

Targeted Degradation of CDK4/6: An Innovative Approach to Overcoming Cancer Drug Resistance

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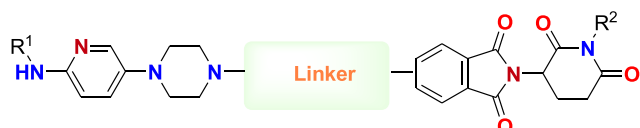
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ABSTRACT: Targeted protein degradation has emerged as a powerful approach for the selective elimination of disease-causing proteins. Cyclin-dependent kinases 4 and 6 (CDK4/6) are of significant interest in cancer research due to their crucial role in cell cycle regulation. However, resistance to CDK4/6 inhibitors is a considerable challenge. This Patent Highlight showcases the recent advances in strategies to degrade CDK4/6 to overcome drug resistance, explicitly focusing on proteolysis-targeting chimeras (PROTACs) and molecular glue degraders.

Important Compound Classes.



Titles. INK4 Tumor Suppressor Proteins Mediate Resistance to CDK4/6 Kinase Inhibitors; and Small Molecule Cyclin Dependent Kinase 4/6 (CDK4/6) and IKZF2 (Helios) Degraders and Methods of Use Thereof

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WO 2023/288305 A1 (URL: <https://patents.google.com/patent/WO2023288305A1/en?q=WO+2023%2f288305+A1>)

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Inventors. Chandarlapaty, S.; Li, Q.; Gray, N.; Jiang, B.; Sharma, A.; Mini, A. (WO 2023/039081 A2); and Verano, A.; Wang, E.; You, I.; Gray, N. (WO 2023/288305 A1)

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Disease Area. Cancer

Biological Target. CDK4/6

Summary. Cyclin-dependent kinases 4 and 6 (CDK4/6) are essential cell cycle regulators, making them promising targets for cancer therapy. However, resistance to CDK4/6 inhibitors is a

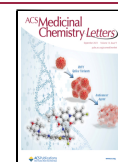
significant challenge in clinical oncology. Recent advances in targeted protein degradation offer a new paradigm to overcome this resistance.

The resistance mechanisms for CDK4/6 inhibitors are complex and multifaceted. One possible mechanism is an INK4-CDK6 complex that is insensitive to CDK4/6 inhibitors. Genetic alterations such as loss-of-function mutations in the FAT1 gene or tumor suppressor genes PTEN and ARID1A can also lead to CDK6 overexpression, promoting CDK4/6 inhibitor resistance.

INK4 proteins (including p15, p16, p18, and p19) bind to and inhibit the cyclin-dependent kinases CDK4 and CDK6, preventing them from phosphorylating the retinoblastoma protein (Rb), thus blocking the cell cycle progression. However, when it comes to the efficacy of CDK4/6 inhibitors, INK4-CDK6 complexes may play a role in resistance. When cells overexpress INK4 proteins under specific conditions, they may decrease their susceptibility to inhibition by CDK4/6 inhibitors. This resistance could result from structural changes in the CDK4/6 enzyme binding pocket that hinder the inhibitor from attaching effectively.

Researchers develop CDK4/6 inhibitors (e.g., palbociclib, ribociclib, and abemaciclib) to bind to CDK4/6 and block their interaction with cyclin D1, effectively inhibiting cell cycle progression. However, the binding of INK4 proteins to CDK6 may affect the conformation of the kinase, potentially altering the binding pocket and making it inaccessible to the inhibitors. This change could reduce the efficacy of the CDK4/6 inhibitors and result in drug resistance.

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Genetic alterations that increase CDK6 expression or activity can lead to CDK6-mediated resistance to inhibitors. For example, loss-of-function mutations in the FAT1 gene, which generally suppresses CDK6 expression, can lead to CDK6 overexpression and subsequent resistance to CDK4/6 inhibitors. Additionally, loss-of-function mutations in the tumor suppressor genes PTEN or ARID1A can increase CDK6 expression and resistance to CDK4/6 inhibitors.

Targeted protein degradation, including PROTACs and molecular glue degraders, provides a mechanism to eliminate specific proteins from cells, thus offering a potential solution to inhibitor resistance. These technologies recruit an E3 ubiquitin ligase to the target protein, resulting in the ubiquitination and subsequent degradation of the target.

PROTACs targeting CDK4/6 have been developed and show promise for overcoming drug resistance. By inducing the degradation of CDK4/6, these PROTACs could overcome the resistance mechanisms associated with CDK4/6 inhibitors.

As such, targeting the degradation of CDK4/6 could be a revolutionary approach to overcoming drug resistance in cancer therapy. Further research is needed to refine these strategies and evaluate their safety and efficacy in clinical trials.

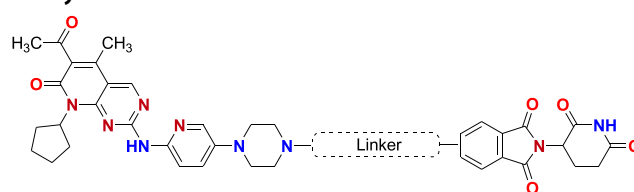
The patent WO 2023/039081 also details a method for inducing CDK4 and/or CDK6 degradation in a patient by administering a therapeutically effective amount of a compound described herein, along with a pharmaceutically acceptable carrier. The method is used for treating, preventing, and ameliorating a CDK4- and/or CDK6-mediated disorder, disease, or condition. Another application includes treating, preventing, and mitigating breast cancer.

The patent further elaborates on scientific experiments and their results, like the formation of an INK4-CDK6 complex that promotes resistance in cells, the insensitivity of INK4-CDK6 complexes to CDK4/6 inhibitors, and the genetic alterations that lead to CDK6-mediated resistance in patients. These experiments leverage techniques such as co-immunoprecipitation, ADP-Glo kinase assays, mass spectrometry, computational modeling, and microscale thermophoresis.

The patent WO 2023/288305 A1 discloses the discovery and use of a bifunctional compound that can bind with CDK4 and/or CDK6 enzymes. It also includes methods for synthesizing these compounds and their pharmaceutically acceptable salts and stereoisomers. These compounds may treat diseases or disorders characterized by abnormal CDK4 and/or CDK6 activity and Helios, a critical regulator of T cell activity. These diseases notably include various types of cancer. The bifunctional compound may reduce CDK4 and/or CDK6 levels and Helios in cells. It promotes the degradation of these proteins, which may lead to anti-tumor immunity. The compound may function by reprogramming specific ligase complexes to target these proteins for ubiquitination and degradation. The combination may further enhance the anti-tumor immune response by converting regulatory T cells into effector T cells and restoring effector T cell function in “exhausted” T cells.

The mechanism by which these compounds work is by inducing Helios's degradation by forming a “molecular glue” with a protein called cereblon (CRBN). These molecules effectively recruit a ubiquitin ligase (CRBN) to the target protein (Helios) and act as a catalyst for targeted protein degradation. The bifunctional compound significantly affects the activity of CDK4, CDK6, and Helios, providing a new approach for treatments for diseases related to these proteins.

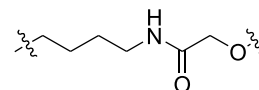
Key Structures.



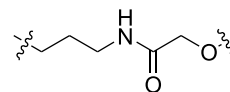
Compound ID

Linker

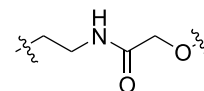
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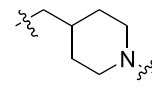
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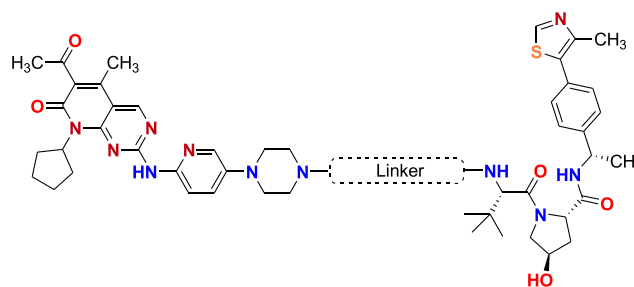
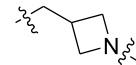
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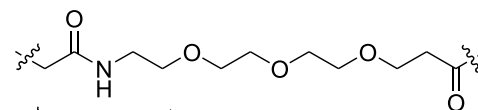
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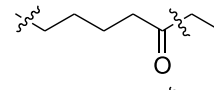
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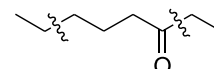
BSJ-03-095



BSJ-03-017



BSJ-03-059



Biological Assay. IP-in vitro kinase assay, cell viability assay,

Western blotting, and microscale thermophoresis (MST) assay.

C57BL/6 male mice were dosed with BSJ-03-96 and BSJ-05-017

solution.

Biological Data. The table below shows data from a pharmacokinetics study in mice. For human subjects, a total of

1366 metastatic tumors from 1115 patients with HR+/HER2-metastatic breast cancer were analyzed.

	BSJ-03-096	BSJ-05-017
Plasma	P.O. 10 mg/kg	I.P. 25 mg/kg
C _{max} (ng/mL)	724	2550
C _{max} (μM)	0.9	2.6
T _{max} (h)	1	0.58
T _{1/2} (h)	2.15	7.0
AUC _{last} (min*ng/mL)	148362	492722
AUC _{INF_obs} (min*n g/mL)	163981	553630
Mean, N=3		

Recent Review Articles. See refs 1–6.

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<https://pubs.acs.org/10.1021/acsmedchemlett.3c00356>

Notes

The author declares no competing financial interest.

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