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A Versatile and Sustainable Multicomponent Platform for the Synthesis of Protein Degraders: Proof-of-Concept Application to BRD4-Degrading PROTACs

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resulting in the discovery of a set of degraders endowed with high degradation efficiency. Compared to the existing methods, our approach offers a versatile and cost-effective means to access libraries of protein degraders and increase the chance of identifying successful candidates.

INTRODUCTION

Protein degraders represent an attractive tool to control levels of disease-related proteins and are set to revolutionize drug discovery.^{1,2} Among them, PROTACs (Proteolysis TArgeting Chimeras) have drawn attention as one of the most appealing approaches.³ Being composed of a ligand for the protein of interest (POI) (referred to as warhead) and a recruiting moiety for the E3 ubiquitin ligase (referred to as anchor) connected through a suitable linker, PROTACs are able to hijack the ubiquitin-proteasome cascade and force the degradation of the POI (Figure 1). They effectively act as catalysts, being available for successive rounds of degradation upon completion of their cycle, thus minimizing the need for a continuous drug administration.⁴ Around 100 different proteins have been targeted so far by applying the PROTAC technology, which also proved to be an effective modality for accessing the "undruggable genome".⁵ However, a grand total of more than 1000 proteins might be considered as PROTACtable, offering opportunities for future efforts based on protein degraders." Compared to the occupancy-driven approaches, PROTACs utilize an event-driven mechanism of action, which allows the knock down of disease-related POIs, offering an extraordinary strategy to overcome issues often experienced in classic drug discovery approaches, including resistance mechanisms (e.g., target protein overexpression and resistance mutations).^{5,}

highly decorated PROTACs. As a proof of principle, our platform has been applied to the preparation of potential BRD4-degrading PROTACs,

The great potential of the approach, which has grown exponentially in the past few years, has been confirmed by the first degraders that have reached clinical trials, including Arvinas degraders ARV-110, ARV-470, and ARV-766,^{10,11} and DT2216 developed by Dialectic Therapeutic, whose structures have been recently disclosed (Figure 2).^{8,10}

While the POIs targeted by PROTACs currently under clinical investigation show a high degree of heterogeneity, bromodomain and extraterminal domain (BET) proteins have emerged as a model POI in many pilot studies probing PROTACs. BET proteins include BRD2, BRD3, BRD4, and BRDT and recruit transcriptional complexes by binding to acetylated lysine residues on histones, thereby controlling genes involved in cellular proliferation and cell cycle progression. Because alterations in the regulation of activities by BET proteins, especially BRD4,¹² are associated with different inflammatory diseases and cancer, several BET degraders, including dBET1,¹³ MZ1,¹⁴ dBET21,^{15,16} and ARV-771¹⁷

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Figure 1. Structural components of PROTACs. Schematic illustration of their two-component reaction (2-CR)-based synthesis and of our multicomponent reaction (MCR)-based platform.



Figure 2. Disclosed structures of PROTACs advanced into clinical investigations. And rogen receptor (AR); estrogen receptor (ER); B-cell lymphomaextra large (BCL- x_L).

(Figure 3) bearing (+)-JQ1^{18,19} as a warhead, have been developed, with the ultimate goal of finding more effective treatments compared to small-molecule inhibitors.^{18–20}

From a structural point of view, while a large variety of warheads have been explored to bind different POIs,

independently from their mechanism of action, cereblon (CRBN) targeting ligands (i.e., immunomodulatory imide drug (IMiD)-based ligands, such as thalidomide and its structural analogues)²¹ and Von Hippel-Lindau (VHL) targeting ligands (i.e., hydroxyproline-based molecules) are



Figure 3. Structures of representative BET-degrading PROTACs based on (+)-JQ1 as a warhead.

the most commonly used anchors to recruit E3 ligases (Figure 1).⁴ In contrast with the limited room of maneuver on both warhead and anchor, the linker and the linkage points represent important sites of diversification and are the focus of several medicinal chemistry programs currently. Linkers of different length and chemical nature (i.e., linear aliphatic and polyethylene glycol (PEG) chains and extended glycol chains) have been investigated with the main goal to modulate the degradation efficiency induced by PROTACs. On the same note, diverse linkage points of the linker to both the anchor and the warhead have been explored so far and their chemical nature is mainly driven by the chemical feasibility of PROTACs synthesis rather than by a rational choice (i.e., amine, ether, amide moieties, Figure 1). The sites of attachment on the anchor and the warhead are critical and are usually selected by analyzing the solvent-exposed regions, to minimize the interference of the linker with the binding of the PROTAC to the E3 ligase and the POI, respectively (Figure 1).

An optimal combination of all different PROTAC structural elements is of pivotal importance in triggering the productive formation of the ternary E3 ligase–PROTAC–POI complex and the following proteasome cascade activation.^{22,23} The rational design of effective PROTACs, which require cooperativity in assembling the ternary complex (TC) and establishing adventitious favorable interactions between the POI and the E3 ligase, is still a key challenge. However, progress in structural biology, X-ray crystallography, molecular modeling, and dynamics simulations has been recently made to rationalize the formation and stability of PROTAC-mediated TC and to understand the PROTAC mechanism of action.^{19,24–26}

Besides degradation efficiency, drug-like properties such as aqueous solubility, metabolic stability, and cell permeability represent additional challenges to face for achieving in vivo efficacy, with far more reason because PROTACs lie outside the rule-of-five space.^{27–29} Hence, massive efforts have been devoted to rationally explore the role of the different structural elements in conferring favorable in vitro ADME properties.³⁰

Because it has become increasingly clear that the length and chemical nature of the linker influence both the bioactivity and physicochemical properties of PROTACs, a shift from synthetically tractable linear aliphatic and PEG-based chains to more sophisticated and rigid motifs (e.g., piperazines, piperidines) is currently ongoing.^{27,31,32}

Despite the advances made so far, the rational conversion of a ligand for the POI into an effective degrader in vivo remains rather empirical and based on a "trial and error" approach.³³ Furthermore, the lack of reliable and economically sustainable synthetic methods represents an important caveat that necessarily limits the number of accessible compounds. The assemble of PROTACs has proven to be far from straightforward, as it involves the asymmetric diversification of the two sides of the linker and their chemoselective reactions with the two protein binding motifs. This challenge is usually achieved through orthogonal conditions or protection/deprotection sequences. Linking strategies to couple the linker with the warhead and the anchor include amide bond formation, N- and O-alkylations, nucleophilic aromatic substitution (S_NAr), acylation, and Williamson ether synthesis, transformations that are known to suffer from poor reactivity, low chemoselectivity, and lack of atom economy (Figure 1).³⁴ Thus, the preparation of PROTACs often requires low-yielding and cumbersome multistep synthetic routes, makes extensive use of protecting groups, and relies on highly functionalized and costly building blocks.3

To overcome these limitations, advances aimed at simplifying the access to protein degraders and their screening have been reported³² and include the use of orthogonally protected bifunctional linkers,³⁶ solid-phase synthesis,³⁷ click chemistry,³⁸ Staudinger ligation,³⁹ a direct-to-biology (D2B) approach,⁴⁰ and others.^{41,42} Nevertheless, none of these strategies has established itself as the ideal method so far, due to existing limitations. For instance, among the recent methods reported to ease the preparation of PROTACs, the click chemistry platform allows the preparation of both CRBN- and VHL-recruiting



"Reagents and conditions: (a) MeOH, 0 °C, 2 h, then rt, 1 h; (b) MeOH, 40 °C, 1 h, then 24 h rt; (c) TFA, CH_2Cl_2 , 0 °C, 30 min, then rt, 3 h; (d) MeOH, reflux, 2 h; (e) CH_2Cl_2 , 40 °C, 4 h. Color code does not refer to the MCR mechanism but is intended to represent the synthons and their assembly into the protein degraders.

degraders, but the introduction of the triazole ring is well-known for leading to poorly soluble products and for its reluctance to scale-up.⁴³ In an effort to streamline the discovery of successful PROTACs and expand the chemical space explored so far, we report a modular synthetic platform that capitalizes on the versatility of multicomponent reactions (MCRs) to assemble heterobifunctional protein degraders (Figure 1).

RESULTS AND DISCUSSION

Selection of Synthons for the Preparation of PROTACs via MCR Platform. In designing our platform we initially





"Reagents and conditions: (a) ethyl formate, reflux, 6 h; (b) TsCl, TEA, dry CH₂Cl₂, rt, 5 h; (c) NaHCO₃, dry DMF, 65 °C, 24 h.

focused on Ugi44,45 and Passerini46 reactions, as these MCRs stand out for their efficiency, versatility, high atom economy, and simple and environmentally friendly experimental procedures, where the only byproduct, when present, is one water molecule per molecule of product. As a proof of concept, we decided to use the following synthons as a model system: five different isocyanides bearing chains of different length and the thalidomide-based CRBN-recruiting anchor (1-5), a carboxylic acid (6) bearing the warhead based on (+)-JQ1,¹⁹ formaldehyde (7) as the carbonyl compound to minimize the interference of the linker with the binding to the POI, and three different commercially available amines (methylamine 8, tritylamine 9, and piperazine 10). Scheme 1 provides an overview of the synthesized PROTACs by using the aforementioned synthons and by exploiting the reactions as discussed in further detail below.

Preparation of Isocyanide-Based CRBN-Recruiting Anchors. While (+)-JQ1-based carboxylic acid (6), formaldehyde (7), and all the selected amines (8, 9, and 10) are commercially available, we applied a sustainable multistep protocol as depicted in Scheme 2 to access the desired thalidomide-based CRBN-recruiting isocyanides (1-5). The first step is the formylation of the proper amino alcohol (11-15)by using ethyl formate, followed by evaporation of the formylating agent. Further treatment of the N-formamide intermediate with tosyl chloride in the presence of triethylamine as base allowed us to obtain the corresponding linker (16-20)bearing both the isocyanide and the tosylate functionalities in 59-73% yield. According to a recent report by Meier et al. published in Green Chemistry, tosyl chloride represents an efficient and practical alternative to toxic and hazardous dehydrating agents such as phosphorus oxychloride in the preparation of aliphatic isocyanides.⁴⁷ This reagent proved to be suitable to our purpose, as it can simultaneously dehydrate the formamide group and convert the hydroxy function of 11-15 to an excellent leaving group. In the last step, an O-alkylation reaction takes place between derivatives 16-20 and 4hydroxythalidomide (21) in the presence of sodium bicarbonate to afford both linear aliphatic (1-3) and PEG (4 and 5) linkers in yields ranging from 46% to 85%. Notably, our optimized and mild experimental conditions avoided both the N-alkylation at the secondary nitrogen atom and the opening of the phthalimide/glutarimide ring of 21, two undesirable events that would result in the loss of E3 ligase recruitment.⁴⁸ Furthermore, all isolated isocyanides proved not to suffer from the unpleasant smell typical of these synthons, due to the high molecular weight and solid state at room temperature, and showed stability upon long-term storage.⁴⁹

Exploration of Different Amines To Probe the Warhead Linkage Point. With all isocyanide-base linkers in our hands, we turned our efforts into the variation of the amine component. We began with methylamine (8) as a model for conventional primary amines and PROTACs 22-26 (Scheme 1A) synthesized in 54-75% yields. To further reduce steric hindrance at the attachment point and minimally influence affinity of the warhead for its target protein, we then explored the use of tritylamine (9), that our laboratory had reported as an effective surrogate of ammonia,⁵⁰ especially when coupled with formaldehyde as the carbonyl component. The reaction was performed in methanol and, upon completion, the trityl group was cleaved by adding trifluoroacetic acid to give the desired PROTACs 27-31 in 55-82% yields (Scheme 1B). Finally, we carried out the split Ugi reaction, which we reported in 2006.⁵¹ In this variant of the Ugi reaction a secondary amine is used, instead of a primary one, leading to the acylation of one nitrogen atom, while the other one is alkylated. This transformation particularly suited our purposes, as it allowed the one-pot incorporation of the piperazine ring, a privileged substructure of several PROTACs, including ARV-110 and ARV-471 (Figure 2), 11,52 that helps to improve solubility and metabolic stability 27 and promotes the formation of stable TC.53-55 Noteworthy is the use of piperazine as the amine component, which allows an increase in the sp³ character and reduces the number of HBDs and HBAs that are associated with oral bioavailability improvement.³¹ Piperazine-based PROTACs 32-36 were successfully synthesized in yields of 63-83% (Scheme 1C).

Due to the well-known configurational instability that affects thalidomide substructures, all PROTACs were obtained as a mixture of diastereomers, which include a racemate at the IMiD stereocenter and an enantiopure (+)-configuration at the JQ1 stereocenter. To explore the scalability of our synthetic approach, the preparation of isocyanide 3 was performed on a 6.00 mmol scale and the split Ugi reaction was scaled up to 1.50 mmol to afford 1 g of 34 in 83% yield.

Implementation of the Platform with the Passerini MCR. Very recently, Ciulli et al. have reported the bioisosteric replacement of an amide function with an ester at the linkage point between the warhead and the linker as a successful strategy to enhance cell permeability, without significantly affecting metabolic stability.⁵⁶ This report encouraged us to include the Passerini reaction⁵⁷ in our MCR platform. To this aim, we selected the isocyanide-bearing linker 2, which reacted with (+)-JQ1-based carboxylic acid 6 and formaldehyde 7 in the absence of the amine component in dichloromethane to afford PROTAC 37 in 60% yield (Scheme 1D).

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Scheme 3. Synthesis of VHL-Recruiting Isocyanides 43 and 44 and PROTACs 45-47^a



^{*a*}Reagents and conditions: (a) KOH, MeOH, 0 $^{\circ}$ C to rt, 2 h and 30 min; (b) HATU, DIPEA, dry DMF, rt, 7 h. A and C refer to reaction conditions reported in Scheme 1A and 1C. Color code does not refer to the MCR mechanism but is intended to represent the synthons and their assembly into the protein degraders.



Figure 4. BRD4 degradation after treatment with CRBN-recruiting compounds. Western blot analysis of BRD4 degradation induced by CRBN-recruiting (A–D) compounds at the concentration of 1 μ M in MDA-MB-231 cells after 4, 8, 16, and 24 h of treatment. Treatments with 1 μ M of (+)-JQ1 or dBET1 were used as negative and positive controls for BRD4 degradation. α -Tubulin was used as internal control for equal loading. Degradation activity is reported below each lane as % of BRD4-immunoreactive band densitometry relative to DMSO vehicle. Western blot results shown in Figure 4 are representative of three (n = 3) independent experiments.

Preparation of VHL-Recruiting PROTACs To Probe the Anchor. To further investigate the versatility of our approach, we synthesized the isocyanide-based linkers bearing the anchor for the VHL proteins 43 and 44 via a two-step procedure depicted in Scheme 3. Saponification of isocyanide methyl esters 38 and 39 gave the corresponding potassium salts 40 and 41, which reacted with the commercially available hydrochloride amine 42 in the presence of HATU as coupling agent and DIPEA as base.

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Figure 5. BRD4 degradation after treatment with VHL-recruiting compounds. Western blot analysis of BRD4 degradation induced by VHL-recruiting (E) compounds at the concentration of 1 μ M in MDA-MB-231 cells after 4, 8, 16, and 24 h of treatment. Treatments with 1 μ M of cis-MZ1 or MZ1 were used as negative and positive controls for BRD4 degradation. α -Tubulin was used as internal control for equal loading. Degradation activity is reported below each lane as % of BRD4-immunoreactive band densitometry relative to DMSO vehicle. Western blot results shown in Figure 5 are representative of three (n = 3) independent experiments.

A final Ugi reaction using methylamine 8 and isocyanide 43 afforded VHL-recruiting PROTAC 45 in 40% yield, as well as two split Ugi reactions employing piperazine 10, and isocyanides 43 and 44 gave 46 and 47, respectively, in 42% and 87% yields (Scheme 3).

In Vitro Biological Validation of PROTACs and Preliminary Assessment of Thermodynamic Aqueous Solubility and Metabolic Stability of Selected BRD4-Degrading PROTACs. With the aim of proving the utility of our platform and the compatibility of the substructures accessible by the MCR strategy with a proximity-based degradation, the ability of our PROTAC candidates to effectively act as BRD4 degraders was tested in the human breast cancer cell line MDA-MB-231, known to express detectable levels of this protein. To this aim, BRD4 protein levels were assessed by Western blot analysis after exposing cells to each of the selected compounds. dBET1 PROTAC was used as the benchmark reference for comparison of CRBN-recruiting compounds, while (+)-JQ1 was used as negative control of protein degradation. Alternatively, MZ1 was identified as reference and cis-MZ1 as negative control for VHLrecruiting PROTAC candidates 45-47.

To preliminarily identify which of these molecules showed effective BRD4 degradation, we ran time-course experiments (4 up to 24 h) using 1 μ M concentration of each compound (Figures 4 and 5). As shown in Figure 4A–D, CRBN-recruiting compounds 27, 28, 29, and 34 led to a decreased expression of BRD4 at all the time-points tested. Also compounds 32 and 33 were able to induce BRD4 degradation, although the onset of this effect was delayed (after 8 h). Conversely, compound 30 caused a modest and transitory degradation of BRD4 (at 8 and 16 h of treatment), as well as compound 31 (at 16 h of treatment), while none of the VHL-recruiting compounds (45–47), with the exception of MZ1, acted as a BRD4 degrader (Figure 5).

According to time-course experiments, compounds 27, 28, 29, 32, 33, and 34 resulted in the most active degraders, albeit with different times of action. Therefore, these PROTACs were selected for further characterization of dose–response and proteasome-dependence of BRD4 degradation. To define whether their activity is dose-dependent, MDA-MB-231 cells were treated with increasing concentrations of each compound (from 100 nM up to 10 μ M) for 8 h, because at this time-point

they were found effective, as shown in Figure 4. By this approach, compounds 27, 28, and 32 showed dose-dependency, with an effect peaking between 1 and 10 μ M concentrations. Conversely, compounds 29 and 33 displayed a narrower window of active concentrations because degradation was observed only at 1 and 3 μ M concentrations (Figure 7). Notably, 34 was the most potent compound, was able to cause maximal degradation of BRD4 at 300 nM, and showed the well-known "hook effect".³³

To assess the dependency of BRD4 degradation on the proteasome machinery, we studied the effect of our best compounds in the presence of the proteasome inhibitor bortezomib. As shown in Figure 6, cotreatment with 5 nM



Figure 6. PROTAC activity is inhibited by the proteasome inhibitor bortezomib. BRD4 protein levels after 8 h treatments with DMSO (vehicle) or 1 μ M concentrations of compounds **27**, **28**, **29**, **32**, **33**, and **34** alone or in combinations with 5 nM bortezomib. α -Tubulin was used as internal control for equal loading. Western blot results shown in Figure 6 are representative of three (n = 3) independent experiments.

bortezomib was able, after 8 h, to abolish BRD4 degradation induced by compounds 27, 28, 29, 32, 33, and 34 (all used at the 1 μ M concentration), demonstrating the specificity of their molecular activity.

Of note, compound 37, which was synthesized using the Passerini reaction and is characterized by a α -acyloxy amide instead of a bis-amide, caused degradation of BRD4, although this effect was evident only after 16 and 24 h of incubation at the 1 μ M concentration (Figure 8A). Similarly to 34, when dose–

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Figure 7. Dose–response of BRD4 degradation induced by PROTAC candidates. Western blot analysis (left) of BRD4 protein levels in MDA-MB-231 cells after 8 h treatments with increasing concentrations (100 nM to 10 μ M) of compounds **27**, **28**, **29**, **32**, **33**, and **34** or DMSO (vehicle). α -Tubulin was used as internal control for equal loading. Results were normalized using densitometry of α -tubulin immunoreactive-bands as internal control for equal loading. Western blot shown in Figure 7 are representative of three (n = 3) independent experiments.



Figure 8. BRD4 degradation after treatment with compound 37. Western blot analysis of BRD4 protein levels in MDA-MB-231 cells treated with 1 μ M 37 for 4, 8, 16, and 24 h (A). Treatments with 1 μ M (+)-JQ1 or dBET1 were used as negative and positive controls for BRD4 degradation (A). Western blot analysis (B) of BRD4 protein levels in MDA-MB-231 cells after 8 h treatments with increasing concentrations (100 nM to 10 μ M) of compound 37 or DMSO (vehicle).

response degradation was investigated after 8 h of treatment, the maximal effect was seen at 300 nM (Figure 8B).

 α -Tubulin was used as internal control for equal loading. Degradation activity is reported below each lane as % of BRD4immunoreactive bands densitometry relative to DMSO vehicle. Western blot results shown in Figure 8 are representative of three (n = 3) independent experiments. DC₅₀ (nM) and D_{max} (%) of PROTACs 27–29, 32–34, and 37 were calculated and are reported in Table 1.

Overall, the SAR study suggests that, for the CRBN-targeting PROTACs, the substructure including the *N*-methyl-substituted tertiary amide (22-26) prevents in all cases BRD4 degradation, while the secondary bis-amide moiety as well as the acetyloxy amide favors the efficiency (27-30, 37), especially when the total linker length is between 10 and 12 atoms, irrespective of its hydrocarbon (27-29, 37) or PEG composition (29). Finally,

Table 1. DC_{50} (nM) and D_{max} (%) Values of Compounds 27–29, 32–34, and 37 after 8 h Treatments^{*a*}

compound	$DC_{50}(nM)$	D_{\max} (%)
27	97.1	88
28	134.0	79
29	184.0	86
32	239.0	81
33	89.9	78
34	60.0	94
37	62.0	86

"Values were calculated using densitometry of Western blots shown in Figures 7 and 8B. $D_{\rm max}$ is expressed as maximal % of reduction toward DMSO control lane.

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the **CONCLUSIONS**

In this study, we present an unprecedented MCR-based platform that streamlines the synthetic entry into the chemical space of protein degraders. By varying the amine component in the MCR, the platform allows extensive structure–activity and structure–property relationship studies around not only the linker and its length and composition but also the linkage point to the warhead that strongly influences the affinity of the PROTAC for the POI and might be responsible for metabolic instability.^{27,59}

When compared to the existing 2-CR methods, our approach stands out for being reliable, high yielding, versatile, protecting group-free, stereoconservative, and sustainable. While in this paper we have focused only on BRD4 degrading PROTACs, the platform might be used for targeting different POIs, provided that the warhead bearing a carboxylic group moiety is accessible. Our approach obviates the need to perform protection/ deprotection steps typically required in the synthesis of PROTACs, except for the use of tritylamine as a surrogate of ammonia. An additional feature of this strategy is given by the well-known stereoconservative nature of Ugi and Passerini reactions that is fundamental when chiral warheads and anchors are involved in the assembly of the protein degrader.^{60,61} Finally, the process is characterized by a low environmental footprint. Except for the (+)-JQ1-based carboxylic acid, all the other substrates required for the preparation of both the linkers and the final degraders are inexpensive and easy to handle, in stark contrast with the cost associated with most of the synthons that are currently commercially available.⁶²

In this paper, we have applied the platform mainly to BRD4degrading PROTACs and, by using a model toolkit, we have demonstrated the compatibility of the substructures accessible by the MCRs and placed at the attachment point between the warhead and the linker with degradation activity, aqueous solubility, and in vitro metabolic stability, despite the presence of additional HBAs and HBDs and potential soft spots. The full characterization and development of BRD4-degrading PRO-TACs are beyond the scope of the present report; however, we have identified a cluster of CRBN-binding PROTACs (27, 28, 29, 34, and 37) that cause a significant and proteasomaldependent reduction of BRD4 levels. At this stage, compound 34, displaying the privileged piperazine ring, and 37, endowed with an α -acyloxy amide, stand out for higher degrading activity, with DC₅₀ values of 60 and 62 nM, respectively, in MDA-MB-231 cells, and properties compatible with further preclinical development.

For the time being, we have only scraped the surface of the potential of the platform in the protein degraders field: the described variants have many applications and can be easily applied to other proteins for which ligands are available. Making protein degraders easily accessible to chemists at a reasonable cost must be a major objective in the time to come, and we envision that our method will help to achieve this goal.

EXPERIMENTAL SECTION

Chemistry. Commercially available reagents and solvents were used as purchased without further purification. When needed, solvents were distilled and stored on molecular sieves. Reactions were monitored by thin layer chromatography (TLC) carried out on 5 cm \times 20 cm silica gel plates with a layer thickness of 0.25 mm, using UV light as a visualizing agent. When necessary, TLC plates were visualized with aqueous KMnO₄ or aqueous acidic solution of cerium sulfate and ammonium molybdate reagent. Column chromatography was performed on flash

PROTACs displaying the piperazine ring are effective when the length is in the range of 13 to 15 (32-34), while longer chains are not tolerated (36).

Having demonstrated that the introduction of additional HBAs and HBDs does not prevent cell membrane crossing and subsequent BRD4 degradation, we were curious to preliminarily assess the aqueous solubility of structures potentially accessible through our platform and, to this aim, we selected compounds 27-29 as representative of Ugi-like products with different lengths, 34 that exemplifies the split-Ugi products and 37 as the result of the Passerini reaction. Being aware of thalidomide hydrolytic degradation at physiological pH (about 30% degradation in 3 h),²⁷ we determined the thermodynamic solubility for the selected compounds for 2 h to limit the permanence in aqueous media and monitored hydrolysis by high performance liquid chromatography (HPLC) analysis. At physiological pH, except for 27, all tested PROTACs showed a poor solubility (<30 μ M), which is comparable with the solubility of the majority of commercial PROTACs.³⁰ As expected, the Passerini product 37 that has one less HBD compared to 27-29 was the least soluble. Under acidic conditions, the solubility significantly increased, especially for compound 34 that displays an ionizable piperazine ring, further confirming the utility of introducing this substructure in PROTAC design (Table 2).

Table 2. Preliminary Data of Thermodynamic AqueousSolubility (μ M) and Metabolic Stability (Residual Substrate%) of Selected PROTACs

compound	PBS 0.1 M (pH = 7.4)	HCl 0.01 N (pH = 2.0)	metabolic stability ^{a,b} (%)
27	48	145	36
28	17	48	40
29	15	39	88
34	11	5073	60
37	8	18	89
dBET1	60 ^c	171 ^c	66

"Residual substrate after 1 h incubation in MLM in the presence of NADPH. ^bThe residual substrate in control incubations without MLMs was >99%. ^cLiterature data.⁵⁸

Finally, we were interested in ruling out the possibility that additional amides and/or esters may represent potential soft spots with a negative impact on metabolic stability. To this aim, we evaluated the metabolic stability of the five selected PROTACs in mouse liver microsomes (MLM). Interestingly, metabolic biotransformation only occurred when the microsomal monooxygenase system was activated by NADPH, suggesting susceptibility to oxidative metabolism but not to hydrolysis.²⁷ Contrary to what happened in phosphate saline buffer, no significant degradation of thalidomide was observed in control incubations carried out in Tris-HCl (see Experimental Section). Concerning the active compounds of the Ugi-like products series (27-29), the stability increased depending on the linker length: indeed, the residual substrate percentage after 1 h incubation raised from 36% for derivative 27 to 88% for 29. Also PROTAC 37, synthesized by the Passerini reaction, proved to be highly stable in MLM, in accordance with previous findings on PROTACs displaying esters in plasma (Table 2).⁵⁶

silica gel using Kieselgel 60 silica gel (particle size 0.040-0.063 mm, 230-400 mesh). Melting points were determined in open glass capillary with a Stuart Scientific SMP3 apparatus. All the target compounds were checked by IR (FT-IR Bruker Alpha II), ¹H NMR and ¹³C NMR (Bruker Avance Neo 400 MHz), and HRMS (Thermo Fisher Q-Exactive Plus) equipped with an Orbitrap (ion trap) mass analyzer. Chemical shifts are reported in parts per million (ppm) with residual solvent signals as internal standard (CDCl₃, δ = 7.26 ppm for ¹H NMR, δ = 77.16 ppm for ¹³C NMR; CD₃OD, δ = 3.31 ppm for ¹H NMR, δ = 49.00 ppm for ¹³C NMR; (CD₃)₂CO, δ = 2.05 ppm for ¹H NMR, δ = 29.84, 206.26 ppm for ¹³C NMR; $(CD_3)_2SO$, $\delta = 2.50$ ppm for ¹H NMR, $\delta = 39.52$ ppm for ¹³C NMR). Coupling constants (1) are quoted in hertz (Hz). Abbreviations used for multiplicity are as follows: s, singlet; d, doublet; t, triplet; q, quartet; quint, quintet; dd, doublet of doublets; dt, doublet of triplets; br, broad; m, multiplet. All PROTACs based on JQ1 structure were obtained as a mixture of diastereomers which include the racemic mixture at the IMiD stereocenter with enantiopure (+)-configuration at the JQ1 stereocenter.

The purity of selected compounds was determined by HPLC using a Shimadzu HPLC system (Shimadzu, Kyoto, Japan) equipped with a Kinetex C18 ($150 \times 4.6 \text{ mm}$, 5 μ m d.p., Phenomenex Torrance, CA) and using 0.2% formic acid in water and 0.2% formic acid in acetonitrile as eluents (for further details, see Supporting Information, SI). The purity of all tested compounds is 95% or higher.

General Procedure A for the Synthesis of Isocyanides 16–20. Amino alcohol (7.50 mmol, 1 equiv) **1–5** and ethyl formate (5.00 mL) were heated under reflux for 6 h. Then the volatile was removed in vacuo. The corresponding formamide (7.50 mmol, 1 equiv) was solubilized in dry CH_2Cl_2 (14.0 mL) under nitrogen and, after adding TEA (45.0 mmol, 6 equiv), the reaction mixture was cooled to 0 °C. Then TsCl (22.5 mmol, 3 equiv) was added, and the reaction was stirred at room temperature for 5 h. The mixture was quenched with a saturated aqueous Na_2CO_3 solution and stirred at 0 °C for 30 min. Water was added, and the aqueous phase was extracted with CH_2Cl_2 (×3). The combined organic layers were dried over sodium sulfate and evaporated. The crude product was purified by column chromatography using the eluent indicated below.

3-Isocyanopropyl 4-Methylbenzenesulfonate (16). The title compound was synthesized following the general procedure A. The crude material was purified by column chromatography using PE/ EtOAc 95:5 as eluent, affording compound 16 (1.31 g, 73%) as a yellow oil. ¹H NMR (400 MHz; CDCl₃): δ 7.82 (d, *J* = 8.4 Hz, 2H), 7.39 (d, *J* = 8.4 Hz, 2H), 4.19 (t, *J* = 5.8 Hz, 2H), 3.51 (t, *J* = 5.8 Hz, 2H), 2.48 (s, 3H), 2.05 (quint, *J* = 5.8 Hz, 2H). ¹³C NMR (101 MHz; CDCl₃): δ 157.5, 145.3, 132.4, 130.1, 127.9, 65.9, 37.8, 28.7, 21.7. HRMS (ESI) m/z (M + Na)⁺ calcd for C₁₁H₁₃NO₃SNa 262.0503, found 262.0504. IR (neat): $\tilde{\nu}$ = 2969, 2149, 1598, 1356, 1174, 1020, 944, 814, 664, 554 cm⁻¹.

4-Isocyanobutyl 4-Methylbenzenesulfonate (17). The title compound was synthesized following the general procedure A. The crude material was purified by column chromatography using PE/EtOAc 95:5 as eluent, affording compound 17 (1.23 g, 65%) as a yellow oil. ¹H NMR (400 MHz; CDCl₃): δ 7.80 (d, *J* = 8.4 Hz, 2H), 7.38 (d, *J* = 8.4 Hz, 2H), 4.09 (t, *J* = 5.9 Hz, 2H), 3.41 (t, *J* = 5.9 Hz, 2H), 2.48 (s, 3H), 1.87–1.82 (m, 2H), 1.78–1.74 (m, 2H). ¹³C NMR (101 MHz; CDCl₃): δ 156.8, 145.0, 132.9, 130.0, 127.9, 69.5, 40.9, 25.8, 25.2, 21.6. HRMS (ESI) *m*/*z* (M + Na)⁺ calcd for C₁₂H₁₅NO₃SNa 276.0659, found 276.0663. IR (neat): $\tilde{\nu}$ = 2923, 2148, 1597, 1354, 1174, 1018, 929, 814, 663, 555 cm⁻¹.

5-Isocyanopentyl 4-Methylbenzenesulfonate (18). The title compound was synthesized following the general procedure A. The crude material was purified by column chromatography using PE/ EtOAc 9:1 as eluent, affording compound 18 (1.14 g, 57%) as a yellow oil. ¹H NMR (400 MHz; CDCl₃): δ 7.81 (d, *J* = 8.4 Hz, 2H), 7.38 (d, *J* = 8.4 Hz, 2H), 4.06 (t, *J* = 6.3 Hz, 2H), 3.37 (dt, *J*_s = 6.6, 1.9 Hz, 2H), 2.48 (s, 3H), 1.75–1.63 (m, 4H), 1.53–1.47 (m, 2H). ¹³C NMR (101 MHz; CDCl₃): δ 155.9, 145.0, 132.8, 130.0, 127.8, 70.1, 41.3, 28.3, 27.9, 22.3, 21.6. HRMS (ESI) *m*/*z* (M + H)⁺ calcd for C₁₃H₁₈NO₃S 268.1002, found 268.1001. IR (neat): $\tilde{\nu}$ = 2950, 2148, 1597, 1353, 1174, 1019, 948, 815, 664, 555 cm⁻¹.

2-(2-Isocyanoethoxy)ethyl 4-Methylbenzenesulfonate (19). The title compound was synthesized following the general procedure A. The crude material was purified by column chromatography using PE/ EtOAc 9:1 as eluent, affording compound 19 (1.31 g, 65%) as a yellow oil. ¹H NMR (400 MHz; CDCl₃): δ 7.82 (d, *J* = 8.4 Hz, 2H), 7.38 (d, *J* = 8.4 Hz, 2H), 4.19 (t, *J* = 4.6 Hz, 2H), 3.78 (t, *J* = 4.7 Hz, 2H), 3.64 (t, *J* = 4.6 Hz, 2H), 3.51 (t, *J* = 4.7 Hz, 2H), 2.47 (s, 3H). ¹³C NMR (101 MHz; CDCl₃): δ 157.7, 145.0, 132.9, 129.9, 127.9, 69.0, 68.9, 68.6, 41.6, 21.6. HRMS (ESI) *m*/*z* (M + Na)⁺ calcd for C₁₂H₁₅NO₄SNa 292.0609, found 292.0609. IR (neat): $\tilde{\nu}$ = 2958, 2152, 1597, 1352, 1174, 1010, 919, 815, 663, 554 cm⁻¹.

2-(2-(2-lsocyanoethoxy)ethoxy)ethyl 4-Methylbenzenesulfonate (**20**). The title compound was synthesized following the general procedure A. The crude material was purified by column chromatography using PE/EtOAc 6:4 as eluent, affording compound **20** (1.67 g, 71%) as a yellow oil. ¹H NMR (400 MHz; CDCl₃): δ 7.82 (d, *J* = 8.2 Hz, 2H), 7.37 (d, *J* = 8.2 Hz, 2H), 4.19 (t, *J* = 4.7 Hz, 2H), 3.72 (t, *J* = 4.8 Hz, 2H), 3.70–3.67 (m, 2H), 3.65–3.62 (m, 4H), 3.58–3.54 (m, 2H), 2.47 (s, 3H). ¹³C NMR (101 MHz; CDCl₃): δ 157.4, 144.9, 133.1, 129.9, 127.9, 70.8 (2C), 69.2, 68.8, 68.7, 41.8, 21.6. HRMS (ESI) *m/z* (M + Na)⁺ calcd for C₁₄H₁₉NO₅SNa 336.0871, found 336.0872. IR (neat): $\tilde{\nu}$ = 2874, 2152, 1701, 1612, 1350, 1257, 1194, 1052, 734, 574 cm⁻¹.

General Procedure B for the Synthesis of Thalidomide-Bearing Isocyanides 1–5. To a solution of thalidomide derivative 21 (2.20 mmol, 1 equiv) in dry DMF (6.00 mL) was added NaHCO₃ (3.30 mmol, 1.5 equiv) under nitrogen, and the reaction mixture was heated at 65 °C. After 15 min, a solution of isocyanide 16–20 (2.64 mmol, 1.2 equiv) in dry DMF (500 μ L) was added dropwise and the resulting mixture was stirred at 65 °C overnight. The reaction mixture was diluted with CH₂Cl₂ and washed with water (×3). The organic layer was dried over sodium sulfate and the volatile solvent was removed in vacuo. The crude product was purified by column chromatography using the eluent indicated below.

2-(2,6-Dioxopiperidin-3-yl)-4-(3-isocyanopropoxy)isoindoline-1,3-dione (1). The title compound was synthesized following the general procedure B, starting from isocyanide 16. The crude material was purified by column chromatography using PE/EtOAc 6:4 as eluent, affording compound 1 (616 mg, 82%) as a yellow solid. ¹H NMR (400 MHz; CDCl₃): δ 8.38 (br s, 1H), 7.73 (t, *J* = 7.4 Hz, 1H), 7.51 (d, *J* = 7.4, 1H), 7.27 (d, *J* = 7.4 Hz, 1H), 4.97 (dd, *J*_s = 12.3, 5.5 Hz, 1H), 4.34 (t, *J* = 5.6 Hz, 2H), 3.77 (t, *J* = 5.6 Hz, 2H), 2.93–2.88 (m, 1H), 2.85– 2.75 (m, 2H), 2.25 (quint, *J* = 5.6 Hz, 2H), 2.17–2.14 (m, 1H). ¹³C NMR (101 MHz; CDCl₃): δ 171.0, 168.1, 166.9, 166.1, 157.1, 155.9, 136.7, 133.8, 119.1, 117.6, 116.5, 64.9, 49.2, 38.2, 31.4, 28.8, 22.6. HRMS (ESI) *m*/*z* (M + Na)⁺ calcd for C₁₇H₁₅N₃O₅Na 364.0898, found 364.0899. IR (neat): $\tilde{\nu}$ = 3094, 2153, 1698, 1614, 1396, 1200, 1051, 744, 610, 471 cm⁻¹. Mp: 199.7–201.1 °C, dec.

2-(2,6-Dioxopiperidin-3-yĪ)-4-(4-isocyanobutoxy)isoindoline-1,3dione (2). The title compound was synthesized following the general procedure B, starting from isocyanide 17. The crude material was purified by column chromatography using PE/EtOAc 6:4 as eluent, affording compound 2 (539 mg, 69%) as a white solid. ¹H NMR (400 MHz; DMSO-d₆): δ 11.10 (br s, 1H), 7.82 (t, *J* = 8.4 Hz, 1H), 7.52 (d, *J* = 8.4 Hz, 1H), 7.46 (d, *J* = 7.2 Hz, 1H), 5.09 (dd, *J*_s = 12.8, 5.4 Hz, 1H), 4.27 (t, *J* = 5.7 Hz, 2H), 3.65 (t, *J* = 5.7 Hz, 2H), 2.93–2.84 (m, 1H), 2.63–2.61 (m, 1H), 2.59–2.57 (m, 1H), 2.06–2.00 (m, 1H), 1.89– 1.82 (m, 4H). ¹³C NMR (101 MHz; DMSO-d₆): δ 173.2, 170.4, 167.3, 165.9, 156.3, 156.0, 137.5, 133.7, 120.3, 116.8, 115.8, 68.6, 49.3, 41.5, 31.4, 25.9, 25.8, 22.5. HRMS (ESI) *m*/*z* (M + Na)⁺ calcd for C₁₈H₁₇N₃O₅Na 378.1055, found 378.1058. IR (neat): $\tilde{\nu}$ = 3085, 2149, 1704, 1612, 1349, 1202, 1047, 747, 609, 464 cm⁻¹. Mp: 201.7–202.0 °C, dec.

2-(2,6-Dioxopiperidin-3-yl)-4-((5-isocyanopentyl)oxy)isoindoline-1,3-dione (3). The title compound was synthesized following the general procedure B, starting from isocyanide 18. The crude material was purified by column chromatography using PE/ EtOAc 5:5 as eluent, affording compound 3 (674 mg, 83%) as a white solid. ¹H NMR (400 MHz; CDCl₃): δ 8.25 (br s, 1H), 7.70 (t, *J* = 8.4 Hz, 1H), 7.48 (d, *J* = 8.4 Hz, 1H), 7.23 (d, *J* = 8.4 Hz, 1H), 4.97 (dd, *J*_s = 12.3, 5.3 Hz, 1H), 4.22 (t, J = 6.2 Hz, 2H), 3.47 (tt, $J_s = 6.2$, 1.9 Hz, 2H), 2.93–2.89 (m, 1H), 2.86–2.82 (m, 1H), 2.79–2.74 (m, 1H), 2.18–2.12 (m, 1H), 1.95 (quint, J = 6.2 Hz, 2H), 1.85–1.82 (m, 2H), 1.76–1.70 (m, 2H). ¹³C NMR (101 MHz; CDCl₃): δ 171.1, 168.2, 167.0, 165.6, 156.4, 156.2, 136.5, 133.8, 119.0, 117.3, 116.0, 69.1, 49.1, 41.4, 31.4, 28.7, 28.0, 23.0, 22.6. HRMS (ESI) m/z (M + Na)⁺ calcd for C₁₉H₁₉N₃O₅Na 392.1211, found 392.1208. IR (neat): $\tilde{\nu} = 3100, 2145$, 1697, 1613, 1365, 1195, 1049, 745, 613, 469 cm⁻¹. Mp: 174.6–175.9 °C, dec.

2-(2,6-Dioxopiperidin-3-yl)-4-(2-(2-isocyanoethoxy)ethoxy)isoindoline-1,3-dione (4). The title compound was synthesized following the general procedure B, starting from isocyanide 19. The crude material was purified by column chromatography using PE/ EtOAc 5:5 as eluent, affording compound 4 (588 mg, 72%) as a yellow solid. ¹H NMR (400 MHz; CDCl₃): δ 8.33 (br s, 1H), 7.71 (t, *J* = 7.2 Hz, 1H), 7.50 (d, *J* = 7.2 Hz, 1H), 7.29 (d, *J* = 7.2 Hz, 1H), 4.98 (dd, *J*_s = 12.1, 5.3 Hz, 1H), 4.38 (t, *J* = 4.5 Hz, 2H), 4.01 (t, *J* = 4.5 Hz, 2H), 3.91–3.88 (m, 2H), 3.62 (t, *J* = 5.5 Hz, 2H), 2.93–2.85 (m, 2H), 2.81– 2.75 (m, 1H), 2.18–2.13 (m, 1H). ¹³C NMR (101 MHz; DMSO-*d*₆): δ 173.2, 170.4, 167.3, 165.8, 156.8, 156.2, 137.5, 133.7, 120.6, 116.9, 116.0, 69.4, 69.1, 68.8, 49.3, 42.1, 31.4, 22.5. HRMS (ESI) *m/z* (M + H)⁺ calcd for C₁₈H₁₈N₃O₆ 372.1190, found 372.1193. IR (neat): $\tilde{\nu}$ = 3082, 2156, 1680, 1613, 1365, 1198, 1054, 744, 613, 467 cm⁻¹. Mp: 183.7–185.2 °C, dec.

2-(2,6-Dioxopiperidin-3-yl)-4-(2-(2-(2-isocyanoethoxy)ethoxy)ethoxy)isoindoline-1,3-dione (5). The title compound was synthesized following the general procedure B, starting from isocyanide 20. The crude material was purified by column chromatography using PE/ EtOAc 4:6 as eluent, affording compound 5 (420 mg, 46%) as a white solid. ¹H NMR (400 MHz; CDCl₃): δ 8.29 (br s, 1H), 7.70 (t, *J* = 7.8 Hz, 1H), 7.49 (d, *J* = 7.8 Hz, 1H), 7.28 (d, *J* = 7.8 Hz, 1H), 4.98 (dd, *J*_s = 12.2, 5.2 Hz, 1H), 4.37 (t, *J* = 4.6 Hz, 2H), 3.98 (t, *J* = 4.6 Hz, 2H), 3.84–3.81 (m, 2H), 3.74–3.71 (m, 4H), 3.58 (t, *J* = 5.8 Hz, 2H), 2.92– 2.82 (m, 2H), 2.79–74 (m, 1H), 2.16–2.13 (m, 1H). ¹³C NMR (101 MHz; DMSO-*d*₆): δ 173.2, 170.4, 167.3, 165.8, 156.3, 153.4, 137.5, 133.7, 120.5, 116.8, 115.9, 70.6, 70.2, 69.4, 69.2, 68.5, 49.3, 42.0, 31.1, 22.5. HRMS (ESI) *m*/*z* (M + Na)⁺ calcd for C₂₀H₂₁N₃O₇Na 438.1266, found 438.1270. IR (neat): $\tilde{\nu}$ = 3110, 2152, 1701, 1613, 1350, 1194, 1052, 747, 608, 467 cm⁻¹. Mp: 188.9–190.3 °C, dec.

General Procedure C for Methylamine-Mediated Ugi Reactions 22–26. To a solution of methylamine 8 (40% aqueous solution, 0.240 mmol, 2 equiv) in MeOH (500 μ L) were added formaldehyde 7 (37% aqueous solution, 0.360 mmol, 3 equiv), isocyanide 1–5 (0.180 mmol, 1.5 equiv), and carboxylic acid 6 (0.120 mmol, 1 equiv) at 0 °C and stirred for 2 h. The reaction mixture was left to reach room temperature and stirred for another 1 h. The volatile solvent was removed in vacuo, and the crude product was purified by column chromatography using the eluent indicated below.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((3-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)propyl)amino)-2-oxoethyl)-N-methylacetamide (22). The title compound was synthesized following the general procedure C, starting from thalidomide-bearing isocyanide 1. The crude material was purified by column chromatography using CH₃CN/MeOH 95:5 as eluent, affording compound 22 (70.6 mg, 75%) as a white solid. ¹H NMR (400 MHz; DMSO- d_{61} 353 K, *: refers to the main rotamer): δ 10.78 (br s, 1H), 8.08 (br s, 1H), 7.75 (t, J = 8.1 Hz, 1H), 7.47–7.44 (m, 5H), 7.42 (d, J = 7.1 Hz, 1H), 5.05 $(dd, J_s = 12.5, 5.4 Hz, 1H), 4.59 (t, J = 6.7 Hz, 1H), 4.26 (t, J = 6.2 Hz, 1H)$ 2H), 4.22-4.19 (m, 1H), 3.98-3.95 (m, 1H), 3.84-3.80 (m, 1H)*, 3.42–3.37 (m, 1H), 3.33 (t, J = 6.3 Hz, 2H), 3.21 (s, 3H), 2.92–2.83 (m, 2H), 2.65-2-63 (m, 1H)*, 2.60 (s, 3H), 2.42 (s, 3H), 2.09-2.03 (m, 1H), 1.96 (quint, J = 6.3 Hz, 2H), 1.66 (s, 3H). ¹³C NMR (101 MHz; CD₃OD, *: refers to the main rotamer): $\delta = 173.1^*$, 171.8, 169.9, 169.3, 167.2*, 164.8*, 156.1, 155.8, 150.7, 136.8, 136.6 (2C), 133.5, 133.4, 132.1, 131.8*, 130.7, 130.6, 130.0*, 128.4, 119.1*, 116.8, 115.2, 67.7*, 54.0*, 51.0, 49.1*, 36.6*, 36.1, 34.9, 30.8, 28.4*, 22.3, 13.0, 11.6, 10.3. HRMS (ESI) m/z (M + H)⁺ calcd for C₃₈H₃₈ClN₈O₇S 785.2267, found 785.2255. IR (neat): $\tilde{\nu}$ = 3368, 2924, 1708, 1643, 1355, 1196,

1049, 747, 545, 466 cm⁻¹. Mp: 211.5–213.8 °C, dec. See SI Figure S1 for NMR spectra of compound **22**.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((4-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)butyl)amino)-2-oxoethyl)-N-methylacetamide (23). The title compound was synthesized following the general procedure C, starting from thalidomide-bearing isocyanide 2. The crude material was purified by column chromatography using CH₃CN/MeOH 98:2 as eluent, affording compound 23 (69.9 mg, 73%) as a white solid. ¹H NMR (400 MHz; DMSO-d₆, 353 K): δ 7.78 (t, J = 7.9 Hz, 1H), 7.48–7.42 (m, 6H), 5.04 (dd, J_s = 12.6, 5.3 Hz, 1H), 4.59 (t, J = 6.7 Hz, 1H), 4.28 (t, J = 6.7 Hz, 2H), 4.21-4.19 (m, 1H), 3.97-3.95 (m, 1H), 3.23-3.18 (m, 5H), 3.05-3.03 (s, 2H), 2.92-2.84 (m, 2H), 2.65-2.63 (m, 1H), 2.42 (s, 3H), 2.60 (s, 3H), 2.09-2.03 (m, 1H), 1.82-1.78 (m, 4H), 1.66 (s, 3H). ¹³C NMR (101 MHz; CD₃OD, *: refers to the main rotamer): δ 173.1*, 171.8*, 169.6, 169.2, 167.2*, 166.1*, 164.8*, 156.4*, 155.9*, 150.8*, 136.8, 136.6, 136.5, 133.6*, 132.1, 131.8*, 130.7, 130.5, 130.0, 128.4, 119.2*, 116.7*, 115.0, 69.0, 54.0*, 51.0, 49.0, 38.5*, 36.1, 34.9, 34.2, 30.8, 25.7*, 22.3, 13.0, 11.6, 10.3. HRMS (ESI) m/z (M + H)⁺ calcd for C₃₉H₄₀ClN₈O₇S 799.2424, found 799.2408. IR (neat): $\tilde{\nu} = 3339$, 2922, 1706, 1648, 1393, 1194, 1048, 747, 558, 466 cm⁻¹. Mp: 190.5–192.0 °C, dec. See SI Figure S2 for the NMR spectra of compound 23.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((5-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)pentyl)amino)-2-oxoethyl)-N-methylacetamide (24). The title compound was synthesized following the general procedure C, starting from thalidomide-bearing isocyanide 3. The crude material was purified by column chromatography using CH₃CN/MeOH 98:2 as eluent, affording compound 24 (63.4 mg, 65%) as a yellow solid. ¹H NMR (400 MHz; DMSO- d_{61} 353 K): δ 7.77 (t, J = 7.9 Hz, 1H), 7.48–7.44 (m, 5H), 7.43–7.41 (m, 1H), 5.03 (dd, J_s = 12.3, 5.4 Hz, 1H), 4.62 (t, J = 6.7 Hz, 1H), 4.24 (t, J = 6.4 Hz, 2H), 4-07-4.03 (m, 1H), 3.68-3.55 (m, 2H), 3.45-3.41 (m, 1H), 3.23 (s, 3H), 3.18 (q, J = 6.4 Hz, 2H), 2.69–2.65 (m, 2H), 2.61 (s, 3H), 2.58-2.57 (m, 1H), 2.43 (s, 3H), 2.12-2.06 (m, 1H), 1.81 (quint, J = 6.4 Hz, 2H), 1.68 (s, 3H), 1.59–1.49 (m, 4H). ¹³C NMR (101 MHz; CD₃OD, *: refers to the main rotamer): δ 173.2, 171.8*, 170.0, 169.6, 169.1, 167.3, 166.0, 165.0*, 156.6*, 155.9*, 150.8, 136.8, 136.5*, 133.7, 132.1, 131.8, 130.7, 130.6, 129.4*, 128.4, 119.1, 116.8, 114.9, 69.0, 54.0*, 51.0, 49.0, 38.8*, 36.0, 34.9*, 30.8, 28.6*, 28.2*, 22.8, 22.3, 13.0, 11.5, 10.2. HRMS (ESI) m/z (M + Na)⁺ calcd for $C_{40}H_{41}ClN_8O_7SNa$ 835.2394, found 835.2381. IR (neat): $\tilde{\nu} = 3294$, 2924, 1709, 1644, 1355, 1196, 1048, 747, 545, 466 cm⁻¹. Mp: 175.2-176.8 °C, dec. See SI Figure S3 for NMR spectra of compound 24.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-peridin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)ethoxy)ethyl)amino)-2oxoethyl)-N-methylacetamide (25). The title compound was synthesized following the general procedure C, starting from thalidomide-bearing isocyanide 4. The crude material was purified by column chromatography using CH₃CN/MeOH 98:2 as eluent, affording compound 25 (52.8 mg, 54%) as a yellow solid. ¹H NMR (400 MHz; DMSO-d₆, 353 K): δ 10.81 (br s, 1H), 8.01 (br s, 1H), 7.78 $(t, J = 7.8 \text{ Hz}, 1\text{H}), 7.54 - 7.51 \text{ (m, 1H)}, 7.49 - 7.43 \text{ (m, 5H)}, 5.05 \text{ (dd, } J_s$ = 12.4, 5.3 Hz, 1H), 4.58 (t, J = 6.7 Hz, 1H), 4.36 (t, J = 4.7 Hz, 2H), 4.22-4.17 (m, 1H), 3.98-3.94 (m, 1H), 3.83-3.80 (m, 2H), 3.64-3.62 (m, 1H), 3.60-3.55 (m, 2H), 3.46-3.44 (m, 1H), 3.32-3.23 (m, 2H), 3.21 (s, 3H), 2.91-2.83 (m, 2H), 2.68-2.63 (m, 1H), 2.60 (s, 3H), 2.42 (s, 3H), 2.08-2.04 (m, 1H), 1.66 (s, 3H). ¹³C NMR (101 MHz; CD₂OD, *: refers to the main rotamer): $\delta = 173.1, 171.8^*, 170.0, \delta = 173.1, 170.0,$ 169.8, 167.1, 166.1, 164.8*, 156.4*, 155.8*, 150.8, 136.8, 136.6, 136.5, 133.6, 132.1, 131.8, 130.7, 130.6, 130.0*, 128.4, 119.5*, 116.9, 115.3, 69.3, 68.9, 68.6, 54.0*, 50.7, 49.1, 39.0*, 35.9, 34.9*, 30.8, 22.3, 13.0, 11.6, 10.3. HRMS (ESI) m/z (M + Na)⁺calcd for C₃₉H₃₉ClN₈O₈SNa 837.2187, found 837.2184. IR (neat): $\tilde{\nu} = 3274$, 2923, 1707, 1644, 1354, 1196, 1051, 747, 545, 466 cm⁻¹. Mp: 155.6–156.8 °C, dec. See SI Figure S4 for NMR spectra of compound 25.

amino)-2-oxoethyl)-N-methylacetamide (26). The title compound was synthesized following the general procedure C, starting from thalidomide-bearing isocyanide 5. The crude material was purified by column chromatography using CH₃CN/MeOH 97:3 as eluent, affording compound 26 (73.1 mg, 71%) as a yellow solid. ¹H NMR (400 MHz; DMSO-d₆, 353 K): δ 10.81 (br s, 1H), 7.98 (br s, 1H), 7.79 $(t, J = 7.8 \text{ Hz}, 1\text{H}), 7.52 - 7.49 \text{ (m, 2H)}, 7.47 - 7.44 \text{ (m, 4H)}, 5.05 \text{ (dd, } J_s$ = 12.5, 5.4 Hz, 1H), 4.59 (t, J = 6.7 Hz, 1H), 4.36 (t, J = 4.7 Hz, 2H), 4.22-4.18 (m, 1H), 3.98-3.94 (m, 1H), 3.82 (d, J = 4.7 Hz, 2H), 3.65 (t, J = 4.4 Hz, 2H), 3.57–3.53 (m, 4H), 3.47 (t, J = 5.9 Hz, 2H), 3.29– 3.26 (m, 2H), 3.21 (s, 3H), 2.91-2.83 (m, 2H), 2.66-2.63 (m, 1H), 2.60 (s, 3H), 2.42 (s, 3H), 2.10-2.04 (m, 1H), 1.66 (m, 3H). ¹³C NMR (101 MHz; CD₃OD, *: refers to the main rotamer): δ 173.1*, 171.8*, 170.0, 169.7, 167.2, 165.9, 164.8*, 156.4, 155.8, 150.7, 136.8, 136.5, 133.7, 132.1*, 131.8, 130.7, 130.6 (2C), 130.0, 128.4, 119.6, 116.9, 115.3, 70.6*, 70.0*, 69.1 (2C), 69.0, 54.0*, 50.8*, 49.1, 39.0*, 35.9, 34.9*, 30.8, 22.3, 13.0, 11.6, 10.2. HRMS (ESI) m/z (M + Na)⁺ calcd for $C_{41}H_{43}ClN_8O_9SNa$ 881.2449, found 881.2442. IR (neat): $\tilde{\nu} = 3294$, 2923, 1707, 1645, 1353, 1196, 1051, 747, 546, 465 cm⁻¹. Mp: 152.5-153.5 °C, dec. See SI Figure S5 for NMR spectra of compound 26.

General Procedure D for Tritylamine-Mediated Ugi Reactions 27–31. A solution of tritylamine 9 (0.120 mmol, 1 equiv) and paraformaldehyde 7 (0.240 mmol, 2 equiv) in MeOH (500 μ L) was stirred at 40 °C for 1 h. The reaction was then cooled to room temperature, and isocyanide 1–5 (0.120 mmol, 1 equiv) and carboxylic acid 6 (0.120 mmol, 1 equiv) were added. The resulting mixture was stirred overnight. The following day, the volatile solvent was removed in vacuo and the crude product was solubilized in CH₂Cl₂ (256 μ L). At 0 °C, TFA (256 μ L) was added and, after 30 min, the mixture was allowed to reach room temperature and stirred for 3 h. The crude material was subjected to column chromatography using Cy/EtOAc 5:5 to remove TFA and then CH₃CN/MeOH in the proportion indicated for each compound to elute the desired product.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((3-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)propyl)amino)-2oxoethyl)acetamide (27). The title compound was synthesized following the general procedure D, starting from thalidomide-bearing isocyanide 1. The crude material was purified by column chromatography using Cy/EtOAc 5:5 and CH₃CN/MeOH 9:1 as eluents, affording compound 27 (75.8 mg, 82%) as a white solid. 1 H NMR (400 MHz; $(CD_3)_2CO$: δ 10.02 (br s, 1H), 8.03 (br t, J = 5.8 Hz, 1H), 7.78–7.76 (m, 1H), 7.73 (t, J = 7.9 Hz, 1H), 7.53 (d, J = 8.2 Hz, 2H), $7.44-7.40 (m, 3H), 7.38 (d, J = 7.9 Hz, 1H), 5.12 (dd, J_s = 13.5, 4.4 Hz, 10.5)$ 1H), 4.66 (t, J = 6.2 Hz, 1H), 4.28 (t, J = 6.5 Hz, 2H), 4.11-4.04 (m, 1H), 3.78 (dt, $J_s = 16.8$, 4.5 Hz, 1H), 3.58-3.52 (m, 1H), 3.47 (t, J = 5.3Hz, 2H), 3.36 (dd, J_s = 14.9, 6.5 Hz, 1H), 2.91–2.88 (m, 1H), 2.78– 2.71 (m, 2H), 2.62 (s, 3H), 2.43 (s, 3H), 2.25-2.19 (m, 1H), 2.03-2.01 (m, 2H), 1.70 (s, 3H). ¹³C NMR (101 MHz; (CD₃)₂CO, *: refers to the main rotamer): δ 171.8*, 170.4*, 169.3, 169.2*, 166.9, 166.0, 163.8, 156.4, 155.8, 150.0, 137.3, 136.7, 135.9, 133.8, 132.6, 130.9*, 130.5, 130.3, 130.2, 128.4, 119.4, 117.0, 115.1, 67.9, 54.4, 49.3, 42.8*, 38.5, 36.4*, 31.1, 29.7, 22.5, 13.6, 12.1, 10.9. HRMS (ESI) m/z (M + Na)⁺ calcd for C₃₇H₃₅ClN₈O₇SNa 793.1925, found 793.1920. IR (neat): $\tilde{\nu}$ = 3351, 2922, 1707, 1656, 1394, 1196, 1049, 748, 529, 467 cm⁻¹. Mp: 221.3–223.6 °C, dec HPLC purity: > 99%. See SI Figure S6 for NMR spectra of compound 27 and Figure S20 for the HPLC chromatogram.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((4-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)butyl)amino)-2oxoethyl)acetamide (**28**). The title compound was synthesized following the general procedure D, starting from thalidomide-bearing isocyanide **2**. The crude material was purified by column chromatography using Cy/EtOAc 5:5 and CH₃CN/MeOH 9:1 as eluents, affording compound **28** (51.8 mg, 55%) as a white solid. ¹H NMR (400 MHz; (CD₃)₂CO): δ 10.03 (br d, J = 9.3 Hz, 1H), 8.17 (br t, J = 6.3 Hz, 1H), 7.76–7.72 (m, 2H), 7.53 (d, J = 8.2 Hz, 2H), 7.44–7.41 (m, 3H), 7.37 (d, J = 7.2 Hz, 1H), 5.11 (dd, J_s = 12.5, 5.4 Hz, 1H), 4.67 (t, J = 8.5 Hz, 1H), 4.20 (t, J = 6.4 Hz, 2H), 4.16–4.10 (m, 1H), 3.70–3.66 (m, 1H), 3.60–3.59 (m, 1H), 3.38–3.36 (m, 1H), 3.34–3.30 (m, 2H), 2.92–2.89 (m, 1H), 2.79–2.73 (m, 2H), 2.62 (s, 3H), 2.42 (s, 3H), 2.23–2.18 (m, 1H), 1.84–1.80 (m, 2H), 1.75–1.72 (m, 2H), 1.69 (s, 3H). ¹³C NMR (101 MHz; (CD₃)₂CO): δ 171.9, 170.4, 169.3, 169.2, 167.0, 165.6, 163.9 (2C), 156.6, 137.3 (2C), 136.7, 135.9, 133.9, 132.6, 131.0, 130.6, 130.3, 130.2, 128.4, 119.5, 117.0, 114.9, 69.1, 54.5, 49.2, 42.7, 38.5, 38.4, 31.1, 26.0, 25.9, 22.4, 13.6, 12.1, 10.9. HRMS (ESI) *m*/*z* (M + H)⁺ calcd for C₃₈H₃₈ClN₈O₇S 785.2267, found 785.2260. IR (neat): $\tilde{\nu}$ = 3323, 2929, 1709, 1651, 1394, 1195, 1048, 748, 528, 465 cm⁻¹. Mp: 231.9–232.6 °C, dec HPLC purity: 95.6%. See SI Figure S7 for NMR spectra of compound **28** and Figure S21 for the HPLC chromatogram.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((5-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)pentyl)amino)-2oxoethyl)acetamide (29). The title compound was synthesized following the general procedure D, starting from thalidomide-bearing isocyanide 3. The crude material was purified by column chromatography using Cy/EtOAc 5:5 and CH₃CN/MeOH 8:2 as eluents, affording compound 29 (67.1 mg, 70%) as a white solid. ¹H NMR (400 MHz; $(CD_3)_2CO$: δ 10.08 (br s, 1H), 8.22 (br t, J = 6.9 Hz, 1H), 7.77 (t, J = 8.4 Hz, 1H), 7.64 (br t, J = 5.8 Hz, 1H), 7.54 (d, J = 8.5 Hz, 2H),7.47–7.39 (m, 4H), 5.12 (ddd, J_s = 12.6, 5.4, 2.3 Hz, 1H), 4.68 (dd, J_s = 8.6, 6.1 Hz, 1H), 4.22 (t, J = 6.3 Hz, 2H), 4.09 (dd, $J_c = 18.5$, 6.7 Hz, 1H), 3.71 (dd, J_s = 16.8, 5.0 Hz, 1H), 3.58 (ddd, J_s = 15.0, 8.6, 3.4 Hz, 1H), 3.34 (ddd, J_s = 14.9, 6.1, 2.0 Hz, 1H), 3.25 (q, J = 6.3 Hz, 2H), 3.01-2.91 (m, 1H), 2.83-2.73 (m, 2H), 2.64 (s, 3H), 2.43 (s, 3H), 2.24-2.19 (m, 1H), 1.83 (quint, J = 6.3 Hz, 2H), 1.71 (s, 3H), 1.58-1.49 (m, 4H). $^{13}{\rm C}$ NMR (101 MHz; (CD₃)₂CO): δ 172.1, 170.6, 169.4 (2C), 167.0, 165.6, 163.9, 156.6, 155.8, 150.1, 137.3, 136.7, 135.9, 133.9, 132.5, 131.0, 130.5, 130.3, 130.2, 128.4, 119.5, 117.0, 115.0, 69.1, 54.5, 49.2, 42.7, 38.6, 38.4, 31.1, 28.8, 28.3, 22.7, 22.4, 13.6, 12.1, 10.9. HRMS (ESI) m/z (M + Na)⁺ calcd for C₃₉H₃₉ClN₈O₇SNa 821.2238, found 821.2231. IR (neat): $\tilde{\nu}$ = 3402, 2927, 1706, 1674, 1365, 1183, 1050, 750, 516, 467 cm⁻¹. Mp: 223.4–225.1 °C, dec HPLC purity: 97.8%. See SI Figure S8 for NMR spectra of compound 29 and Figure S22 for the HPLC chromatogram.

2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((2-(2-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)ethoxy)ethyl)amino)-2oxoethyl)acetamide (30). The title compound was synthesized following the general procedure D, starting from thalidomide-bearing isocyanide 4. The crude material was purified by column chromatography using Cy/EtOAc 5:5 and CH₃CN/MeOH 9:1 as eluents, affording compound 30 (56.7 mg, 59%) as a yellow solid. ¹H NMR (400 MHz; CD₃OD): δ 7.72 (t, J = 8.4 Hz, 1H), 7.46 (d, J = 8.6 Hz, 2H), 7.43–7.40 (m, 3H), 7.39 (d, J = 8.4 Hz, 1H), 5.09 (dd, $J_s = 12.8$, 5.8 Hz, 1H), 4.63 (t, J = 7.1 Hz, 1H), 4.33 (t, J = 4.1 Hz, 2H), 4.01-3.96 (m, 1H), 3.93-3.91 (m, 1H), 3.89-3.87 (m, 2H), 3.69 (t, J = 5.4 Hz, 2H), 3.47-3.44 (m, 4H), 2.88-2.78 (m, 2H), 2.73-2.71 (m, 1H), 2.69 (s, 3H), 2.43 (s, 3H), 2.13–2.09 (m, 1H), 1.68 (s, 3H). ¹³C NMR (101 MHz; CD₃OD): δ 173.1, 171.8, 169.9, 169.3, 167.2, 164.8, 156.1, 155.8, 150.7, 136.8, 136.6 (2C), 133.5, 133.4, 132.1, 131.8, 130.7, 130.6, 130.0, 128.4, 119.1, 116.8, 115.2, 67.7, 51.0, 37.2, 36.6, 36.1, 34.9, 34.2, 30.8, 28.4, 22.3, 13.0, 11.6, 10.3. HRMS (ESI) m/z (M + Na)⁺ calcd for $C_{38}H_{37}ClN_8O_8SNa$ 823.2030, found 823.2026. IR (neat): $\tilde{\nu} = 3328, 2970, 1710, 1656, 1365, 1200, 1050, 747, 529, 466$ cm⁻¹. Mp: 224.4–226.6 °C, dec. See SI Figure S9 for NMR spectra of compound 30.

2⁻((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-N-(2-((2-(2-(2-(2-(2,6-diax-opiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)ethoxy)ethoxy)ethoxy)ethyl)amino)-2-oxoethyl)acetamide (**31**). The title compound was synthesized following the general procedure D, starting from thalidomide-bearing isocyanide **5**. The crude material was purified by column chromatography using Cy/EtOAc 5:5 and CH₃CN/MeOH 9:1 as eluents, affording compound **31** (48.6 mg, 58%) as a yellow solid. ¹H NMR (400 MHz; (CD₃)₂CO): δ 10.00 (br s, 1H), 8.04 (br t, *J* = 6.1 Hz, 1H), 7.78 (t, *J* = 8.5 Hz, 1H), 7.58 (br t, *J* = 5.5 Hz, 1H), 7.54 (d, *J* = 8.5 Hz, 2H), 7.50 (d, *J* = 8.5 Hz, 1H), 7.44–7.42 (m, 3H), 5.11 (dd, *J*_s = 12.4, 5.3 Hz, 1H), 4.66 (t, *J* = 7.3 Hz, 1H), 4.39 (t, *J* = 4.6 Hz, 2H), 4.06 (dd, $J_s = 16.8, 6.9$ Hz, 1H), 3.90 (t, J = 4.6 Hz, 2H), 3.71 (t, J = 4.1 Hz, 2H), 3.68–3.66 (m, 1H), 3.60 (t, J = 5.4 Hz, 2H), 3.56–3.54 (m, 1H), 3.53–3.51 (m, 2H), 3.44–3.41 (m, 1H), 3.39–3.32 (m, 2H), 2.83–2.73 (m, 3H), 2.64 (s, 3H), 2.44 (s, 3H), 2.25–2.19 (m, 1H), 1.71 (s, 3H). ¹³C NMR (101 MHz; (CD₃)₂CO): δ 171.9, 170.4, 169.3 (2C), 166.9, 165.5, 163.8, 156.4, 155.8, 150.0, 137.3, 136.6, 135.9, 133.9, 132.6, 130.9, 130.5, 130.3, 130.2, 128.4, 119.9, 117.2, 115.3, 70.7, 70.1, 69.4, 69.3, 69.2, 54.4, 49.2, 42.7, 38.8, 38.4, 31.1, 22.4, 13.6, 12.1, 10.9. HRMS (ESI) m/z (M + H)⁺ calcd for C₄₀H₄₂ClN₈O₉S 845.2478, found 845.2465. IR (neat): $\tilde{\nu} = 3308, 2970, 1709, 1659, 1354, 1198, 1051, 747, 528, 466 cm⁻¹. Mp: 201.2–202.9 °C, dec. See SI Figure S10 for NMR spectra of compound 31.$

General Procedure E for Piperazine-Mediated Split Ugi Reactions 32–36. To a solution of piperazine 10 (0.120 mmol, 1 equiv) in MeOH (500 μ L) were added paraformaldehyde 7 (0.120 mmol, 1 equiv), isocyanide 1–5 (0.120 mmol, 1 equiv), and carboxylic acid 6 (0.120 mmol, 1 equiv) sequentially. The reaction mixture was heated at reflux for 2 h, the volatile was removed in vacuo, and the crude product was purified by column chromatography using the eluent indicated below.

2-(4-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetyl)piperazin-1-yl)-N-(3-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)propyl)acetamide (32). The title compound was synthesized following the general procedure E, starting from thalidomide-bearing isocyanide 1. The crude material was purified by column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound 32 (68.5 mg, 68%) as a white solid. ¹H NMR (400 MHz; CDCl₃): δ 10.12 (br s, 1H), 7.84-7.81 (m, 1H), 7.79-7.72 (m, 1H), 7.50 (d, J = 8.5 Hz, 2H), 7.49-7.46 (m, 1H), 7.44-7.42 (m, 2H), 7.42-7.40 (m, 1H), 5.13 (dd, $J_s = 12.4, 5.3 \text{ Hz}, 1\text{H}$, 4.72 (t, J = 6.6 Hz, 1H), 4.33 (t, J = 6.0 Hz, 2H), 3.76-3.73 (m, 2H), 3.66-3.60 (m, 2H), 3.51 (q, J = 6.6 Hz, 2H), 3.47-3.45 (m, 1H), 3.43-3.41 (m, 1H), 3.04 (s, 2H), 2.93-2.89 (m, 2H), 2.83-2.78 (m, 2H), 2.75-2.73 (m, 1H), 2.62 (s, 3H), 2.51-2.48 (m, 2H), 2.44 (s, 3H), 2.26–2.20 (m, 1H), 2.11–2.08 (m, 2H), 1.70 (s, 3H). ¹³C NMR (101 MHz; (CD₃)₂CO, *: refers to the main rotamer): δ 171.9*, 169.4*, 169.3, 168.5, 166.9*, 165.7*, 163.1, 156.4, 155.8*, 149.6, 137.4, 136.8*, 135.8, 133.9*, 132.7, 130.7, 130.4, 130.2 (2C), 128.4, 119.7*, 117.2*, 115.3*, 67.8*, 61.5, 54.6, 53.5, 53.0, 49.3, 45.4, 41.4, 36.0, 35.0, 31.2, 30.7, 22.5*, 13.7, 12.2, 10.9. HRMS (ESI) m/z $(M + H)^+$ calcd for $C_{41}H_{43}ClN_9O_7S$ 840.2689, found 840.2677. IR (neat): $\tilde{\nu} = 3366, 2927, 1709, 1639, 1357, 1198, 1048, 747, 559, 467$ cm⁻¹. Mp: 205.1–207.4 °C, dec. See SI Figure S11 for NMR spectra of compound 32.

2-(4-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetyl)piperazin-1-yl)-N-(4-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)butyl)acetamide (33). The title compound was synthesized following the general procedure E, starting from thalidomide-bearing isocyanide 2. The crude material was purified by column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound 33 (84.0 mg, 82%) as a white solid. ¹H NMR (400 MHz; $(CD_3)_2CO$): δ 10.16 (br d, J = 15.5 Hz, 1H), 7.81–7.74 (m, 2H), 7.51 (d, J = 8.3 Hz, 2H), 7.48–7.46 (m, 1H), 7.44–7.41 (m, 3H), 5.12 (dd, J_s = 14.2, 6.9 Hz, 1H), 4.73 (t, J = 8.0 Hz, 1H), 4.29 (t, J = 6.5 Hz, 2H), 3.79 - 3.76 (m, 2H), 3.66 (dd, J_s = 16.2, 6.9 Hz, 2H), 3.48–3.46 (m, 1H), 3.43–3.41 (m, 1H), 3.37 (q, J = 6.5 Hz, 2H), 3.01 (s, 2H), 2.96-2.91 (m, 2H), 2.82-2.75 (m, 2H), 2.71-2.67 (m, 1H), 2.62 (s, 3H), 2.51-2.48 (m, 2H), 2.44 (s, 3H), 2.24–2.20 (m, 1H), 1.90 (quint, J = 6.5 Hz, 2H), 1.78 (quint, J = 6.5 Hz, 2H), 1.70 (s, 3H). ¹³C NMR (101 MHz; (CD₃)₂CO, *: refers to the main rotamer): δ 172.0, 169.4 (2C)*, 168.6, 167.0*, 165.6*, 163.2, 156.5*, 155.8, 149.7, 137.4, 136.7, 135.8, 133.9, 132.7, 130.8, 130.4, 130.3, 130.2, 128.4, 119.4*, 117.0*, 115.1*, 69.1, 61.5, 54.6, 53.5, 53.1, 49.3*, 45.4, 41.5, 38.1, 35.1, 31.2*, 26.4, 26.2, 22.5, 13.7, 12.2, 11.0. HRMS (ESI) m/z (M + H)⁺ calcd for C₄₂H₄₅ClN₉O₇S 854.2846, found 854.2837. IR (neat): $\tilde{\nu}$ = 3344, 2946, 1711, 1643, 1362, 1197, 1048, 747, 558, 466 cm⁻¹. Mp: 229.4–230.0 °C, dec. See SI Figure S12 for NMR spectra of compound 33.

2-(4-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetyl)piperazin-1-yl)-N-(5-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)pentyl)- acetamide (34). The title compound was synthesized following the general procedure E, starting from thalidomide-bearing isocyanide 3. The crude material was purified by column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound 34 (78.1 mg, 75%) as a white solid. ¹H NMR (400 MHz; CD₃OD): δ 7.77 (t, J = 8.2 Hz, 1H), 7.47 (d, J = 8.7 Hz, 2H), 7.45–7.42 (m, 4H), 5.10 (dd, $J_s = 13.6$, 5.3 Hz, 1H), 4.70 (t, J = 6.8 Hz, 1H), 4.26 (t, J = 6.0 Hz, 2H), 3.82-3.79 (m, 2H), 3.70 (t, J = 6.0 Hz, 2H), 3.61-3.57 (m, 2H), 3.38-3.35 (m, 2H)2H), 3.09 (s, 2H), 2.92-2.80 (m, 2H), 2.76-2.73 (m, 1H), 2.72 (s, 3H), 2.69-2.65 (m, 2H), 2.56-2.53 (m, 2H), 2.47 (s, 3H), 2.13-2.09 (m, 1H), 1.92 (quint, J = 6.0 Hz, 2H), 1.72 (s, 3H), 1.68–1.60 (m, 4H). ^{13}C NMR (101 MHz; CD₃OD): δ 173.1, 171.1, 170.0, 169.3, 167.3, 165.9, 164.7, 156.6, 155.8, 150.7, 136.8, 136.6, 136.5, 133.7, 132.1, 131.8, 130.7, 130.6, 130.0, 128.4, 119.1, 116.7, 115.0, 69.0, 60.8, 54.0, 53.0, 52.7, 49.1, 45.3, 41.5, 38.5, 34.6, 30.8, 28.8, 28.3, 23.2, 22.3, 13.0, 11.6, 10.2. HRMS (ESI) m/z (M + H)⁺ calcd for C₄₃H₄₇ClN₉O₇S 868.3002, found 868.2990. IR (neat): $\tilde{\nu} = 3363$, 2924, 1711, 1640, 1364, 1197, 1047, 747, 558, 466 cm⁻¹. Mp: 194.8–196.8 °C, dec HPLC purity: 95.7%. See SI Figure S13 for NMR spectra of compound 34 and Figure S23 for the HPLC chromatogram.

2-(4-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetyl)piperazin-1-yl)-N-(2-(2-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)ethoxy)ethyl)acetamide (35). The title compound was synthesized following the general procedure E, starting from thalidomide-bearing isocyanide 4. The crude material was purified by column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound 35 (65.7 mg, 63%) as a yellow solid. ¹H NMR (400 MHz; $(CD_3)_2CO$): δ 9.96 (br s, 1H), 7.85-7.81 (m, 1H), 7.63-7.60 (m, 1H), 7.52 (d, J 8.6 Hz, 2H), 7.45-7.37 (m, 4H), 5.12 (dd, $J_s = 12.5$, 5.5 Hz, 1H), 4.70 (t, $J_s = 12.5$, 5.5 Hz, 10.5 H = 6.4 Hz, 1H), 4.46-4.43 (m, 2H), 3.94 (t, J = 6.0 Hz, 2H), 3.74-3.71 (m, 2H), 3.63–3.59 (m, 2H), 3.56–3.54 (m, 2H), 3.46 (q, J = 5.6 Hz, 2H), 3.41-3.89 (m, 1H), 3.37-3.35 (m, 1H), 2.99 (s, 2H), 2.94-2.90 (m, 1H), 2.79–2.77 (m, 2H), 2.75–2.73 (m, 2H), 2.63 (s, 3H), 2.48 (s, 3H), 2.39–2.36 (m, 2H), 2.25–2.20 (m, 1H), 1.74 (s, 3H). ¹³C NMR (101 MHz; DMSO-d₆): δ 173.1, 170.3, 169.6, 168.6, 167.7, 166.2, 163.4, 156.1, 155.7, 150.2, 137.3, 136.9, 135.7, 134.3, 132.7, 131.2, 130.6, 130.4, 130.1, 128.9, 120.0, 117.4, 115.6, 69.8, 69.3, 68.9, 61.4, 54.6, 53.4, 53.0, 51.7, 44.5, 41.6, 38.6, 35.2, 31.0, 23.6, 14.5, 13.1, 11.7. HRMS (ESI) m/z (M + H)⁺ calcd for C₄₂H₄₅ClN₉O₈S 870.2795, found 870.2788. IR (neat): $\tilde{\nu}$ = 3338, 2925, 1712, 1639, 1365, 1173, 1051, 748, 558, 465 cm⁻¹. Mp: 173.8–174.5 °C, dec. See SI Figure S14 for NMR spectra of compound 35.

2-(4-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f]-[1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetyl)piperazin-1-yl)-N-(2-(2-(2-((2-(2,6-dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)ethoxy)ethoxy)ethyl)acetamide (36). The title compound was synthesized following the general procedure E, starting from thalidomide-bearing isocyanide 5. The crude material was purified by column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound 36 (80.0 mg, 73%) as a yellow solid. ¹H NMR (400 MHz; $(CD_3)_2CO$: δ 10.02 (br s, 1H), 7.78 (t, J = 8.6 Hz, 1H), 7.60 (br t, J = 6.1 Hz, 1H), 7.53-7.49 (m, 3H), 7.45-7.42 (m, 3H), 5.11 (dd, $J_s =$ 13.9, 6.7 Hz, 1H), 4.70 (t, J = 6.7 Hz, 1H), 4.43 (t, J = 4.5 Hz, 2H), 3.95 (t, J = 4.5 Hz, 2H), 3.79 (t, J = 4.6 Hz, 2H), 3.67-3.62 (m, 5H), 3.58-3.54 (m, 5H), 3.43-3.38 (m, 2H), 2.99 (s, 2H), 2.95-2.93 (m, 2H), 2.79-2.74 (m, 2H), 2.68-2.65 (m, 1H), 2.63 (s, 3H), 2.54-2.49 (m, 2H), 2.45 (s, 3H), 2.25–2.19 (m, 1H), 1.73 (s, 3H). ¹³C NMR (101 MHz; $(CD_3)_2CO_1$ *: refers to the main rotamer): δ 171.2, 169.2, 169.0, 168.5, 166.9, 165.4, 163.1, 156.5*, 155.8, 149.6, 137.5, 136.6*, 135.8, 134.0, 132.7, 130.7, 130.4, 130.3, 130.2, 128.4, 120.0, 117.3, 115.3, 71.0, 70.2, 69.6, 69.5, 69.3, 61.4, 54.7, 53.4, 53.0, 49.3*, 45.5, 41.5, 38.5, 35.0, 31.1*, 22.4*, 13.6, 12.1, 10.9. HRMS (ESI) m/z (M + H)⁺ calcd for $C_{44}H_{49}ClN_9O_9S$ 914.3057, found 914.3053. IR (neat): $\tilde{\nu} = 3351, 2925,$ 1711, 1643, 1364, 1199, 1051, 748, 558, 466 cm⁻¹. Mp 194.5–195.8 °C, dec. See SI Figure S15 for NMR spectra of compound 36.

Synthesis of 2-((4-((2-(2,6-Dioxopiperidin-3-yl)-1,3-dioxoisoindolin-4-yl)oxy)butyl)amino)-2-oxoethyl2-((S)-4-(4-chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f][1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetate (**37**). To a solution of thalidomide-bearing isocyanide **2** (0.240 mmol, 2 equiv) in CH₂Cl₂ (1.30 mL) were added formaldehyde 7 (37% aqueous solution, 0.480 mmol, 4 equiv) and (+)-JQ1 carboxylic acid 6 (0.120 mmol, 1 equiv), and the resulting mixture was stirred at 40 °C for 4 h. Then the volatile was removed in vacuo, and the crude product was purified by column chromatography using Cy/EtOAc 1:9 as eluent, affording compound 37 (56.5 mg, 60%) as a white solid. ¹H NMR (400 MHz; DMSO- d_6): δ 11.10 (br s, 1H), 8.09 (t, J = 5.8 Hz, 1H), 7.80 (t, J = 6.0 Hz, 1H), 7.53–7.48 (m, 3H), 7.45–7.43 (m, 3H), 5.45 (t, J = 5.8 Hz, 1H), 5.08 (dd, $J_s = 13.0$, 5.3 Hz, 1H), 4.52 (dd, $J_s =$ 24.2, 14.8 Hz, 2H), 4.21 (t, J = 5.7 Hz, 2H), 3.57 (dd, J_s = 6.7, 4.0 Hz, 2H), 3.19 (q, J = 5.7 Hz, 2H), 2.92-2.83 (m, 1H), 2.61 (s, 3H), 2.57-2.54 (m, 2H), 2.40 (s, 3H), 2.06–2.00 (m, 1H), 1.76 (quint, J = 5.7 Hz, 2H), 1.66–1.64 (m, 2H), 1.62 (s, 3H). ¹³C NMR (101 MHz; DMSO d_{6i} *: refers to the main rotamer): δ 173.1, 170.6, 170.3, 167.3, 166.9, 165.8, 163.9, 156.5, 155.2, 150.5, 137.5, 137.2, 135.8, 133.8, 132.7, 131.3, 130.7, 130.5, 130.1, 128.9, 120.4, 116.9, 115.7, 69.2, 63.0*, 53.9, 49.3*, 38.4*, 36.9, 31.4*, 26.3, 26.0, 22.5*, 14.4, 13.1, 11.7. HRMS (ESI) m/z (M + H)⁺ calcd for C₃₈H₃₇ClN₇O₈S 786.2107, found 786.2097. IR (neat): $\tilde{\nu}$ = 3370, 2921, 1708, 1554, 1391, 1195, 1048, 747, 559, 467 cm⁻¹. Mp: 105.2–107.0 °C, dec HPLC purity: 97.6%. See SI Figure S16 for NMR spectra of compound 37 and Figure S24 for the HPLC chromatogram.

General Procedure F of Saponification Reaction for 40 and 41. Isocyanide **38** or **39** (1.00 mmol, 1 equiv) was solubilized in dry MeOH (1.00 mL) under nitrogen, and the resulting mixture was cooled to 0 °C. Then KOH (1.00 mmol, 1 equiv) was added, and the reaction was stirred for 30 min. The mixture was allowed to reach room temperature and stirred for additional 2 h. Then the volatile solvent was removed in vacuo, and the obtained products **40** or **41** was used, without further purification and characterization, in the following step.

General Procedure G for the Synthesis of Isocyanides 43 and 44. To a solution of **40** or **41** (0.500 mmol, 1 equiv) in dry DMF were sequentially added (2S,4R)-1-((S)-2-amino-3,3-dimethylbutanoyl)-4hydroxy-*N*-(4-(4-methylthiazol-5-yl)benzyl)pyrrolidine-2-carboxamide hydrochloride **42** (0.500 mmol, 1 equiv), DIPEA (2.00 mmol, 4 equiv), and HATU (0.630 mmol, 1.25 equiv) under nitrogen. The reaction was stirred at room temperature for 7 h. EtOAc was added, and the organic layer was washed with water (×3), dried over sodium sulfate, and evaporated. The crude material was purified by column chromatography using the eluent indicated below.

(2S,4R)-4-Hydroxy-1-((S)-2-(2-isocyanoacetamido)-3,3-dimethylbutanoyl)-N-(4-(4-methyl thiazol-5-yl)benzyl)pyrrolidine-2-carboxamide (43). The title compound was synthesized following the general procedure G, using isocyanide 40. The crude material was purified by column chromatography using Cy/EtOAc 1:9 as eluent, affording compound 43 (99.4 mg, 40%) as a yellow solid. ¹H NMR (400 MHz; CD₃OD): δ 8.94 (s, 1H), 8.18 (br s, 1H), 7.91 (br d, J = 8.6 Hz, 1H), 7.48 (d, J = 4.2 Hz, 2H), 7.44 (d, J = 4.2 Hz, 2H), 4.67 (t, J = 8.0, 1H), 4.58–4.54 (m, 1H), 4.52 (s, 2H), 4.38 (dd, *J*_s = 15.1, 5.5 Hz, 1H), 3.98 $(d, J = 11.1 \text{ Hz}, 1\text{H}), 3.91 (dd, J_s = 11.1, 3.8 \text{ Hz}, 1\text{H}), 3.83 (s, 2\text{H}), 2.50$ (s, 3H), 2.25–2.21 (m, 1H), 2.13–2.07 (m, 1H), 1.06 (s, 9H). ¹³C NMR (101 MHz; CD₃OD, *: refers to the main rotamer): δ 173.0*, 170.6*, 169.3, 162.8, 147.5, 138.9 (2C), 132.0, 130.0, 129.0, 127.6, 70.0, 59.4, 57.7*, 56.6, 42.3, 40.5*, 37.5*, 35.4*, 25.6, 14.3. HRMS (ESI) m/z (M-H)⁻ calcd for C₂₅H₃₀N₅O₄S 496.2013, found 496.2027. IR (neat): $\tilde{\nu} = 3292, 2922, 2159, 1664, 1535, 1416, 1229, 1084, 728,$ 550 cm⁻¹. Mp: 63.1–64.6 °C.

(25,4R)-4-Hydroxy-1-((S)-2-(4-isocyanobutanamido)-3,3-dimethylbutanoyl)-N-(4-(4-methylthiazol-5-yl)benzyl)pyrrolidine-2-carboxamide (44). The title compound was synthesized following the general procedure G, using isocyanide 41. The crude material was purified by column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound 44 (86.7 mg, 33%) as a white solid. ¹H NMR (400 MHz; CDCl₃): δ 8.68 (s, 1H), 7.36 (d, *J* = 8.3 Hz, 2H), 7.32 (d, *J* = 8.3 Hz, 2H), 6.70 (br d, *J* = 8.8 Hz, 1H), 4.67 (t, *J* = 8.0, 1H), 4.56 (d, *J* = 8.9 Hz, 1H), 4.54–4.48 (m, 2H), 4.34 (dd, *J*_s = 15.1, 5.4 Hz, 1H), 3.99 (d, *J* = 11.4 Hz, 1H), 3.67 (dd, *J* = 11.2, 3.7 Hz, 1H), 3.45–3.38 (m, 2H), 2.50 (s, 3H), 2.41 (t, *J* = 7.4 Hz, 2H), 0.96 (s, 9H). HRMS (ESI) m/z (M + H)⁺ calcd for C₂₇H₃₆N₅O₄S 526.2483, found

526.2481. IR (neat): $\tilde{\nu}$ = 3295, 2953, 2148, 1621, 1531, 1416, 1224, 1085, 850, 569 cm⁻¹. Mp: 65.7–66.8 °C.

(2S,4R)-1-((S)-2-(2-(2-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f][1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)-Nmethylacetamido)acetamido)acetamido)-3,3-dimethylbutanoyl)-4-hydroxy-N-(4-(4-methylthiazol-5-yl)benzyl)pyrrolidine-2-carboxamide (45). To a solution of methylamine 8 (40% aqueous solution, 0.240 mmol, 2 equiv) in MeOH (500 μ L) were added formaldehyde 7 (37% aqueous solution, 0.360 mmol, 3 equiv), isocyanide 43 (0.180 mmol, 1.5 equiv), and carboxylic acid 6 (0.120 mmol, 1 equiv) at 0 °C, and the mixture was stirred for 2 h at the same temperature. The volatile solvent was removed in vacuo, and the crude product was purified by column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound **45** (45.1 mg, 40%) as a yellow solid. ¹H NMR (400 MHz; CD₃OD, *: refers to the main rotamer): δ 8.84 (s, 1H), 8.58 (br s, 1H), 7.47-7.38 (m, 8H), 4.73 (t, J = 7.0, 1H), 4.65-4.63 (m, 1H), 4.55 (s, 2H), 4.51 (s, 2H), 4.36 (d, J = 15.3 Hz, 1H), 4.25 (d, J = 15.3 Hz, 1H), 4.07 (d, J = 16.4 Hz, 1H), 4.02-3.98 (m, 1H), 3.92-3.85 (m, 1H), 3.80 $(d, J = 3.9 \text{ Hz}, 1\text{H}), 3.76 (dd, J_s = 14.5, 7.0 \text{ Hz}, 1\text{H}), 3.64 (dd, J_s = 16.1)$ 7.0 Hz, 1H), 3.37 (s, 3H), 2.70 (s, 3H), 2.47 (s, 3H), 2.44 (s, 3H), 2.19-2.10 (m, 2H), 1.69 (s, 3H)*, 1.00 (s, 9H)*. ¹³C NMR (101 MHz; CD₃OD, *: refers to the main rotamer): δ 173.0*, 171.9*, 170.5*, 170.4*, 169.8*, 165.0, 159.8, 155.9*, 150.8*, 147.6, 138.9, 136.8, 136.6*, 132.1, 132.0, 131.8*, 130.7*, 130.6, 130.1, 130.0*, 129.0, 128.4*, 127.6*, 69.7, 59.4, 57.7*, 56.5*, 53.9*, 52.5, 51.0, 42.3*, 37.5, 36.1, 35.5*, 34.8*, 25.6, 14.4, 13.0*, 11.5, 10.2*. HRMS (ESI) $m/z \, ({\rm M}$ + H)⁺ calcd for C₄₆H₅₄ClN₁₀O₆S₂ 941.3352, found 941.3347. IR (neat): $\tilde{\nu} = 3294$, 2925, 1626, 1532, 1415, 1348, 1194, 747, 548, 445 cm⁻¹. Mp: 114.6-116.0 °C. See SI Figure S17 for NMR spectra of compound 45.

General Procedure H for Piperazine-Mediated Split Ugi Reactions 46 and 47. To a solution of piperazine 10 (0.120 mmol, 1 equiv) in MeOH (500 μ L) were added paraformaldehyde 7 (0.120 mmol, 1 equiv), isocyanide 43 or 44 (0.240 mmol, 2 equiv), and carboxylic acid 6 (0.120 mmol, 1 equiv) sequentially. The reaction mixture was heated at reflux for 2 h, the volatile solvent was removed in vacuo, and the crude product was purified by column chromatography using the eluent indicated below.

(2S,4R)-1-((S)-2-(2-(2-(4-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f][1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetyl)piperazin-1-yl)acetamido)acetamido)-3,3-dimethylbutanoyl)-4-hydroxy-N-(4-(4-methylthiazol-5-yl)benzyl)pyrrolidine-2-carboxamide (46). The title compound was synthesized following the general procedure H, using (2S,4R)-4-hydroxy-1-((S)-2-(2-isocyanoacetamido)-3,3-dimethylbutanoyl)-N-(4-(4-methylthiazol-5-yl)benzyl) pyrrolidine-2-carboxamide 43. The crude material was purified by column chromatography using CH₃CN/MeOH 8:2 as eluent, affording compound 46 (50.2 mg, 42%) as a yellow solid. ¹H NMR (400 MHz; CD₃OD): δ 8.86 (s, 1H), 8.57 (br s, 1H), 7.45–7.43 (m, 2H), 7.42-7.39 (m, 4H), 7.37 (d, J = 8.2 Hz, 2H), 4.69 (t, J = 8.0, 1H), 4.68-4.66 (m, 1H), 4.63-4.61 (m, 1H), 4.59-4.56 (m, 1H), 4.52 (s, 2H), 4.36 (dd, J_s = 15.1, 5.5 Hz, 1H), 4.01 (d, J = 3.4 Hz, 1H), 3.90 (d, J = 11.1 Hz, 1H), 3.84–3.82 (m, 4H), 3.79–3.75 (m, 1H), 3.74–3.66 (m, 2H), 3.58 (d, J = 6.9 Hz, 2H), 3.17–3.15 (m, 2H), 2.70 (s, 3H), 2.62– 2.59 (m, 2H), 2.46 (s, 3H), 2.45 (s, 3H), 2.27-2.22 (m, 1H), 2.14-2.09 (m, 1H), 1.71 (s, 3H), 1.06 (s, 9H). ¹³C NMR (101 MHz; CD₃OD): δ 173.1, 172.0, 170.5, 169.7, 169.3, 168.1, 164.7, 155.8, 150.7, 147.6, 138.9, 136.8, 136.5, 132.1, 132.0, 131.8, 130.7 (2C), 130.1, 129.9, 128.9, 128.4, 127.6, 69.7, 60.7, 59.5, 57.6, 56.6, 54.0, 53.0, 52.7, 45.4, 42.3, 41.8, 41.6, 37.6, 35.6, 34.6, 25.6, 14.5, 13.0, 11.6, 10.2. HRMS (ESI) m/z (M + H)⁺ calcd for C₄₉H₅₉ClN₁₁O₆S₂ 996.3774, found 996.3765. IR (neat): $\tilde{\nu}$ = 3293, 2921, 1625, 1528, 1417, 1228, 1088, 731, 548, 444 cm⁻¹. Mp: 183.3–184.8 °C. See SI Figure S18 for NMR spectra of compound 46.

 $(2S, \hat{4R})$ -1-((S)-2-($\hat{4}$ -(2-(4-(2-((S)-4-(4-Chlorophenyl)-2,3,9-trimethyl-6H-thieno[3,2-f][1,2,4]triazolo[4,3-a][1,4]diazepin-6-yl)acetyl)piperazin-1-yl)acetamido)butanamido)-3,3-dimethylbutanoyl)-4hydroxy-N-(4-(4-methylthiazol-5-yl)benzyl)pyrrolidine-2-carboxamide (47). The title compound was synthesized following the general procedure H, using (2S,4R)-4-hydroxy-1-((S)-2-(4-isocyanobutanamido)-3,3-dimethylbutanoyl)-N-(4-(4-methylthiazol-5-yl)benzyl) pyrrolidine-2-carboxamide 44. The crude material was purified by column chromatography using CH₃CN/MeOH 8:2 as eluent, affording compound 47 (107 mg, 87%) as a white solid. ¹H NMR (400 MHz; CD₃OD): δ 8.87 (s, 1H), 7.47–7.44 (m, 4H), 7.43–7.39 (m, 4H), 4.70 (t, J = 6.9 Hz, 1H), 4.66-4.64 (m, 1H), 4.61 (t, J = 8.3 Hz, 1H), 4.52 (s, 2H), 4.36 (d, J = 15.5 Hz, 1H), 3.94 (d, J = 11.0 Hz, 1H), 3.84-3.82 (m, 3H), 3.73-3.70 (m, 2H), 3.64 (d, J = 9.1 Hz, 1H), 3.60-3.58 (m, 1H), 3.30-3.28 (m, 2H), 3.10 (s, 2H), 2.70 (s, 3H), 2.68-2.65 (m, 2H), 2.55 (t, J = 5.1 Hz, 2H), 2.47 (s, 3H), 2.45 (s, 3H), 2.37–2.32 (m, 2H), 2.25-2.21 (m, 1H), 2.13-2.10 (m, 1H), 1.85 (quint, J = 7.8 Hz, 2H), 1.70 (s, 3H), 1.07 (s, 9H). ¹³C NMR (101 MHz; CD₃OD): δ 173.8, 173.1, 171.3, 170.9, 169.3, 164.7, 155.8, 151.4, 150.7, 147.6, 138.8, 136.8, 136.5, 132.1, 132.0, 131.8, 130.7, 130.6, 130.0 (2C), 128.9, 128.4, 127.6, 69.7, 60.8, 59.4, 57.8, 56.6, 54.0, 53.0, 52.7, 45.3, 42.3, 41.5, 38.2, 37.6, 35.1, 34.6, 32.5, 25.7, 25.5, 14.5, 13.0, 11.6, 10.2. HRMS (ESI) m/z (M + H)⁺ calcd for C₅₁H₆₃ClN₁₁O₆S₂ 1024.4087, found 1024.4085. IR (neat): $\tilde{\nu}$ = 3302, 2921, 1628, 1526, 1416, 1227, 1088, 803, 549, 445 cm⁻¹. Mp: 164.3–165.5 °C. See SI Figure S19 for NMR spectra of compound 47.

Synthesis of Isocyanide 18 on a Multigram Scale. Amino alcohol 13 (2.10 g, 20.0 mmol, 1 equiv) and ethyl formate (13.0 mL) were heated under reflux for 6 h. The volatile solvent was removed in vacuo. The corresponding formamide (2.62 g, 20.0 mmol, 1 equiv) was solubilized in dry CH_2Cl_2 (30.0 mL) under nitrogen and, after adding TEA (16.7 mL, 120 mmol, 6 equiv), the reaction mixture was cooled at 0 °C. TsCl (11.4 g, 60.0 mmol, 3 equiv) was added, and the reaction was stirred at room temperature for 5 h. The reaction mixture was quenched with a saturated aqueous Na_2CO_3 solution and stirred under cooling at 0 °C for 30 min. Water was added, and the aqueous phase was extracted with CH_2Cl_2 (×3). The organic layer was dried over sodium sulfate and evaporated. The crude product was purified by silica gel column chromatography using PE/EtOAc 9:1, affording compound 18 (3.15 g, 59%) as a yellow oil.

Synthesis of Isocyanide-Based Library of CRBN-Recruiting Anchor 3 on a Multigram Scale. To a solution of thalidomide derivative 21 (1.65 g, 6.00 mmol, 1 equiv) in dry DMF (25.5 mL) was added NaHCO₃ (756 mg, 9.00 mmol, 1.5 equiv) under nitrogen, and the reaction mixture was heated at 65 °C. After 15 min, a solution of isocyanide 18 (1.92 g, 7.20 mmol, 1.2 equiv) in dry DMF (1.50 mL) was added dropwise and the resulting mixture was stirred at 65 °C overnight. The reaction mixture was diluted with CH_2Cl_2 and washed with water (×3). The organic layer was dried over sodium sulfate, and the volatile solvent was removed in vacuo. The crude product was purified by silica gel column chromatography using PE/EtOAc 5:5 as eluent, affording compound 3 (1.88 g, 85%) as a white solid.

Scaled-up Synthesis of PROTAC 34 by Split Ugi Reaction on a One-Gram Scale. To a solution of piperazine 10 (129 mg, 1.50 mmol, 1 equiv) in MeOH (6.00 mL) were added paraformaldehyde (72.1 mg, 1.50 mmol, 1 equiv), isocyanide 3 (554 mg, 1.50 mmol, 1 equiv), and carboxylic acid 6 (601 mg, 1.50 mmol, 1 equiv) sequentially. The reaction mixture was heated at reflux for 2 h, the volatile solvent was removed in vacuo, and the crude product was purified by silica gel column chromatography using CH₃CN/MeOH 9:1 as eluent, affording compound 34 (1.08 g, 83%) as a white solid.

Cell Culturing and Western Blot Analysis. The human breast adenocarcinoma MDA-MB-231 cell line was cultured in DMEM (Sigma-Aldrich) supplemented with 10% fetal bovine serum (FBS), 1% glutamine, and 1% penicillin/streptomycin. For Western blot analysis, 15×10^5 cells/well were plated into 12-well plates containing DMEM complete growth medium, to be treated with the synthesized compounds, at the indicated concentrations, for different time lengths. After treatments, cells were scraped and lysed to obtain protein extract using RIPA lysis buffer supplemented with protease (PIC, Merck-Millipore) and phosphatase inhibitors (NaF 1 M, Na₃VO₄ 1 M, PMSF 100 nM, Sigma-Aldrich). Cells lysates were centrifuged at 13 000 rpm for 10 min at 4 °C to remove membranes and insoluble fractions. Protein quantification of the supernatant fraction was performed with Braford protein assay (Sigma-Aldrich). Thirty milligram aliquots of protein extracts were used for 12% SDS-PAGE. After transfer into nitrocellulose membranes, BRD4 and tubulin protein levels were

evaluated by immunoblotting using a rabbit polyclonal anti-BRD4 (Bethyl Laboratories) and a mouse monoclonal α -tubulin (Sigma-Aldrich) IgG as primary antibodies and an antirabbit or antimouse HRP-conjugated secondary antibody (Bio-Rad Laboratories). Protein levels were detected by chemiluminescent conversion of the HRP substrate LumiGLO (Thermo Fisher Scientific) using the ChemiDoc imaging system (Bio-Rad). Band densitometry was assessed, normalized to α -tubulin immunoreactive bands, and reported as percentage of the 0.3% DMSO control lane.

Thermodynamic Solubility. Saturated solutions were prepared by dissolving the tested compounds in 0.1 M phosphate-buffered saline (pH = 7.4) or 0.01 N HCl solutions and sonicated for 30 s. After equilibration on an orbital shaker at 25 °C for 2 h, the solutions were centrifuged at 13 000 rpm for 5 min and filtered through a 0.2 μ m regenerated cellulose (RC) syringe filter. A 400 μ L aliquot of each filtrate was diluted with 100 μ L of acetonitrile and analyzed by LC-UV. When the measured peak area was out of the calibration range (5–100 μ M), the sample was further diluted in water–acetonitrile 1:1 mixture.

Metabolic Stability. Mouse liver microsomes (MLM) (pooled male mouse CD-1, protein concentration: 20 mg/mL) were purchased from Corning B.V. Life Sciences (Amsterdam, The Netherlands). The standard incubation mixture (250 μ L final volume) was carried out in a 50 mM Tris (tris[hydroxymethyl]aminomethane) buffer (pH 7.4) containing 150 mM KCl, 1.5 mM, 3.3 mM MgCl₂, 1.3 mM NADPNa₂, 3.3 mM glucose 6-phosphate, 0.4 units/mL glucose 6-phosphate dehydrogenase, acetonitrile as cosolvent (1% of total volume), and the substrate (50 μ M). After pre-equilibration of the mixture, an appropriate volume of MLM suspension was added to give a final protein concentration of 1.0 mg/mL. The mixture was shaken for 60 min at 37 °C. Control incubations were carried out without the presence of an NADPH-regenerating system or microsomes. Each incubation was stopped by the addition of 250 μ L of ice-cold acetonitrile, vortexed, and centrifuged at 13 000 rpm for 5 min. The supernatants were analyzed by LC-UV.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.jmedchem.2c01218.

¹H and ¹³C NMR spectra of final compounds; LC-UV methods for determination of purity of selected active compounds, aqueous solubility, and metabolic stability; HPLC chromatograms (PDF)

Molecular formula strings (CSV)

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Author Contributions

Regarding the synthesis, M.S. and T.P. conceived and designed the experiments. I.P.B., A.R., and M.S. performed the experiments. I.P.B., A.R., R.M.C.D.M., and T.P. analyzed the experimental data. Regarding the biochemistry evaluation, F.C. designed the experiments and F.C.B. performed them. Regarding the evaluation of metabolic stability and solubility, S.A. designed and performed the experiments. T.P. and R.M.C.D.M. wrote the manuscript with input from all the authors. All the authors have approved the final version of the manuscript.

Notes

The authors declare no competing financial interest.

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ABBREVIATIONS USED

2-CR, two-component reaction; AR, androgen receptor; BCLx₁, B-cell lymphoma-extra large; BET, bromodomain and extraterminal domain; CRBN, cereblon; D2B, direct-to-biology; DIPEA, N,N-diisopropylethylamine; DMEM, Dulbecco's Modified Eagle's Medium; EtOAc, ethyl acetate; ER, estrogen receptor; FBS, fetal bovine serum; HATU, 1-[bis-(dimethylamino)methylene]-1H-1,2,3-triazolo[4,5-b]pyridinium 3-oxide hexafluorophosphate; IMiD, immunomodulatory imide drug; MCR, multicomponent reaction; MDA-MB-231 Cells, MD Anderson-metastatic breast-231 cells; MeOH, methanol; MLM, mouse liver microsomes; PE, petroleum ether; POI, protein of interest; PROTAC, Proteolysis TArgeting Chimera; RC, regenerate cellulose; RIPA, radioimmunoprecipitation assay; S_NAr , nucleophilic aromatic substitution; TC, ternary complex; TEA, triethylamine; TFE, trifluoroethanol; TsCl, tosyl chloride; VHL, von Hippel-Lindau.

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