

Supplementary Fig. 1. Expression levels of CDKs in various cancers. Comparison of the expression of CDKs between tumor (red) and normal (blue) tissues. For the boxplots, the center line of the box indicates the median. The upper boundary of the box represents the upper quantile, while the bottom boundary of the box represents the lower quartile. The top and bottom ends of the whiskers indicate the maximum and minimum values, respectively. (\* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001, NS: no significant difference).

Supplementary Table 1. Expression and overall survival rate of patients expressing various CDKs in diverse cancers

Cancer type	Acronym	Specific CDK(s) overexpressed (p < 0.05)	Specific CDKs affecting overall survival rate
			(p < 0.05)
adrenocortical carcinoma	ACC	no data	CDK1, CDK2, CDK3, CDK6, CDK10, CDK17
acute myeloid leukemia	LAML	no data	CDK14, CDK18
bladder urothelial carcinoma	BLCA	CDK1, CDK2, CDK3, CDK4, CDK5, CDK7, CDK10, CDK16, CDK17, CDK18, CDK20	CDK2, CDK6, CDK10, CDK15, CDK17
brain lower grade glioma	LGG	no data	CDK1, CDK,2, CDK3, CDK6, CDK7, CDK8, CDK12, CDK13, CDK16, CDK20
breast invasive carcinoma	BRCA	CDK1, CDK2, CDK3, CDK4, CDK5, CDK6, CDK7, CDK8, CDK9, CDK12, CDK14, CDK15, CDK16, CDK18, CDK19, CDK20	CDK8
cervical squamous cell carcinoma	CESC	CDK1, CDK2, CDK5, CDK7, CDK13, CDK16, CDK18,	-
cholangiocarcinoma	CHOL	CDK2, CDK3, CDK4, CDK5, CDK6, CDK7, CDK8, CDK9, CDK10, CDK12,	CDK15

		CDK13, CDK14, CDK16, CDK17,	
		CDK18, CDK19, CDK20	
colon adenocarcinoma	COAD	CDK1, CDK2, CDK3, CDK4, CDK5,	CDK10
		CDK6, CDK7, CDK8, CDK9, CDK10,	
		CDK12, CDK13, CDK15, CDK16,	
		CDK18, CDK20	
esophageal carcinoma	ESCA	CDK1, CDK2, CDK3, CDK4, CDK5,	-
		CDK6, CDK7, CDK8, CDK9, CDK10,	
		CDK12, CDK13, CDK16, CDK19,	
glioblastoma multiforme	GBM	CDK1, CDK2, CDK4, CDK6, CDK13,	CDK5, CDK7
		CDK15, CDK17, CDK20	
head and neck squamous cell	HNSC	CDK1, CDK2, CDK3, CDK4, CDK5,	CDK3, CDK19
carcinoma		CDK6, CDK8, CDK9, CDK10, CDK12,	
		CDK14, CDK17, CDK18, CDK19,	
kidney chromophobe	KICH	CDK2, CDK3, CDK5, CDK6, CDK7,	CDK1, CDK2, CDK4, CDK5, CDK7, CDK8,
		CDK10, CDK12, CDK13, CDK16,	CDK12
		CDK18, CDK19, CDK20	
kidney renal clear cell carcinoma	KIRC	CDK1, CDK2, CDK3, CDK4, CDK5,	CDK1, CDK3, CDK4, CDK5, CDK9, CDK10,
		CDK6, CDK7, CDK9, CDK10, CDK13,	CDK12 CDK14, CDK15 CDK17, CDK19, CDK20
		CDK16, CDK17, CDK18, CDK19,	

kidney renal papillary cell	KIRP	CDK1, CDK2, CDK3, CDK4, CDK5,	CDK1, CDK2, CDK3, CDK4, CDK7, CDK12
carcinoma		CDK6, CDK7, CDK9, CDK10, CDK13,	CDK18, CDK19
		CDK18, CDK20	
liver hepatocellular carcinoma	LIHC	CDK1, CDK2, CDK3, CDK4, CDK5,	CDK1, CDK2, CDK4, CDK5, CDK7, CDK8
		CDK6, CDK7, CDK8, CDK10, CDK12,	CDK13, CDK16, CDK19, CDK20
		CDK13, CDK14, CDK16, CDK17,	
		CDK18, CDK19, CDK20	
lung adenocarcinoma	LUAD	CDK1, CDK2, CDK3, CDK4, CDK5,	CDK1, CDK6
		CDK7, CDK8, CDK9, CDK10, CDK12,	
		CDK13, CDK14, CDK15, CDK16,	
		CDK17, CDK18,	
lung squamous cell carcinoma	LUSC	CDK1, CDK2, CDK3, CDK4, CDK5,	-
		CDK6, CDK7, CDK8, CDK10, CDK12,	
		CDK14, CDK15, CDK16, CDK17,	
		CDK18, CDK19, CDK20	
lymphoid neoplasm diffuse large	DLBC	no data	-
B-cell lymphoma			
mesothelioma	MESO	no data	CDK1, CDK2, CDK4, CDK6, CDK8, CDK16,
			CDK19
ovarian serous cystadenocarcinoma	OV	no data	CDK8, CDK14, CDK15, CDK19

pancreatic adenocarcinoma	PAAD	CDK12, CDK15, CDK19,	CDK1, CDK2 CDK3, CDK6, CDK7, CDK10
prostate adenocarcinoma	PRAD	CDK1, CDK3, CDK4, CDK4, CDK5, CDK7, CDK8, CDK9, CDK10, CDK14, CDK15, CDK16, CDK17, CDK18, CDK19,	CDK7
pheochromocytoma and paraganglioma	PCPG	CDK5, CDK13, CDK14,	CDK1, CDK3, CDK16, CDK19
rectum adenocarcinoma	READ	CDK1, CDK2, CDK4, CDK5, CDK7, CDK8, CDK9, CDK10, CDK12, CDK14, CDK15, CDK16, CDK17, CDK18, CDK20	CDK12
sarcoma	SARC	CDK4, CDK9,	CDK6, CDK9, CDK14, CDK18
skin cutaneous melanoma	SKCM	CDK13,	CDK1, CDK2, CDK4
testicular germ cell tumors	TGCT	no data	-
thyroid carcinoma	THCA	CDK4, CDK5, CDK6, CDK7, CDK8, CDK9, CDK12, CDK14, CDK16, CDK18, CDK20	CDK15, CDK20
thymoma	THYM	CDK16,	CDK2, CDK7, CDK14, CDK18
stomach adenocarcinoma	STAD	CDK1, CDK2, CDK3, CDK4, CDK6, CDK7, CDK8, CDK9, CDK10, CDK12,	no data

		CDK13, CDK16, CDK17, CDK18,	
		CDK19, CDK20	
UCEC and uveal melanoma	UVM	no data	CDK2, CDK5, CDK10, CDK15
uterine corpus endometrial	UCEC	CDK1, CDK2, CDK4, CDK6, CDK7,	CDK6, CDK14 CDK16, CDK19, CDK20
carcinoma		CDK8, CDK9, CDK13, CDK14, CDK16,	
		CDK17, CDK18, CDK19,	
uterine carcinosarcoma	UCS	no data	CDK7

Positive media	ators of CDK1		
Mediator	Function	Mechanism	Ref
CDK7	Activates CDK1	Increases phosphorylation of CDK1 at Thr161	27
LAR	Activates CDK1	LAR protein tyrosine phosphatase mediates focal adhesion by enhancing	28
		phosphorylation of CDK1 at Thr161	
NOL11	Activates CDK1	Depletion of NOL11 delays entry into the M phase owing to increased	29
		inhibitory phosphorylation of CDK1 at Tyr15	
C53	Activates CDK1	C53 promotes CDK1 activation and mitotic entry in both unperturbed cell	30
		cycle progression and DNA damage response	
CDC25A	Activates CDK1	CDC25A mediates the activity of CDK1–2/cyclin A and CDK1/cyclin B in	27 32
		G2 phase	
CDC25B	Activates CDK1	CDC25B activates human CDK1 kinase by dephosphorylating both Thr14	33
		and Tyr15	
CDC25C	Activates CDK1	CDC25C positively mediates activation of CDK1/cyclin B	34,35
CDK2	Activates CDK1	Cyclin A/CDK2 complexes are key mediators of CDK1 activation in	36
		human cells	

Activates CDK1	BECN1 promotes radiation-induced G2/M arrest through mediation of	40
	CDK1 activity: a potential role for autophagy at the G2/M checkpoint	
Activates CDK1	KCTD12 binds to CDC25B and activates CDK1 and Aurora kinase A to	41
	facilitate the G2/M transition and promote tumorigenesis	
Activates CDK1	Phosphorylation at Ser10 and Ser70 of NPM plays a role in mediating	42
	CDK1 kinase activity during the G2/M transition	
Activates CDK1	MCM10 activates CDK1 in G2 phase	43
Activates CDK1	Aurora kinase A is required for initial activation of CDK1 at centrosomes	39 44,194
Activates CDK1	PFKFB3 promotes cell cycle progression and suppresses apoptosis through	45
	CDK1-mediated phosphorylation of p27	
Activates CDK1	UCH-L1 physically interacts with CDK1 enhancing its kinase activity and	46
	promoting cell proliferation	
Activates CDK1	MASTL enhances cyclin B1/CDK1-dependent mitotic phosphorylation	47
	events, directing mitotic entry, anaphase and cytokinesis in human cells	
Activates CDK1	TSPY enhances the cyclin B1-CDK1 phosphorylation activity in	48
	modulating cell-cycle progression at the G2/M stage	
Activates CDK1	KPNB1 promotes CDK1/cyclin B1 translocation from the cytoplasm to the	49
	nucleus	
Activates CDK1	Knockdown of SIL in cancer cells <i>in vitro</i> delays entrance into mitosis and	50
	decreases activation of the CDK1/cyclin B complex	
	Activates CDK1         Activates CDK1	CDK1CDK1 activity: a potential role for autophagy at the G2/M checkpointActivates CDK1KCTD12 binds to CDC25B and activates CDK1 and Aurora kinase A to facilitate the G2/M transition and promote tumorigenesisActivates CDK1Phosphorylation at Ser10 and Ser70 of NPM plays a role in mediating CDK1 kinase activity during the G2/M transitionActivates CDK1MCM10 activates CDK1 in G2 phaseActivates CDK1Aurora kinase A is required for initial activation of CDK1 at centrosomesActivates CDK1PFKFB3 promotes cell cycle progression and suppresses apoptosis through CDK1-mediated phosphorylation of p27Activates CDK1UCH-L1 physically interacts with CDK1 enhancing its kinase activity and promoting cell proliferationActivates CDK1MASTL enhances cyclin B1/CDK1-dependent mitotic phosphorylation events, directing mitotic entry, anaphase and cytokinesis in human cellsActivates CDK1TSPY enhances the cyclin B1-CDK1 phosphorylation activity in modulating cell-cycle progression at the G2/M stageActivates CDK1KPNB1 promotes CDK1/cyclin B1 translocation from the cytoplasm to the nucleusActivates CDK1KPNB1 promotes CDK1/cyclin B1 translocation from the cytoplasm to the nucleus

E2F1-3	Promote CDK1	E2F1, E2F2, and E2F3 bind to the CDK1 promoter promoting its	51
	transcription	transcription	
MRG15	Promotes CDK1	MRG15 acts in a HAT complex involving Tip60 to modify chromatin	52
	transcription	through the acetylation of histone H4 at the CDK1 promoter to activate	
		transcription	
LIN54	Promotes CDK1	LIN54 is an essential core subunit of the DREAM/LINC complex that binds	53
	transcription	to the CDK1 promoter and promotes CDK transcription	
CENPE	Promotes CDK1	Overexpression of CENPE significantly increases CDK1 promoter activity	54
	transcription		
SP1	Promotes CDK1	Overexpression of SP1 significantly decreases G2/M arrest in cervical	56
	transcription	cancer cells, which is related to upregulation of CDK1 expression	
RPS9	Promotes CDK1	Knockdown of RPS9 inhibits the growth of human colon cancer cells at the	57
	transcription	G2/M phase by downregulating CDK1 expression at the promoter level	
CHPF	Enhances CDK1	CHPF promotes development of malignant melanoma through mediation of	58
	expression	CDK1	
CD276	Enhances CDK1	CD276 maintains proliferation and increases differentiation by enhancing	59
	expression	CDK1 expression in endothelial progenitor cells	
NSUN2	Enhances CDK1	NSUN2 methylates CDK1 mRNA enhancing CDK1 translation	60,61
	translation		

DAP5	Enhances CDK1	The translation initiation factor DAP5 is a mediator of cell survival during	62
	translation	mitosis by enhancing CDK1 translation	
RBM7	Stabilizes CDK1 mRNA	Oncogenic action of the exosome cofactor RBM7 by stabilization of CDK1	63
		mRNA in breast cancer	
HDAC3	Stabilizes CDK1	HDAC3 controls G2/M phase progression mainly through post translational	64
	expression	stabilization of CDK1	
DNMT3A	Enhances CDK1	DNMT3A mutation-induced CDK1 overexpression promotes	66
	expression	leukemogenesis	
RPS15A	Enhances CDK1	RPS15A overexpression induces CDK1 expression	195
	expression		
TPX2	Enhances CDK1	Knocking down TPX2 inhibits CDK1 expression and represses	196
	expression	phosphorylation of the ERK/GSK3 $\beta$ /SNAIL signaling pathway, thereby	
		inhibiting the tumor epithelial-mesenchymal transition	
MYC	Enhances CDK1	MYC regulates the CDK1/cyclin B1 dependent-G2/M cell cycle	197
	expression	progression	
ZIC5	Enhances CDK1	ZIC5 promotes colorectal cancer cell proliferation and cell cycle	198
	expression	progression by enhancing CDK1 signaling	
KPNA2	Enhances CDK1	KPNA2 promotes tumor cell proliferation by increasing the expression of	199
	expression	CDK1	

Enhances CDK1	UBAP2L knockdown decreases CDK1 expression and plays an oncogenic	200
expression	role in HCC	
Enhances CDK1	NUSAP1 enhances proliferation, migration and invasion in breast cancer	201
expression	cells by increasing CDK1 expression	
Enhances CDK1	Activation of PLK1 enhanced CDK1 expression in cells	202,203
expression		
Enhances CDK1	IFITM3 activates the STAT3/CDK1 pathway	204
expression		
Enhances CDK1	KLF8 knockdown induces cell phase arrest decreasing the expression of	205
expression	CDK1	
Enhances CDK1	LOX knockdown reduces CDK1 expression in cancer cell mitosis	206
expression		
Enhances CDK1	Knockdown of RTKN affects proliferation and metastasis of colon cancer	207
expression	by reducing expression of CDK1	
Enhances CDK1	Overexpression of PTTG1 causes upregulation of CDK1 expression	208
expression	resulting in enhanced proliferation	
tors of CDK1	1	1
Function	Mechanism	Ref
Inactivates CDK1	WEE1 kinase phosphorylates the CDK1-cyclin B complex and suppresses	67-69
	the CDK1-cyclin B kinase activity	
	<ul> <li>expression</li> <li>Enhances CDK1</li> <li>expression</li> </ul>	expression       role in HCC         Enhances CDK1       NUSAP1 enhances proliferation, migration and invasion in breast cancer         expression       cells by increasing CDK1 expression         Enhances CDK1       Activation of PLK1 enhanced CDK1 expression in cells         expression       intervention of PLK1 enhanced CDK1 expression in cells         expression       IFITM3 activates the STAT3/CDK1 pathway         expression       CDK1         Enhances CDK1       KLF8 knockdown induces cell phase arrest decreasing the expression of expression         Enhances CDK1       LOX knockdown reduces CDK1 expression in cancer cell mitosis         expression       CDK1         Enhances CDK1       LOX knockdown reduces CDK1 expression in cancer cell mitosis         expression       CDK1         Enhances CDK1       Knockdown of RTKN affects proliferation and metastasis of colon cancer         expression       by reducing expression of CDK1         Enhances CDK1       Overexpression of CDK1         expression       resulting in enhanced proliferation         tors of CDK1       Mechanism         Inactivates CDK1       WEE1 kinase phosphorylates the CDK1-cyclin B complex and suppresses

MYT1	Inactivates CDK1	The human MYT1 kinase preferentially phosphorylates CDK1 (Thr14) and	70 71 72
		localizes to the endoplasmic reticulum and Golgi complex	
PTEN	Inhibits CDK1	PTEN decreases phosphorylation of CDK1 (Tyr15), resulting in decreased	74
	activation	G2/M cell cycle arrest	
DYRK1A	Inactivates CDK1	Inhibition of DYRK1A results in massive cyclin B accumulation and	75
		saturation of CDK1 activity	
USP7	Inactivates CDK1	USP7 limits CDK1 activity throughout the cell cycle	209
p17	Inactivates CDK1	p17 suppresses CDK1 action and disrupts vimentin phosphorylation,	210
		causing G2/M cell cycle arrest	
IFI27	Inactivates CDK1	IFI27 knockdown increases CDK1 phosphorylation (Tyr15) involved in cell	211
		proliferation.	
CDKN3	Inactivates CDK1	CDKN3 controls mitosis by inhibiting CDK1 activation by	212
		dephosphorylation of CDK1 (Thr161)	
BUBR1	Inhibits CDK1	Deletion of BURR1 activates cyclin B/CDK1 kinase, leading to premature	213
	activation	mitotic entry	
CDH1	Inhibits CDK1	UV-induced degradation of CDH1 results in accumulation of cyclin B1 and	214
	activation	thus increased CDK1 activity	
CSNK2B	Inhibits CDK1	Knockdown of CSNK2B causes stabilization of WEE1 and increases	215
	activation	phosphorylation of CDK1 (Tyr15)	

PRKCD	Inactivates CDK1	PRKCD induces phosphorylation of CDK1 (Tyr15), a critical event in the	216
		G2/M checkpoint	
BDNF	Inhibits both CDK1	Inhibition of both CDK1 expression and activity through a BDNF-	217
	expression & activity	dependent mechanism contributes to the maintenance of tetraploid RGCs in	
		a G2-like state	
DEDD	Inactivates CDK1	DEDD is an inhibitor of mitotic CDK1/cyclin B1	79,80
CHEK1	Inhibits CDK1	CHEK1 prevents entry into mitosis by inhibiting CDK1 activity	76,77
	activation		
KIF22	Inactivates CDK1	Inhibition of KIF22 increases CDC25C expression and CDK1 activity,	78
		resulting in delayed mitotic exit	
p21	Inactivates CDK1	Loss of p21CIP1 unleashes CDK1 activity	218 219
SET	Inactivates CDK1	SET inhibits cyclin B/CDK1 activity	220,221
DUX4	Inactivates CDK1	DUX4 functions as a direct inhibitor of CDK1, and decreases colon cancer	222
		cell proliferation	
ESR2	Inactivates CDK1	Unliganded ESR2 causes a G2 cell cycle arrest by inactivating CDK1	223
PP4C	Inactivates CDK1	Loss of PP4C leads to an unscheduled activation of CDK1 in interphase	224
CDC14B	Inhibits CDK1	CDC14B counteracts CDK1 action and causes mitotic arrest in mouse pre-	225
	activation	implantation embryo models	
HDAC4	Inhibits CDK1	HDAC4 inhibits the CDK1 activity and the progression of proliferating	226
	activation	HEK293T and HT22 cells through the cell cycle.	

TSPX	Inhibits CDK1	TSPX represses the cyclin B1/CDK1 phosphorylation activity	48
	activation		
p57	Inhibits CDK1	p57 is required to trigger endoreduplication by inhibiting CDK1	227
	activation		
SUMO1	Inhibits CDK1	Inhibits CDK1 activity by SUMOylation	228
	activation		
FGF1	Inhibits CDK1	FGF1 inhibits the cyclin B1/CDK1 kinase activity to induce a transient G2	229
	activation	arrest	
CDC6	Inhibits CDK1	Phosphorylation of CDC6 promotes the interaction of CDC6 and CDK1,	230
	activation	leading to attenuated CDK1 activity	
PKR	Induces CDK1	PKR mediates Tyr4 phosphorylation facilitating CDK1 ubiquitination and	82
	degradation	triggering CDK1 downregulation	
BTRC	Induces CDK1	CDK1 accumulation in patients' tumors shows a negative correlation with	83
	degradation	BTRC and a positive correlation with the degree of tumor malignancy	
HDAC6; p62	Induces CDK1	p62/SQSTM1/HDAC6-dependent autophagy mediates CDK1 degradation	87
	degradation	in human breast cancer	
TWEAK	Induces CDK1	TWEAK induces CDK1 phosphorylation and degradation, resulting in the	231
	degradation	arrest of cell growth at the G2/M phase	
HEI10	Inhibits CDK1	HEI10 negatively mediates cell invasion by inhibiting cyclin B/CDK1	89
	expression	expression	

SFN	Inhibits CDK1 nuclear	SFN is required to sequester CDK1/cyclin B1 complexes in the cytoplasm	90
	translocation	and prevent mitotic catastrophes	

CDK1 tumor promot	or substrates		
Target substrate	Phosphorylation site	Biological function	Ref
BRAF	Ser144	CDK1/cyclin B directly phosphorylates BRAF at Ser144, which is	91
		required for mitotic activation of BRAF and subsequent activation of	
		the MAPK cascade	
ERK3	Thr698	CDK1 phosphorylates ERK3 at Thr698, which operates in a cell-cycle-	92
		dependent manner	
AR	Ser81, Ser 515	CDK1 phosphorylates the AR at Ser81 and leads to AR-mediated	93-95
		transactivation in prostate cancer; phosphorylation of AR at Ser515 by	
		CDK1 may be an independent prognostic marker	
HIF1A	Ser668	CDK1 stabilizes HIF1A through direct phosphorylation of Ser668 to	98
		promote tumor growth	
YAP	Thr119, Ser289, Ser367	CDK1 phosphorylates YAP, promoting mitotic defects and cell motility	99
TAZ	Ser90, Ser105, Thr326,	CDK1 phosphorylation of TAZ in mitosis inhibits its oncogenic activity	100
	Thr346		
AJUBA	Ser119, Ser175	Adaptor protein AJUBA is phosphorylated by CDK1 at Ser119 and	101
		Ser175 during the G2/M phase of the cell cycle, which promotes cell	
		proliferation and tumorigenesis.	

Thr172	CDK1-mediated BCL9 phosphorylation in controlling mitotic Wnt	232
	signaling to promote cell division and growth	
Thr24, Ser32, Se59	CDK1-mediated mitotic phosphorylation of PBK is involved in	233
	cytokinesis and inhibits its oncogenic activity	
Thr34, Thr46, Ser72	SRC phosphorylation by CDK1 has been demonstrated to increase SRC	234
	kinase activity	
Ser131, Thr146	CDK1 phosphorylation mediates the activity of NET1 towards RhoA	235
	during mitosis.	
Thr596, Ser251	CDK1 phosphorylates FOXM1B, which is essential for recruitment of	102,236
	CREB binding protein coactivator proteins.	
Ser269	Phosphorylation of ISL1 at Ser269 by CDK1 increases its	103
	transcriptional activity and promotes cell proliferation in gastric cancer	
Ser83	CDK1 phosphorylation of mRNA translation regulator 4E-BP1 at	104
	Ser83, which contributes to cell transformation	
Thr51, Thr53	CDK1-mediated phosphorylation of human ATF7 at Thr51 and Thr53	106
	promotes cell-cycle progression into M phase	
Ser48, Ser303,	Phosphorylation of RUNX1 by CDK1 reduces direct interaction with	107
Ser424	HDAC1 and HDAC3	
Ser451	Mediates G2 and M phases and promotes tumor angiogenesis	108
		237
	Image:	signaling to promote cell division and growthThr24, Ser32, Se59CDK1-mediated mitotic phosphorylation of PBK is involved in cytokinesis and inhibits its oncogenic activityThr34, Thr46, Ser72SRC phosphorylation by CDK1 has been demonstrated to increase SRC kinase activitySer131, Thr146CDK1 phosphorylation mediates the activity of NET1 towards RhoA during mitosis.Thr596, Ser251CDK1 phosphorylates FOXM1B, which is essential for recruitment of CREB binding protein coactivator proteins.Ser269Phosphorylation of ISL1 at Ser269 by CDK1 increases its transcriptional activity and promotes cell proliferation in gastric cancerSer83CDK1 phosphorylation of mRNA translation regulator 4E-BP1 at Ser83, which contributes to cell transformationThr51, Thr53CDK1-mediated phosphorylation of human ATF7 at Thr51 and Thr53 promotes cell-cycle progression into M phaseSer424HDAC1 and HDAC3

RXRA	Ser265	CDK1 phosphorylates RXRA by activating c-Jun NH <sub>2</sub> -terminal kinases	109
CEBPA	Ser21	CDK1 phosphorylates CEBPA on Ser21, which inhibits its	110
		differentiation-inducing function	
OCT4	Ser229	CDK1 interplays with OCT4 to repress differentiation of embryonic	113
		stem cells into trophectoderm	
TFCP2L1	Thr177	TFCP2L1 is phosphorylated at Thr177 by CDK1, which affects	111
		embryonic stem cell cycle progression, pluripotency and differentiation	
BCL2	Ser70	Mediates BCL2 phosphorylation and link the coupling of mitotic arrest	114,115,117,118
		and apoptosis	
BCL2L1	Ser62	Mediates BCL2L1 phosphorylation and links coupling mitotic arrest	114,116,118
		and apoptosis	
DRP1	Ser585; Ser616	DRP1 is specifically phosphorylated in mitosis by CDK1/cyclin B on	119-121
		Ser585; DRP1 mediates mitochondrial fusion and apoptosis in human	
		cancer cells	
CPD	Thr176	The CPD kinase can mediate CDK phosphorylation at Thr176	123
FBXO28	Ser344	CDK1 mediates activation of the FBXO28 ubiquitin ligase promoting	122
		MYC-driven transcription and tumorigenesis in breast cancer	
EIF4G1	Ser1232	Phosphorylation of EIF4G1 (Ser1232) in the mitotic translation	238
		initiation shift	

CEP55	Ser425, Ser428	CEP55 is triggered by CDK1-dependent phosphorylation at Ser425 and	239
		Ser428, which is required for its recruitment to midbody and	
		cytokinesis	
MDM2	Thr216	MDM2 is phosphorylated by cyclin A/CDK2 or cyclin A/CDK1 at	240
		Thr216, which is most prevalent at the onset of S phase	
SIRT3	Thr150, Ser159	SIRT3 enzymatic activity is further enhanced through of	241
		Thr150/Ser159 phosphorylation by cyclin B1-CDK1, which is also	
		induced by radiation and relocated to mitochondria together with	
		SIRT3	
NCOA3	Ser728, Ser867	Phosphorylation of NCOA3 at mitosis is mediated by CDK1	242
EZH2	Thr487	CDK1-dependent phosphorylation of EZH2 suppresses methylation of	243
		H3K27 and promotes osteogenic differentiation of human	
		mesenchymal stem cells	
CDK1 tumor suppre	essor substrates		
Target substrate	Phosphorylation site	Biological function	Ref
p53	Ser315	Anti-apoptotic function of mitochondrial p53 mediated by cyclin	125
		B1/CDK1 Ser315 phosphorylation in p53-wildtype tumor cells	
p73	Thr86	p73 proteins are targets of CDK complexes and phosphorylation on	126
		Thr86 by CDK1 mediates p73 functions	

FOXO1	Ser249	CDK1 contributes to tumorigenesis by promoting cell proliferation and	127,128
		survival through phosphorylation and inhibition of FOXO1	
CASP8	Ser387	Inhibits apoptosis during mitosis and contributes to tumorigenesis	129,244
CASP9	Thr125	Cyclin/CDK1 phosphorylates CASP9 on Thr125 and protects mitotic	130
		cells from apoptosis	
EMI1	Ser145, Ser149	EMI1 is phosphorylated by CDK1 providing a delay for APC activation	132
DLG1	Ser158, Ser442	DLG1 is phosphorylated by CDK1 on Ser158 and Ser442, which affect	131
		the nuclear localization of DLG1.	
P62	Thr269, Ser272	Phosphorylation of p62 by CDK1 controls the timely transit of cells	133
		through mitosis and tumor cell proliferation	
EPHA2	Ser897	EPHA2 phosphorylation on Ser897 by the CDK1/MEK/ERK/RSK	245
		pathway mediates M-phase progression by maintaining cortical rigidity.	
VGLL4	Ser58, Ser155,	CDK1 mediates mitotic phosphorylation of the transcriptional co-	136
	Thr159, Ser280	repressor VGLL4 and inhibits its tumor-suppressing activity	
RAP80	Ser677	CDK1 mediates phosphorylation of RAP80 at Ser677 and modulates	137
		DNA damage-induced G2/M checkpoint and cell survival	
ING1	Ser126	CDK1 phosphorylates Ser126 of ING1, which plays a key role in	246
		mediating the expression of cyclin B1 and proliferation of melanoma	
		cells	

EML2	Thr84	CDK1 phosphorylates EML2, modulating cell proliferation and	138
		migration.	
CDK1 cell cycle subs	strates		
Target substrate	Phosphorylation site	Biological function	Ref
BORA	Thr12, Thr15, Ser25,	Phosphorylation of the N terminus of BORA for PLK1 activation and	140
	Ser41, Thr52, Ser112,	mitotic entry	
	Ser137, Ser183,		
	Thr243, Ser252,		
	Ser271, Ser274,		
	Ser278, Thr287,		
	Ser375, Thr381		
MLL5	Thr912	Phosphorylation of MLL5 by CDK1 affects its cellular distribution and	141
		is required for mitotic entry	
GWL	T193, T206	CDK1 phosphorylates GWL during mitotic entry	142
KIF4A	Ser1186	CDK1-dependent KIF4A phosphorylation at Ser1186 is a trigger for	247
		chromosomal organization during early mitosis	
VCIP135	Ser130	In early mitosis, phosphorylation of VCIP135 by CDK1 at a single	248
		residue, Ser130, is sufficient to inactivate the enzyme and inhibit	
		p97/p47-mediated Golgi membrane fusion	

Thr242	Mitotic exit function of Polo-like kinase CDC5 is dependent on	143
	sequential activation by CDK1	
Thr320	CDK1 inhibits PP1-mediated dephosphorylation at mitotic exit	249
Ser68	Eliminates the binding of CENPA to the assembly factor HJURP, thus	144
	preventing the premature loading of CENPA to the centromere prior to	
	mitotic exit	
Ser727	Phosphorylation of STAT3 (Ser727) by CDK1 is critical for	250
	nocodazole-induced mitotic arrest	
Thr92	Phosphorylation of MCL1 by CDK1-cyclin B1 during mitotic arrest	251
Thr1415	Thr1415 of the CAP-D3 subunit is a CDK1 phosphorylation site, which	145
	is required for timely chromosome condensation during prophase	
Thr210	CDK1 is necessary for NEK9 activation for centrosome separation	252
	during mitosis.	
Ser774	CDK1 phosphorylates CHK2 kinase in metaphase, influencing cellular	146
	morphogenesis	
T358, T360	SKA3 is phosphorylated by CDK1 in mitosis to promote anaphase	148
	onset	
Ser71, Ser86	CDK1 activity prevents premature nuclear envelope assembly and	253
	phosphorylation of the inner nuclear membrane protein LBR by CDK1	
	contributes to the temporal control	
	Thr320         Ser68         Ser727         Thr92         Thr1415         Thr210         Ser774         T358, T360	sequential activation by CDK1Thr320CDK1 inhibits PP1-mediated dephosphorylation at mitotic exitSer68Eliminates the binding of CENPA to the assembly factor HJURP, thus preventing the premature loading of CENPA to the centromere prior to mitotic exitSer727Phosphorylation of STAT3 (Ser727) by CDK1 is critical for nocodazole-induced mitotic arrestThr92Phosphorylation of MCL1 by CDK1-cyclin B1 during mitotic arrestThr1415Thr1415 of the CAP-D3 subunit is a CDK1 phosphorylation site, which is required for timely chromosome condensation during prophaseThr210CDK1 is necessary for NEK9 activation for centrosome separation during mitosis.Ser774CDK1 phosphorylates CHK2 kinase in metaphase, influencing cellular morphogenesisT358, T360SKA3 is phosphorylated by CDK1 in mitosis to promote anaphase onsetSer71, Ser86CDK1 activity prevents premature nuclear envelope assembly and phosphorylation of the inner nuclear membrane protein LBR by CDK1

LMN1	Ser22, Ser392	The nuclear envelope is a dynamic structure that completely	254
		disassembles in response to CDK1 activity in mitosis	
EML3	Thr881	CDK1 phosphorylates EML3 regulating mitotic spindle assembly and	255
		the kinetochore-microtubules connection	
CLASP2	Ser1234	CDK1 mediates CLASP2 stabilizing kinetochore-microtubule	256
		attachments	
NUMA	Thr2055	CDK1-mediated phosphorylation at T2055 negatively modulates	257
		NUMA cortical localization mediating mitotic progression with proper	
		spindle function.	
DIAPH1	Ser629, Ser640,	CDK1-mediated DIAPH1 phosphorylation maintains metaphase	258
	Ser665	cortical tension and inactivates the spindle assembly checkpoint at	
		anaphase	
CRM1	Ser391	Phosphorylation of CRM1 by CDK1/cyclin B promotes Ran-dependent	259
		mitotic spindle assembly	
TIAM1	Ser1466	S1466 of TIAM1 is a novel CDK1 site for which phosphorylation is	260
		required for the mitotic function of TIAM1 to promote mitotic spindle	
		formation	
KIF11	Thr926	CDK1 phosphorylates KIF11 at Thr926 during mitosis, which is crucial	261
		for the mitotic function of KIF11	

NDEL1	Thr219	Silencing of CDK1 clearly suppresses the phosphorylation of NDEL1	224
		at Thr219 mediating microtubule organization	
NEDD1	Thr550	Promotes the interaction of NEDD1 with the $\gamma$ TuRC component $\gamma$ -	262
		tubulin during mitosis	
TUBB	Ser172	Mediation of microtubule dynamics during mitosis.	263
TPPP1	Thr14, Ser18, Ser45,	CDK1 phosphorylation of TPPP1 inhibits its modulation of cell cycle	264
	Ser160	to	
		increase cell proliferation	
CEP169	Ser451	Mediates microtubule dynamics of mitotic spindle	265
TMAP	Thr622	CDK1/cyclin B1-mediated phosphorylation of TMAP contributes to	266
		proper regulation of microtubule dynamics and establishment of	
		functional bipolar spindles during mitosis	
RPL12	Ser38	Phosphorylation of the ribosomal protein RPL12 by CDK1 affects	267
		translation during mitosis	
CREB	Ser270, Ser271	CDK1-dependent phosphorylation of CREB at Ser270 and Ser271	268
		facilitates	
		its dissociation from chromatin during mitosis by reducing its intrinsic	
		DNA binding potential	
SP1	Ser720, Thr723,	CDK1-mediated SP1 phosphorylation facilitates cell cycle progression	269
	Thr737	during mitosis	

CUX1	Ser1237, Ser1270	Hyperphosphorylation of CUX1 by CDK1/cyclin B inhibits its DNA	270
		binding activity in mitosis	
TOP1	Ser112, Ser394	TOP1 is phosphorylated during mitosis by CDK1, which enhances	271
		DNA relaxation activity	
NCAPG	Thr308, Thr332	Phosphorylation of NCAPG is required for its chromosomal DNA	272
		localization during mitosis	
XIAP	Ser40	Phosphorylation of XIAP by CDK1-cyclin-B1 controls mitotic cell	273
		death	
UBC9	Ser71	Phosphorylation of UBC9 by CDK1 enhances SUMOylation activity in	274
		cell cycle mediation	
EBP50	Ser279, Ser301	CDK1/cyclin B-mediated EBP50 phosphorylation plays a role in the	275
		modulation of various cell functions by affecting actin cytoskeleton	
		reorganization	
FLNa	Ser1436	CDK1/cyclin B1 phosphorylates Ser1436 of FLNa mediating actin	276
		remodeling	
cGAS	Ser305 (human)	DNA sensor cGAS is post-translationally modulated by cell cycle-	277
	Ser291 (mouse)	dependent enzymes to ensure its proper activation for host defense of	
		cytosolic DNA in interphase and is inert to self-DNA in mitosis	
ARHGAP19	Thr404, Thr476	Phosphorylation of ARHGAP19 by CDK1 mediates its subcellular	278
		localization and function during mitosis	

ARHGEF12	Ser190, Ser1176	CDK1 phosphorylates ARHGEF12 at Ser190 and Ser1176 during	279
		mitosis	
CHK1	Ser286, Ser301	CHK1 is phosphorylated at Ser286 and Ser301 by CDK1 during	76
		mitosis	
CC2D1A	Ser208	CDK1 is a kinase of CC2D1A during mitosis and its phosphorylation of	280
		CC2D1A mediates mitotic function	
GM130	Ser25	GM130 is phosphorylated in early mitosis on Ser25 by CDK1/cyclin B.	281
CENPA	Ser68	Dynamic phosphorylation of CENPA at Ser68 by CDK1 orchestrates	144
		its cell-cycle-dependent deposition at centromeres	
MCM3	Ser112	CDK1-dependent phosphorylation of Ser112 triggers the assembly of	282
		MCM3 with the remaining MCM subunits and subsequent chromatin	
		loading of MCMs	
CDCA5	Ser21, Thr48, Ser75,	phosphorylation of CDCA5 by CDK1 influences sister chromatid	283,284
	Ser79, Ser83, Thr111,	cohesion	
	Thr115, Thr159,		
	Ser181, Ser209		
S6K1	Ser371, Ser411,	CDK1 triggers inactivation of S6K1 in mitosis, serving to spare energy	285
	Thr421, Ser424	for costly mitotic processes at the expense of ribosomal protein	
		synthesis	

NIPA	Ser395	CDK1/cyclin B1 amplifies phosphorylation of NIPA, which may	286
		contribute to the regulation of its own abundance in early mitosis	
SEPT9	Thr24	Mitotic regulation of SEPT9 by CDK1 is important for the completion	287
		of cytokinesis	
FAR1	Ser87	Phosphorylation by CDK1 primes FAR1 for ubiquitin-mediated	149
		proteolysis	
WRN	Ser1133	CDK1 phosphorylates WRN at collapsed replication forks	150
CDC7	Ser16, Ser302,	CDK1-mediated phosphorylation of CDC7 suppresses DNA re-	151
	Thr376, Thr472,	replication	
	Thr503		
BRCA1	Ser1497, Ser1189,	CDK1 participates in BRCA1-dependent S phase checkpoint control in	152
	Ser1191	response to DNA damage	
RAD9	Ser11	Dynamics of RAD9 chromatin binding and checkpoint function are	153
		mediated by CDK1 activity	
SLBP	Thr61	Phosphorylation of Thr61 by cyclin A/CDK1 triggers degradation of	288
		stem-loop binding protein at the end of S phase	
TRF1	Thr371	CDK1 phosphorylates TRF1, which is recruited to sites of DNA	154
		damage to facilitate homologous recombination and checkpoint	
		activation	

ELAVL1	Ser202	CDK1 phosphorylates ELAVL1 during G2, thereby helping to retain it	155
		in the nucleus hindering its post-transcriptional function and anti-	
		apoptotic influence	
VIM	Ser55	Mediates proliferation of mammalian cell lines	289
EZH2	Thr350	Link between the cell-cycle machinery and epigenetic gene silencing	290
CLIP1	Thr287	CDK1-mediated phosphorylation of CLIP1is essential for the normal	291
		function of this protein during cell cycle progression	
VPS34	Thr159	Negative regulation of VPS34 by CDK1 mediates cell-cycle	292
		progression	
CDC25A	Ser17, Ser115	Mitotic stabilization of CDC25A reflects its phosphorylation on Ser17	293,294
	Ser115, Ser320	and Ser115 by CDK1/cyclin B, which is required to uncouple CDC25A	
		from its ubiquitin-proteasome-mediated turnover	
SIRT1	Thr530, Ser540	Mutation of these two residues phosphorylated by CDK1/cyclin B	295,296
		disturbs normal cell cycle progression and fails to rescue proliferation	
		defects in SIRT1-deficient cells	
HHR6A	Ser120	CDK1-mediated phosphorylation of HHR6A, which is an important	297
		regulatory event in the control of cell cycle progression	
PRKAR2A	Thr54	Thr54 phosphorylation of PRKAR2A by CDK1 might serve to mediate	298
		PRKACA during cell cycle	

CDC25C	Ser214	The cell cycle control protein CDC25C is present, and phosphorylated	299
		on Ser214 by CDK1 in the transition from germinal vesicle to	
		metaphase II in human oocyte meiosis	

Compound	Structure	Target (IC50)	Cell base potency (IC50 or	Animal model (type and	Ref
name			effective inhibitory dose)	dose and route of	
				administration)	
RO-3306	°	CDK1 (35 nM)	<i>IC</i> <sub>50:</sub>	Patient derived xenograft	157-159
	S NH S		WAC2 (1.8 µM)	(PDX) tumor model-	
			SHEP (1.3 µM)	RO3306 4 mg/kg every 2	
	N		NB69 (2.3 µM)	days, oral gavage	
			SK-N-FI (4 μM)	administration	
			NLF (3.8 μM)		
			RH-41 (> 8 μM)		
			IMR32 (1.6 µM)		
			IMR-32 LV-h-p53		
			(2.4 µM)		
			NGP (4 µM)		
			HDMB-3 (1.8 μM)		

CGP-74514A		CDK1 (25 nM)	<i>IC</i> <sub>50</sub> :	NA	160,161
	СІ		MCF-7 (5 µM)		
	N N		T47D (5 μM)		
	NH <sub>2</sub> NH <sub>2</sub> NH <sub>2</sub> NH <sub>2</sub> NH <sub>2</sub> NH <sub>2</sub> N		SK-BR-3 (5 μM)		
	NH2		NCI-H720 (2.2 µM)		
			NC1-H727 (3.2 µM)		
			BON-1 (1.9 μM)		
			hTERT-RPE1 (13 µM)		
Alsterpaullone		CDK1 (35 nM)	Effective inhibitory dose	D458 cerebellar xenograft	300,301
	N N	CDK2 (15 nM)	D425 (5 µM)	model- alsterpaullone 30	
	-O-N+ O	CDK5 (40 nM)	D458 (5 µM)	mg/kg, SC, daily for 2	
	0			weeks;	
CVT-313	NH	CDK1 (4.2 µM)	A2058 (16 µM)	Rat restenosis model—	302,303
		CDK2 (500 nM)	Caco-2 (4.5 µM)	Rats were injected with	
		CDK4 (215 µM)	Capan-1 (10 µM)	CBT-313 (0.75 and 0.25	
	ОН		L1210 (4.2 µM)	mg/kg) intraperitoneally	
			MCF-7 (6.25 µM)		
			MRC-5 (20 µM)		
			Neonatal (5 µM)		
			RVSMC (1.25 μM)		

			P-388D1 (4 μM)		
			Panc1 (8 μM)		
Indirubin	HZ	CDK1 (10 µM)	Effective inhibitory dose:	Full-thickness wounds	304,305
		CDK2 (2.2 µM)	HaCaT (100 nM)	were created on the dorsal	
	HN	CDK4 (12 µM)		skin of BALB/c mice,	
		CDK5 (5.5 µM)		which were then treated	
				with vaseline containing	
				DMSO (1%) or indirubin	
				(262.26 ng/g vaseline).	
Indirubin-5-	Na <sup>*</sup> SO <sub>3</sub> *	CDK1 (55 nM)	Effective inhibitory dose:	Human prostate xenograft	304
sulfonic acid.	8	CDK2 (35 nM)	HUVEC (25 µM)	mouse model-indirubin	
		CDK4 (300 nM)		solution was intralesional	
	н // о	CDK5 (65 nM)		injected at a dose of 10	
				mg/kg daily	
Staurosporine	H N VO	CDK1 (5 nM)	<i>IC</i> <sub>50</sub> :	Bladder T-24 xenograft	306-308
		CDK2 (7 nM)	Primary hOB (11.63 µM)	model-athymic nude mice	
		CDK4 (<10 µM)	U2OS (6.69 µM)	were treated with	
	H <sub>3</sub> CO	CDK5 (4 nM)	HOS (6 µM)	staurosporine 0.1 mg/kg	
	NHCH <sub>3</sub>		Saos-2 (6.16 µM)	daily by intraperitoneal	
			Τ-24 (0.029 μΜ)	injection	

			HL-60 (0.13 μM)		
			CEC (0.022 µM)		
JNJ-7706621	√−F	CDK1 (9 nM)	<i>IC</i> <sub>50</sub> :	A375 melanoma human	158,309
		CDK2 (4 nM)	WAC2	tumor xenograft model-	
	N N NH2		(10.6 µM)	JNJ-7706621 was	
			SHEP (1 µM)	administered to mice	
			NB69 (2.3 µM)	orally at 100 or 125	
			SK-N-FI (4.1 μM)	mg/kg daily	
			NLF (3.4 µM)		
			RH-41 (2.8 μM)		
			IMR-32 (1.3 µM)		
			IMR-32 LV-h-p53		
			(2.1 µM)		
			NGP (3.1 µM)		
			HDMB-3 (1.6 µM)		
Olomoucine		CDK1 (7 µM)	IC <sub>50</sub> :	HCT116 colon xenograft	310,311
		CDK2 (7 µM)	HL-60 (40 µM)	model-	
		CDK5 (3 µM)	Α2780 (30 μΜ)	Olomoucine was	
	H CH₃		A2780Cis <sup>R</sup> (45 μM)	administered in a volume	

CH1 (48 µM)	of 0.1 mL/10 g
CH1Cis <sup>R</sup> (78 µM)	bodyweight in 50 mmol/L
CH1Cox <sup>R</sup> (86 μM)	by IV
SKOV-3 (> 50 μM)	
BE (> 50 μM)	
HT29 (58 μM)	
Mawi (> 50 μM)	
Lovo (> 50 µM)	
SW620 (> 50 μM)	
HCT116 (52 μM)	
COLO-205 (41 µM)	
KM12 (90 μM)	
SA-OS2 (78 μM)	
U2-OS (101.5 µM)	
MCF 7 (64 µM)	
MB-MDA231 (101.5 μM)	
Α549 (54.5 μΜ)	
MOR (66 µM)	
HX147 (106.5 μM)	
CORL23 (54 µM)	

			GCT (27 µM)		
AT-7519		CDK1 (220 nM)	<i>IC</i> <sub>50</sub> :	Xenograft tumor model-	312,313
		CDK2 (44 nM)	HeLa (0.325 μM)	AT7519 was	
	N N H	CDK4 (67 nM)	ME-180 (0.599 μM)	administrated at 15 mg/kg	
		CDK5 (11 nM)		body weight once a day	
				for 5 days for a 2-week	
				duration by IV.	
R547	NH <sub>2</sub> Q F	CDK1 (0.001 µM)	<i>IC</i> <sub>50</sub> :	Xenograft models–R547	314
	N F	CDK2 (0.003 µM)	MDA-MB-468 (0.11 µM)	was administered at 40,	
	HN N O	CDK4 (0.001 µM)	MDA-MB-435 (0.08 µM)	60 or 80 mg/kg with	
	Ņ		MCF-7 (0.06 µM)	formulated as a	
	O=S=O		HCT116 (0.08 μM)	suspension in Klucel	
			SW480 (0.07 µM)	LF/Tween 80 or R547	
			RKO (0.05 μM)	was formulated as a	
			HT-29 (0.17 μM)	solution in	
			HCT15 (0.61 µM)	hydroxylpropyl h-	
			H460a (0.06 µM)	cyclodextrin, sodium	
			C33A (0.32 µM)	hydroxide, and water for	
			DU145 (0.08 μM)	IV injection at 3.75, 7.5,	
			OSA-CL (0.19 μM)	15, or 30 mg/kg.	
				10, 01 00 mg/kg.	

			LOX (0.05 µM)		
			JEKO-1 (0.08 µM)		
			REC-1 (0.09 µM)		
SU9516		CDK1 (0.04 µM)	<i>IC</i> <sub>50</sub> :	Duchenne muscular	315,316
		CDK2 (0.022 µM)	Α431 (2.2 μΜ)	dystrophy mouse model-	
	V N H		H460 (3.6 µM)	mice were administered 5	
			Colo-205 (2.6 µM)	mg/kg SU9516 by oral	
			RKO (5.6 μM)	gavage from 3 to 10	
			SW480 (6.4 µM)	weeks of age	
AZD5438		CDK1 (16 nM)	<i>IC</i> <sub>50</sub> :	BT474c, CoLo-205,	317
		CDK2 (6 nM)	MCF-7 (0.22 µM)	HX147, PC-3, SW620	
		CDK9 (20 nM)	MCF-7Adr (0.31 µM)	cell line xenografts-	
			MDA-MB-23 (0.46 µM)	The mice were orally	
			HCT-116 (0.32 µM)	administered	
			ΗCT-15 (1.13 μΜ)	AZD5438 at 25, 50, or	
			ΗΤ29 (1.05 μΜ)	100 mg/kg, daily	
			LoVo (0.63 μM)		
			SW620 (0.58 µM)		
			Colo-205 (0.7 µM)		
			Α549 (0.57 μΜ)		

			H322 (0.4 µM)		
			H460 (0.87 μM)		
			PC-3 (0.2 μM)		
			DU145 (0.42 µM)		
			A2780 (1.26 μM)		
			HeLa (1.1 μM)		
			IM-9 (1 µM)		
			MOLP-8 (1.1 µM)		
			AMO-1 (1 μM)		
			ARH-77 (1.7 μM)		
			KARPAS-620 (0.74 µM)		
			JJN-3 (0.94 µM)		
			L-363 (0.5 µM)		
CDKi277	n n h	CDK1 (0.008 µM)	<i>IC</i> <sub>50</sub> :	Colo205 and PC-3	318
	N N N N N N N N N N N N N N N N N N N	CDK2 (0.004 µM)	Daudi (0.171 µM)	xenograft models-mice	
		CDK5 (0.005 µM)	HL60 (0.129 μM)	were administered	
		CDK6 (0.051 µM)	K562 (0.438 µM)	CDKi277 at 25 or 12.5	
			Jurkat (0.14 µM)	mg/kg for 4 days on and 1	
			HS294 (0.221 μM)	day off, for three	
			Α375 (0.12 μΜ)		

			MiaPaca2 (0.116 µM)	consecutive cycles by IP	
			PC-3 (0.251 µM)	injection	
			SaSO2 -Rb (0.32 µM)		
			U2OS+Rb (0.28 μM)		
			Colo205 (0.278 µM)		
			Colo320 (0.241 µM)		
			HCT116 p53 <sup>+</sup> (0.29 μM,		
			HCT116 p53 <sup>-</sup> (0.31 μM)		
			MDA-MB-231 (0.137 µM)		
			MCF-7 p53(+) (0.18 µM)		
			MCF-7 p53(-) (0.14 µM)		
Dinaciclib		CDK1 (3 nM)	Effective inhibitory dose:	PDX mouse model-	319,320
		CDK2 (1 nM)	BT474 (10 nM)	Dinaciclib alone was	
	он ни	CDK5 (1 nM)	MCF7 (1 nM)	administered by	
		CDK9 (4 nM)	T47D (1 nM)	intraperitoneal injection at	
	ОН			a dose of 50 mg/kg/day, 5	
				days a week for total of 4	
				weeks	

Bohemine		CDK1 (3.89 µM)	<i>IC</i> <sub>50</sub> :	HCT116 xenograft	311,321
		CDK2 (4.6 µM)	Α2780 (12.5 μΜ),	model-	
		CDK9 (2.7 µM)	CH1 (21.5 µM)	Bohemine were given i.v.	
	1130 0113		CH1Cis <sup>R</sup> (22 µM)	in a volume of 0.1 mL/10	
			CH1Cox <sup>R</sup> (19 µM)	g bodyweight in 50 mM	
			SKOV-3 (81 μM)	HCl/saline at 50 mg/kg	
			BE (25 μM)		
			ΗΤ29 (27.7 μΜ)		
			Mawi (28.5 μM)		
			Lovo (25 μM)		
			SW620 (27 µM)		
			НСТ116 (17 μМ)		
			COLO-205 (21 µM)		
			КМ12 (32 μМ)		
			SA-OS2 (38 μM)		
			U2-OS (36 µM)		
			MCF 7 (21.5 µM)		
			MB-MDA231 (40 µM)		
			Α549 (20 μΜ)		
			MOR (36 µM)		

		HX147 (38 μM)		
		CORL23 (29.5 µM)		
		GCT (14.5 µM)		
CYC202	CDK1 (0.65 µM)	<i>IC</i> <sub>50</sub> :	HCT116 tumor	311,322
	CDK2 (0.7 µM)	A2780 (4.9 μM)	xenograft-mice were	
	CDK5 (0.16 µM)	A2780Cis <sup>R</sup> (8.4 µM)	orally administered	
	CDK7 (0.46 µM)	CH1 (7.7 µM)	CYC202 at 50, 500, or	
	CDK9 (0.6 µM)	CH1Cis <sup>R</sup> (9.3 µM)	2,000 mg/kg, daily	
		CH1Cox <sup>R</sup> (7.4 μM)		
		SKOV-3 (31 μM)		
		BE (17.5 μM)		
		HT29 (20.3 μM)		
		Mawi (18 μM)		
		Lovo (20 μM)		
		SW620 (23 µM)		
		HCT116 (6.9 μM)		
		COLO-205 (8.5 µM)		
		KM12 (15 μM)		
		SA-OS2 (16.5 μM)		
		U2-OS (15 μM)		

			MCF 7 (7.8 µM)		
			MB-MDA231 (15 µM)		
			A549 (9.3 µM)		
			MOR (12.5 µM)		
			HX147 (19 μM)		
			CORL23 (10.5 µM)		
			GCT (5.2 µM)		
3-ATA	s s	CDK1	Effective inhibitory dose:	NA	323,324
		(88 µM)	cerebellar granule neurons		
	V NH <sub>2</sub>	CDK2 (>100 µM)	(10 µM)		
		CDK4 (3.1 µM)			
Kenpaullone	H	CDK1 (0.4 µM)	Effective inhibitory dose:	NA	12,325
		CDK2 (0.7 µM)	MDA-MB-231 (5 µM)		
	Br	CDK5 (0.9 µM)			
NU 12050			10		326,327
NU2058	ν ν ν ν ν ν	CDK1 (26 µM)	IC <sub>50</sub> :	NA	520,527
		CDK2 (17 µM)	LNCaP (15 µM)		
	$H_2N$ N H		LNCaP-cdxR (17 µM)		
			LNCaP-AI (10 µM)		
			ΡC3 (38 μΜ)		

		DU145 (14 μM) CWR22Rv1 (16 μM)		
Purvalanol A	CDK1 (4 μM) CDK2 (35 μM)	<i>Effective inhibitory dose:</i> MCF-7 (25 μM)	NA	328-332
	CDK4 (850 μM) CDK5 (75 μM)	MDA-MB-231 (25 μM) HCT 116 (10 μM)		
		HT29 (5 μM) SW480 (5 μM)		
BMS-265246	CDK1 (6 nM) CDK2 (9 nM)	<i>IC</i> 50: Hep-3B (2.84 µM) Hep-G2 (1.73µM)	NA	333

Drug Name	Structure	Major Targets	Phase	Clinical Trial	Cancer type	Ref
				ID		
BEY1107		CDK1	1, 2	NCT03579836	Locally Advanced or Metastatic Pancreatic Cancer	162
Flavopiridol		CDK1/2/4/7/9	1, 2	NCT02520011	Leukemia, Multiple Myeloma,	163,164
	OH O			NCT00112723	Sarcoma, Gastrointestinal Stromal	
	но			NCT00005974	Tumor, Lymphoma, Solid Tumors	
	HO CI			NCT00098579		
	N N			NCT00007917		
				NCT00324480		
Roniciclib	NH NH	CDK1/2/3/4/7/9	1, 2	NCT01188252	Neoplasms, Lung Cancer	170,171
				NCT02047890		
	NH NN =			NCT01335256		
				NCT02161419		
	F F OH			NCT01573338		

P276-00		CDK1/4/9	1, 2	NCT00408018	Neoplasm,	173
	ОН			NCT00407498	Melanoma,	334
	но			NCT00835419	Breast Cancer,	13
	CI			NCT01333137	Multiple Myeloma,	174
	о́н о́ н−сі			NCT00824343	Advanced Head and Neck Cancer,	
				NCT00882063	Lymphoma	
Dinaciclib	-0, N+ ~	CDK1/2/5/9	1, 2,	NCT01783171	Pancreatic cancer,	176-178
			3, 4	NCT00732810	Non-Small-Cell Lung cancer,	
	HN			NCT03484520	Neoplasms,	
				NCT01624441	Leukemia, Breast cancer, Myeloma,	
	ОН			NCT01676753	Lymphoma,	
				NCT01096342	Melanoma	
				NCT02684617		
				NCT01711528		
				NCT01650727		
				NCT00937937		
				NCT01515176		
				NCT01434316		
				NCT01624441		

AT7519	$\square$	CDK1/2/4/6/9	1, 2	NCT00390117	Lymphoma,	181,182,312
				NCT01652144	Unspecified Adult Solid Tumor,	
				NCT01627054	Multiple Myeloma,	
	N/IN H			NCT01183949	Lymphoma	
Seliciclib		CDK1/2/5/7	1,2	NCT00999401	Non-small Cell Lung Cancer, Breast	335
				NCT00372073	Cancer, Advanced Solid Tumors	
	NH			NCT01333423		
AG-024322	N H	CDK1/2/4	1	NCT00147485	Neoplasms,	13
	F N N-NH				Non-Hodgkin Lymphoma,	
PHA-793887		CDK1/2/4/5	1	NCT00996255	Advanced/Metastatic Solid Tumors	334
	× N					
R547		CDK1/2/4/7	1	NCT00400296	Neoplasms	13
	0=\$=0					

RGB-286638	H-CI H-CI	CDK1/2/3/4/5/9	1	NCT01168882	Hematological Malignancies	
AZD-5438		CDK1/2/9	1	NCT00088790	Neoplasms	
Indirubin		CDK1/2/4/5	4	NCT02200978	Childhood Acute Promyelocytic Leukemia	336