

The background features a 3D rendering of several blue capsules and green molecular structures. One capsule is lying on its side on the left, another is on the right, and a third is in the center, partially obscured by the title. Green molecular models, consisting of spheres and connecting lines, are scattered around the capsules. The entire scene is set against a light gray background with a subtle grid pattern.

# Targeted Protein Degradation as Emerging Antiviral Therapeutics

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1<sup>st</sup> year PHD student  
16/12/2021

Department of Microbiology



# *Outline*

## **PART 01 Protein degradation**

## **PART 02 Targeted protein degradation(TPD)**

- Principal
- Advantages
- Applications

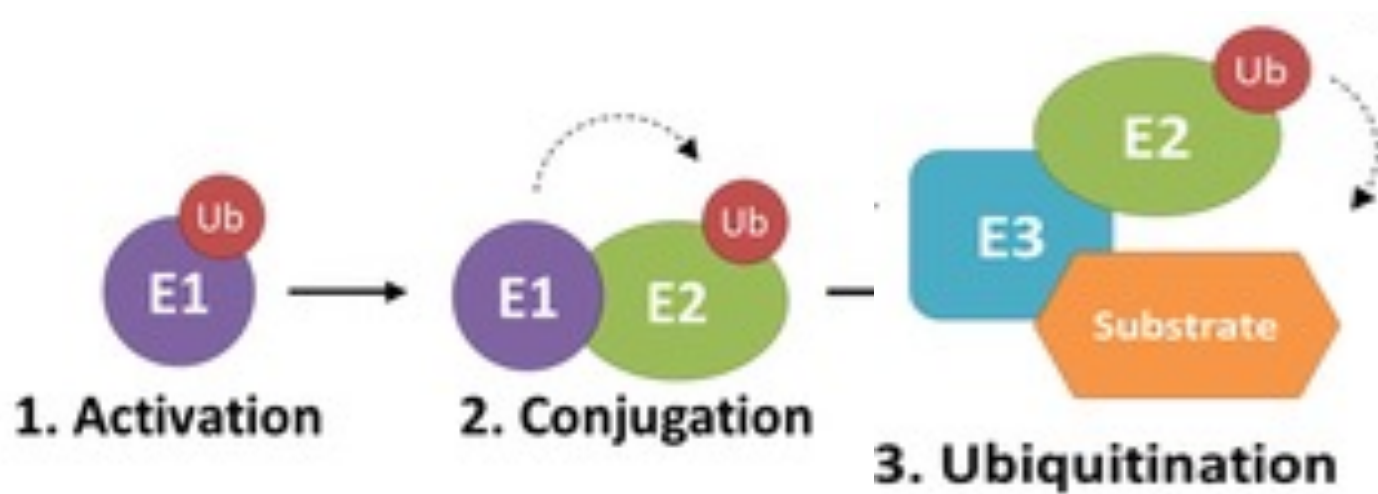
## **PART 03 TPD as antiviral therapeutics**

- Anti-HBV
- Anti-HCV
- Anti-Cov

## **PART 04 Challenges of TPD**

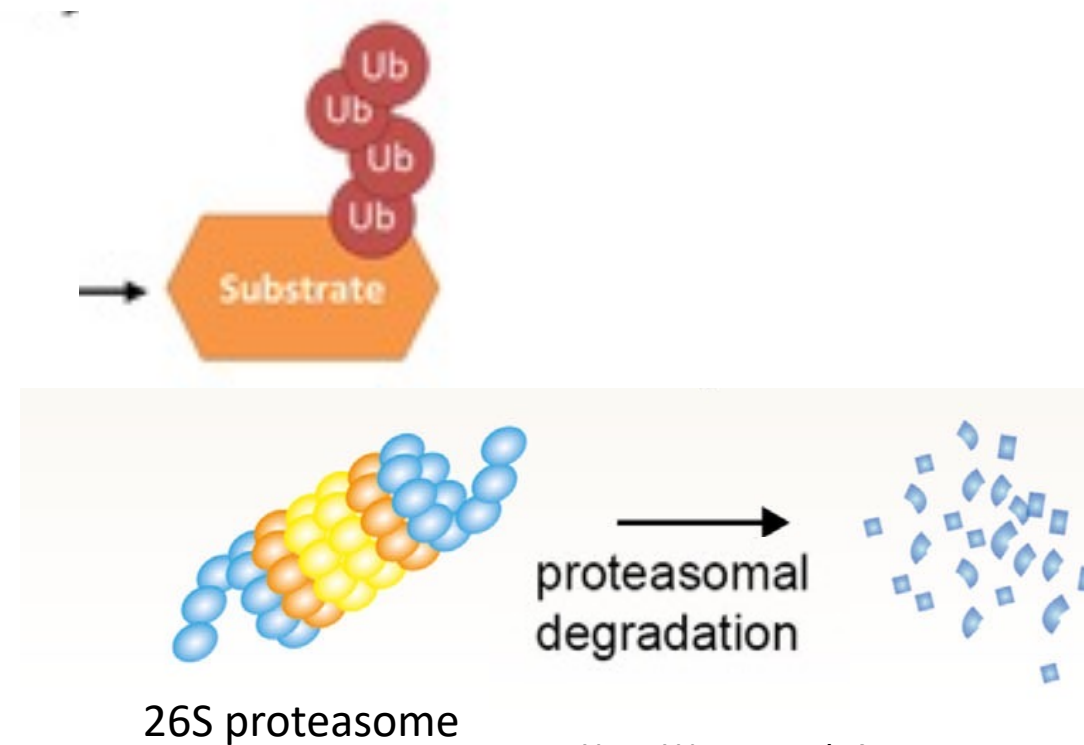
# ❖ Protein degradation

- Misfolded proteins
- Maintaining protein homeostasis



## Ubiquitin-proteasome System (UPS)

- 26S proteasome mediated



# ❖ Protein degradation

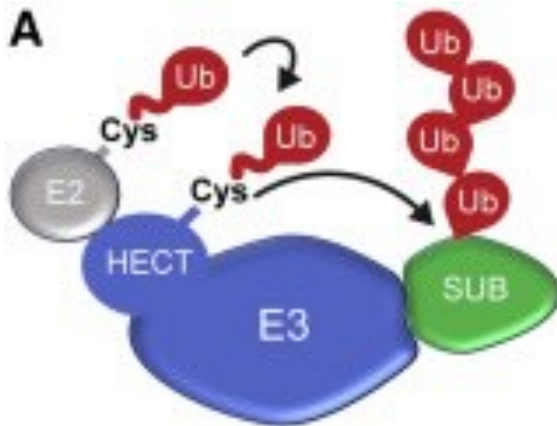
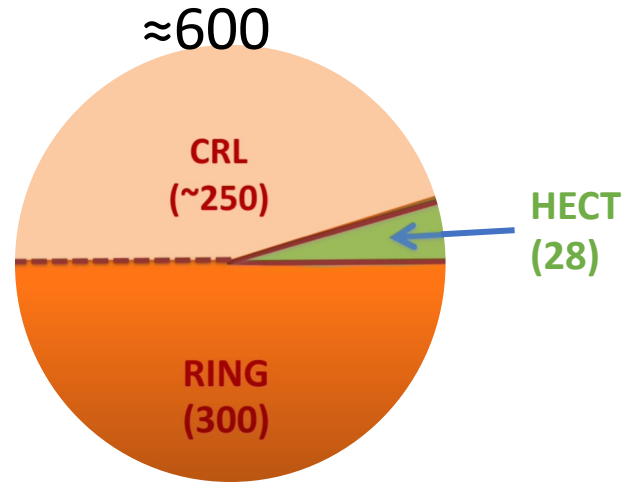
## E3-ubiquitin ligase

### (N)Target recognizing domain

- Specific recognition
- Specificity and versatility

### (C)E2 interacting domain

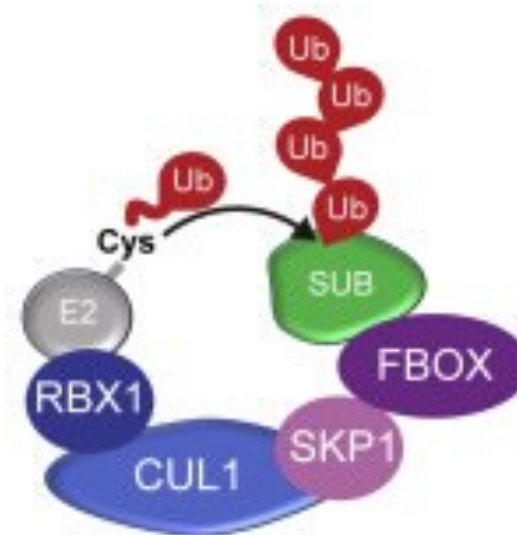
- HECT
- RING-finger protein



**HECT E3 ligase**  
*i.e. UBR5, NEDD4*



**RING E3 ligase**  
*i.e. MDM2, ciAP1, UBR1*

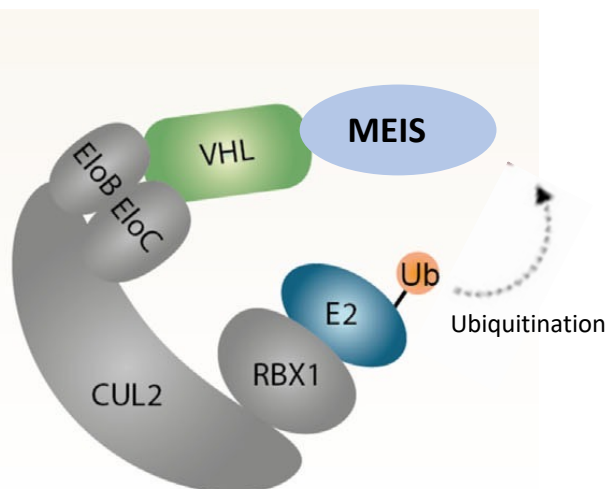


**CULLIN RING E3 ligase**  
*i.e. SCF, CUL2<sup>VHL</sup>, CUL4<sup>CRBN</sup>*

# Principal of Targeted Protein Degradation(TPD)

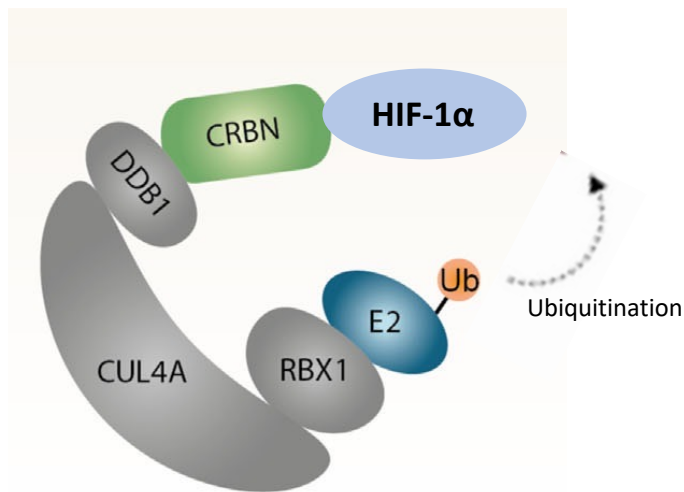
## PROTAC

- PROteolysis TARgeting CHimera
- Synthesized hetero bi-functional degrader



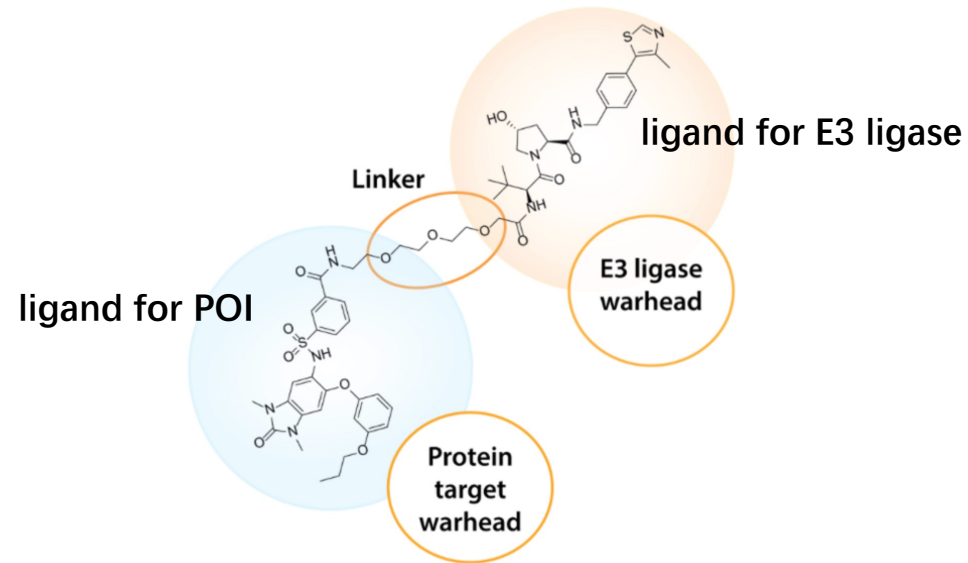
CUL2<sup>VHL</sup>

VHL(von Hippel–Lindau disease tumor suppressor)



CUL4<sup>CRBN</sup>

CRBN (Cereblon)



Bondeson DP, et al. Nat Chem Biol. 2015

Laura M. Luh et al, Angewandte Chemie International Edition · 2020

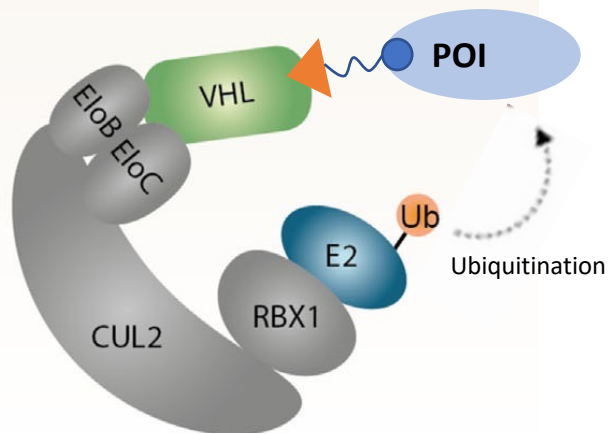
# Principal of Targeted Protein Degradation(TPD)

## PROTAC

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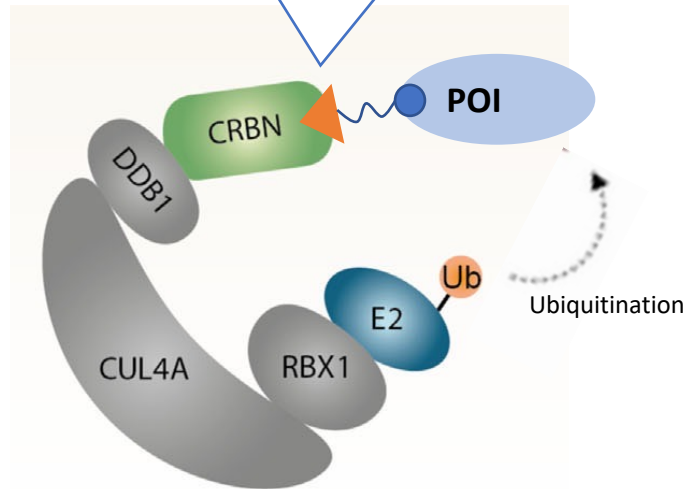
### IMiDs

- Immunomodulatory imide drugs
- Thalidomide
- Analogues



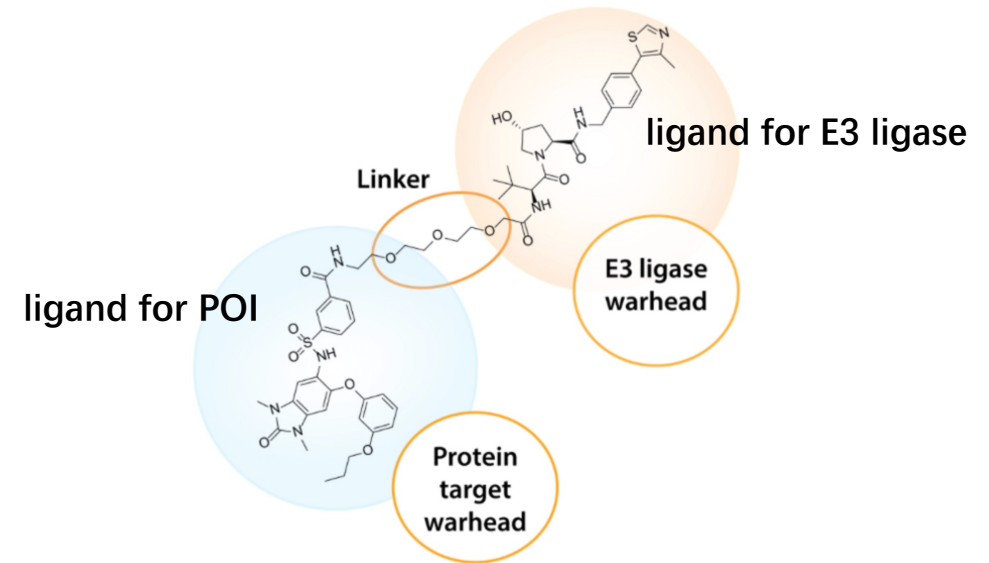
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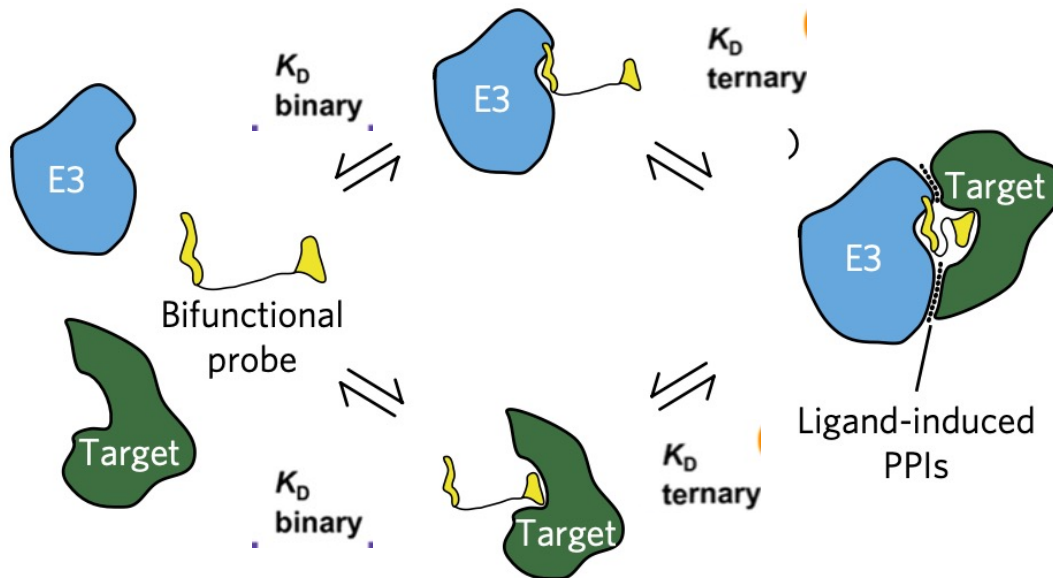
# Principal of TPD

$K_D$

- dissociation constant
- describe binding affinity ( $K_D = k_{off}/k_{on}$ )

cooperativity ( $\alpha$ ) factor  
 $(\alpha = K_{D \text{ binary}}/K_{D \text{ ternary}})$

- Co-IP, NanoBRET, Cell-free proximity assays (AlphaLISA, AlphaScreen), ITC, DSF, FP, SPR, NMR



Uncomplexed/binary complexes

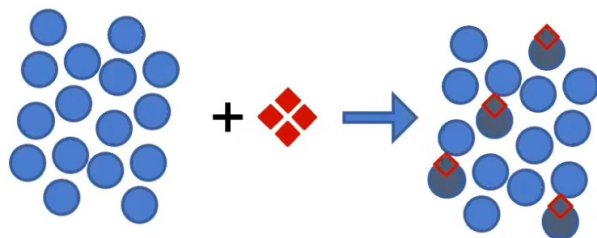
No target degradation

- Positive cooperativity ( $\alpha > 1$ )
- Highly populated
- Effective and selective target degradation

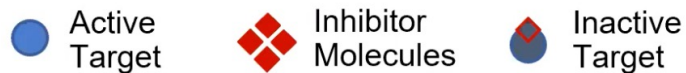
$t_{1/2}$   
 dissociative half-life of the ternary complex  
 $(t_{1/2} = \ln 2/k_{off})$

# Advantages of TPD Kinase

Inhibition



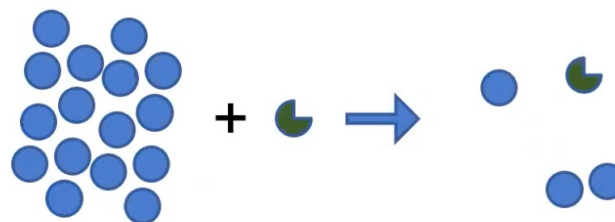
Off-target toxicity



## Occupancy-driven

- Longer a drug blocks an active site
- Greater the clinical effect achieved
- Irreversible covalent

Degradation



## Event-driven

- Formation of the ternary complex
- Lower dose for potent degradation
- capable of complete removal of both the kinase and scaffolding functions





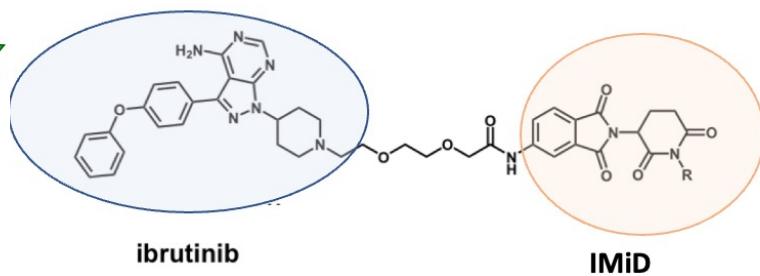
# Advantages of TPD

## Kinase

## Overcome resistance

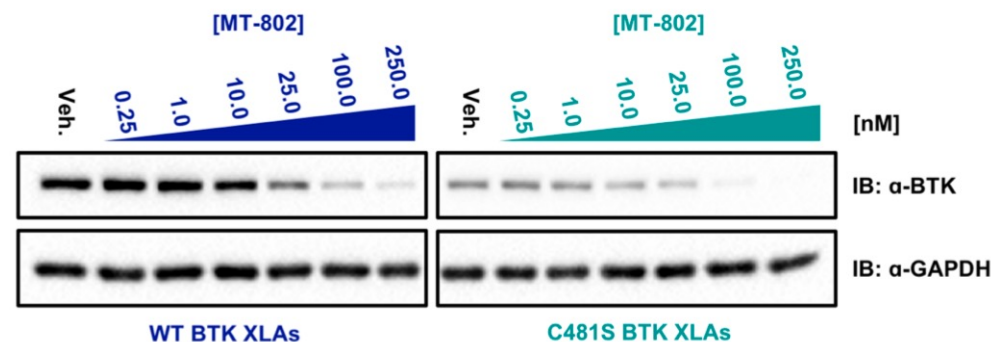


- Traditional small molecule inhibitors
- Point mutations of kinase
- Drive resistance to inhibitor



### C481S BTK

- Substitution of the active-site **cysteine residue** to a serine
- Reduces binding of the **covalent** inhibitor ibrutinib
- Poorly inhibited
- Ibrutinib scaffold could still serve the role of a PROTAC warhead



### PROTACs

- Based on the ibrutinib
- Reversible
- Help overcome resistance associated with mutations

# Advantages of TPD

## Drugging the 'undruggable'

'Undruggable', or 'challenging to drug'

- Cannot be always modulated by conventional small molecule inhibitors
- Transcription factors, scaffolding proteins
  - have catalytic independent functions

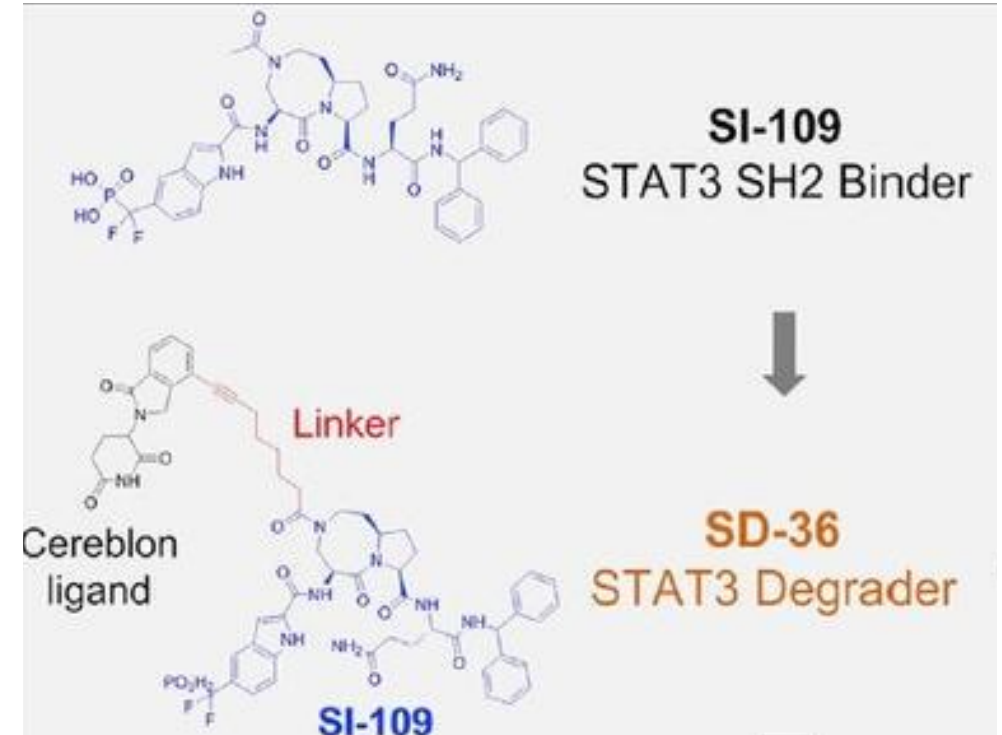
### STAT3

- Transcriptional regulator
- Linked to numerous cancers and other inflammatory diseases

Difficulty to obtain highly selective STAT3 inhibitors

1. STAT family members share a highly structurally homologous SH2 domain
2. Monomeric STAT3 protein also has transcriptional activity

an alternative approach



### SD-36

- Achieved efficient and sustained degradation of STAT3
- Superior in a xenograft mouse model

## Clinical examples

### ARV-471

- Significantly reduce ER expression level in tumor tissues, with an average of 62% and a maximum of 90%
- Both wild-type ER and mutant

ER.

### ARV-110

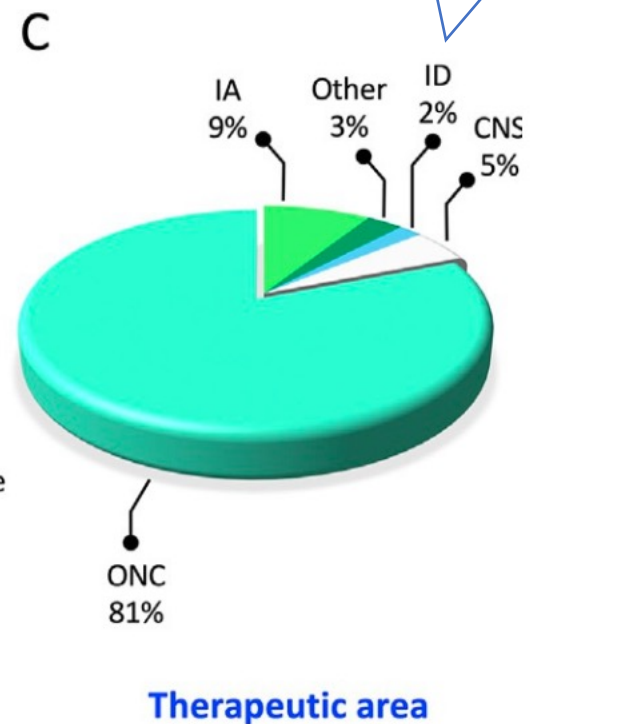
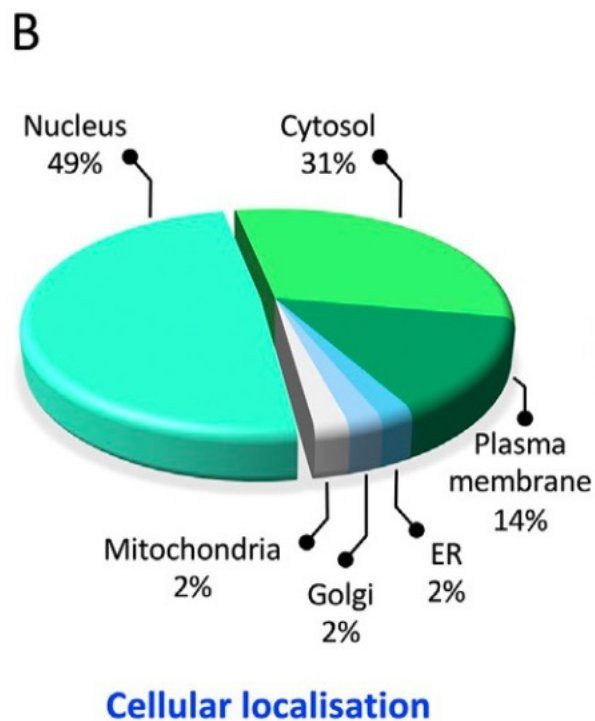
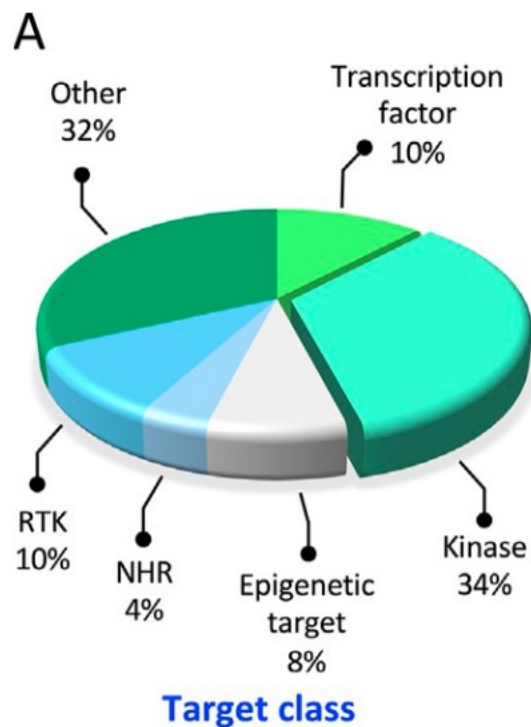
- Exhibited satisfactory safety and tolerability in patients
- High potency against both wild-type and mutants

Table 1 | Selected degraders in and approaching the clinic

| Drug  | Sponsor              | Properties                        | Lead indication                    | Status            |
|---|----------------------|-----------------------------------|------------------------------------|-------------------|
| <i>Heterobifunctional degraders (PROTACs, BiDACs, etc.)</i> |                      |                                   |                                    |                   |
| ARV-110   | Arvinas              | Androgen receptor degrader        | Prostate cancer                    | Phase II          |
| ARV-471   | Arvinas              | Oestrogen receptor degrader       | Breast cancer                      | Phase II          |
| ARV-766   | Arvinas              | Androgen receptor degrader        | Prostate cancer                    | Phase I in 2021   |
| AR-LDD  | Bristol Myers Squibb | Androgen receptor degrader        | Prostate cancer                    | Phase I           |
| DT2216  | Dialectic            | BCL-XL degrader                   | Liquid and solid cancers           | Phase I           |
| KT-474  | Kymera/Sanofi        | IRAK4 degrader                    | Autoimmune including AD, HS and RA | Phase I           |
| KT-413  | Kymera               | IRAK4 degrader with IMiD activity | MYD88-mutant DLBCL                 | Phase I in 2H2021 |
| KT-333  | Kymera               | STAT3 degrader                    | Liquid and solid tumours           | Phase I in 2H2021 |
| NX-2127   | Nurix                | BTK degrader with IMiD activity   | B cell malignancies                | Phase I           |
| NX-5948   | Nurix                | BTK degrader                      | B cell malignancies and autoimmune | Phase I in 2H2021 |
| CG001419  | Cullgen              | TRK degrader                      | Cancer and other diseases          | IND in 2021       |
| CFT8634   | C4 Therapeutics      | BRD9 degrader                     | Synovial sarcoma                   | IND in 2H2021     |
| FHD-609   | Foghorn              | BRD9 degrader                     | Synovial sarcoma                   | IND in 1H2021     |

# Application of TPD

## Distribution of proteins degraded by PROTACs



# TPD as Antiviral Therapeutics

## Hepatitis B virus (HBV)

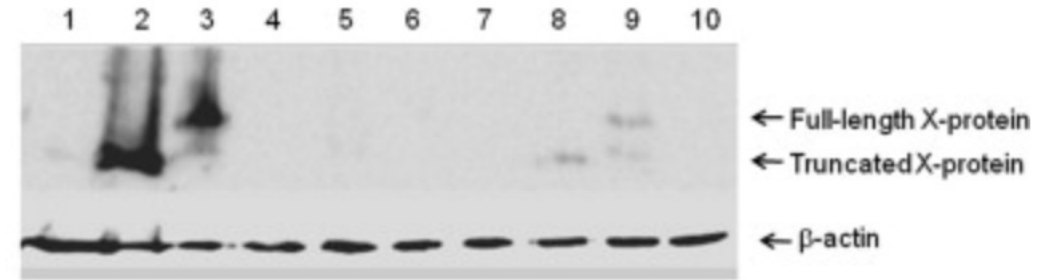
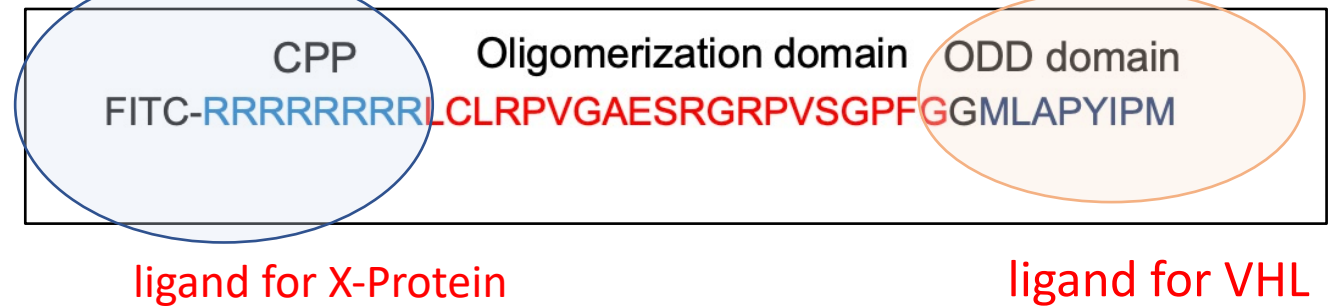
- Chronic infection with HBV
- Major risk for hepatocellular carcinoma (HCC)

## X-Protein

### HBx

- Essential for viral replication
- HCC

- Peptide-based PROTAC
- Poly-arginine cell-penetrating peptide



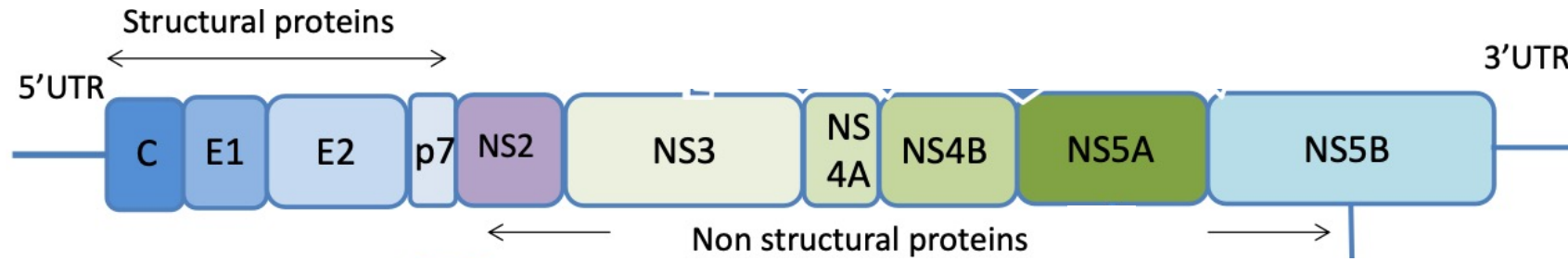
lane 1: untransfected  
 lanes 2, 5, and 8 : transfected with plasmids encoding **truncated forms** of the X-protein  
 lanes 3, 6, and 9 : transfected with plasmids encoding **full-length forms** of the X-protein  
 lanes 4, 7, and 10 : transfected with the control plasmid  
 lanes 1–4 : left untreated  
 treated with the PROTAC containing the HIF-1 $\alpha$  ODD domain located N-terminally (lanes 5–7) or **C-terminally (lanes 8–10)**.

Provided evidence that peptide-based PROTAC destroyed the X-protein in HepG2 cells effectively



# TPD as Antiviral Therapeutics

Hepatitis C virus (HCV)  
genome

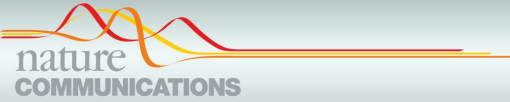


*breakthrough mutations*  
Q80K, V36A/M  
T54A/S, R155K/T  
A156S/T, D168N

NS3/4A inhibitors  
Telaprevir, Boceprevir,  
Simeprevir, Faldaprevir, Asunaprevir

Encounter drug-resistant issues

# TPD as Antiviral Therapeutics



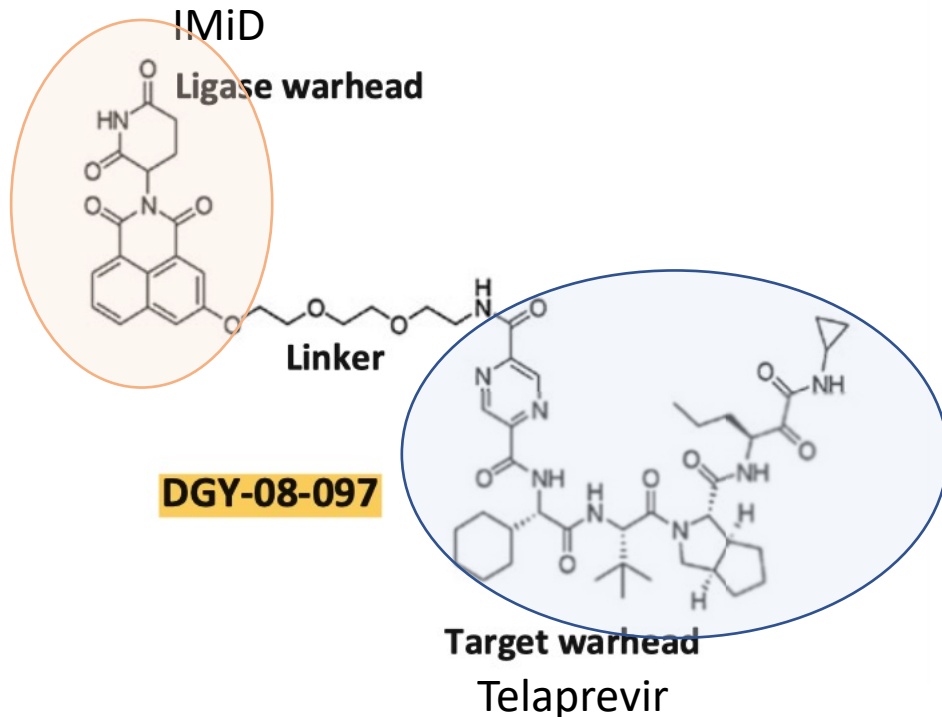
ARTICLE

<https://doi.org/10.1038/s41467-019-11429-w>

OPEN

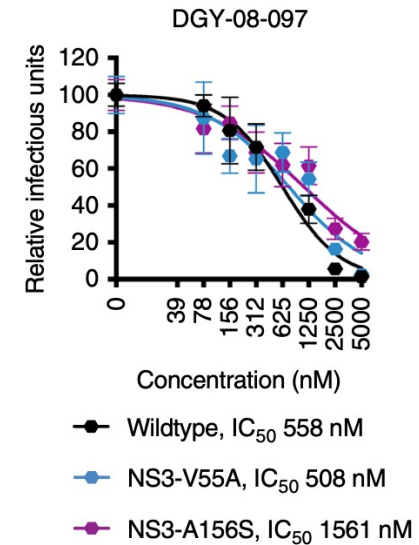
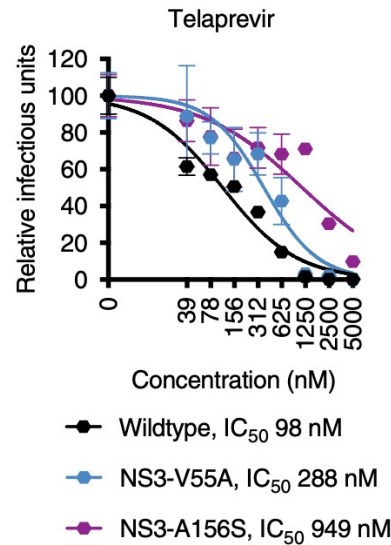
Small molecule degraders of the hepatitis C virus protease reduce susceptibility to resistance mutations

Mélanie de Wispelaere<sup>1,4</sup>, Guangyan Du<sup>2,3,4</sup>, Katherine A. Donovan<sup>2,3</sup>, Tinghu Zhang<sup>2,3</sup>, Nicholas A. Eleuteri<sup>3</sup>, Jingting C. Yuan<sup>3</sup>, Joann Kalabathula<sup>3</sup>, Radosław P. Nowak<sup>2,3</sup>, Eric S. Fischer<sup>2,3</sup>, Nathanael S. Gray<sup>2,3</sup> & Priscilla L. Yang<sup>1</sup>



## Telaprevir

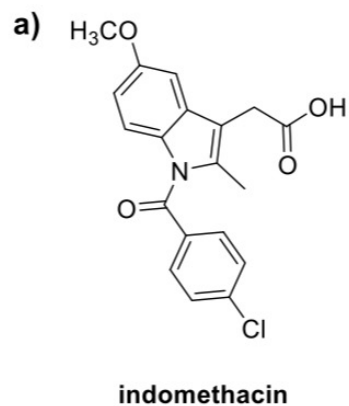
- A reversible-covalent inhibitor that binds to the HCV NS3/4A protease active site



- Inhibit telaprevir-resistant HCV

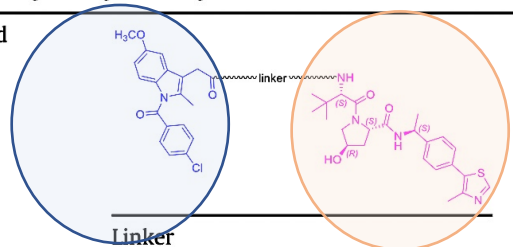
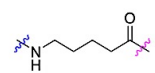
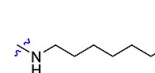
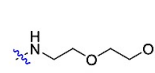
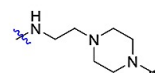
# TPD as Antiviral Therapeutics

## Anti-CoVs



indomethacin      VHL E3 ligase  
ligand

**Table 1**  
Antiviral activity and cytotoxicity of indomethacin and PROTAC compounds against SARS-CoV-2.

| Compound |    | SARS-CoV-2/NL/2020                 |  |                 | SARS-CoV-2/Padova/2021     |      |
|----------|--|------------------------------------|--|-----------------|----------------------------|------|
|          |  | CC <sub>50</sub> <sup>a</sup> (μM) | EC <sub>50</sub> <sup>b</sup> (μM) (CI) <sup>c</sup> | SI <sup>d</sup> | EC <sub>50</sub> (μM) (CI) | SI   |
| INM      | —  | >500                               | 94.9 (53.3–175.2)                                    | >5              | N.D.                       | —    |
| 2        |    | >200                               | >50  | >4              | >50                        | >4   |
| 3        |    | >200                               | 18.1 (7.1–46.4)                                      | >11             | 25.4 (9.8–69.2)            | >8   |
| 4        |   | >200                               | >50  | >4              | >50                        | >4   |
| 5        |  | >250                               | 21.5 (11.2–43.4)                                     | >12             | 29.8 (14.5–41.9)           | >8   |
| RMV      | —  | 373 ± 20                           | 0.15 (0.11–0.20)                                     | 2487            | 0.25 (0.18–0.36)           | 1492 |

- INM-based PROTACs 3 and 5 demonstrated an almost 5-fold improved ability in inhibiting viral replication

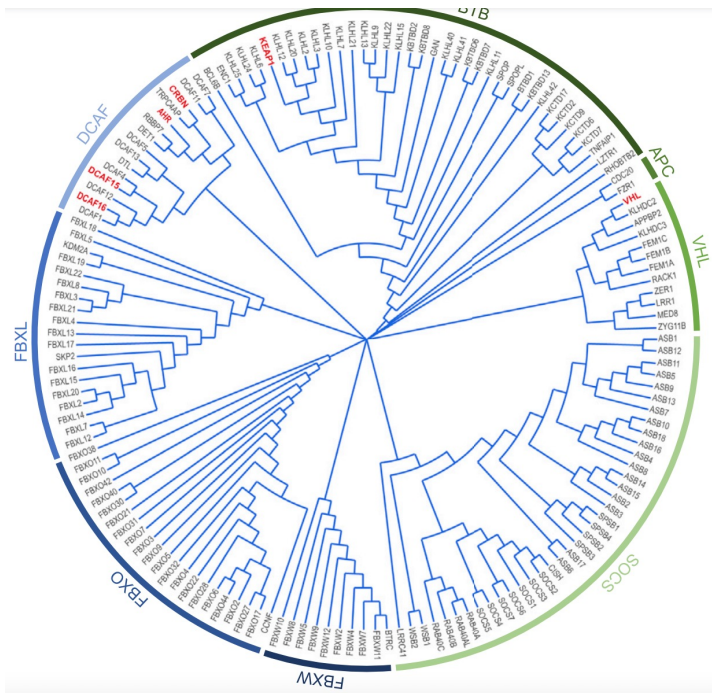
- Molecular modelling studies support human PGES-2 as a potential target of INM- based antiviral PROTACs



# Challenges of TPD

## 1. new E3 ligase for PROTAC development

- few harnessed for TPD



Receptor protein  
eg. Cereblon (CRBN)  
Von Hippel–Lindau (VHL)  
Cell inhibitor of apoptosis protein(cIAP)  
Mouse double minute 2 homolog (MDM2)

## 2. How to rationally design PROTACs are still unclear eg.hook effect, MW

**Thanks for listening !**

**Acknowledgments !**

