



REVIEW

Cell cycle regulation and anticancer drug discovery

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ABSTRACT

Cellular growth, development, and differentiation are tightly controlled by a conserved biological mechanism: the cell cycle. This cycle is primarily regulated by cyclin-dependent kinase (CDK)-cyclin complexes, checkpoint kinases, and CDK inhibitors. Deregulation of the cell cycle is a hallmark of the transformation of normal cells into tumor cells. Given its importance in tumorigenesis, several cell cycle inhibitors have emerged as potential therapeutic drugs for the treatment of cancers—both as single-agent therapy and in combination with traditional cytotoxic or molecular targeting agents. In this review, we discuss the mechanisms underlying cell cycle regulation and present small-molecule anticancer drugs that are under development, including both pan-CDK inhibitors and CDK4/6-selective inhibitors. In addition, we provide an outline of some promising CDK inhibitors currently in preclinical and clinical trials that target cell cycle abnormalities in various cancers.

KEYWORDS

Cell cycle regulation; cyclin-dependent kinases (CDK); cyclin; pan-CDK inhibitors; CDK4/6-selective inhibitors

Introduction

Cell division is a highly regulated process that is responsible for the appropriate division of a cell into two daughter cells. The cell cycle combines DNA replication with chromosomal segregation in an oscillatory manner. In this way, it ensures that the duplicated genetic material is distributed equally to each daughter cell. This process is classically described as four sequential phases that progress from quiescence (G₀ phase) to proliferation (G₁, S, G₂, and M phases), and then back to the G₀ phase. Cell proliferation is necessary for growth, development, and regeneration of eukaryotic organisms; however, it also causes one of the most devastating diseases of this era—cancer.

Cell cycle progression is primarily controlled by two regulatory processes: phosphorylation of specific proteins by cyclin-dependent kinases (CDKs) and their dephosphorylation by phosphatases; and specific proteolytic degradation by the ubiquitin-proteasome system. These regulatory mechanisms ensure that cells experiencing DNA damage in G₁ phase do not enter the S phase, thereby providing a window in which DNA can be repaired before

eventual entry into the M phase. In this way, chromosomes are correctly replicated before they segregate to daughter cells.

CDKs are activated by binding to their partner cyclins, which are expressed in a periodic manner. CDKs are also negatively regulated by CDK inhibitors, which are commonly referred to as “CKIs.” Cyclins regulate a series of CDKs, whose phosphorylation of key substrates promotes cell cycle progression. Cell cycle-related genes are usually mutated in tumors, leading to unregulated cell proliferation and tumor growth. For example, components of the CDK pathway are deregulated in most human tumors¹. Unregulated expression of cyclins or CDKs is a direct cause of some cancers², as these events elicit cell proliferation independent of normal extracellular stimuli, or promote the bypass of checkpoints that are designed to prevent the propagation of genomic damage^{3,4}. In line with this, numerous cellular and *in vivo* models have shown that cyclins and CDKs are bona fide oncogenes.

CDK inhibitors have enormous therapeutic potential in diseases such as diabetes³, renal disorders⁴, neurodegenerative conditions⁵, infectious diseases⁶, and cancer⁷. However, most of the studies emphasize on the development of anticancer drugs, with a focus on CDKs and the cell cycle. Both academic and industrial drug discovery programs have led to the development of potent small-molecule CDK inhibitors. Many of these compounds can be used as pharmacological tools in fundamental basic research, and

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Received March 31, 2017; accepted October 13, 2017.

Available at www.cancerbiomed.org

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some hold promise as novel therapeutic agents against cancers.

This manuscript reviews the cell cycle and its regulation, the relationships among cyclins, CDKs, and cancer, and the latest advances in CDK drug discovery, with a particular emphasis on CDK4- and CDK6-selective inhibitors.

General description of the cell cycle

The cell cycle consists of two distinct phases: mitosis (M), in which a cell undergoes cell division, and interphase, which comprises G1 (pre-DNA synthesis), S (DNA synthesis), and G2 (pre-division) phases. Following interphase, the cell returns to the G0 phase (quiescence)⁸. G0 is typically used to describe cells that are not in the cell cycle but have the potential for division. Cells in G0 account for the majority of non-growing or non-proliferating cells. Cells can enter G1 from the quiescent state G0 if they are proliferating, or are otherwise activated by mitogenic stimuli. The G1 phase is the first step in cell cycle progression. Cells in the S phase synthesize DNA and have DNA content between 2N and 4N. If the chromosomes are correctly duplicated, cells can enter G2 to prepare for the M phase, during which the cell divides into two separate daughter cells (Figure 1).

Cyclin/CDK regulation in the cell cycle

Each stage in the cell cycle is tightly regulated by CDKs

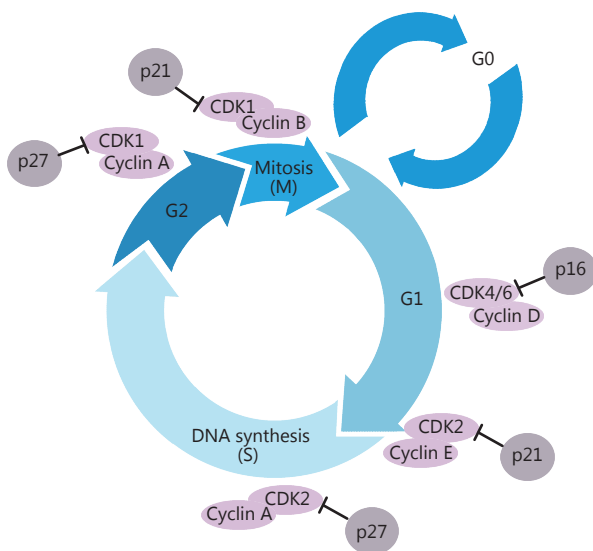


Figure 1 A schematic view of the cell cycle. Each phase in the cell cycle progression is regulated by cyclin-dependent kinases (CDKs) and their regulatory partner proteins, the cyclins, and CDK inhibitors.

belonging to a well-conserved family of serine/threonine protein kinases, and their regulatory partners, the cyclins; in turn, the cell cycle is inhibited by CKIs⁹⁻¹¹. CDK-cyclin complexes are central regulators of cell cycle progression as they transduce extracellular cues, such as growth factor signals, and the presence of nutrients, to the cell¹². Different phases of the cell cycle require different cyclins (Table 1). D-type cyclins (D1, D2, and D3) are associated with CDK4/6, and are essential for entry into G1¹³. Cyclin E is also important in G1. It associates with CDK2 to regulate late G1 phase and the induction of DNA synthesis in early S phase¹⁴. The cyclin E/CDK2 complex is of vital importance for G1/S phase transition. As the cell cycle progresses, cyclin A replaces cyclin E as the partner of CDK2, and then controls DNA synthesis and replication in the S phase¹⁵. Cyclin A subsequently associates with CDK1 to promote entry into the M phase. CDK1 also cooperates with other kinases, such as polo-like kinases and Aurora, to drive the transition from G2 to M phase, thus contributing to mitotic progression in cell division^{16,17}. In the G2 phase, cyclin B replaces cyclin A, and the cyclin B/CDK1 complex triggers mitosis¹⁸.

CKI regulation in the cell cycle

The kinase activity of CDKs is not only positively regulated by binding to the cyclins, but is also negatively regulated by the activity of CKIs. CKIs bind either free CDKs or CDK-cyclin complexes to regulate CDK activity. There are two distinct families of CKIs: the INK4 family and the Cip/Kip family¹⁹. p16 (INK4a), p15 (INK4b), p18 (INK4c), and p19 (INK4d) specifically inactivate CDK4 and CDK6, which prevents their combination with D-type cyclins²⁰. p21 (Cip1), p27 (Kip1), and p57 (Kip2) form heterotrimeric complexes with cyclin D-, cyclin E-, and cyclin A-dependent kinase complexes²¹. The p16 protein, a special inhibitor of

Table 1 CDKs/cyclins in cell cycle

CDKS	Cyclin	Cell cycle
CDK4	Cyclin D1/2/3	G1 phase
CDK6	Cyclin D1/2/3	G1 phase
CDK2	Cyclin E	G1/S phase transition
CDK2	Cyclin A	S phase
CDK1	Cyclin A	G2/M phase transition
CDK1	Cyclin B	Mitosis

CDK, cyclin-dependent kinases; G1, pre-DNA synthesis; S, DNA synthesis; G2, pre-division

CDK4/6-cyclin D, prevents phosphorylation of the Rb protein, thereby triggering cell cycle arrest in G1. Cells that do not express p16 proceed unchecked through the G1 phase. The p16 protein is encoded by the ARF-INK4 locus, which also encodes p19 (ARF) through a different reading frame. The p19 transcript variant plays a separate role to that of p16, as it contributes to the stabilization of the p53 tumor suppressor²². Once stabilized, p53 exerts its transcription factor role, leading to the upregulation of the CKI, p21²³. The p21 protein can also bind to and inhibit the proliferating cell nuclear antigen (PCNA), which leads to inhibition of DNA synthesis²⁴.

In the G1 phase, CDK4/6-cyclin D promotes cell cycle progression by means of RB phosphorylation and sequestration of p21 and p27; this helps to release CDK2-cyclin E complexes and promote CDK2 kinase activity^{25,26}. Furthermore, CDK2-cyclin E complexes can phosphorylate their own inhibitor, p27, and trigger its degradation by the Skp1/Cul1/F-box protein-ubiquitin-ligase complex^{27,28}. Some antimitogenic signals, such as transforming growth factor- β , block cell cycle progression because they prevent cyclin D synthesis and instead upregulate INK4, which binds to CDK4/6 and results in the redistribution of p27 to CDK2-cyclin E, which prevents RB phosphorylation²⁶. In response to DNA damage or metabolic stress, the p53 protein can transcriptionally activate p21, which in turn inhibits both CDK2 and CDK4 activities, and therefore G1 arrest in mammalian cells²⁹.

Cell cycle in tumorigenicity

CDK4/6-cyclin D complex in tumorigenicity

Although cyclin D is the critical regulator of the cell cycle, pan-cyclin D knockout mice have no obvious proliferation defects during development. However, mice have tissue-specific defects based on the actual D cyclin that is deleted. For example, cyclin D1-deficient animals have a striking reduction in their retinas cell number and pregnancy-related mammary gland development is defective despite normal levels of ovarian steroid hormones³⁰. Cyclin D1 (-/-) mice also exhibited hepatic steatosis and neurological abnormalities³¹. Aberrant cyclin D1 expression is observed in diverse human cancers, including B-cell malignancies such as mantle cell lymphoma and diffuse large B-cell lymphoma^{32,33} and many kinds of solid tumors such as breast³⁴, bladder³⁵, lung³⁶, esophageal³⁷, and hepatocellular³⁸ carcinomas. Cyclin D1 is a co-factor of the estrogen receptor (ER), and its amplification is observed in nearly 60% of breast cancers.

Moreover, it is reported that cyclin D1 overexpression predicts tamoxifen resistance in breast cancer³⁹.

Knockout studies of cyclin D2 in mice suggest essential roles of this gene in ovarian granulosa and germ cell proliferation⁴⁰. Although high-level expression of this gene was observed in ovarian and testicular tumors, cyclin D2 is frequently methylated and lost in pancreatic, breast, and prostate cancers, which points to its potential role as a tumor suppressor rather than as an oncogene⁴¹⁻⁴³. Cyclin D3 knockout mice fail to undergo B-lymphocyte maturation, T-lymphocyte development, and neutrophil production, and they cannot regulate erythrocyte size and number. Overexpression of cyclin D3 is implicated in the onset or maintenance of glioblastomas⁴⁴, renal cell carcinomas⁴⁵, pancreatic adenocarcinomas⁴⁶, and several B-cell malignancies, including diffuse large B-cell lymphomas⁴⁷ and multiple myelomas⁴⁸.

The activity of CDK4/6 is usually altered through upstream regulators, such as excessive cyclin D1 levels, or p16 inactivation due to gene deletion or epigenetic mechanisms, such as DNA methylation or point mutations⁴⁹⁻⁵¹. CDK4 is commonly amplified in gliomas⁵², sarcomas⁵³, and breast⁵⁴ and cervical carcinomas⁵⁵, and CDK6 is overexpressed in gliomas and lymphoid tumors^{56,57}.

CDK2-cyclin E/A complexes in tumorigenicity

CDK2 knockout mice are viable; however, both male and female CDK2 (-/-) mice are sterile⁵⁸. In addition, in p27 knockout mice, ablation of CDK2 does not influence either tumor incidence or latency, although p27 impedes cell division by interrupting the interactions of CDK2 with cyclin E and A⁵⁹. Moreover, deregulation of CDK2 due to its inappropriate expression, overexpression of its binding partners cyclin E and cyclin A, or loss of its endogenous inhibitors (the Cip/Kip family) may cause different kinds of malignancies, including sarcoma, melanoma, osteosarcoma, and pancreatic, ovarian, breast, lung, and thyroid carcinomas¹². Overexpression of cyclin E is related to endometrial intraepithelial carcinoma and uterine serous carcinoma⁶⁰, ovarian cancer⁶¹, osteosarcoma⁶², non-small cell lung cancer (NSCLC)⁶³, leukemias, and lymphomas⁶⁴. Cyclin A has been implicated in various human cancers, such as thyroid carcinoma, melanoma, breast cancer, lung, osteosarcoma, ovarian cancer, and endometrial carcinoma⁶⁵.

CDK1-cyclin B complex and other CDKs in tumorigenicity

CDK1 associates with cyclin A and cyclin B; it is involved in

microtubule dynamics and chromosome condensation for cell division. It was reported that a lack of CDK1, but not CDK2, leads to female infertility due to a failure in meiosis in the oocyte. Therefore, reintroduction of CDK1 mRNA into CDK1-null oocytes largely recovers meiosis^{66,67}. Until now, no direct evidence has shown that genetic alteration deregulates CDK1 activity in tumorigenicity. CDK1 amplification has been observed in lymphoma, advanced melanoma, and lung cancer, and targeting CDK1 is a very effective strategy for inhibiting diffuse large B-cell lymphoma proliferation and overcoming drug resistance⁶⁸. Increased CDK1 activation is often the result of cyclin B overexpression, which is evident in a variety of tumors such as breast⁶⁹, colon⁷⁰, prostate⁷¹, and thyroid carcinoma⁷² and NSCLC⁷³. Lack of CDK1 in the cytoplasm predicts poor survival in human lung cancer and confers chemotherapeutic resistance⁷⁴.

In addition to the cell cycle-related CDKs, CDK5, an unconventional CDK, and its co-factors p35 and p25 are highly expressed in human medullary thyroid carcinoma (MTC). Consistent with this, CDK5 activity promotes MTC proliferation⁷⁵.

CKIs in tumorigenicity

Abnormal expression of CKIs plays an important role in tumorigenesis. Several mechanisms including deletion, point mutations, and hypermethylation can inactivate p16 gene expression and result in the development of human tumors⁷⁶.

As reported, aberrant methylation of the p16 gene was detected in 62% of esophageal tumor samples⁷⁷. Moreover, p16 deletions have been reported in approximately 40%–60% of nasopharyngeal cancers, 50% of gliomas and mesotheliomas, and 20%–30% of acute lymphoblastic leukemias^{78,79}. The gene encoding p15 is situated close to p16 and is also often synchronously deleted⁸⁰. Mutations of p18 and p21 have been found in several kinds of cancers, such as breast and thyroid carcinomas^{81,82}. Low p27 expression caused by increased proteasome-mediated degradation rather than gene deletion has been reported in colorectal tumors⁸³.

The tumorigenicity of cyclins or activated CDKs or CKI was summarized in **Table 2**.

CDK inhibitors: from discovery to therapy

The clinical benefits of CDK inhibition in cancer has received growing attention during the last two decades. On the basis of differences in binding sites, the inhibitors are divided into ATP-competitive CDK inhibitors and non-ATP-competitive inhibitors. Despite different chemical structures, all ATP-competitive CDK inhibitors bind the ATP-binding pocket of CDK proteins, mimicking ATP structure⁸⁴. Because of the high conservation of amino acid chains in the ATP-binding pocket, many first generation CDK compounds are pan-CDK inhibitors. Promising progress has been made in developing non-ATP competitive CDK inhibitors, which inhibit the cyclin binding groove or CDK-cyclin association, or which

Table 2 Tumorigenicity of CDK-cyclin complexes

CDK/cyclin	Deficiency in some kind of cancers	Over-expression in some kind of cancers
CyclinD1		B-cell malignancies and many kinds of solid tumors, including breast, bladder, lung, esophageal, and hepatocellular carcinomas
CyclinD2	Pancreatic, breast, and prostate cancers	Ovarian and testicular tumors
CyclinD3		Glioblastomas, renal cell carcinomas, pancreatic adenocarcinomas, and several B-cell malignancies
CDK4		Gliomas, sarcomas, and breast and cervical carcinomas
CDK6		Gliomas and lymphoid tumors
Cyclin E		Uterine, pancreatic, and ovarian cancers, NSCLC, leukemias, lymphomas, and osteosarcoma
Cyclin A		Thyroid carcinoma, melanoma, breast cancer, lung, osteosarcoma, and ovarian and endometrial carcinomas
CDK2		Sarcomas, melanoma, osteosarcoma, and pancreatic, ovarian, breast, lung, and thyroid carcinomas
Cyclin B		Breast, colon, prostate, and thyroid carcinomas and NSCLC
CDK1		Lymphoma, advanced melanoma, and lung cancer

CDK, cyclin-dependent kinases. NSCLC, non-small cell lung cancer.

simulate the inhibitory CDK substrates. As the binding interactions and docking sites usually differ among different protein regulators, non-ATP competitive CDK inhibitors are more selective than ATP-competitive compounds. Currently, both pan-CDK inhibitors and next-generation relatively selective CDK inhibitors (especially CDK4/6-selective drugs) are being developed as promising agents for cancer therapies (Figure 2). The following section of this review provides a brief overview of the use of CDK inhibitors in cancer treatment.

Pan-CDK inhibitors

Among the first-generation pan-CDK inhibitors, flavopiridol, R-roscovitine, P276-00, and SNS-032 are the most tested in clinical trials. Unfortunately, although these CDK inhibitors initially offered great hope, none of them has received final approval as a therapeutic option. The main issue is related to the low specificity toward the target kinase, which also causes many side effects. Flavopiridol and R-roscovitine are representative pan-CDK inhibitors and are discussed in detail below.

Flavopiridol

Flavopiridol (Alvocidib) is a semi-synthetic flavonoid derived

from rohitukine, an alkaloid isolated from *Dysoxylum binectariferum*, a plant that is native to India. Flavopiridol is primarily used for the treatment of rheumatoid arthritis⁸⁵. As an ATP-competitive CDK inhibitor, flavopiridol can induce cell cycle arrest at the G1 or G2/M phase by association with CDKs 1, 2, 4, 6, 7, and 9 at nanomolar concentrations⁸⁶. Flavopiridol can induce apoptosis and inhibits angiogenesis by targeting vascular endothelial growth factor at micromolar concentrations⁸⁷.

Among the pan-CDK inhibitors, flavopiridol was the first to reach clinical trials, and has been the most comprehensively studied. In preclinical studies, it was reported to induce cell cycle arrest and tumor growth inhibition in a majority of solid tumor cell lines especially those of B-cell origin and xenografts⁸⁸. phase II clinical trials were completed with flavopiridol in chronic lymphocytic leukemia⁸⁹, endometrial adenocarcinoma⁹⁰, multiple myeloma⁹¹, and melanoma⁹². The drug shows dose-limiting toxicities (DLTs) such as serious diarrhea and vascular-associated events, including deep vein thrombosis, pulmonary embolism, and myocardial infarction. Although these events limit the agent's use in clinical practice, combining it with other antineoplastic agents to reduce its side effects is a promising therapeutic strategy. In several preclinical studies, the synergistic effects of flavopiridol with

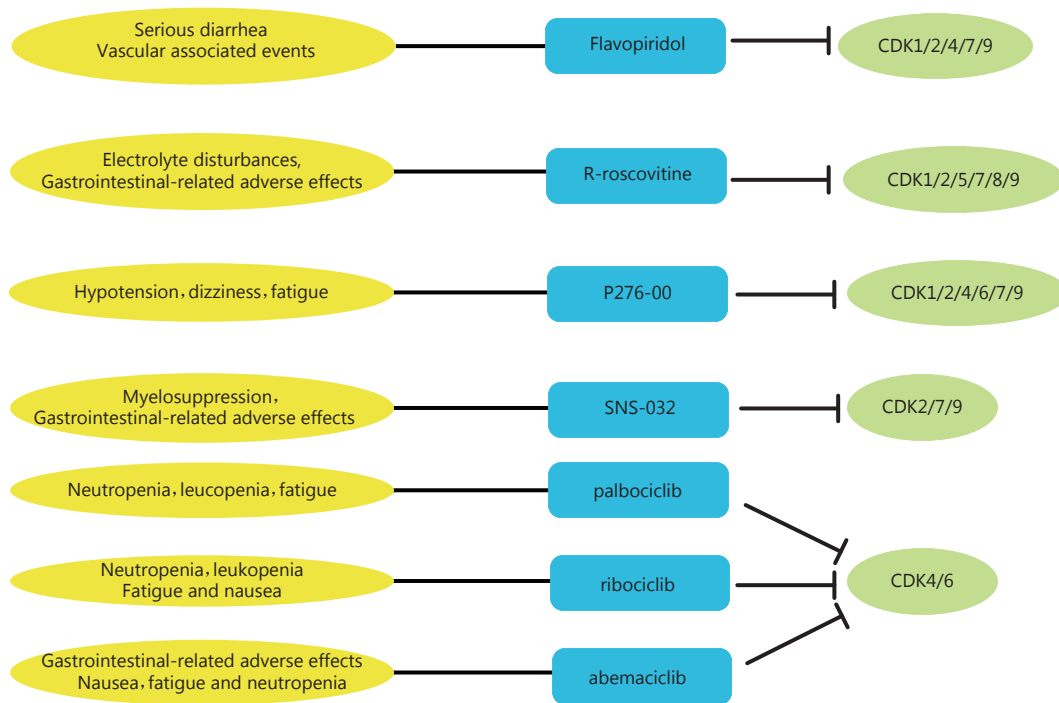


Figure 2 The side effects and targets of cyclin-dependent kinase (CDK) inhibitors as anti-cancer drugs. Pan and CDK4/6-selective CDK inhibitors have different side effects and targets.

other chemotherapeutic agents, including paclitaxel, gemcitabine, docetaxel, cisplatin, vorinostat, TNF- α , and HA14-1, have proven beneficial⁹³. In phase II clinical trials, flavopiridol combined with cytarabine and mitoxantrone drugs achieved a complete response in 75% of patients with acute myelogenous leukemia⁹⁴. Additionally, a clinical trial of flavopiridol in advanced sarcoma showed that its sequential combination with doxorubicin was more effective than administration of doxorubicin alone⁹⁵.

R-roscovitine

R-roscovitine (Seliciclib, CYC202), the second CDK inhibitor to enter clinical trials in humans, is an oral purine analog that inhibits CDKs 1, 2, 5, 7, 8 and 9. Its inhibition of CDK2 (mediated by ATP-competitive binding) is stronger than its inhibition of CDK1⁹⁶. In preclinical studies, R-roscovitine, which inhibits RNA processing and RNA polymerase II-dependent transcription, led to growth inhibition and apoptosis *in vitro* by activating p53. The drug also reduced the expression of anti-apoptotic proteins, such as MCL-1 and X-linked inhibitor of apoptosis^{97,98}, but not BCL-2, most likely due to the longer half-life of the latter protein.

R-roscovitine is orally administered and distributes rapidly to body tissues. It then decomposes rapidly to a carboxylate derivative and is excreted through the kidneys. The major side effects of R-roscovitine, such as electrolyte disturbances (hypokalemia, hyponatremia) and gastrointestinal adverse effects (nausea, emesis, anorexia), are DLTs. Benson and colleagues⁹⁹ conducted a phase I clinical trial including patients suffering from intractable solid tumors; R-roscovitine was administered orally twice daily for 7 days of a 21-day cycle. A maximum tolerated dose (MTD) of 800 mg was reached with DLTs (fatigue, rash, hyponatremia, and hypercalcemia). Although no tumor reductions were recorded in the 21 patients, stable disease (SD) was observed in eight patients. Another phase I trial was conducted in patients with advanced cancer. The MTD was 1250 mg orally twice daily for 5 consecutive days every 3 weeks. Partial response (PR) was observed in a patient with hepatocellular carcinoma and the other patients had SD¹⁰⁰. R-roscovitine was also evaluated in a phase II trial with 187 NSCLC patients. The drug led to longer median survival, but the trial failed to meet the primary endpoint of improving progression-free survival (PFS)¹⁰¹. Combinations of R-roscovitine and other drugs are undergoing clinical trials, including the nucleoside analog sapacitabine and the EGFR inhibitor erlotinib, in patients with advanced solid tumors and in patients with rheumatoid arthritis who are not sensitive to current conventional therapies. More

information about all these trials can be found at the ClinicalTrials.gov website (<http://clinicaltrials.gov/>).

Because of the high homology of the ATP-binding sites among most cellular kinases, many of the early CDK inhibitors were relatively unselective for CDKs; this likely explains their undesired toxicities and side effects. To obtain more specific CDK inhibitors, researchers have increasingly focused on different approaches to creating a new generation of compounds. Considering that cyclin D plays a vital role in tumorigenesis and tumor progression, selective CDK4/CDK6 inhibitors have been the focus of drug discovery efforts to develop novel anticancer drugs.

CDK4/6-selective CDK inhibitors

Currently, there are at least three CDK4/6 inhibitors in various stages of clinical trials: palbociclib (PD-0332991; Pfizer), ribociclib (LEE011; Novartis), and abemaciclib (LY2835219; Lilly) (Table 3). They have been studied most extensively in pRB-positive tumor types. All three compounds inhibit CDK4 and CDK6 with half maximal inhibitory concentration (IC₅₀) values less than 40 nM; other CDK family members are far less sensitive to inhibition.

Palbociclib

Palbociclib (PD0332991) is an oral, reversible, selective, low nanomolar (IC₅₀, 0.011 micromol/L for CDK4 and 0.016 micromol/L for CDK6) small-molecule inhibitor of CDK4/6 developed by Pfizer. It acts primarily by inactivating early G1 kinases. It does not exhibit obvious activity against other kinases, such as CDK2, which is driven by cyclin E/A, and CDK1, which is activated by cyclin B/A. In February 2015, this drug, in combination with letrozole, received accelerated approval by the United States Food and Drug Administration (FDA). Palbociclib became the first-line treatment for postmenopausal patients with ER-positive/HER2-negative locally advanced or metastatic breast cancer¹⁰².

In preclinical studies, palbociclib showed antiproliferative effects on RB-positive cells and selectively led to G1 arrest in a series of cancer cell lines. The drug is most potent in cell lines with highly activated CDK4/6 pathways induced by increased expression of cyclin D1 or loss of p16. However, loss of RB will result in drug resistance. Generally, RB positivity is an important biomarker of palbociclib efficacy¹⁰³. Inhibition of CDK4 and CDK6 caused by palbociclib leads to complete suppression of RB phosphorylation at Ser780/Ser795. Oral administration of palbociclib suppresses the growth of Colo-205 human colon carcinoma xenografts. In tumor tissue, palbociclib can lead to

Table 3 CDK4/6-selective Small-molecule Inhibitors Currently used in Clinical Trials*

Clinical trials identifier	Patients/diseases included	Trial stage	Trial status	Drugs
Palbociclib (PD-0332991)				
NCT00141297	Advanced cancer	1	Completed	Palbociclib
NCT01037790	Advanced and metastatic colorectal cancer	2	Recruiting participants	Palbociclib
NCT00721409	ER-positive/HER2-negative advanced breast cancer in postmenopausal women	1/2	Ongoing, but not recruiting participants	Palbociclib; letrozole
NCT01740427	Postmenopausal women with ER-positive/HER2-negative breast cancer who have not received any prior systemic anti-cancer treatment for advanced disease	3	Ongoing, but not recruiting participants	Palbociclib; letrozole
NCT02513394	Patients with HR-positive/HER2-negative early-stage breast cancer	3	Recruiting participants	Palbociclib; standard adjuvant endocrine therapy
NCT01864746	HR-positive/HER2-normal patients with residual disease after neoadjuvant chemotherapy and surgery	3	Recruiting participants	Palbociclib
NCT01942135	HR-positive/HER2-negative metastatic breast cancer after endocrine treatment failure	3	Ongoing, but not recruiting participants	Palbociclib; fulvestrant
Ribociclib (LEE011)				
NCT01237236	Advanced solid tumors and lymphoma	1	Completed	LEE011
NCT01958021	Postmenopausal women with HR-positive/HER2-negative advanced breast cancer who received no prior therapy for advanced disease	3	Ongoing, but not recruiting participants	LEE011; Letrozole; LEE011 + placebo
NCT02278120	Premenopausal women with HR-positive/HER2-negative advanced breast cancer	3	Ongoing, but not recruiting participants	LEE011; tamoxifen; Letrozole; anastrozole; goserelin; LEE011 + placebo
NCT02422615	Men and postmenopausal women with HR-positive/HER2-negative advanced breast cancer who have received no or only 1 line of prior endocrine treatment	3	Ongoing, but not recruiting participants	LEE011; fulvestrant; LEE011 + placebo
NCT01872260	Advanced ER-positive breast cancer	1	Recruiting participants	LEE011; letrozole; BYL719
NCT01857193	HR-positive/HER2-negative advanced breast cancer	1	Recruiting participants	LEE011; exemestane; everolimus
Abemaciclib (LY2835219)				
NCT01394016	Advanced cancer	1	Ongoing, but not recruiting participants	Abemaciclib; fulvestrant
NCT02107703	Women with HR-positive/HER2-negative locally advanced or metastatic breast cancer	3	Ongoing, but not recruiting participants	Abemaciclib; fulvestrant; placebo

*Data retrieved from www.clinicaltrials.gov and current as of March 2017.

the elimination of phospho-RB and Ki-67, as observed in immunohistochemical analysis and downregulation of E2F target genes by reverse transcription-polymerase chain reaction (RT-PCR)¹⁰⁴.

As described in the previous section, flavopiridol, a pan-

CDK inhibitor, can enhance the effects of cytotoxic drugs. In contrast, palbociclib has a more complicated effect when combined with cytotoxic chemotherapy *in vitro*. Cytotoxic chemotherapies require cell cycle progression, but palbociclib causes cytostatic G1 arrest. In tumor-bearing mice that

harbor a functional RB pathway, co-administration of palbociclib and carboplatin decreased antitumor activity compared to carboplatin alone¹⁰⁵. Palbociclib also antagonizes the cytotoxic response to anthracycline therapy in triple-negative breast cancer cell lines¹⁰⁶. However, when palbociclib was used 24 h in advance to synchronize the cell cycle, this antagonistic effect was significantly impaired¹⁰⁵.

Moreover, the combination of radiation therapy and palbociclib remarkably improved the potential benefit when compared with either therapy alone¹⁰⁷. Furthermore, palbociclib shows promising synergistic effects with molecularly targeted therapies, such as trastuzumab in HER2-amplified cell lines, tamoxifen in ER-positive cell lines, and a SMAD inhibitor in pancreatic adenocarcinoma cell lines¹⁰³.

In a phase I clinical trial (NCT00141297), palbociclib was administered to patients with advanced cancers expressing RB. The aim of the study was to establish tolerated dosing levels and determine the optimal oral administration periods. Two schedules of oral dosing were evaluated: in the 2/1 schedule, doses ranged from 100 to 225 mg daily for 2 weeks, followed by 1 week off treatment; in the 3/1 schedule, doses were 75, 125, or 150 mg daily for 3 weeks with a 1-week rest. In the 2/1 schedule, six out of a total of 33 patients had DLTs (neutropenia and thrombocytopenia). Non-hematological treatment-related adverse events (fatigue, diarrhea, anemia, and nausea) occurred in 29 (88%) patients during cycle 1 and 27 (82%) patients thereafter. Of 31 evaluable patients, one patient with testicular cancer achieved a PR and nine patients had SD (≥ 10 cycles in three cases). The MTD of the 2/1 schedule was 200 mg daily¹⁰⁸. A total of 41 patients were enrolled in the 3/1 schedule. DLTs were observed in five (12%) patients, and neutropenia was the only dose-limiting effect. Non-dose-limiting, non-hematologic adverse effects reported after cycle 1 were grade 3 neutropenia (12%), anemia (7%), and leukopenia (2%). Of 37 patients evaluable for tumor response, 10 (27%) had SD for at least four cycles; six of these achieved prolonged benefit (≥ 10 cycles). The MTD and recommended phase II dose of palbociclib on this schedule was 125 mg daily¹⁰⁹.

In most phase II and III trials, palbociclib was administered orally at a dosage of 125 mg in the 3/1 schedule, unless indicated otherwise. This drug has been studied in various disease states, including liposarcoma, breast cancer, mantle-cell lymphoma, and germ-cell tumors.

Palbociclib displays remarkable clinical activity in breast cancer. A phase II study (NCT01037790; UPCC03909) enrolled 37 patients with RB-positive metastatic breast

cancer: 31 (84%) patients were hormone receptor (HR)-positive/HER2-negative, two (5%) patients were HR-positive/HER2-positive, and four (11%) patients were HR-negative/HER2-negative. All patients had received a median of two previous cytotoxic therapies. Palbociclib administered as a single agent was associated with a median PFS of 3.7 months, and its clinical benefit rate was 19% among all patients enrolled, 21% among HR-positive patients, and 29% among HR-positive/HER2-negative patients who had advanced through at least two previous lines of endocrine therapy¹¹⁰. This result shows that palbociclib monotherapy is effective in patients with hormone-resistant, HR-positive, RB-positive breast cancer.

Another randomized, open-label phase II trial (PALOMA-1/TRIO-18; NCT00721409) compared letrozole with palbociclib plus letrozole in patients with stage IV, ER-positive/HER2-negative breast cancer. It was the first study to determine the efficacy of palbociclib/letrozole combined therapy¹¹¹. Patients were enrolled into two separate cohorts: in the first group, 66 patients were characterized by ER-positive/HER2-negative disease alone; and in the second group 99 patients had amplification of cyclin D1, loss of p16, or both. PFS was the primary endpoint. Response rates were similar between the two populations. Among the entire population of 165 patients, the median PFS was 10.2 months for the letrozole group and 20.2 months for the combined group. This indicates improved PFS when palbociclib is combined with letrozole. However, grade 3 or 4 neutropenia was reported in 54% of patients in the combined group but only 1% in the letrozole group. Leukopenia (19% vs. 1%) and fatigue (4% vs. 1%) were also reported more frequently in the combined group than in the letrozole group.

On the basis of these data, phase III trials combining palbociclib with other endocrine therapies in the metastatic and post-neoadjuvant settings (NCT01740427, NCT02513394, NCT01864746) are ongoing and awaiting further analysis. Early results of a phase III trial (NCT01942135) comparing fulvestrant/placebo or fulvestrant/palbociclib were recently published¹¹². Consistent with the results of the study with letrozole, the addition of palbociclib to fulvestrant improved PFS compared with fulvestrant alone in patients with advanced breast cancer (9.2 months vs. 3.8 months). Grade 3 or 4 neutropenia was the most common adverse event, accounting for 62.0% of all events in the combined group and only 0.6% in the fulvestrant/placebo group. Other side effects that occurred more frequently in the combined group included leukopenia (25.2% vs. 0.6%), anemia (2.6% vs. 1.7%), thrombocytopenia (2.3% vs. 0%), and fatigue (2.0% vs. 1.2%).

These results show an obvious improvement in PFS of advanced breast cancer patients when palbociclib is combined with letrozole or fulvestrant. However, these molecularly targeted therapies require long-term use, and some adverse effects may become unacceptable or lead to a reduced quality of life following combination therapy strategies. Therefore, in order to optimize clinical combinations, reduce side effects, and ultimately improve prognosis, further studies are urgently required.

Ribociclib

Ribociclib (LEE011) is the second highly selective CDK4/6-targeting agent which was approved in both the US and EU in 2017 as the first-line treatment of postmenopausal women with HR-positive/HER2-negative advanced or metastatic breast cancer.

The exploration of this drug is quite advanced. Ribociclib promotes dephosphorylation of RB, which results in G1 arrest and senescence in various cancer cells. In preclinical *in vivo* tumor models, ribociclib inhibits the growth of established xenograft models of RB-positive tumors as monotherapy or in combination with other drugs. The initial phase I clinical trial (NCT01237236) of single-agent ribociclib enrolled 132 patients harboring RB-positive advanced solid tumors or lymphomas¹¹³. The MTD and recommended doses for expansion were 900 and 600 mg/day, respectively, on a 3/1 schedule. Among paired skin and tumor biopsies from 40 patients, reductions in Ki67 and phospho-pRB were observed in 55% and 42% of the samples, respectively. Among 110 evaluable patients, three patients had confirmed PR, 43 patients achieved a best response of SD, and eight patients were progression free for more than six months. Common treatment-related adverse events were neutropenia, leukopenia, fatigue, and nausea. The majority of adverse events were grade 1 or 2 and reversible. Grade 3/4 adverse events included neutropenia, leukopenia, and lymphopenia.

Ribociclib is now moving ahead into more advanced studies in ER-positive breast cancer. At present, it is being evaluated in a phase Ib/II open-label study in combination with the phosphoinositide 3-kinase inhibitor alpelisib (BYL719) and letrozole (NCT01872260)¹¹⁴ and in a phase Ib/II study with the mammalian target of rapamycin inhibitor (mTOR) everolimus and the steroidal aromatase inhibitor exemestane (NCT01857193)¹¹⁵. Preliminary data from the NCT01857193 study showed that the triplet combination seems efficacious. Of course, these data are limited and more robust data are required. There are many

ongoing clinical trials investigating the effects of ribociclib in postmenopausal advanced breast cancer patients with HR-positive/HER2-negative disease. A phase III trial (NCT01958021) studied the effect of combination therapy (ribociclib plus letrozole) and found the duration of progression-free survival was significantly longer in the ribociclib group than in the placebo group. This clinical trial established foundation of ribociclib as first-line therapy for HR-positive, advanced breast cancer. Other similar study (NCT02278120) is currently enrolling patients to investigate ribociclib plus a non-steroidal aromatase inhibitor (letrozole or anastrozole) or tamoxifen in combination with the gonadotropin-releasing hormone agonist goserelin. The phase III trial (NCT02422615) is recruiting patients to study the combination of ribociclib plus fulvestrant in first- and second-line settings.

Ribociclib is also undergoing clinical evaluation, both alone and in combination with chemotherapeutic agents, in other tumors, including advanced NRAS-mutant melanoma, locally advanced or metastatic BRAF-mutant melanoma, advanced melanoma, advanced solid tumors, lymphoma, and neuroblastoma. More information about all these trials can be found at the ClinicalTrials.gov website (<http://clinicaltrials.gov/>).

Abemaciclib

Abemaciclib (LY2835219) is another orally bioavailable CDK4/6 inhibitor in clinical development; it achieves CDK4/6 inhibition at nanomolar concentrations. Preclinical data indicate that it causes G1 arrest in RB-proficient cells and results in antitumor activity in human tumor xenograft models as both monotherapy and in combination with other chemotherapies¹¹⁶. It also inhibits the growth of intracranial tumors, as the drug appears to be able to cross the blood–brain barrier¹¹⁷. Abemaciclib improves the effect of cytotoxic drugs, underscoring the fact that it can be used in combined therapeutic regimens¹¹⁸.

Abemaciclib is currently being tested in various clinical trials against breast cancer and a variety of other tumors. The first phase I study (NCT01394016) involving abemaciclib included 75 patients with advanced solid tumors; the primary results of the study were presented in Invest New Drugs in 2014¹¹⁶. The MTD was defined at 200 mg twice daily. The most common treatment-related toxicities were mild to moderate and included diarrhea, nausea, fatigue, vomiting, and neutropenia. This phase I study showed promising activity in a metastatic breast cancer expansion cohort in which patients were administered abemaciclib after systemic

therapies (median of seven cycles)¹¹⁹. Among 44 evaluable patients, nine (20%) patients achieved PR, 24 (55%) patients had SD, and 11 (25%) patients showed disease progression. Patnaik and colleagues¹¹⁹ reported that the PR rate was 25% and the SD rate was 56% in the HR-positive subgroup. They reported PFS in HR-positive patients was 9.1 months compared to 5.8 months in all patients. The trial also included a small cohort to investigate tolerance to the combination of abemaciclib plus fulvestrant; the assessment concluded that the combination was safe and the toxicities were acceptable. These two drugs are currently undergoing evaluation in another randomized, double-blind, placebo-controlled phase III study (NCT02107703). A total of 550 women with HR-positive/HER2-negative locally advanced or metastatic breast cancer were enrolled. The results have not yet been published.

The other expansion cohorts from NCT01394016 recruited 49 NSCLC patients (19 KRAS wild-type, 26 KRAS mutant, and four unknown KRAS status)¹²⁰. The enrolled patients had advanced NSCLC and were treated with single-agent abemaciclib following a median of four previous systemic therapies. The disease control rate was 51% [37% for KRAS wild-type ($n=19$) and 54% for KRAS mutants ($n=26$)] with one confirmed patient achieving PR. In this clinical study, abemaciclib was associated with acceptable safety and tolerability profiles, with diarrhea (2%), nausea (4%), fatigue (2%), vomiting (2%), and anemia (2%) being the most common adverse effects. There were no grade 4 side effects.

There are other ongoing evaluations of abemaciclib in clinical trials in other disease states, such as squamous cell carcinoma in NSCLC, NSCLC with KRAS mutant, and mantle cell lymphoma. Information about all these trials can be found at the ClinicalTrials.gov website (<http://clinicaltrials.gov/>). Results from the trials that are currently in progress will improve understanding of the activity of abemaciclib and help distinguish and differentiate it from other CDK4/6 inhibitors.

There are some similarities among the three CDK4/6-selective inhibitors in functional and toxicity profiles. As they are RB-dependent, loss of RB will result in drug resistance. However, these three inhibitors have various treatment-related adverse effects, especially when used in combination with other drugs; these effects may limit the uses of the inhibitors as cancer therapies. Several phase III trials involving these three selective CDK4/6 inhibitors are currently under way and may uncover differences among them in terms of niche activity and tolerability. According to the information obtained from those clinical trials, clinical

combinations can hopefully be optimized, treatment-related side effects can be reduced, and, ultimately, the prognoses of various cancers can be improved.

Conclusions

In all organisms, from yeast to humans, it is well documented that the cell cycle is controlled by the sequential activation of a group of serine-threonine kinases called CDKs, which associate with the cyclin proteins. Since the discovery of the key role of CDK1 in driving the cell cycle, significant progress has been made in understanding the controlled processes of the cell cycle. Deregulation of the cell cycle is a hallmark of the transformation of normal cells into tumor cells. Tumor-associated alterations in CDKs or their activators and inhibitors (cyclins and CKIs, respectively) contribute to sustained proliferation that is independent of the external signals, which is a hallmark of cancer.

In the cell cycle, CDKs regulate critical checkpoints, and are thus important antiproliferative drug targets for several diseases, including cancer. Indeed, therapeutic approaches based on CDK inhibition represent a unique opportunity for drug discovery and a promising strategy for cancer treatment.

In the past two decades, the major research focus has been on ATP-competitive small molecules that target several CDK proteins. However, none of these compounds has yet received approval following clinical trials, primarily due to the lack of specificity in targeting CDKs, as well as their accompanying toxicities and side effects. CDK4 and CDK6 play pivotal roles in regulating cell cycle progression at the G1 restriction point, and are therefore important anticancer drug targets. Indeed, the development of CDK4/6-selective inhibitors has made promising progress: palbociclib, abemaciclib, and ribociclib have been approved or are currently being evaluated in various cancers in ongoing clinical trials. Other ongoing evaluations of CDK4/6-selective inhibitors are in clinical trials. Although CDK4/6 inhibitors are showing great promise in many cancers, there are some questions and concerns that have emerged from the preclinical and clinical data. First, some cancers such as melanomas, colorectal cancer, triple-negative breast cancer are resistant to CDK4 and CDK6 inhibition. Second, there remains a strong need to identify predictive markers that can help define the patient subgroup(s) most likely to benefit from CDK4/6 inhibition. Another question about the future direction of anti-cell cycle therapy is how to optimize treatment protocols: by combining CDK4/6 inhibitors with other targeted therapies or cytotoxic chemotherapy. The

preclinical trials are expected to identify the most promising and most well-tolerated combination therapeutic strategies and improve our understanding of novel functioning and resistance mechanisms. In the near future, more efficient and specific CDK selective inhibitors should be available. New strategies based on CDK inhibition will thus accelerate the development of precision cancer therapy.

Conflict of interest statement

No potential conflicts of interest are disclosed.

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Cite this article as: Bai J, Li Y, Zhang G. Cell cycle regulation and anticancer drug discovery. *Cancer Biol Med.* 2017; 14: 348-62. doi: 10.20892/j.issn.2095-3941.2017.0033