easy preparation of azide-terminated E3 ligase binders to be then used in the synthesis of clickable PROTACs.^[114] In this case, the fluorosulfuryl azide (FSO₂N₃) was used as a diazoting reagent to provide rapid access to the azido derivatives, starting from easily accessible amine precursors. Thus, in a one-step reaction, a previously developed library of amine-based immunomodulatory drugs (IMiDs)^[115] was converted to azide-terminated molecules using a procedure reported by Dong and Sharpless (Figure 20A).^[116]

 $\rm FSO_2N_3$ was reacted in a mixture of solvents: methyl *tert*-butyl ether (MTBE), DMF and water with the addition of KHCO_3. After few minutes, nineteen derivatives of pomalidomide and lenalidomide with different alkyl or PEG linkers were obtained, with yields between 45 % and 80 %. ^[114]

To corroborate the versatility of the synthesized IMiD-based azide library, PROTACs targeting two different oncogenic proteins, i.e., BCR-ABL (bromodomain) and BET (extraterminal domain) were prepared. In both cases, previously reported POI ligands^[117,118] were transformed to alkynyl-tagged derivatives and then, PROTACs were synthetized *via* a CuAAC reaction (**35**

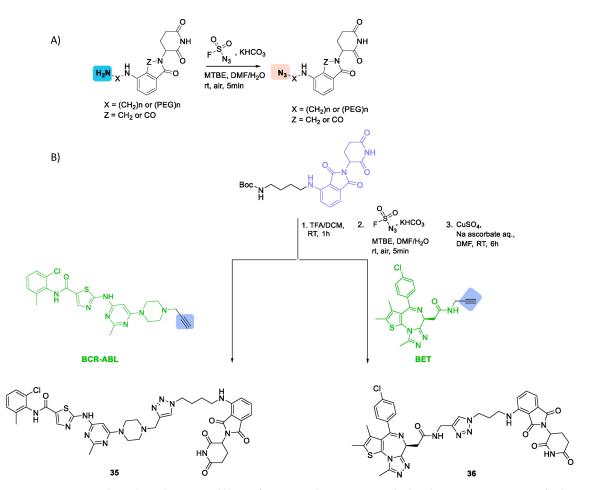


Figure 20. A) One-step reaction to obtain the azide-terminated library of IMiDs according to Dong's method; B) three-step one-pot process for the synthesis of BCR-ABL PROTAC 35 and BCR-ABL PROTAC 36.

8607187,20

5, 20, Downloaded from https://chemistry-europe.onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online Library on [09:01/2024]. See the Terms and Conditions (https://onlinelibrary.wikey.com/doi/10.1002/cndc.202300422 by Area Sistemi Dipart & Document, Wiley Online L

and 36, Figure 20B). To note, the authors optimized a successful three-step one-pot process, including BOC-deprotection, azide formation and click reaction,^[114] with 50-65% yields (Figure 20B). It is worth mentioning that the newly synthesized BCR-ABL and BET PROTACs displayed a stronger degradation effect than reference compounds.[117,118] Ultimately, the developed IMiD-based azide library may serve as a platform for speeding up the construction of large and diverse PROTAC libraries.

3. Summary and Outlook

TPD is a rapidly emerging modality in medicinal chemistry and drug discovery. A degrader is the prototypical example of a heterobifunctional molecule, constituted by a ligand for the target of interest bridged to a scaffold recruiting the degradation system by an appropriate linker. Therefore, medicinal chemists have struggled to rapidly and effectively assemble the different PROTAC elements in a single molecule able to induce the formation of a productive ternary complex. As a result, the Nobel Prize-winning technology of click/bioorthogonal chemistry may be a valuable option to the multi-step synthesis and difficult purification of degraders. The selectivity, ease, rapidity, and modularity of click transformations may speed up degrader synthesis and allow rapid SAR exploration in terms of ligand assembly, attachment points, and linker length/composition.

However, an effective degrader molecule must also possess optimal drug-like properties. So far, it is generally recognized that a limitation of degraders towards clinical translation is their poor drug-like profile, mainly due to the low cell permeability and water solubility. To obtain the right balance between potency and drug-like properties, the exploitation of 1,2,3triazole as an optimal linker in PROTAC development, has attracted significant interest. Its introduction could greatly influence the whole physicochemical properties and rigidity of the degrader. Hence, it may be a suitable strategy to improve cell permeability, solubility, or stability. Here, we provided an overview of the combination of TPD technology with click and bioorthogonal chemistry. We discussed selected examples of click chemistry transformations for the development of both small molecule- and oligonucleotide-based PROTACs. This strategy has been also extended to solid-phase synthesis and combinatorial click chemistry. The possibility to apply these reactions directly in cellulo expanded even more the scope and advantages of click chemistry, especially in terms of efficient delivery. Therefore, an avenue that could be pursued to quickly generate PROTACs endowed with optimal drug-like properties is offered by the cheap and easy click/bioorthogonal chemistry.

Clearly, it is not a panacea. A first issue that has not been enough explored so far is the metabolism of these clickgenerated degraders. It is well-known that triazole-containing drugs, like fungicides, undergo to a rapid cytochrome P450mediated metabolism. This would affect bioavailability and clinical efficacy. In addition, as CYP substrates and inhibitors, triazoles are reported to interact with many drugs, which may lead to toxicity in case of polypharmacy or ineffective treatment.

Furthermore, from a patent perspective, as for September 2023, not so many triazole-based degraders have been patented. A SciFinder site search for patents containing the keywords "click chemistry" and "PROTAC" retrieved just 6 entries.

All in all, far from being an easy task, we are confident that TPD modality together with click/bioorthogonal chemistry may be a winning combination toward clinical translation.

Acknowledgements

This research was supported by MIUR-FISR2019_03796 "Proteolysis targeting chimeras (PROTACs) to treat leishmaniasis"-PROLEISH and by EU funding within the NextGenerationEU-MUR PNRR Extended Partnership initiative on Emerging infectious Diseases (Project no. PE00000007, INF-ACT).

Conflict of Interests

The authors declare no conflict of interest.

Keywords: click chemistry · bioorthogonal chemistry PROTACs · Targeted Protein Degradation · triazoles

- [1] V. V. Rostovtsev, L. G. Green, V. V. Fokin, K. B. Sharpless, Angew. Chem. Int. Ed. 2002, 41, 2596-2599.
- C. W. Tornøe, C. Christensen, M. Meldal, J. Org. Chem. 2002, 67, 3057-[2] 3064.
- [3] C. Bertozzi, ACS Cent. Sci. 2022, 2-3.
- [4] E. Saxon, C. R. Bertozzi, Science 2000, 287, 2007–2010. [5] J. Hou, X. Liu, J. Shen, G. Zhao, P. G. Wang, Expert Opin. Drug Discovery
- 2012, 7, 489-501.
- [6] H. C. Kolb, K. B. Sharpless, Drug Discovery Today 2003, 8, 401-422.
- [7] X. Jiang, X. Hao, L. Jing, G. Wu, D. Kang, X. Liu, P. Zhan, Expert Opin. Drug Discovery 2019, 14, 779-789.
- [8] H. C. Kolb, M. G. Finn, K. B. Sharpless, Angew. Chem. Int. Ed. 2001, 40, 2004-2021.
- [9] M. Meldal, C. W. Tomøe, Chem. Rev. 2008, 108, 2952-3015.
- [10] C. E. Hoyle, C. N. Bowman, Angew. Chem. Int. Ed. Engl. 2010, 49, 1540-1573.
- Y. Méndez, A. V. Vasco, A. R. Humpierre, B. Westermann, ACS Omega [11] 2020, 5, 25505-25510.
- [12] J. Tu, D. Svatunek, S. Parvez, A. C. Liu, B. J. Levandowski, H. J. Eckvahl, R. T. Peterson, K. N. Houk, R. M. Franzini, Angew. Chem. Int. Ed. Engl. 2019, 58, 9043-9048.
- [13] J. Dong, L. Krasnova, M. G. Finn, K. Barry Sharpless, Angew. Chem. Int. Ed. 2014, 53, 9430-9448.
- [14] D. Zeng, W. P. Deng, X. Jiang, Chem. A Eur. J. 2023, 29, e2023005.
- [15] T. Guo, G. Meng, X. Zhan, Q. Yang, T. Ma, L. Xu, K. B. Sharpless, J. Dong, Angew. Chemie-Int. Ed. 2018, 57, 2605-2610.
- [16] H.-L. Qin, Q. Zheng, G. A. L. Bare, P. Wu, K. B. Sharpless, Angew. Chem. Int. Ed. Engl. 2016, 55, 14155-14158.
- [17] Z. Liu, J. Li, S. Li, G. Li, K. B. Sharpless, P. Wu, J. Am. Chem. Soc. 2018, 140, 2919-2925.
- [18] C. Zhao, K. P. Rakesh, L. Ravidar, W.-Y. Fang, H.-L. Qin, Eur. J. Med. Chem. 2019, 162, 679-734.
- [19] A. S. Barrow, C. J. Smedley, Q. Zheng, S. Li, J. Dong, J. E. Moses, Chem. Soc. Rev. 2019, 48, 4731-4758.
- [20] S. Wilson Lucas, R. Zijian Qin, K. P. Rakesh, K. S. Sharath Kumar, H.-L. Qin, Bioorg. Chem. 2023, 130, 106227.
- [21] E. M. Sletten, C. R. Bertozzi, Acc. Chem. Res. 2011, 44, 666-676.



8607187,20

3, 20, Downloaded

from https://chemistry-europe.onlinelibrary.wiley.com/doi/10.1002/cmdc.202300422 by Area Sistemi

Dipart & Document, Wiley Online Library on [09:01/20:24]. See the Terms and Conditions (https://onlinelibrary.wiley.com/terms-and-conditions) on Wiley Online Library for rules of use; OA articles are governed by the applicable Creative Commons

- [22] J. C. Jewett, C. R. Bertozzi, Chem. Soc. Rev. 2010, 39, 1272-1279.
- [23] K. L. Kiick, E. Saxon, D. A. Tirelli, C. R. Bertozzi, PNAS 2002, 99, 19-24. [24] V. O. Rodionov, V. V. Fokin, M. G. Finn, Angew. Chem. Int. Ed. 2005, 44,
- 2210-2215 [25] B. L. Oliveira, Z. Guo, G. J. L. Bernardes, Chem. Soc. Rev. 2017, 46, 4895-4950.
- [26] J.-B. Béquignat, N. Ty, A. Rondon, L. Taiariol, F. Degoul, D. Canitrot, M. Quintana, I. Navarro-Teulon, E. Miot-Noirault, C. Boucheix, J.-M. Chezal, E. Moreau, Eur. J. Med. Chem. 2020, 203, 112574.
- [27] M. L. Blackman, M. Royzen, J. M. Fox, J. Am. Chem. Soc. 2008, 130, 13518-13519.
- [28] C. Bednarek, I. Wehl, N. Jung, U. Schepers, S. Bräse, Chem. Rev. 2020, 120, 4301-4354.
- [29] N. J. Agard, J. A. Prescher, C. R. Bertozzi, J. Am. Chem. Soc. 2004, 126, 15046-15047.
- [30] M. Handula, K.-T. Chen, Y. Seimbille, Molecules 2021, 26, 10.3390/ molecules26154640.
- [31] K. Bozorov, J. Zhao, H. A. Aisa, Bioorg. Med. Chem. 2019, 27, 3511-3531. [32] N. K. Devaraj, ACS Cent. Sci. 2018, 4, 952–959.
- [33] The Nobel Committee for Chemistry, K. Vetenkaps-Akademien 2022, 50005.1-18.
- [34] T. W. Corson, N. Aberle, C. M. Crews, ACS Chem. Biol. 2008, 3, 677–692. [35] A. C. Lai, C. M. Crews, N. Haven, Nat. Rev. Drug Discov. (84.69; Q1) 2017,
- 16, 101–114.
- [36] X. Sun, H. Gao, Y. Yang, M. He, Y. Wu, Y. Song, Y. Tong, Y. Rao, Signal Transduct. Target. Ther. 2019, 4, 10.1038/s41392-019-0101-6.
- [37] M. He, C. Cao, Z. Ni, Y. Liu, P. Song, S. Hao, Y. He, X. Sun, Y. Rao, Signal Transduct. Target. Ther. 2022, 7, 10.1038/s41392-022-00999-9.
- [38] G. Ahn, S. M. Banik, C. L. Miller, N. M. Riley, J. R. Cochran, C. R. Bertozzi, Nat. Chem. Biol. 2021, 17, 937-946.
- [39] J. Pei, X. Pan, A. Wang, W. Shuai, F. Bu, P. Tang, S. Zhang, Y. Zhang, G. Wang, L. Ouyang, Chem. Commun. 2021, 57, 13194-13197.
- [40] D. Takahashi, J. Moriyama, T. Nakamura, E. Miki, E. Takahashi, A. Sato, T. Akaike, K. Itto-Nakama, H. Arimoto, Mol. Cell 2019, 76, 797–810.e10.
- [41] S. B. Alabi, C. M. Crews, J. Biol. Chem. 2021, 296, 100647.
- [42] L. Hua, Q. Zhang, X. Zhu, R. Wang, Q. You, L. Wang, J. Med. Chem. 2022, 65, 8091-8112.
- [43] A. Salerno, F. Seghetti, J. Caciolla, E. Uliassi, E. Testi, M. Guardigni, M. Roberti, A. Milelli, M. L. Bolognesi, J. Med. Chem. 2022, 65, 9507-9530.
- [44] R. M. Espinoza-Chávez, A. Salerno, A. Liuzzi, A. Ilari, A. Milelli, E. Uliassi, M. L. Bolognesi, ACS bio med chem Au 2023, 3, 32-45.
- [45] M. Teng, N. S. Gray, Cell Chem. Biol. 2023, 30, https://doi.org/10.1016/j. chembiol.2023.06.020.
- [46] X. Liu, A. Ciulli, ACS Cent. Sci. 2023, 9, 1269-1284.
- [47] C. Steinebach, I. Sosič, S. Lindner, A. Bricelj, F. Kohl, Y. L. D. Ng, M. Monschke, K.G. Wagner, J. Krönke, M. Gütschow, MedChemComm **2019**, *10*, 1037–1041.
- [48] M. S. Gadd, A. Testa, X. Lucas, K.-H. Chan, W. Chen, D. J. Lamont, M. Zengerle, A. Ciulli, Nat. Chem. Biol. 2017, 13, 514-521.
- [49] M. J. Roy, S. Winkler, S. J. Hughes, C. Whitworth, M. Galant, W. Farnaby, K. Rumpel, A. Ciulli, ACS Chem. Biol. 2019, 14, 361-368.
- [50] H. Lebraud, T. D. Heightman, Essays Biochem. 2017, 61, 517-527.
- [51] V.G. Klein, C.E. Townsend, A. Testa, M. Zengerle, C. Maniaci, S.J. Hughes, K.-H. Chan, A. Ciulli, R. S. Lokey, ACS Med. Chem. Lett. 2020, 11, 1732-1738
- [52] A. Zagidullin, V. Milyukov, A. Rizvanov, E. Bulatov, Explor. Target. Antitumor Ther. 2020, 1, 381-390.
- [53] N. Bai, S. A. Miller, G. V. Andrianov, M. Yates, P. Kirubakaran, J. Karanicolas, J. Chem. Inf. Model. 2021, 61, 1368-1382.
- [54] S. Tomoshige, M. Ishikawa, Bioorg. Med. Chem. 2021, 41, 116221.
- [55] C. M. Crews, J. Med. Chem. 2018, 61, 403-404.
- [56] M. Schiedel, D. Herp, S. Hammelmann, S. Swyter, A. Lehotzky, D. Robaa, J. Oláh, J. Ovádi, W. Sippl, M. Jung, J. Med. Chem. 2018, 61, 482-491.
- [57] Q.-J. Wu, T.-N. Zhang, H.-H. Chen, X.-F. Yu, J.-L. Lv, Y.-Y. Liu, Y.-S. Liu, G. Zheng, J.-Q. Zhao, Y.-F. Wei, J.-Y. Guo, F.-H. Liu, Q. Chang, Y.-X. Zhang, C.-G. Liu, Y.-H. Zhao, Signal Transduct. Target. Ther. 2022, 7, 402.
- [58] T. Rumpf, M. Schiedel, B. Karaman, C. Roessler, B. J. North, A. Lehotzky, J. Oláh, K. I. Ladwein, K. Schmidtkunz, M. Gajer, M. Pannek, C Steegborn, D. A. Sinclair, S. Gerhardt, J. Ovádi, M. Schutkowski, W. Sippl, O. Einsle, M. Jung, Nat. Commun. 2015, 6, 6263.
- [59] R. P. Wurz, K. Dellamaggiore, H. Dou, N. Javier, M. C. Lo, J. D. McCarter, D. Mohl, C. Sastri, J. R. Lipford, V. J. Cee, J. Med. Chem. 2018, 61, 453-461.
- [60] P. Filippakopoulos, J. Qi, S. Picaud, Y. Shen, W. B. Smith, O. Fedorov, E. M. Morse, T. Keates, T. T. Hickman, I. Felletar, M. Philpott, S. Munro,

ChemMedChem 2023, 18, e202300422 (16 of 17)

M. R. McKeown, Y. Wang, A. L. Christie, N. West, M. J. Cameron, B. Schwartz, T. D. Heightman, N. La Thangue, C. A. French, O. Wiest, A. L. Kung, S. Knapp, J. E. Bradner, Nature 2010, 468, 1067-1073.

- [61] L. Wang, X. Shao, T. Zhong, Y. Wu, A. Xu, X. Sun, H. Gao, Y. Liu, T. Lan, Y. Tong, X. Tao, W. Du, W. Wang, Y. Chen, T. Li, X. Meng, H. Deng, B. Yang, Q. He, M. Ying, Y. Rao, Nat. Chem. Biol. 2021, 17, 567-575.
- [62] H. Döhner, D. J. Weisdorf, C. D. Bloomfield, N. Engl. J. Med. 2015, 373, 1136-1152.
- [63] O. R. Ayinde, C. Sharpe, E. Stahl, R. J. T. li, J. R. Lerma, N. Muthusamy, J. C. Byrd, J. R. Fuchs, ACS Med. Chem. Lett. 2023, DOI 10.1021/ acsmedchemlett.3c00082.
- [64] M. S. Squires, R. E. Feltell, N. G. Wallis, E. J. Lewis, D.-M. Smith, D. M. Cross, J. F. Lyons, N. T. Thompson, Mol. Cancer Ther. 2009, 8, 324–332.
- [65] L. Qu, S. Li, L. Ji, S. Luo, M. Ding, F. Yin, C. Wang, H. Luo, D. Lu, X. Liu, W. Peng, L. Kong, X. Wang, Eur. J. Med. Chem. 2021, 226, 10.1016/ j.ejmech.2021.113889.
- X. Jiang, J. Zhou, Y. Wang, X. Liu, K. Xu, J. Xu, F. Feng, H. Sun, Eur. J. [66] Med. Chem. 2021, 210, 112949.
- [67] P. Sivaprakasam, X. Han, R. L. Civiello, S. Jacutin-Porte, K. Kish, M. Pokross, H. A. Lewis, N. Ahmed, N. Szapiel, J. A. Newitt, E. T. Baldwin, H. Xiao, C. M. Krause, H. Park, M. Nophsker, J. S. Lippy, C. R. Burton, D. R. Langley, J. E. Macor, G. M. Dubowchik, Bioorg. Med. Chem. Lett. 2015, 25, 1856-1863.
- [68] K. Yang, Y. Song, H. Xie, H. Wu, Y. T. Wu, E. D. Leisten, W. Tang, Bioorg. Med. Chem. Lett. 2018, 28, 2493-2497.
- [69] C. Boyault, K. Sadoul, M. Pabion, S. Khochbin, Oncogene 2007, 26, 5468-5476.
- [70] T. Hideshima, J. Qi, R. M. Paranal, W. Tang, E. Greenberg, N. West, M. E. Colling, G. Estiu, R. Mazitschek, J. A. Perry, H. Ohguchi, F. Cottini, N. Mimura, G. Görgün, Y.-T. Tai, P. G. Richardson, R. D. Carrasco, O. Wiest, S. L. Schreiber, K. C. Anderson, J. E. Bradner, Proc. Natl. Acad. Sci. USA **2016**, *113*, 13162–13167.
- [71] S. Neumann, M. Biewend, S. Rana, W. H. Binder, Macromol, Rapid Commun. 2020, 41, 10.1002/marc.201900359.
- [72] H. Wu, K. Yang, Z. Zhang, E. D. Leisten, Z. Li, H. Xie, J. Liu, K. A. Smith, Z. Novakova, C. Barinka, W. Tang, J. Med. Chem. 2019, 62, 7042-7057.
- [73] E. Munoz, G. Chen, A. Hossain, S. Wu, E. Ocequera Nava, J. Hang, T. Lee, Q. Zhang, G. Wang, Q. H. Chen, Bioorg. Med. Chem. Lett. 2022, 72, 128870.
- [74] G. Loren, I. Espuny, A. Llorente, C. Donoghue, X. Verdaguer, R. R. Gomis, A. Riera, Eur. J. Med. Chem. 2022, 243, 114770.
- Y. Chen, X. Yuan, M. Tang, M. Shi, T. Yang, K. Liu, D. Deng, L. Chen, [75] Bioorg. Chem. 2022, 119, 105508.
- [76] S. Su, Z. Yang, H. Gao, H. Yang, S. Zhu, Z. An, J. Wang, Q. Li, S. Chandarlapaty, H. Deng, W. Wu, Y. Rao, J. Med. Chem. 2019, 62, 7575-7582.
- [77] M. Zheng, J. Huo, X. Gu, Y. Wang, C. Wu, Q. Zhang, W. Wang, Y. Liu, Y. Liu, X. Zhou, L. Chen, Y. Zhou, H. Li, J. Med. Chem. 2021, 64, 7839-7852.
- [78] T. K. Neklesa, H. S. Tae, A. R. Schneekloth, M. J. Stulberg, T. W. Corson, T. B. Sundberg, K. Raina, S. A. Holley, C. M. Crews, Nat. Chem. Biol. 2011, 7, 538-543.
- [79] H. Xie, J. J. Liang, Y. L. Wang, T. X. Hu, J. Y. Wang, R. H. Yang, J. K. Yan, Q. R. Zhang, X. Xu, H. M. Liu, Y. Ke, Eur. J. Med. Chem. 2020, 204, 10.1016/i.eimech.2020.112512.
- [80] M. E. Van Dort, D. M. Robins, B. Wayburn, J. Med. Chem. 2000, 43, 3344-3347.
- [81] H. S. Tae, T. B. Sundberg, T. K. Neklesa, D. J. Noblin, J. L. Gustafson, A. G. Roth, K. Raina, C. M. Crews, ChemBioChem 2012, 13, 538-541.
- [82] M. Naganuma, N. Ohoka, G. Tsuji, H. Tsujimura, K. Matsuno, T. Inoue,
- M. Naito, Y. Demizu, ACS Med. Chem. Lett. 2022, 13, 134-139. [83] G. M. Burslem, C. M. Crews, Chem. Rev. 2017, 117, 11269-11301.
- [84] X. Zhang, Z. Zhang, X. Xue, T. Fan, C. Tan, F. Liu, Y. Tan, Y. Jiang, ACS
- Pharmacol. Transl. Sci. 2022, 5, 1109-1118.
- [85] C. M. Klinge, Nucleic Acids Res. 2001, 29, 2905-2919.
- [86] J. S. Disch, J. M. Duffy, E. C. Y. Lee, D. Gikunju, B. Chan, B. Levin, M. I. Monteiro, S. A. Talcott, A. C. Lau, F. Zhou, A. Kozhushnyan, N. E. Westlund, P. B. Mullins, Y. Yu, M. Von Rechenberg, J. Zhang, Y. A. Arnautova, Y. Liu, Y. Zhang, A. J. McRiner, A. D. Keefe, A. Kohlmann, M. A. Clark, J. W. Cuozzo, C. Huguet, S. Arora, J. Med. Chem. 2021, 64, 5049-5066
- M. A. Clark, R. A. Acharya, C. C. Arico-Muendel, S. L. Belyanskaya, D. R. [87] Benjamin, N. R. Carlson, P. A. Centrella, C. H. Chiu, S. P. Creaser, J. W. Cuozzo, C. P. Davie, Y. Ding, G. J. Franklin, K. D. Franzen, M. L. Gefter, S. P. Hale, N. J. V. Hansen, D. I. Israel, J. Jiang, M. J. Kavarana, M. S. Kelley, C. S. Kollmann, F. Li, K. Lind, S. Mataruse, P. F. Medeiros, J. A.

Messer, P. Myers, H. O'Keefe, M. C. Oliff, C. E. Rise, A. L. Satz, S. R. Skinner, J. L. Svendsen, L. Tang, K. van Vloten, R. W. Wagner, G. Yao, B. Zhao, B. A. Morgan, *Nat. Chem. Biol.* **2009**, *5*, 647–654.

- [88] N. Cankařová, E. Schütznerová, V. Krchňák, Chem. Rev. 2019, 119, 12089–12207.
- [89] H. Xu, T. Kurohara, R. Takano, H. Yokoo, N. Shibata, N. Ohoka, T. Inoue, M. Naito, Y. Demizu, *ChemistryOpen* **2022**, *11*, 1–6.
- [90] S. Krajcovicova, R. Jorda, D. Hendrychova, V. Krystof, M. Soural, Chem. Commun. 2019, 55, 929–932.
- [91] C. A. Lipinski, F. Lombardo, B. W. Dominy, P. J. Feeney, Adv. Drug Delivery Rev. 2001, 46, 3–26.
- [92] P. Matsson, J. Kihlberg, J. Med. Chem. 2017, 60, 1662-1664.
- [93] S. D. Edmondson, B. Yang, C. Fallan, Bioorg. Med. Chem. Lett. 2019, 29, 1555–1564.
- [94] H. Lebraud, D. J. Wright, C. N. Johnson, T. D. Heightman, ACS Cent. Sci. 2016, 2, 927–934.
- [95] M. Zengerle, K.-H. Chan, A. Ciulli, ACS Chem. Biol. 2015, 10, 1770–1777.
- [96] R. Si, H. Zhu, J. Wang, Q. Zhang, Y. Li, X. Pan, J. Zhang, *Bioorg. Chem.* 2023, 134, 10.1016/j.bioorg.2023.106439.
- [97] R. Si, H. Zhu, J. Wang, Q. Zhang, Y. Li, X. Pan, J. Zhang, *Bioorg. Chem.* 2023, 135, 10.1016/j.bioorg.2023.106497.
- [98] J. Chen, J. Guo, Z. Chen, J. Wang, M. Liu, X. Pang, Sci. Rep. 2016, 6, 29382.
- [99] Y. Shan, H. Gao, X. Shao, J. Wang, X. Pan, J. Zhang, Eur. J. Med. Chem. 2015, 103, 80–90.
- [100] J. Liu, H. Chen, H. Ü. Kaniskan, L. Xie, X. Chen, J. Jin, W. Wei, J. Am. Chem. Soc. 2021, 143, 8902–8910.
- [101] S. A. Lambert, A. Jolma, L. F. Campitelli, P. K. Das, Y. Yin, M. Albu, X. Chen, J. Taipale, T. R. Hughes, M. T. Weirauch, *Cell* **2018**, *172*, 650–665.
 [102] J. H. Burburghter, *Nat. Phys. Comm.* **2020**, *10*, 611 (2018).
- [102] J. H. Bushweller, *Nat. Rev. Cancer* 2019, *19*, 611–624.
 [103] R. Si, P. Hai, Y. Zheng, J. Wang, Q. Zhang, Y. Li, X. Pan, J. Zhang, *Eur. J. Med. Chem.* 2023, *257*, 115497.
- [104] S. M. Wilhelm, C. Carter, L. Y. Tang, D. Wilkie, A. McNabola, H. Rong, C. Chen, X. Zhang, P. Vincent, M. McHugh, Y. Cao, J. Shujath, S. Gawlak, D. Eveleigh, B. Rowley, L. Liu, L. Adnane, M. Lynch, D. Auclair, I. Taylor, R. Gedrich, A. Voznesensky, B. Riedl, L. E. Post, G. Bollag, P. A. Trail, *Cancer Res.* 2004, 64, 7099–7109.

- [105] T. Kamiya, J. Clin. Biochem. Nutr. 2022, 71, 22–28.
- [107] Z. Luo, X. Yuan, Y. Yu, Q. Zhang, D. T. Leong, J. Y. Lee, J. Xie, J. Am. Chem. Soc. 2012, 134, 16662–16670.
 [108] Z. Wang, M. Tan, W. Su, W. Hunger, J. Zhu, S. F. K. C. C. M. Kin, W. Kuman, J. Zhu, S. K. Kuman, J. Zhu, S. K. Kuman, J. Zhu, S. Kuman, J. Zhu, J. Zhu, S. Kuman, J. Zhu, S. Kuman, J. Zhu, J. Zhu, J. Zhu, S. Kuman, J. Zhu, S. Kuman, J. Zhu, S. Kuman, J. Zhu, S. Kuman, J. Zhu, J. Zhu, S. Kuman, J. Z
- [108] Z. Wang, M. Tan, W. Su, W. Huang, J. Zhang, F. Jia, G. Cao, X. Liu, H. Song, H. Ran, G. Nie, H. Wang, J. Med. Chem. 2023, 66(9), 6263–6273.
 [100] V. Jia, J. San, P. Wang, Y. Wang, J. Hui, G. Yu, J. Zhang, G. Cao, X. Liu, H. Song, H. Rang, C. Standard, Standard, C. Standard, Standard, C. Standard, C. Standard,
- [109] Y. Jin, J. Fan, R. Wang, X. Wang, N. Li, Q. You, Z. Jiang, J. Am. Chem. Soc. 2022, 145(13), 7218–7229.
 [110] H. Kharaldina, O. Bachid, A. M. Ushib, A. F. Al Maustafa, J. F. Burti, C. S. Karaldina, A. M. Ushib, A. F. Al Maustafa, J. F. Burti, C. S. Karaldina, A. M. Ushib, A. F. Al Maustafa, J. F. Burti, C. S. Karaldina, A. M. Ushib, A. F. Al Maustafa, J. F. Burti, C. S. Karaldina, A. M. Ushib, A. F. Al Maustafa, J. F. Burti, C. S. Karaldina, S. Karaldina, A. Karaldina, A. M. Ushib, A. F. Al Maustafa, J. F. Burti, C. S. Karaldina, A. M. Ushib, A. F. Al Maustafa, J. F. Burti, C. S. Karaldina, A. Karaldina, A.
- [110] H. Kheraldine, O. Rachid, A. M. Habib, A.-E. Al Moustafa, I. F. Benter, S. Akhtar, Adv. Drug Delivery Rev. 2021, 178, 113908.
- [111] L. Garnar-Wortzel, T. R. Bishop, S. Kitamura, N. Milosevich, J. N. Asiaban, X. Zhang, Q. Zheng, E. Chen, A. R. Ramos, C. J. Ackerman, E. N. Hampton, A. K. Chatterjee, T. S. Young, M. V. Hull, K. B. Sharpless, B. F. Cravatt, D. W. Wolan, M. A. Erb, ACS Cent. Sci. 2021, 7, 815–830.
- [112] M. A. Erb, T. G. Scott, B. E. Li, H. Xie, J. Paulk, H.-S. Seo, A. Souza, J. M. Roberts, S. Dastjerdi, D. L. Buckley, N. E. Sanjana, O. Shalem, B. Nabet, R. Zeid, N. K. Offei-Addo, S. Dhe-Paganon, F. Zhang, S. H. Orkin, G. E. Winter, J. E. Bradner, *Nature* 2017, *543*, 270–274.
- [113] S. Kitamura, Q. Zheng, J. L. Woehl, A. Solania, E. Chen, N. Dillon, M. V. Hull, M. Kotaniguchi, J. R. Cappiello, S. Kitamura, V. Nizet, K. B. Sharpless, D. W. Wolan, J. Am. Chem. Soc. 2020, 142, 10899–10904.
- [114] H. Liu, R. Sun, C. Ren, X. Qiu, X. Yang, B. Jiang, Org. Biomol. Chem. 2021, 19, 166–170.
- [115] X. Qiu, N. Sun, Y. Kong, Y. Li, X. Yang, B. Jiang, Org. Lett. 2019, 21, 3838–3841.
- [116] G. Meng, T. Guo, T. Ma, J. Zhang, Y. Shen, K. B. Sharpless, J. Dong, *Nature* **2019**, *574*, 86–89.
- [117] G. E. Winter, D. L. Buckley, J. Paulk, J. M. Roberts, A. Souza, S. Dhe-Paganon, J. E. Bradner, *Science* 2015, 348, 1376–1381.
- [118] A. C. Lai, M. Toure, D. Hellerschmied, J. Salami, S. Jaime-Figueroa, E. Ko, J. Hines, C. M. Crews, Angew. Chem. Int. Ed. Engl. 2016, 55, 807–810.

Revised manuscript received: September 1, 2023 Version of record online: September 14, 2023

