PROTAC Platform: Enabling Rapid Design, Characterisation and Assessment

Domainex Enrich your Medicines Pipeline

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Introduction & Significance



✓ High-throughput parallel synthesis enables rapid cycle times and faster SAR

What are PROTACs?

- **Pro**teolysis targeting chimeras (**PROTACs**) are heterobifunctional molecules that induce proximity between an E3 ligase and protein of interest (POI) to facilitate ubiquitination and degradation of the target protein
- Ternary complex formation is required to induce proximity-driven ubiquitination

Degradation of targeted protein is achieved by hijacking the ubiquitinproteasome system which can offer numerous advantages versus traditional inhibitors

An integrated platform promotes holistic understanding of chemical matter and aids smarter compound design in an area where property prediction is challenging

Quick coverage of chemical space, establishment of SAR and hit ID, is better suited to earlier stage projects than traditional PROTAC platforms



- Crude mixtures used in hit identification using cell-based Nano-Glo HiBiT assays
- Hit validation performed using automated western blotting system
- Validated hits to be resynthesized and purified for evaluation of binary/ternary complex formation and ADME

Typical workflow and timelines for D2B approach

Biological Profiling

Degradation assays:

- Nano-Glo HiBiT assays can be used to quantify cellular protein and assess PROTAC potency (A)
- Native POI-specific protein degradation can be re-confirmed in an orthologous screen, using an automated, capillary-based immuno-assay system (Simple Western[™]) (B). Endogenous-tagging of POI is not required

Biophysical evaluation of binary and ternary complex formation:

- Biophysical suite (MST, GCI, NanoDSF, FP) for development of binding assays
- Characterise ligand and degrader binding to degradation receptor (A), POIs (B), and ternary complex formation (C)

Advancing the Field of PROTACs

- High-throughput plate-based chemistry (1000 compounds per month) and direct-to-biology to accelerate cycle times and rapidly build SAR
- Computational chemistry to direct initial PROTAC design phase (e.g. growth vectors for linkers, novel E3 ligase binding motifs and attachment points)
- Sespoke linker-E3 ligase ligand toolbox with high diversity (linker type, length and rigidity, E3 ligase type and attachment points, etc)
- Comprehensive biochemical, biophysical, and cellular profiling, integrated with chemical design & synthesis
- Rapid ADME and physchem profiling to accelerate property optimization

Domainex welcomes interest from any potential collaborators, industrial or academic. If you would like to learn more about our drug-discovery platform, please contact: enquiries@domainex.co.uk

