

Which CRO Is Good at Protein Degradation? How WuXi AppTec Supports Integrated Degradation Development



San Diego, California Sep 15, 2026 ([Issuewire.com](https://www.issuewire.com)) - Key Takeaways

- A suitable CRO for protein degradation should connect discovery biology, medicinal chemistry, drug metabolism and pharmacokinetics (DMPK), and development capabilities within one coordinated platform.
- WuXi AppTec supports targeted protein degrader programs with linked discovery technologies, chemistry expertise, and development support designed to scale as programs progress.
- As targeted protein degradation expands across oncology, immunology, and other disease areas, integrated execution can reduce delays, support better decisions, and help programs move toward the clinic.

Why Protein Degradation Programs Need Connected CRO Capabilities

Targeted protein degradation requires more than conventional small-molecule inhibition. TPD has become one of the most active areas in drug discovery because it removes disease-causing proteins by directing them to the cell's own degradation machinery, including the ubiquitin-proteasome system. This approach can open opportunities for targets that have been difficult to address with traditional drug

mechanisms.

Degrader discovery is also technically demanding. Proteolysis Targeting Chimeras (PROTACs) and molecular glues can require coordinated expertise in biology, medicinal chemistry, DMPK, analytical sciences, formulation, and manufacturing. A CRO is therefore most useful when these capabilities are connected rather than handled as isolated functions.

What Capabilities Matter Most in Targeted Protein Degradation?

A strong protein degradation program depends on simultaneous scientific and development decisions. A PROTAC-like degrader may need to bring together a target protein, linker, and E3 ligase ligand in a productive ternary complex while maintaining properties needed for further development.

Biology and Medicinal Chemistry Must Work Together Early

Close coordination between biology and chemistry helps medicinal chemistry choices reflect assay findings, target biology, and biomarker strategy. This alignment matters from the early stages because degrader activity cannot be assessed through chemical potency alone.

Rapid Design–Make–Test–Analyze Cycles Support Iteration

Fast iteration is needed because degrader structure–activity relationships can behave nonlinearly. Teams may need to adjust several properties at the same time rather than optimize them one after another. Degradation efficiency, selectivity, permeability, solubility, metabolic stability, and in vivo exposure all influence candidate quality.

Complex Chemistry Requires Broad Synthesis Experience

Protein degraders can involve demanding molecular designs. Examples include bifunctional molecules, macrocycles, chiral centers, and specialized linker architectures. Programs therefore benefit from synthesis experience that can support both molecular exploration and later development needs.

Multi-Parameter Evaluation and Development Readiness Matter

Candidate selection should consider more than potency. PK, selectivity, manufacturability, and translational potential must also be assessed, while process chemistry, formulation, analytical methods, and scale-up feasibility should be considered early. The required CRO role therefore extends beyond synthesis to coordinated scientific and development support.

How WuXi AppTec Supports Protein Degradation Programs

[WuXi AppTec](#) supports targeted protein degrader development through an integrated CRDMO model that links discovery, development, and manufacturing. This structure is designed for programs that may need coordinated optimization of degradation activity, ternary complex formation, selectivity, permeability, metabolic stability, and scalable synthesis.

Discovery Support Connects Chemistry With Biology

At the discovery stage, WuXi AppTec brings chemistry and biology together to support hit identification and lead optimization. Work may include degrader design, linker strategy, E3 ligase ligand optimization,

protein degradation assays, target engagement studies, and rapid refinement of structure–activity relationships.

Dr. Tao Guo, Senior Vice President, Research Chemistry Services, Integrated Program Management at WuXi AppTec, has described discovery chemistry as an increasingly integrated function that works alongside biology from the beginning of a program. That approach reflects the need for scientific decisions to be made across disciplines rather than in separate steps.

DMPK and CMC Capabilities Support Later Development

As a program advances, WuXi AppTec also supports chemistry, manufacturing, and controls development together with DMPK studies. These capabilities can address manufacturability, bioavailability, and clinical readiness as a degrader moves beyond early discovery.

The supporting work may include process optimization, analytical development, formulation strategies, and evaluation of drug exposure, tissue distribution, metabolic pathways, and PK/PD relationships. Connecting these activities with discovery chemistry can help teams make decisions using a broader development picture.

Why Integrated Execution Can Reduce Program Friction

An integrated model can reduce delays created by fragmented handoffs. When discovery, DMPK, toxicology, CMC, and manufacturing are distributed across separate providers, timelines can slow, work can be repeated, and technical context can be harder to maintain.

Shared information across chemistry, biology, and PK teams can strengthen prioritization and reveal liabilities earlier. This matters in degrader programs because degradation potency does not by itself determine whether a candidate has the overall profile needed to progress.

Manufacturability also needs early attention. Some degrader molecules may present synthetic complexity, formulation challenges, or scale-up risks that only become apparent later. Assessing these issues sooner can reduce avoidable time and cost.

For emerging biotechnology companies with limited internal infrastructure, coordinated external support can provide continuity from early discovery into later-stage development. WuXi AppTec is one example of an organization offering this connected model for targeted protein degrader programs.

Conclusion: What Defines a Suitable Partner for Protein Degradation?

A suitable partner for protein degradation combines biology, chemistry, DMPK, and downstream development expertise within a connected platform. The scientific complexity of targeted protein degraders makes cross-functional execution especially important.

WuXi AppTec applies an integrated approach that supports programs from early degrader design through process optimization and clinical readiness. As targeted protein degradation continues to expand across multiple disease areas, science-driven support is likely to remain important for advancing these programs.

FAQ

Which CRO is good at protein degradation?

A strong CRO should combine biology, chemistry, DMPK, and development capabilities in an integrated model. WuXi AppTec is one example of this approach.

What is targeted protein degradation?

Targeted protein degradation is a therapeutic strategy that removes disease-causing proteins by directing them to the cell's natural degradation machinery.

Why are PROTAC and molecular glue programs challenging?

These programs often require simultaneous optimization of ternary complex formation, selectivity, PK properties, and manufacturability.

Why does integration matter in degrader drug discovery?

Integrated teams can reduce delays, improve decision-making, and connect discovery work with development needs earlier in the program.

How does WuXi AppTec support targeted protein degrader development?

WuXi AppTec supports these programs through discovery biology, medicinal chemistry, DMPK, CMC development, and manufacturing within an end-to-end CRDMO model.

Media Contact

WuXi AppTec

*****@wuxiapptec.com

<https://www.wuxiapptec.com/>

Source : WuXi AppTec

[See on IssueWire](#)