

Mechanisms of cisplatin sensitivity and resistance in testicular germ cell tumors and potential therapeutic agents (Review)

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Abstract. Testicular germ cell tumors (TGCTs) are the most common tumors in men aged 20-40 years and are primarily treated with cisplatin-based drugs. Although TGCTs are highly sensitive to DNA damage induced by cisplatin and show a hypersensitive apoptotic response, cisplatin resistance still exists. Emerging evidence shows that cisplatin resistance in TGCTs is mainly related to the inhibition of apoptotic pathways such as MDM2/p53, OCT4/NOXA, PDGFR/PI3K/AKT, inhibition of cell cycle checkpoints, increased methylation or neddylation and DNA repair balance. In this review, recent advances regarding the mechanisms of TGCTs' sensitivity and resistance to cisplatin were summarized and potential therapeutic agents for cisplatin-resistant TGCTs were presented, providing a new therapeutic strategy for drug-resistant TGCTs.

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1. Introduction

Testicular germ cell tumors (TGCTs) are a heterogeneous group of tumors that occur primarily in children and young men and are the most common tumor types in men aged 20-40 years. Based on histology, TGCTs are classified as seminoma (SE-TGCT) or non-seminoma (NSE-TGCT) tumors (1,2). The latter can be further classified as undifferentiated embryonal carcinoma (EC), choriocarcinoma, yolk sac tumor, differentiated teratoma or mixed tumors (a mixture of two or more components) (3-5). TGCTs develop from precancerous germ cell tumors in the renal tubule, mainly because of the failure of lymphocytes to mature properly during fetal or postnatal development. It progresses to aggressive TGCTs such as seminomas and non-seminomas after puberty, and environmental and genetic risk factors are significant factors in the susceptibility to TGCTs (6). Studies have shown that most testicular germ cell tumors originate from germ cell neoplasia *in situ*, which is thought to be due to the stagnation and transformation of the original germ cells. Seminomas have the same characteristics as germ cells or primordial germ cell formation *in situ*, whereas non-seminomas exhibit differential differentiation. Neoplasms and embryonic cell carcinomas are pluripotent and are thought to be responsible for the histological heterogeneity and mixed pathology of testicular germ cell tumors (7).

Platinum-based therapies are often used as first-line treatment for TGCTs in children and adults. TGCTs have a very high cure rate and are highly sensitive to cisplatin chemotherapy (8,9). The sensitivity of TGCTs to cisplatin may correlate with two key reactions: Inadequate repair of cisplatin-induced DNA damage and a hypersensitive apoptotic response (10). Various sources of endogenous and exogenous damage constantly assault the genome, among which DNA double-strand breaks (DSBs) are the most cytotoxic DNA lesions (11-13). To maintain chromatin stability, cells have developed a complex system of biochemical pathways called the DNA damage response (DDR) (14). DNA repair mechanisms have an indispensable role in cisplatin-induced cytotoxicity. DNA repair systems, including nucleotide excision repair (NER), homologous recombination (HR), nonhomologous end joining (NHEJ) and mismatch repair (15).

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For metastatic disease, the overall disease-free survival rate of TGCTs is ~80%. Even in patients with advanced metastatic disease, complete remission can be achieved with systemic therapy and secondary removal of residual mass. However, ~10-20% of advanced TGCTs are resistant to platinum-based chemotherapy and have a poor prognosis or recurrence, and the treatments for these patients are poorly understood. Studies have shown that cisplatin resistance is mainly related to the apoptotic pathway, tumor cell cycle regulation-related factors, DNA methylation and DNA damage repair pathways. Among them, the regulated death pathway is mainly related to murine double minute 2 (MDM2)/p53, octamer-binding transcription factor 4 (OCT4)/phorbol-12-myristate-13-acetate-induced protein 1 (NOXA), platelet-derived growth factor receptor (PDGFR)/PI3K/AKT and insulin-like growth factor 1 receptor (IGF1R)/AKT, and certain related microRNA (miRNA) pathways. The main cell cycle regulatory factors were cyclin-dependent kinase (CDK)4/6 inhibitors and CDK2 inhibitor p21. During DNA methylation, hypermethylation can promote cisplatin resistance; therefore, since vir-like N6-methyladenosine (m⁶A) methyltransferase associated (VIRMA) is positively correlated with m⁶A in TGCTs, it stimulates the methylation process and thus cisplatin resistance. However, the DNA hypomethylating agent 5-azacytidine (5-aza) and the demethylating agent guadecitabine (SGI-110) promoted cisplatin sensitivity and inhibited cisplatin resistance. In the process of DNA damage repair, cisplatin resistance is positively correlated with NHEJ inhibition and increased HR, and a decrease in DNA damage response protein p53-binding protein 1 (53BP1) and DNA-dependent protein kinases (DNA-PKCs) can inhibit NHEJ and promote HR, thus promoting cisplatin resistance.

This review describes the sensitivity of TGCTs to cisplatin (also referred to as CDDP), the underlying mechanisms related to CDDP and the potential therapeutic agents for cisplatin-resistant TGCTs.

2. Molecular mechanisms involved in the sensitivity of TGCTs to cisplatin

In cisplatin-sensitive TGCT cells, cisplatin interacts with the DNA base to produce different forms of cisplatin-DNA adducts, of which in-chain crosslinking (>90%) is the dominant DNA adduct (15,16). The NER pathway can repair intra-chain crosslinking that can destroy DNA structures induced by cisplatin, and intra-chain crosslinking can be recognized by proteins associated with DNA repair (17). Certain proteins also act as inhibitors. For instance, high-mobility-group (HMG) proteins can inhibit NER by binding to DNA adducts. HMG box protein 4 (HMGB4) enhances the sensitivity of TGCTs to cisplatin by blocking the excision repair of cisplatin-DNA adducts (18). Furthermore, numerous essential proteins involved in NER have low expression levels in TGCTs, including xeroderma pigmentosum group A (XPA), xeroderma pigmentosum group F (XPF) and excision repair cross-complementing 1 (ERCC1) (19). A study showed that high levels of XPF and ERCC could decrease the sensitivity to cisplatin in TGCT cell lines, indicating that XPF and ERCC are rate-limiting factors for the repair of cisplatin-DNA cross-links in TGCTs,

which contributes to remarkable cisplatin sensitivity (20). Furthermore, homologous recombination repair defects of DSB also cause high cisplatin sensitivity in TGCT cell lines (18). Of note, a related study showed that compared to normal testes, poly(ADP-ribose) polymerase (PARP) is overexpressed in TGCT and is involved in the repair of DNA single-strand breaks by base excision repair (21). Meanwhile, PARP inhibitors can promote the response of resistant EC cells to cisplatin by reducing their ability to repair damage, providing a potential therapeutic approach for resistant TGCTs (22).

The high sensitivity of TGCTs to cisplatin is due to the various apoptotic pathways induced by cisplatin, in which wild-type p53 plays a major role (23). p53 is frequently mutated in cancer as a tumor suppressor, but is not mutated in TGCTs, retains its wild-type configuration and is activated following exposure to chemotherapeutic agents (8). Cisplatin treatment increases the expression of p53 transcriptional target FAS death receptors and subsequently activates exogenous apoptotic pathways through the interaction of FAS with its ligand (24,25). It has also been shown that cisplatin treatment can upregulate the pro-apoptotic protein p53 upregulated modulator of apoptosis (PUMA) and NOXA (26). As a target protein of p53, NOXA is a Bcl-2 family protein that belongs to a subclass of BH3-only proteins that induce apoptosis via p53-dependent and/or p53-independent mechanisms (27), thus playing a crucial role in cisplatin-induced apoptosis of TGCTs cells (28). In addition, high expression of the pluripotent factor OCT4 (also known as POU5F1) is positively correlated with NOXA, leading to a hypersensitive apoptotic reaction in EC cells (23). OCT4 is also related to the regulation of p21 expression and p21 is associated with cisplatin sensitivity both *in vivo* and *in vitro* (28). Furthermore, crosstalk between exogenous and intrinsic apoptotic pathways may further enhance the apoptotic response (10) (Fig. 1).

3. Mechanisms of cisplatin resistance in TGCTs

Cisplatin resistance is a major clinical challenge and a thorough understanding of the mechanisms underlying the development of TGCT resistance will help improve the efficacy of TGCT therapy. Several mechanisms of cisplatin resistance have been identified, mainly related to apoptotic pathways, tumor cell cycle regulation (Fig. 2), methylation and DNA repair systems (Fig. 3). Specific mechanisms are outlined and potential treatments that currently exist are discussed in this chapter.

Apoptotic pathways

MDM2/p53 pathway. Signal changes in response to apoptotic transduction pathways mediated by DNA damage are important factors in cisplatin resistance in TGCTs. Overexpression of anti-apoptotic factors as well as decreased expression or dysfunction of pro-apoptotic factors can alter the induction of apoptosis. MDM2, a ubiquitin ligase, mediates ubiquitination and degradation of p53, inhibiting apoptosis and cell-cycle arrest for DNA repair (10,24) and recruit transcriptional co-repressors of p53 (24). Loss of p53 function results in a lack of cell-cycle regulation and contributes to more aggressive tumors and chemotherapy resistance (29). p53 is a tumor suppressor and inactivation of p53 at the genetic or protein level is ascribed to cisplatin resistance in

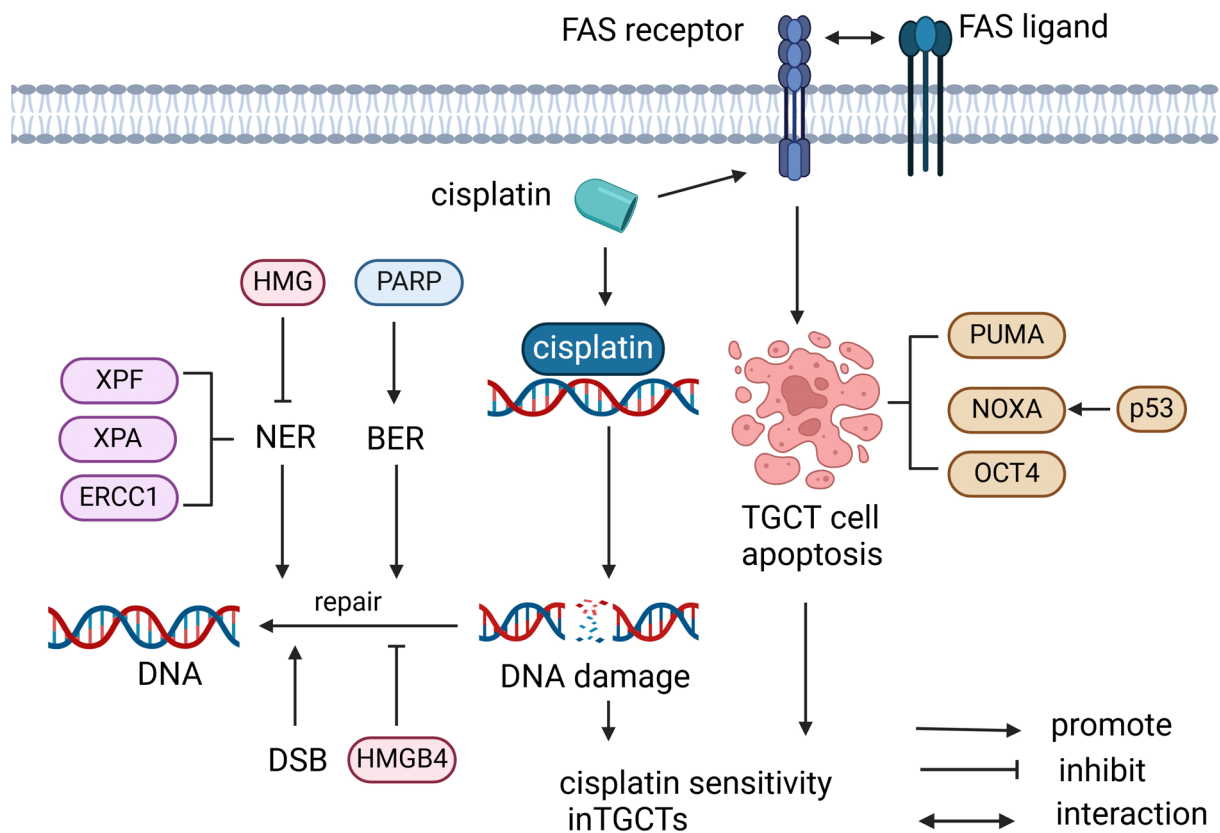


Figure 1. Mechanism of cisplatin sensitivity of TGCTs. DNA damage and tumor cell apoptosis can promote TGCT sensitivity to cisplatin. On the one hand, NER, BER and DSB can repair DNA damage. ERCC1, XPF and XPA are the main components of NER, HMG can inhibit NER and PARP can promote BER. However, HMGB4 can inhibit damage repair. On the other hand, FAS ligand-receptor interaction promotes apoptosis of TGCT cells. Furthermore, the upregulation of p53, PUMA, OCT4 and NOXA can also promote the apoptosis of TGCT cells, thus promoting the sensitivity of TGCTs to cisplatin. TGCTs, testicular germ cell tumors; NER, nucleotide excision repair; BER, base excision repair; DSB, DNA double-strand break; ERCC1, excision repair cross-complementing group 1; XPF, *Xeroderma pigmentosum* complementation group F; PARP, poly(ADP-ribose) polymerase; HMGB4, high mobility group box-4; PUMA, pro-apoptotic protein p53 upregulated modulator of apoptosis; OCT4, octamer-binding transcription factor 4; NOXA, phorbol-12-myristate-13-acetate-induced protein 1.

numerous tumor types, particularly in a subset of refractory TGCTs (28). In addition, p53 regulates MDM2 expression by binding to and positively regulating its promoter. More than 17% of human tumors exhibit MDM2 gene amplification, leading to cisplatin resistance (30). Compared with cisplatin-sensitive tumors, chemotherapy-resistant tumors showed a higher positive tendency for p53 and MDM2. However, inhibition of p53 activity was mediated by MDM2. A study on cisplatin-sensitive and resistant testicular carcinoma (TC) cell lines showed that the interaction between p53 and MDM2 requires higher doses of cisplatin to be disrupted in resistant cell lines (24). Besides, it has been shown that the reduced p53-induced apoptotic response detected in different TGCT cell lines in their relative cisplatin resistance is different, which is associated with the formation of MDM2-p53 complex (28).

OCT4/NOXA pathway. OCT4, a key stem cell transcription factor, is highly expressed in TGCTs and pluripotent stem cells (31,32). However, OCT4 and NOXA are underexpressed in cisplatin-resistant TGCT cells (28). OCT4 is a key regulator of cisplatin resistance (33). Knockdown of OCT4 reduces NOXA transcript levels, leading to the loss of NOXA protein, which decreases cisplatin hypersensitivity (23). In addition, it

has been indicated that hypoxia induces resistance to TGCT, which may be related to the hypoxia-induced downregulation of OCT4. Under hypoxic conditions, overexpression of SUMO1 increases the sumoylation of OCT4, thereby reducing OCT4 protein stability (34).

PDGFR/PI3K/AKT pathway. Several studies have demonstrated that the PDGFR/PI3K/AKT signaling pathway has an important role in cisplatin resistance. PDGF has been shown to bind and phosphorylate its receptor PDGFR, subsequently activating the downstream PI3K/AKT pathway (35-38). In the activation of the PDGFR/PI3K/AKT signaling pathway, in which activated PI3K is recruited to the plasma membrane and phosphorylates phosphatidylinositol 4,5-diphosphate (PIP2), thus producing the second messenger phosphatidylinositol 3,4,5-triphosphate (PIP3), which can activate the serine/threonine kinase AKT through 3-phosphoinositide dependent protein kinase-1 (PDPK1) (39-41). Of note, this signal transduction process can be inhibited by phosphatase and tensin homolog deleted on chromosome 10 (PTEN) (42). Di Vizio *et al* (43) found that the tumor suppressor gene *PTEN* is widely expressed in germ cells and germ-cell neoplasia *in situ* but is virtually absent in 56% of seminomas, 86% of embryonal carcinomas and all teratomas. In cisplatin-resistant

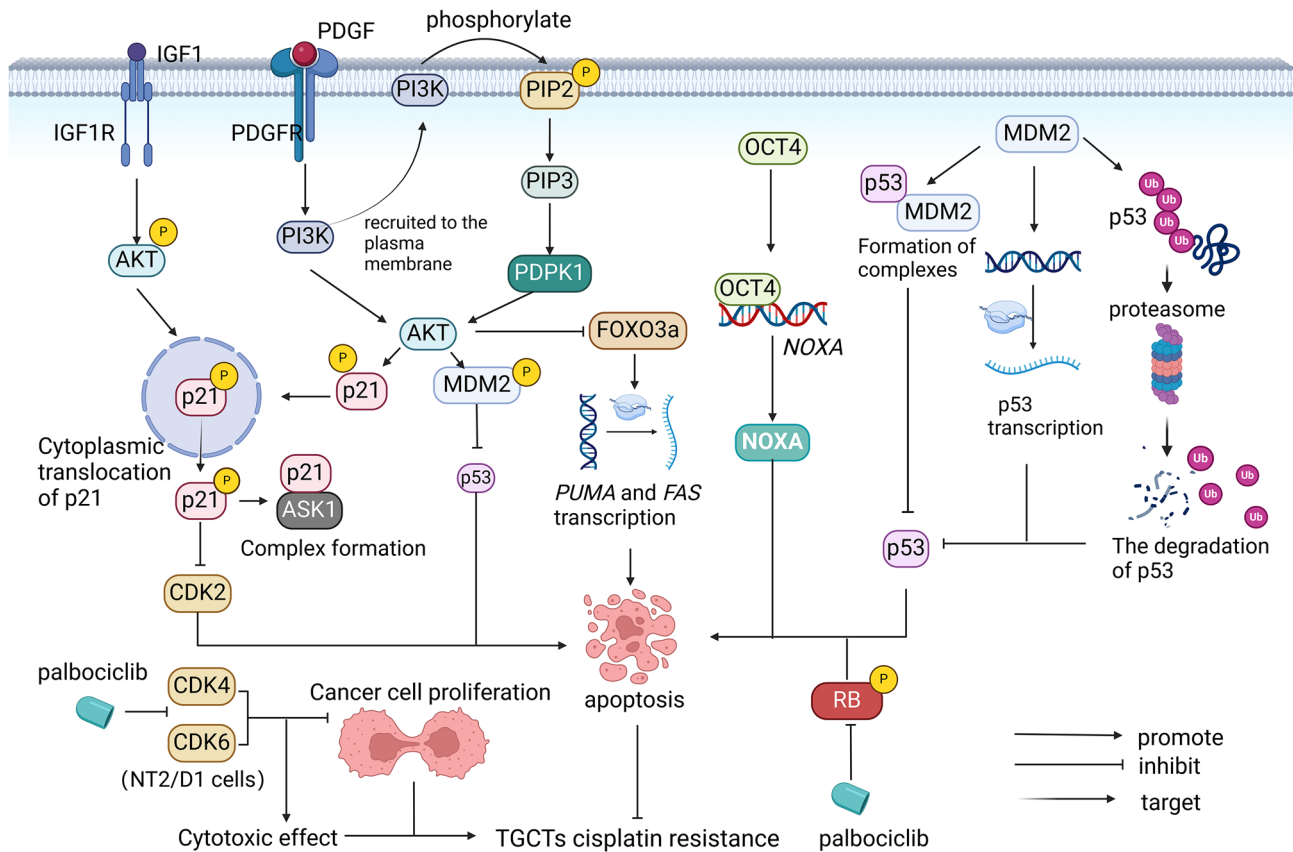


Figure 2. Mechanism of cisplatin resistance of TGCTs by influencing apoptosis and tumor cell cycle regulation. MDM2 can form a complex with p53 and the upregulation of MDM2 can promote the ubiquitination and degradation of p53 and the formation of p53 transcription inhibitors, all of which lead to the downregulation of p53. The interaction between PDGF and PDGFR could enhance the recruitment of PI3K to the cytoplasm membrane, resulting in the phosphorylation of PIP2 and the formation of PDKP1 in PIP3, thus contributing to the phosphorylation of AKT. The IGF1R pathway also promotes AKT phosphorylation. The increase of AKT phosphorylation may promote the phosphorylation of MDM2 leading to the ubiquitination and degradation of p53 and triggering the cytoplasmic translocation of p21, inhibiting the expression of FOXO3a. All of the above processes inhibit apoptosis and promote TGCT resistance to cisplatin. Palbociclib, an inhibitor of CDK4/6, reduces the phosphorylation of RB, thereby promoting cisplatin resistance. Palbociclib also increased CDK4/6 in NT2/D1 cell lines, promoting cancer cell proliferation and inducing cisplatin resistance. Phosphorylation of AKT promotes the transfer of p21 from nuclear ectopic space to the cytoplasm, while high cytoplasmic p21 expression promotes the formation of complexes between p21 and ASK1, reducing the expression of ASK1, thereby inhibiting the regulated death of TGCT cells and promoting cisplatin resistance. TGCTs, testicular germ cell tumors. MDM2, murine double minute 2; PDGFR, platelet-derived growth factor receptor; PIP2, phosphatidylinositol 4,5-diphosphate; PDKP1, 3-phosphoinositide dependent protein kinase-1; IGF1R, insulin-like growth factor 1 receptor; FOXO3, forkhead box O3; RB, retinoblastoma protein.

TGCTs cells, AKT is overactivated due to increased levels of PDGFRb and PDGF-b ligands at both the mRNA and protein levels. Subsequently, the PDGFR/PI3K/AKT pathway is over-activated, leading to the phosphorylation of p21 and its cytoplasmic accumulation, while increasing the phosphorylation of MDM2 which leads to the inhibition of P53-mediated apoptosis (44,45). Furthermore, phosphorylation of AKT directly inhibits forkhead box O3 (FOXO3a), which is responsible for the transcription of pro-apoptotic proteins such as PUMA and FAS ligands; thus, AKT indirectly negatively affects apoptosis (46).

IGF1R/AKT pathway. IGF1R has been implicated in numerous carcinogenic processes and can activate anti-apoptotic proteins and improve cell survival by promoting the PI3K/AKT pathway. Furthermore, the MAPK (Ras/Raf/MEK/ERK) pathway can be activated to promote proteins involved in the cell cycle and drive cell proliferation (47). IGF1R signaling is also associated with numerous cellular processes that promote metastasis. For instance, it is involved in the integrin pathway to induce RhoA-dependent

movement through focal adhesion kinase and receptor for activated C kinase 1, and is associated with TGCT survival and migration. A study has shown that IGF1R is highly expressed in TGCT cell lines and *IGF1R* knockdown can reduce the growth of non-seminoma cells, leading to apoptosis (48). In cisplatin-resistant TGCTs, IGF1R levels are upregulated and the upregulation of IGF1R could promote the overexpression of AKT and phospho-AKT, leading to increased translocation of p21 in the cytoplasm, thereby inhibiting apoptosis and promoting cisplatin resistance (49). However, *IGF1R* silencing leads to decreased phosphorylated AKT levels, which allows the translocation of p21 to the cytoplasm for transfer to the nucleus, thereby restoring cisplatin sensitivity. Therefore, inhibition of the IGF1R in combination with chemotherapy may promote re-sensitization in the treatment of chemotherapy-resistant diseases (47).

Factors associated with tumor cell cycle regulation

CDK4/CDK6. CDKs are key enzymes that control cell cycle progression, and CDK4/6 are targets of cell cycle checkpoint

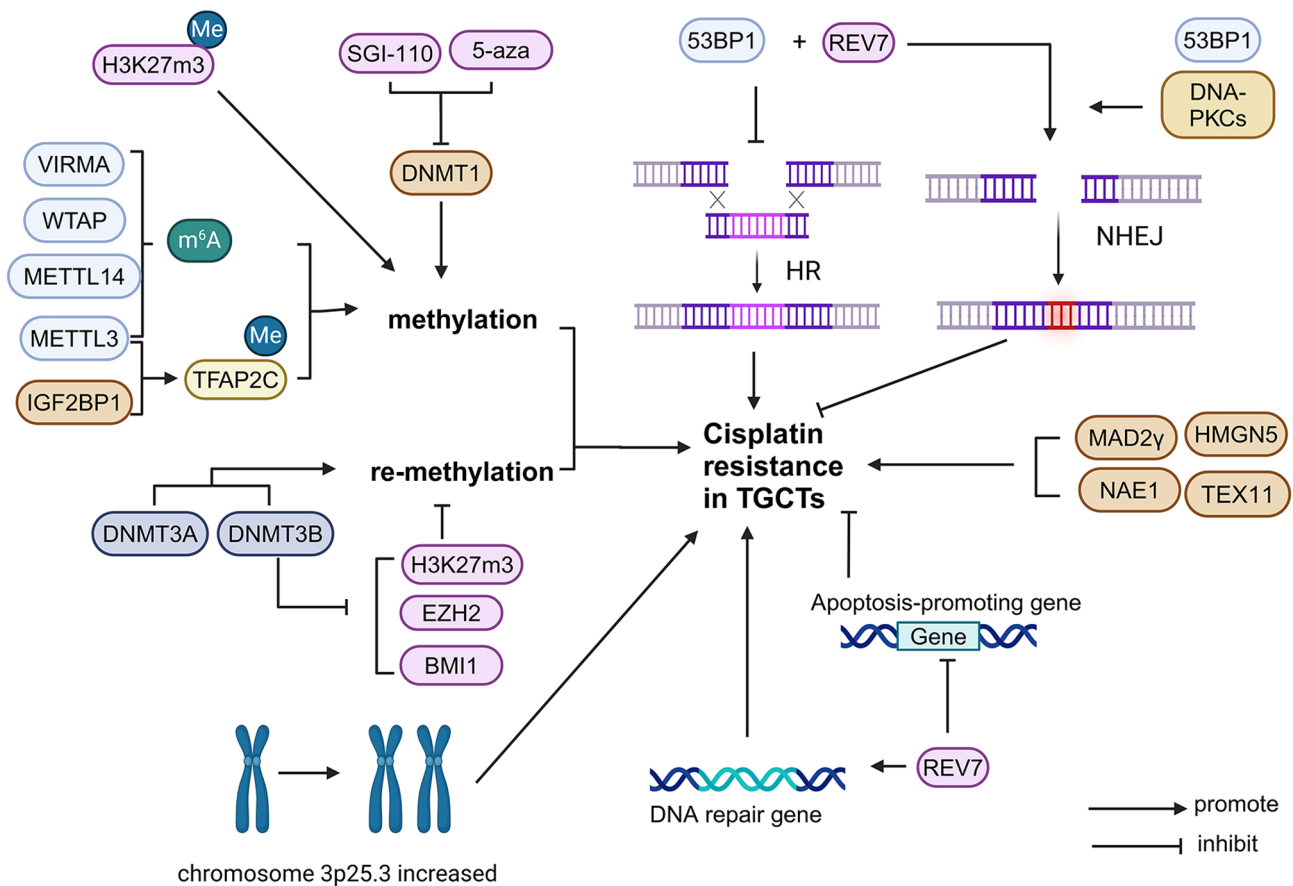


Figure 3. Mechanism of cisplatin resistance of TGCTs by influencing methylation, DNA repair systems and others. In terms of methylation, VIRMA WTAP, METTL3, METTL4 and IGF2BP1 can promote the increase of protein methylation levels, thus promoting TGCTs of cisplatin resistance, and methylation inhibitors such as 5-aza and SGI-110 reduce methylation levels, thereby reducing cisplatin resistance. Furthermore, DNMT3A/B and other remethylation-promoting enzymes can promote cisplatin resistance and DNMT3B can inhibit methylation inhibitors such as H3K27m3. In DNA repair systems, HR upregulation and NHEJ downregulation promote cisplatin resistance. By preventing DNA repair away from HR, REV7 and 53BP1 work together to promote NHEJ, both leading to inhibition of cisplatin resistance. However, REV7 upregulation can promote the expression of DNA repair-related genes and inhibit the expression of apoptosis genes, leading to the promotion of drug resistance. In addition, the increase of proteins such as MAD2γ and 3P2.3 chromatin also promoted the development of drug resistance. TGCTs, testicular germ cell tumors; VIRMA, vir-like N6-methyladenosine (m⁶A) methyltransferase associated; WTAP, Wilms' tumor 1-associating protein; METTL3, methyltransferase-like protein 3; IGF2BP1, IGF2 mRNA-binding protein 1; 5-aza, 5-azacytidine; SGI-110, demethylating agent guadecitabine; DNMT3A, DNA methyltransferase 3A; H3K27m3, histone H3 lysine 27; NHEJ, nonhomologous end joining; 53BP1, p53-binding protein 1.

inhibitors such as palbociclib (50). It has been shown that CDK4/6 inhibitors can induce cell cycle arrest and apoptosis in different types of TGCTs, indicating that CDK4/6 inhibitors have therapeutic potential for TGCTs. Palbociclib has a concentration-dependent inhibitory effect on TGCT cell survival and can induce apoptosis by decreasing the phosphorylation of retinoblastoma protein (RB). However, palbociclib can induce a rapid increase in CDK4/6 protein levels in NT2/D1 cells, a TGCT cell line, thus limiting its cytotoxic effects. It was demonstrated that the NT2/D1 cell line could overcome the effect of palbociclib and develop resistance, but the specific mechanism remains elusive and requires further investigation (51).

p21. A previous study showed that high cytoplasmic expression of the CDK inhibitor p21 is a critical factor in cisplatin resistance in testicular EC. Nucleic p21 controls the cell cycle and DNA replication, whereas cytoplasmic p21 is involved in the inhibition of apoptosis. In most patients with refractory testicular cancer, cytoplasmic p21 is highly expressed and high cytoplasmic p21 can form complexes with CDK2 and

apoptosis signal-regulating kinase 1 (ASK1). The formation of complexes inhibits the pro-apoptotic functions of CDK2 and ASK1, thereby protecting EC cells from cisplatin-induced apoptosis, leading to cisplatin resistance. Phosphorylated (P-) AKT-mediated p21 phosphorylation is critical for the localization of p21 in the cytoplasm, whereas p-AKT inhibition (LY294002) promotes p21 translocation to the nucleus, thus reducing p21-CDK2 complex formation and sensitizing EC cells to cisplatin in a CDK2-dependent manner (45).

Different types of TGCT sensitive to cisplatin have different p21 expression levels. For instance, p21 is almost undetectable in cisplatin-sensitive seminomas and EC but is significantly higher in cisplatin-sensitive choriocarcinoma and mature teratoma than in cisplatin-resistant ones (45). In mature teratomas, intrinsic chemotherapy resistance is associated with the upregulation of various factors that affect p21-induced cell cycle arrest (52).

Cisplatin is a highly cytotoxic drug that is effective against EC by inducing apoptosis in numerous cell types. Cisplatin treatment of EC cells increased p53 and MDM2 levels,

activated the Fas apoptotic pathway and induced apoptosis; however, the expression level of p21 was almost unaffected. However, in EC cells, gamma irradiation increased the levels of p53 and MDM2 and significantly induced cytoplasmic p21 without inducing cell cycle arrest or apoptosis. These results suggested that cytoplasmic p21 has an important role in preventing DNA damage-induced apoptosis of EC cells (45). Studies have also shown that high cytoplasmic p21 expression and cisplatin resistance of EC/TC are negatively associated with the expression of OCT4 and miR-106b but positively associated with p53 and MDM2 (28,45,53). Therefore, targeting cytoplasmic p21 may provide a new strategy for the treatment of chemotherapy-resistant testicular cancer (45) (Fig. 2).

Role of epigenetic mechanisms. Epigenetics plays an important role in the development and classification of TGCTs (54). DNA methylation is one of the major epigenetic modifications that have a key role in cisplatin resistance in TGCTs (55,56). Seminomas are undifferentiated tumors that are hypomethylated and exhibit low cisplatin resistance (18). More differentiated non-seminomas have higher DNA methylation levels (18). Embryonic cancers have moderate levels of DNA methylation and are sensitive to cisplatin, but may also acquire cisplatin resistance (18). Teratomas, choriocarcinomas and yolk sac tumors have the highest levels of DNA methylation, which is closely related to cisplatin resistance (18). A study has shown that global remodeling of DNA methylation is a critical factor that mediates TGCT resistance and cisplatin hypersensitivity (56).

m⁶A. m⁶A is the most abundant of modifications in mRNA, and this process includes ‘writers’, ‘erasers’ and ‘readers’ of m⁶A methylation, proteins that can add, remove or recognize m⁶A-modified sites, respectively, and alter biological processes accordingly (57). It can be introduced into the mRNA by different families of enzymes. They are assembled to form methyltransferase complexes that participate in methylation modification, mainly by the catalytically active component Methyltransferase-like protein 3 (METTL3) and other cofactors that recruit the complex, such as VIRMA, Wilms’ tumor 1-associating protein (WTAP) and METTL14. Various m⁶A-related proteins have been shown to be differentially expressed in TGCT subtypes (as biomarkers for the disease) and in cisplatin-resistant and cisplatin-sensitive TGCT cells. VIRMA and METTL3 were found to be more important. In the SE subtype of TGCTs, VIRMA is significantly upregulated at the mRNA and protein levels, promoting the progression of multiple malignancies by regulating cell cycle progression, migration, invasion, apoptotic resistance and tumor growth in an m⁶A-dependent manner. Furthermore, VIRMA introduced m⁶A modifications to TGCTs and contributed to tumor aggressiveness and cisplatin resistance in TGCT cells both *in vitro* and *in vivo* by modulating the DDR (58).

METTL3 and IGF2 mRNA-binding protein 1 (IGF2BP1) play key roles in enhancing cisplatin resistance in seminoma by enhancing transcription factors to activate m⁶A methylation of enhancer binding protein 2C (TFAP2C).

METTL3, as the ‘writer’ of m⁶A, enhanced the m⁶A methylation level of TFAP2C mRNA and improved the stability of TFAP2C mRNA, thus enhancing the viability of TGCT-resistant cells under cisplatin treatment. Furthermore,

IGF2BP1 functions as a TFAP2C m⁶A ‘reader’, enhances TFAP2C mRNA stability and is involved in cisplatin resistance (59).

Regulatory mechanisms of miRNAs in TGCTs. Of note, in addition to the signaling pathways mentioned above, miRNAs also play a critical role in oncogenesis in TGCTs by interacting with different mechanisms. In general, miRNAs are involved in the p53, epidermal growth factor receptor, Wnt/β-catenin and PTEN/AKT/mTOR pathway [see ref. (60) to review the regulatory mechanisms of miRNAs in TGCTs]. In addition, miRNAs play different roles in CDDP-resistant and -sensitive TGCT cells. On the one hand, miR-302a and miR-106b mainly inhibit the resistance to cisplatin by targeting p21, while miR-371-373 increases cisplatin resistance by targeting p53. By contrast, miR-106b-5p reverses resistance to cisplatin by targeting testis development related 1. miR-383 can improve cisplatin sensitivity by targeting histone H2AX, the adaptor protein of PP1 phosphatase and CDK4. miR-514a-3p decreases cisplatin resistance by targeting paternally expressed gene 3. Furthermore, miR-27b-3p, miR-31-5p, miR-125b-5p, miR-218-5p, miR-199a-5p and miR-324 were upregulated in cisplatin-resistant TCGT cells, while miR-374b-5p, miR-320a, miR-20b-5p, miR-375-5p, miR-17-5p, miR-106a, miR-378a-3p and miR-30e-3p were downregulated in cisplatin-resistant TCGT cells. However, miR-378a-3p was downregulated only in the CDDP-resistant TGCT cell line 1411HP, and miR-30e-3p was only down-regulated in the CDDP-resistant TGCT cell line 1777NRpmet (Table I).

DNA repair systems

The balance of DNA repair pathways. Cisplatin can cause several types of DNA damage, including inter-strand cross-linking (ICLs) (61). ICLs covalently join two strands of a DNA double-strand, blocking basic cellular processes such as DNA replication and leading to cell death. Cisplatin can also induce ICL-associated formation of DSB, which can be repaired by HR during the S/G2 phase of the cell cycle using sister chromatids as a repair template. One mechanism of TGCT resistance to cisplatin is its enhanced ability to repair DSB through HR (62).

In addition to HR, NHEJ is also involved in the restoration process and is the main DSB repair pathway throughout the cell cycle, accounting for almost all DSB repair, except in the S and G2 phases. NHEJ relies on Ku binding to the end of DNA, thereby improving the affinity of the NHEJ enzyme component consisting of polymerase (Pol μ and Pol λ), nuclease (Artemis·DNA-PKcs complex) and ligase (XRCC4-like factor-X-ray repair cross-complementing protein 4-DNA ligase IV complex), promoting DNA repair (63). NHEJ acts throughout the cell cycle, mediating the direct reconnection of DSB ends in an error-prone manner (64). In the process of repair, HR and NHEJ compete with each other to repair DSB, and the choice of repair pathway depends on the balance of pro-NHEJ and pro-HR factors at the DSB site. Key pro-NHEJ factors include 53BP1, the ATP-dependent DNA helicase 2 subunit KU70/KU80 and DNA-PKcs. 53BP1 inhibits extensive excision of DNA ends by assembly on DSB-side chromatin, repairing far away from HR, whereas KU70/KU80 and DNA-PKcs are involved in the later steps of DSB blunt

Table I. Effect of miRNA on cisplatin resistance or sensitivity in TGCT cells.

miRNAs	Target	Mechanism	(Refs.)
miR-302a	p21	Improve the sensitivity of cisplatin, up-regulated in TGCTs cisplatin-resistant cells	(147)
miR-106b-5p	TDRG1	Reverses the resistance to cisplatin	(148)
miR-383	H2AX, PNUTS, CDK4	Improve the sensitivity of cisplatin	(6,149)
miR-106b	p21	Inhibits the resistance to cisplatin	(45)
miR-514a-3p	PEG3	Decreases the resistance to cisplatin	(150,151)
miR-371-373	p53	Increase cisplatin resistance	(150)
miR-27b-3p, miR-31-5p, miR-125b-5p, miR-218-5p and miR-199a-5p	-	Up-regulated in TGCTs cisplatin-resistant cells	(152)
miR-374b-5p, miR-320a, miR-20b-5p miR-375-5p, miR-17-5p and miR-106a	-	Down-regulated in TGCTs cisplatin-resistant cells	(28,152)
miR-324	-	Up-regulated in metastatic CDDP resistant TGCTs cell lines (1411HP and 1777NRpmet)	(152)
miR-378a-3p	-	Down-regulated only in CDDP resistant TGCTs cell line 1411HP	(152)
miR-30e-3p	-	Down-regulated only in CDDP resistant TGCTs cell line 1777NRpmet	(152)

miRNA/miR, microRNA; TGCTs, testicular germ cell tumors; TDRG1, testis development related 1; PNUTS, the adaptor protein of PP1 phosphatase; PEG3, paternally expressed gene 3; CDDP, cisplatin.

end ligation. Cisplatin resistance was found to be positively associated with NHEJ inhibition and an increase in HR, and both often occur together (65).

Cisplatin-resistant cell lines have an NHEJ-less phenotype, with a decreased basal expression of 53BP1 and DNA-PKC proteins, thus forming fewer 53BP1 lesions after cisplatin treatment. Similarly, reduced expression of DNA-PKCs and 53BP1 inhibited the repair of DSB through the action of NHEJ, but could be effectively repaired by HR. Therefore, inhibition of 53BP1 protein expression and DNA-PKC activity can antagonize cisplatin cytotoxicity (3).

XPA. NER is an important DNA repair pathway that promotes ICL repair by mediating lesion recognition and incision through >30 DNA repair-associated proteins. Among these, the XPA protein has a pivotal role in DNA damage verification and assembly of the NER incision complex (66). One related study has shown that XPA expression is inversely associated with the prognosis of patients with TGCT and that increased co-expression of ERCC1, XPF and XPA is also associated with unfavorable overall survival. In primary TGCTs, increased XPA expression, and, to a lesser extent, NER co-expression, may promote increased DNA repair capacity; therefore, XPA may be a new promising therapeutic target in TGCTs (67).

REV7. REV7 (also known as MAD2L2, MAD2B or FANCV) is a HORMA protein and is regulated through stable structural rearrangement (68). Under physiological conditions, endogenous and exogenous agent-induced DNA damage is commonly repaired using the DNA repair system (DDR).

If the damage remains unrepaired, the damage-tolerance system is activated. Translesion synthesis (TLS) is a major mechanism underlying DNA damage tolerance (69). REV7 is highly expressed in human TGCT tissues (70) and is a multi-functional protein involved in TLS, DNA repair, cell cycle regulation, gene expression and histone modification through protein-protein interactions. REV7 plays a critical role in the DSB repair pathway. Mechanistically, REV7 and 53BP1 promote non-homologous terminal junctions at DSB sites, thereby promoting DNA damage repair. Knockdown of REV7 could lead to the downregulation of DNA repair-associated genes, as well as cell cycle checkpoint genes in TCGT cells, while promoting the transcription of pro-apoptotic genes. Therefore, REV7 inactivation antagonizes the cisplatin resistance induced by DNA damage-associated DSB accumulation. Collectively, REV7 loss contributes to the effective cytotoxicity of cisplatin in cancer cells, allowing drug-resistant TGCTs to regain their sensitivity to cisplatin, suggesting that targeting REV7 could be a potential treatment for patients with elevated REV7 or cisplatin resistance (71).

Neddylation. Neddylation refers to the process by which the small ubiquitin-like molecule neuronal precursor cell expression and development down-regulated protein 8 (NEDD8) is coupled to a target protein to alter its function, stability or subcellular position. The activation of neddylation increases the degradation of tumor suppressor proteins 21, p27 and p53, thereby promoting tumor progression. It has been observed in lung, breast and pancreatic cancer. A recent study found

that the overexpression of neddylation-related protein NEDD8-activating enzyme E1, which can lead to cisplatin resistance, and the neddylation inhibitor MLN4924 and cisplatin co-induced apoptosis and cell cycle arrest, effectively reducing the viability of TCGTs. Therefore, MLN4924 in combination with cisplatin could be a potential treatment for resistant TGCTs (72).

Other mechanisms. Enhancer of zeste homolog 2 (EZH2) is an epigenetic transcriptional suppressor involved in cell cycle control and cell fate determination (73). Genetic, transcriptional and post-transcriptional dysregulation of EZH2 are frequently observed in numerous cancer types (74). They play different roles in different types of tumor cells. Studies have shown that EZH2 upregulation increases cisplatin resistance in lung, ovarian and breast cancers, and the combination of EZH2 inhibitors and cisplatin may be beneficial for oncotherapy (75-77). However, at 833 K, in the NT2/D1 and 2102EP cell lines of TGCTs, decreased EZH2 expression resulted in cisplatin resistance. EZH2 is a negative regulator of cisplatin resistance in TGCTs. In addition, cisplatin resistance in TGCTs is associated with decreased H3K27 methylation (78), ubiquitination of H2A-K119 and expression of (B cell-specific Moloney murine leukemia virus integration site-1) BMI1 (79). However, DNA methyltransferase 3b knockdown leads to the induction of histone H3 lysine 27 (H3K27m3), EZH2 and BMI1 expression, thereby enhancing the sensitivity of TGCT cells to cisplatin (80). Gain of chromosome 3p25.3 was detected in all cisplatin-resistant TGCT cell lines, and the copy number in this region was associated with the level of cisplatin resistance. Gains in this region were detected at a low frequency in primary tumors and at a higher frequency in relapsed and/or cisplatin-resistant tumors. Increased chromosome 3p25.3 is associated with shorter progression-free and overall survival (81).

Mitotic arrest deficient-2 (MAD2) is a key activator of the mitotic spindle assembly checkpoint (SAC) (82), which defers the start of anaphase during mitosis when microtubules are broken or centromeres are mislocated or unattached. SAC damage causes cells to enter anaphase prematurely, resulting in chromosomal missegregation, aneuploidy or chromosomal instability (83). Abnormal MAD2 expression has a role in chromosomal abnormalities in different types of cancer cells, including TCs. Mad2 γ is a new isomer of MAD2 derived from the alternative splicing of MAD2 pre-mRNA without including exons 2 and 3. Studies have shown that the expression of MAD2 γ is markedly increased in patients with TGCT who are resistant to cisplatin chemotherapy, suggesting that MAD2 γ overexpression is related to cisplatin resistance. Furthermore, the expression of MAD2 γ increased after cisplatin treatment. Research has shown that SAC destruction is associated with resistance to DNA-damaging chemotherapy drugs. The N-terminal domain of MAD2a can interact with proteins that participate in the DNA damage response and is thought to be important for chemotherapy resistance. Similarly, MAD2 γ retains the N-terminal domain of MAD2a and is related to chemotherapy drug resistance; its overexpression promotes cisplatin resistance (84). It was found that the mRNA levels of the meiosis-related gene Testis-expressed gene 11 (TEX11) and mobility-group nucleosome-binding gene High-mobility group nucleosome-binding domain 5 (HMG5) are significantly

upregulated in TGCTs, and knockdown of HMG5 or TEX11 in cisplatin-resistant TGCTs significantly reduces the activity of cisplatin-resistant TGCT cells (85).

4. Potential therapeutic drugs

In the past 50 years, significant progress has been made in the treatment of TGCTs, particularly advanced diseases, and the cure rate has increased from 25% in the 1970s to nearly 80% (86). Radical orchiectomy combined with chemotherapy is currently considered the standard treatment (87,88). As TGCT is highly sensitive to chemotherapy, particularly cisplatin chemotherapy, the combination of cisplatin, etoposide and bleomycin or ifosfamide remains the first-line treatment (10,89,90). These combination chemotherapy regimens have shown high efficacy in the treatment of early and advanced TGCT, resulting in a significant increase in the 5-year survival rate to >95% (91,92). However, owing to the long-term side effects of platinum chemotherapy, patients still experience long-term toxicity after being cured (93,94). Approximately 20-30% of patients cannot be cured by standard treatment, particularly those who develop cisplatin resistance, and the prognosis is still unfavorable (95,96). Patients with recurrent TGCT are usually treated with high-dose chemotherapy, but this treatment results in severe adverse effects and cytotoxicity (97-99). Therefore, there is an urgent need to develop new, less toxic drugs to overcome this challenge and to improve the health status and quality of life of patients.

Numerous studies have shown that epigenetic drugs can induce tumor suppressor gene expression, making cancer sensitive to chemotherapeutic drugs again (100-102). Of note, TGCT tumorigenesis is closely associated with epigenetic and drug resistance mechanisms. Therefore, epigenetic drugs may be a viable treatment option for TGCT treatment (103-105). Numerous agents have been tested in TGCT cell lines and animal models with promising results (103,106-108). DNA methylation is a key epigenetic regulatory mechanism that silences gene expression by adding methyl groups to promoter regions of genes through the action of DNA methyltransferase (DNMT). This phenomenon often leads to loss of tumor suppressor gene function (109-112). DNA methylation is mainly mediated by DNMT1, whereas DNA remethylation is mainly mediated by DNMT3A and DNMT3B (113). Studies have shown that chemical DNA demethylation can restore the expression of silencing genes, e.g., using a first-generation DNMT inhibitor (DNMTI) 5-aza. The cytotoxic activity of 5-aza as a ribonucleoside is based on its wide range of effects on DNA methylation patterns, cellular transcriptomes and gene expression profiles. It can bind to DNA and RNA and is primarily used to treat myelodysplastic syndromes (114,115). A previous study demonstrated that 5-aza could contribute to decreased cisplatin resistance in a seminoma cell line (116). Oing *et al* (107) investigated the effects of 5-aza on two different EC cell lines (NCCIT carrying TP53 mutations and 2102Ep carrying wild-type TP53), using isogenic resistant sublines 2102EP-R and NCCIT-R to establish an acquired cisplatin resistance model system *in vivo* and *in vitro*. They found that nanomolar doses of EC cells are highly sensitive to 5-aza regardless of cisplatin sensitivity and that 5-aza at nanomolar concentrations can overcome cisplatin resistance

and induce an intense and long-lasting apoptotic response in EC cells (107). DNMTIs such as 5-aza and decitabine (DAC) have been approved by the US Food and Drug Administration (FDA), and breakthroughs have been made in the treatment of hematological cancers by blocking the activity of DNMT (117). DNMTIs have also shown potential in preclinical experimental studies for the treatment of TGCT (106,118). Beyrouthy *et al* (118) in their experiments with cisplatin-resistant embryonic cancer cells pretreated with 5-aza, found that the drug not only affected resistant cells, but also through the induction of p53 target genes or through corresponding promoter methylation to the expression of other genes (such as *MGMT*, *RASSF1A* and *HOXA-9*), to restore sensitivity to cisplatin chemotherapy. Collectively, these results suggest that 5-aza can promote apoptosis and overcome cisplatin resistance in non-cysteine germ cell tumor cells. As mentioned previously, 5-aza is a first-generation DNMTI. Notably, SGI-110, a second-generation DNMTI, has been used to treat refractory TGCTs (106). Issa *et al* (119) evaluated the effects of SGI-110 (a demethylating agent) on EC cells and in cisplatin-resistant non-seminoma testicular cancer animal models and found that cisplatin-resistant cells and tumors derived from EC were highly sensitive to SGI-110, which showed potential antitumor effects in EC (106). Notably, very low doses of SGI-110 as a single agent eliminated the progression of cisplatin-resistant EC tumors (106). In addition, they conducted a Phase I trial combining SGI-110 with cisplatin in patients with relapsed cisplatin-resistant TGCT (120), indicating that the combination of SGI-110 and cisplatin displayed good efficacy in patients with platinum-refractory germ cell cancer (120). Furthermore, Lobo *et al* (105) tested a newly synthesized flavonoid-derived compound, MLo1302, in TGCT cell lines and found that it significantly reduced tumor cell activity and induced apoptosis and cell cycle arrest by reducing DNMT expression.

In addition to DNMTIs, histone deacetylase inhibitors (HDACIs) related to histone modifications have also shown promising results in preclinical studies (103,121,122). Lobo *et al* (103) used FDA-approved HDACIs, Belinostat and Panobinostat, and observed that these agents reduced cell viability and induced cell cycle arrest and apoptosis in cisplatin-sensitive and cisplatin-resistant TGCT cell lines. In addition, Steinemann *et al* (122,123) used a novel dual-mode compound called animacroxam, which was conjugated to an HDAC-inhibiting component and a cytoskeletal-disrupting component, to show significant anti-proliferation, cell cycle arrest and apoptosis-inducing effects in TGCT cell lines, and no nonspecific cytotoxicity was observed. Of note, animacroxam also reduced glucose uptake in TGCT and inhibited the expression of glycolytic enzymes, leading to a collapse in glycolytic energy production (122,123). Nettersheim *et al* (124) used the HDACI romidepsin and found that it was highly toxic to cisplatin-resistant TGCT cells. They also showed that combined treatment with the glucocorticoid dexamethasone further enhanced the expression of romidepsin effector factors and reduced TGCT cell activity more significantly than monotherapy (124). Notably, the bromodomain inhibitor JQ1, which acts by interfering with the function of bromodomain and extra terminal

(BET) (125), increased apoptosis and growth arrest in TGCT cells, reduced tumor size and vascular density in mice and was more effective in combination with romidepsin (108). Burmeister *et al* (126) treated cisplatin-resistant TGCT cells with dual HDAC and BET inhibitors, resulting in decreased cell viability and impairment of the cell cycle. Furthermore, in the TGCT xenograft model, the dual inhibitor significantly reduced the tumor burden.

Other targeted agents have also shown potential for the treatment of cisplatin-resistant TGCT. TLS protects cells against DNA damage. Recently, Lengert *et al* (127) found that TLS inhibitors may overcome the resistance of TGCTs to cisplatin, and the targets can be diverse. MG-132 (carbobenzoxyl-L-leucyl-L-leucyl-L-leucine) is a peptide aldehyde that effectively blocks the proteolytic activity of the 26S proteasome complex and prevents TLS in human cancer cells but not in normal cells (128). However, the exact mechanism through which MG-132 inhibits TLS remains elusive. However, MG-132 showed significant cytotoxicity in cisplatin-resistant TGCT cell lines in the nanomolar range (127). In addition, Hasibeder *et al* (129) demonstrated the potential of the phytoestrogens *Belamcanda chinensis* extract and tectorigenin to inhibit TGCT.

Combination therapy is a promising therapeutic strategy. For cisplatin-resistant TGCTs, triple therapy with gemcitabine, oxaliplatin and paclitaxel is the standard of care (130). In addition, the combination of palbociclib and cisplatin greatly reduced TGCT cell viability *in vitro* and in zebrafish embryos and the drug combination also played a positive role in cell recovery after toxic damage (51). In addition, it has been reported that inhibitors specifically targeting PARP, MDM2 or AKT/mTOR combined with cisplatin have been shown to successfully overcome cisplatin resistance, which was also demonstrated in patient-derived xenograft models (10). Therefore, these agents may be promising candidates for the treatment of TGCT, particularly cisplatin-resistant TGCT; however, their application in patients with TGCT still requires further clinical studies and validation (Table II).

5. Conclusions and future perspectives

Cisplatin is the first-line drug for TGCT; therefore, it is necessary to study the mechanisms of its sensitivity and resistance.

In the mechanism of TGCT sensitivity to cisplatin, the expression of the NER essential proteins XPA, XPF and ERCC1 was significantly associated with TGCT sensitivity to cisplatin. In addition, defects in DSB homologous recombination repair also lead to high sensitivity of TGCT cell lines to cisplatin. Furthermore, the expression of the apoptosis-related factor p53 and its related proteins such as NOXA and OCT4 is also closely related to cisplatin sensitivity.

In the mechanism of TGCT resistance to cisplatin, studies have shown that cisplatin-resistant cells are mainly associated with the apoptotic pathway (24), DNA methylation, tumor cell cycle regulation-related factors and DNA damage repair pathways.

In the apoptotic signaling pathway, MDM2 can mediate the ubiquitination and further degradation of p53, thereby

Table II. Potential therapeutic drugs for cisplatin-resistant TGCTs.

Type	Name	Mechanism	(Refs.)
DNMTIs	5-aza	Block the activity of DNMT, and induct of p53 target genes or through corresponding promoter methylation to expression of other genes (such as <i>MGMT</i> , <i>RASSF1A</i> and <i>HOXA-9</i>)	(117,118)
	DAC	Block the activity of DNMT	(117)
	SGI-110	Same as 5-aza	(106)
	MLo1302	Reduce tumor cell activity and induce apoptosis and cell cycle arrest by reducing DNMT expression	(105)
HDACIs	Animacroxam	Reduce cell viability and induce cell cycle arrest and apoptosis. Reduce glucose uptake in TGCT and inhibit the expression of glycolytic enzymes, leading to a collapse in glycolytic energy production	(103,122)
	Romidepsin	Reduce TGCT cell activity	(124)
BET inhibitor	JQ1	Increase apoptosis and growth arrest in TGCT cells and reduce tumor size and vascular density in mice	(108)
TLS inhibitor	MG-132	Block the proteolytic activity of the 26S proteasome complex	(128)
Others	BCE, tectorigenin	NA	(129)

TGCTs, testicular germ cell tumors. DNMTIs, DNA methyltransferase inhibitors; 5-aza, 5-azacytidine; DNMT, DNA methyltransferase; DAC, decitabine; BET, bromodomain and extra terminal; TLS, translesion synthesis; MG-132, carbobenzoxy-L-leucyl-L-leucyl-L-leucine; BCE, belamcanda chinensis extract.

inhibiting apoptosis and cell cycle arrest and promoting cisplatin resistance (24). PDGF phosphorylates PDGFR, thereby promoting the PDGFR/PI3K/AKT signaling pathway, resulting in enhanced AKT phosphorylation. Enhanced phosphorylation of p21 and MDM2 by AKT promotes the degradation of p53. In addition, decreased expression of OCT4 could result in a decrease in NOXA, which inhibits apoptosis and promotes cisplatin resistance (23). By contrast, inhibition of FOXO3 expression leads to the suppression of *PUMA* and *FAS*, thereby inhibiting the process of programmed death (35-37). Similarly, upregulation of IGF1R promotes AKT phosphorylation and inhibits apoptosis (49). Regarding the mechanism of tumor cell cycle-related regulatory factors, p-AKT could lead to the enhancement of p21 phosphorylation, which leads to an increase in cytoplasmic p21, and high expression of cytoplasmic p21 could inhibit cell apoptosis and promote drug resistance. Cytoplasmic p21 is a new therapeutic target or biomarker that also plays an important role in chemotherapy resistance in other tumor cells. For instance, nuclear protein 1 contributes to chemotherapy resistance in breast cancer by inducing Akt-mediated phosphorylation and the subsequent cytoplasmic relocation of p21 (131). In addition, cytoplasmic p21 is a potential predictor of cisplatin sensitivity in ovarian cancer (132). Therefore, whether it is possible to regain sensitivity to chemotherapy by inhibiting the expression of p21 in the cytoplasm or altering its localization needs to be further investigated.

In addition, studies have shown that increased methylation can promote cisplatin resistance, and VIRMA, METTL3 and IGF2BP1 can promote the upregulation of m⁶A and thus promote drug resistance, whereas DNMT1 inhibitors 5-aza

and SGI-110 can inhibit methylation and lead to cisplatin sensitivity, which can be used as a potential treatment for resistant TGCTs. However, its practicability requires further investigation. In the DNA repair system, 53BP1 was far from HR. Upregulation of HR and downregulation of NHEJ were positively correlated with cisplatin resistance, and the downregulation of 53BP1 and DNA-PKs may promote cisplatin resistance (3,133).

In the TCGT NT2/D1 cell line, the CDK4/6 inhibitor palbocinib enhanced CDK4/6 protein expression in NT2/D1 cells, thereby inhibiting the cytotoxic effect of cisplatin and inducing cisplatin resistance by influencing tumor cell cycle regulation-related factors (51). However, the specific mechanisms involved require further investigation. For instance, whether an increase in CDK4/6 protein expression directly leads to cisplatin resistance or whether other factors lead to cisplatin resistance requires further exploration (51). In addition, palbocinib reduced RB phosphorylation in TGCTs, thus promoting apoptosis and inhibiting cisplatin resistance. Whether RB phosphorylation is inhibited in NT2/D1 cells to promote cisplatin resistance should be investigated in the future (134). MG-132 has been proven to promote apoptosis in refractory TGCTs and it can treat resistant TGCTs alone or in combination with cisplatin; however, its mechanism of action remains elusive (127). Study has shown that p53 promotes apoptosis in cisplatin-resistant TGCTs. Notably, MDM2-mediated ubiquitination and degradation of p53 can result in the downregulation of p53 and inhibition of apoptosis, contributing to cisplatin resistance (135). Therefore, it is speculated that MG-132, a proteasome inhibitor, promotes apoptosis by inhibiting the ubiquitination and degradation of

p53, thereby inhibiting cisplatin resistance, which requires further investigation (136).

In DNA repair systems, REV7 can bind to 53BP1 to promote NHEJ, and depletion of REV7 leads to the accumulation of DNA double-strand breaks and activation of cell apoptosis, thereby inhibiting cisplatin resistance (137). However, study has shown that cisplatin resistance is negatively related to NHEJ and that REV7 upregulation could promote NHEJ, which inhibits cisplatin resistance (71). However, study has shown that *REV7* depletion can also inhibit cisplatin resistance, and whether this is a contradiction requires further discussion. In addition, the DNA repair-related factor XPA is highly expressed in TGCTs, and its association with cisplatin resistance in TGCTs should be further studied (138). In cisplatin-sensitive and drug-resistant TGCT cells, there were differential expressions of the *LITD1*, *GAL*, *NTF3*, *NANOG*, *WNT6*, *POU5F1*, *IGFBP2*, *IGFBP7*, *PCP4*, *ZFP42*, *ID2*, *TRIB3* and *SLC40A1* genes (139). However, whether their corresponding proteins are differentially expressed has not yet been studied, and whether these proteins can be used as potential therapeutic targets needs to be studied (134).

In cisplatin-resistant TGCT therapy, studies have shown that methylation is associated with cisplatin-resistant TGCT; therefore, DNMTIs such as 5-aza and DAC SGI-110 can restore sensitivity to cisplatin-resistant chemotherapy by inhibiting methylation. In addition, drugs such as HDACIs, belinostat and panobinostat can exert therapeutic effects by inducing apoptosis. Furthermore, TLS inhibitors such as MG-132 and lomidoxin can enhance the toxic effects on cisplatin-resistant TGCT cells. Although the efficacy of certain drugs has been demonstrated in patient-derived xenotransplantation models, their use in patients with TGCT still requires further clinical research and validation.

In addition, similar to the mechanism of cisplatin resistance in other tumors, such as in ovarian yolk sac tumor (oYST), long-term cisplatin exposure resulted in a 7-fold increase in the IC₅₀ concentration in resistant cells. Drug-resistant cells showed significantly increased expression of prominin-1 (CD133), ATP-binding cassette subfamily G member 2 and aldehyde dehydrogenase 3 subtype A1 (ALDH3A1), decreased gene and promoter methylation, increased expression of ALDH1A3 and increased overall ALDH enzyme activity (140). The study showed that high expression of ALDH1A3 and elevated activity of ALDH were detected in cisplatin-resistant TCGT cells, which is identical to oYST (141). However, in TGCTs, reduced methylation promotes cisplatin resistance, in contrast to that in oYST. However, the role of other factors in the cisplatin resistance mechanism of TGCTs remains elusive and may be considered a potential factor. In medulloblastoma, anti-miR-31 enhances cisplatin resistance through the PI3K/AKT and NF-κB pathways (142). Similarly, in glioblastoma, circPTN (a newly discovered circular RNA) contributes to cisplatin resistance through PI3K/AKT signaling via the miR-3-542p/PIK3R3 pathway (143). The PI3K/AKT pathway also affects cisplatin resistance in TGCTs; therefore, it is worth exploring whether miR-31 and circPTN also play a role in cisplatin resistance in TGCTs. A previous study showed that PD-L1 is highly expressed in TGCTs (144). Studies have shown that in non-small cell lung cancer (NSCLC), NSCLC

cell-derived programmed death-ligand 1 (PD-L1)-containing exosomes promote cell stemness and increase the resistance of NSCLC cells to cisplatin (145), and high expression of PD-L1 in cancer cells drives immune-independent, cell-intrinsic functions, leading to resistance to DNA-damaging therapies, such as cisplatin (146). In NSCLC, the increase in CDK5 could lead to the downregulation of F-box only protein 22, which leads to the downregulation of PD-L1 ubiquitination, and the increase in PD-L1 levels leads to cisplatin resistance (146). In medulloblastoma, CDK5 and PD-L1 also play a role in cisplatin resistance; therefore, the role of CDK5 in cisplatin resistance of TGCTs is also worth discussing (146).

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Authors' contributions

ZZ, XL, JS and LC conceived the study, performed the investigation, and wrote and edited the original draft. XJ and MY wrote, reviewed and edited the manuscript. All authors read and approved the final manuscript. Data authentication is not applicable.

Ethics approval and consent to participate

Not applicable.

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Competing interests

The authors declare that they have no competing interests.

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