Killing cancer with platycodin D through multiple mechanisms

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Abstract

Cancer is a multi-faceted disease comprised of a combination of genetic, epigenetic, metabolic and signalling aberrations which severely disrupt the normal homoeostasis of cell growth and death. Rational developments of highly selective drugs which specifically block only one of the signalling pathways have been associated with limited therapeutic success. Multi-targeted prevention of cancer has emerged as a new paradigm for effective anti-cancer treatment. Platycodin D, a triterpenoid saponin, is one the major active components of the roots of *Platycodon grandi-florum* and possesses multiple biological and pharmacological properties including, anti-nociceptive, anti-atherosclerosis, antiviral, anti-inflammatory, anti-obesity, immunoregulatory, hepatoprotective and anti-tumour activities. Recently, the anti-cancer activity of platycodin D has been extensively studied. The purpose of this review was to give our perspectives on the current status of platycodin D and discuss its anti-cancer activity and molecular mechanisms which may help the further design and conduct of pre-clinical and clinical trials to develop it successfully into a potential lead drug for oncological therapy. Platycodin D has been shown to fight cancer by inducing apoptosis, cell cycle arrest, and autophagy and inhibiting angiogenesis, invasion and metastasis by targeting multiple signalling pathways which are frequently deregulated in cancers suggesting that this multi-target activity rather than a single effect may play an important role in developing platycodin D into potential anti-cancer drug.

Keywords: platycodin D • saponin • apoptosis • autophagy • invasion • metastasis

Introduction

Cancer remains one of the major public health problems in both developed and developing countries and represents second leading cause of death worldwide [1], with approximately 14 million new cases and 8.2 million cancer-related deaths in 2012 [1, 2]. According to World Health Organization estimate, by 2050, 17.5 million cancer deaths are projected to occur in the world [3] Regardless of the factors contributing to the initiation and development of a

*Correspondence to: Tonghui MA E-mail: tonghuima@dlmedu.edu.cn particular cancer, accumulating research evidence suggests that most cancers are caused by dysfunction of many genes coding for proteins such as anti-apoptotic proteins, inhibitors of apoptosis, transcriptional factors, growth factor, growth factor receptors and tumour suppressors, which constituted the targets for cancer treatment [1, 4].

Prevailing chemotherapeutic drugs have limited therapeutic success in cancer; because they are highly toxic, expensive and activate the alternative cell signalling pathways [5]. Rational development of highly selective drugs that specifically block only one of the signalling pathways such as monoclonal antibodies that kill the cancer cells by

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specifically binding to the extracellular domains of receptor tyrosine kinase have been associated with sporadic responses because of emergence of secondary drug resistance [1, 6–8]. As cancer development is a multi-step process which is characterized by multiple abnormalities rather than a single mutation, it is unlikely to inhibit cancer effectively using mono-target agent [9, 10]. A plethora of research reports have shown that unlike pharmaceutical drugs that act as mono-target molecules, plants have multi-target molecules that can regulate cancer growth and progression by interfering with multiple mechanisms which are central to cancer progression [5].

Cancer chemoprevention by natural compound has emerged as a promising and pragmatic approach to reduce the risk of cancer and is gaining increasing attention because it is considered as safe, cost effective and alternative form of modern health care system [4, 11, 12]. Plants have a long history of use in the treatment of cancer and remain the most attractive source of anti-cancer drugs as tremendous chemical diversity is found in millions of plant species [4, 13-15]. There exist several solid reasons for continued interest in exploring plants extensively as a source of new potential therapeutic anti-cancer agents. First, at present more than 60% FDA approved commercially available anti-cancer drugs are derived from natural sources including plants [16]. However, the plants which have been thoroughly investigated for their potential values as a source of drugs make a very small fraction (about 10%) of the total estimated 2.5 to 5 million plant species [17, 18]. As a very small fraction of available plant flora has obviously contributed for more than 60% of prevailing chemotherapeutic anti-cancer drugs, the need for the identification of novel structures from remaining flora and understanding their molecular mechanisms for the development of more effective therapeutic drugs to reduce the burden of cancer is therefore paramount. Second, nature has spent over 3 billion years to create a wonderful and near-perfect complex compound library by performing countless high throughput screens to remove inactive and retain bioactive molecules that no synthetic chemist could ever dream up [19]. Although combinatorial chemistry is one of the most important methodologies developed by chemists to create new leads in anticancer drug discovery, however, only one FDA approved anti-cancer drug 'Sorafenib' has been made available in public domain by this method from 1981 to 2006 [20]. Third, a comparative genomic analvsis revealed that 70% of cancer-related human genes have orthologues in Arabidopsis thaliana indicating that human and plants use similar receptors and signalling pathways in some cases [21, 22]. Given the similarity of many human and plant genes, it can be assumed that some secondary metabolites produced by plants to modulate their own metabolism might be able to interact with molecules which have a role in human cancers. One such example of such a case is multidrug resistance-like proteins that are used by Arabidopsis thaliana to transport auxin have orthlogues in human that play important role in development of multidrug resistance by transporting anti-cancer drugs out of cells. Flavonoids, which are auxin distribution modulators in Arabidopsis, have been shown to overcome multidrug resistance by modulating p-glycoprotein in various human cancers [23]. Fourth, natural anti-cancer compounds fit into mechanism-based approach as perfectly as a hand fits into a

glove. There is solid evidence that plant-derived bioactive compounds inhibit cancer by interfering with multiple mechanisms which are central to cancer progression [1]. Fifth, plants often produce bioactive compounds that exceed the current capacity of synthetic chemistry [24, 25].

Plants' secondary metabolites such as alkaloids, terpenes and polyphenols have now been well characterized for their anti-cancer activity [16, 26-30]. Till now, more than 55,000 terpenes have been isolated [31] but only a small fraction of terpenes is well characterized for their potential value as a source of anti-cancer drugs [32]. Therefore, it is imperative to identify and characterize more and more bioactive compounds with promising anti-cancer activity for faster development of more effective anti-cancer drugs. Diterpenoids, sesquiterpene lactones and saponins are the major classes of terpenes with well-documented anti-cancer activity against a wide range of human cancers [32-34]. Saponins are a large and structurally diverse group of bioactive compounds [32] which are reported to occur in more than 100 families of plants [35]. Chemically, saponins are natural glycosides of steroid or triterpene [36]. The steroidal saponins are important precursor molecules for various steroid drugs such as anti-inflammatory drugs, androgen, oestrogen and progestins [37]. The triterpene saponins have been shown to exhibit a wide range of biological and pharmacological activities including anti-inflammatory, antiviral, anti-fungal, antibacterial, antioxidant, anti-cancer, anti-diarrheal, anti-ulcerogenic, anti-oxytocic, antitussive, anticoagulant. immunomodulatory, hepatoprotective, neuroprotective, expectorant, analgesic, molluscicidal, hypoglycaemic and hypocholesterolaemic [36, 37]. Modern research has shown that saponins are potential anti-cancer agents with different mechanisms of action [35]. To date, about 150 kinds of natural saponins have been found to possess significant anti-cancer activity against various human cancer cells [35]. Saponins are usually considered toxic because of their haemolytic properties. The number of sugar chains and type of sugar residue attached to C-3 and C-28 of advcone play important role in pore formation as well as in adjuvant activity of triterpenoid saponins. Saponins with two sugar chains induce less haemolytic activity and membrane permeability than those with only one side chain [38, 39]. Other studies show that presence of single glucosvl group in sugar chain attached to C-3 of aglycone is responsible for strong adjuvant property of saponins [40, 41]. Platycodin D, an oleanane type triterpenoid saponin possesses two sugar chains; a single glucosyl group attached at position C-3 and other chain of four sugar residues at position C-28 of aglycone. Platycodin D has been shown to exert anti-tumour activity in various cancer cells without affecting normal cells in various in vitro [42] and in vivo [43-45] models. Platycodin D has been extensively investigated and found to exhibit significant anti-cancer activity against a wide range of human cancers types both in vitro and in vivo through multiple mechanisms which are considered to be crucial and central to cancer development and metastasis. Among other oleanane type, triterpenoid saponins with two sugar chains at position C-3 and C-28, tubeimoside I, a bioactive component of Traditional Chinese Medicinal herb Bolbostemma paniculatum, has been extensively studied for its anti-cancer activity against various human cancers including lung cancer, gastric cancer, ovarian cancer, choriocarcinoma cancer, glioma, hepatoma, squamous oesophageal carcinoma and cervical carcinoma [46]. The anti-cancer activity of tubeimoside I has been reported to be associated with apoptosis induction through mitochondrial dysfunction, Bcl-2 familv protein modulation. COX-2 down-regulation. caspases activation. reactive oxygen species (ROS) generation, endoplasmic reticulum stress [46, 47], Fas and FasL activation, c-Jun N-terminal kinase (JNK) phosphorylation, down-regulation of tumour necrosis factor- α , and NF- κ B [48], p38 phosphorylation, Protein kinase B (AKT) and Extracellular Signal Regulating Kinase (ERK) dephosphorylation [49] and G2/M phase arrest by inhibiting tubuline polymerization [50] and down-regulating the expression of cyclin B1/cdc2 complex through p21 activation [51]. The IC₅₀ values of tubeimoside I against various cancer cells were comparable to platycodin D. The aim of this review was to summarize the current knowledge and discuss the natural sources, anti-cancer activity, molecular targets, mechanisms of action and usefulness of platycodin D for anti-cancer drug development.

Natural sources

Platycodin D is one of the major bioactive components of the Traditional Chinese Medicinal herb Platycodon grandiflorum [52, 53] which is called Jiegeng in Chinese [45] and Chinese balloon flower or common balloon flower in English [37]. It is abundantly found in China, Japan and Korea [43, 54]. The content of platycodin D in root and aerial parts of P. grandiflorum is determined to be 0.20% [52] and 0.018% [53] of dry weight respectively. It can also be obtained from platycoside E and platycodin D3 by enzymatic transformation [55]. The yield of platycodin D by enzymatic transformation method is twofold as compared to direct isolation and purification from methanol extract of roots [55]. The root of P. grandiflorum called Platycodonis Radix [56] or Platycodi Radix [44, 57] is a well-known Traditional Chinese Medicine used as an expectorant for pulmonary diseases and a remedy for respiratory disorders including cough, colds, sore throat, tonsillitis and chest congestion [54, 58]. In Korea, in addition to treating diabetes and respiratory and inflammatory diseases, the roots of *P. grandiflorum* are used to prepare salad and many other traditional Korean dishes. Pan fried roots are also served as a popular dish in Korean traditional foods [37].

Killing cancer with platycodin D

Platycodin D (Fig. 1A) is a potential anti-cancer compound that has been shown to exhibit a broad spectrum of cytotoxicity against a wide range of human cancer cell lines of different origin both *in vitro* and *in vivo* through multiple mechanisms of action involving proliferation and survival inhibition, apoptosis induction, non-specified apoptotic cell death stimulation, cell cycle arrest, autophagy, and angiogenesis and metastasis inhibition and transcription factors regulation as shown in Figure 1B.

Targeting apoptosis pathways in cancer cells with platycodin D

Apoptosis is a highly synchronized gene-controlled process during which a series of intracellular events come into play to decommission the unwanted and cancerous cells [1, 59]. It is characterized by typical morphological and biochemical hallmarks, including membrane blebbing, DNA fragmentation, cell shrinkage and caspase-3 activation [60-62]. Apoptosis is of widespread biological significance playing a vital role in a myriad of physiological and pathological processes in a wide variety of tissues. Apoptosis plays a key role in maintaining tissue homoeostasis by selectively eliminating unwanted or damaged cells. Tissue homoeostasis is regulated through a precise balance between apoptosis and cell proliferation. However, disruption of this balance between apoptosis and cell proliferation may promote severe pathological conditions including, tumourigenesis, neurodegeneration, auto-immune diseases and developmental abnormalities [63-66]. Inhibition of apoptosis may lead to tumour development and drug resistance [65, 67]. Tumour cells use a variety of molecular mechanisms to suppress apoptosis [68]. Thus, activation of the molecular mechanisms that regulate the apoptotic machinery in various human malignancies may provide a novel opportunity for cancer drug development. Identification of novel therapeutic agent that have potential to selectively induce apoptosis in cancer cells are the focus of modern anti-cancer drug discovery.

Platycodin D, a triterpenoid saponin has recently been evaluated for its potential value as a source of anti-cancer drug. Accumulating evidence from various studies has shown that platycodin D holds the promise to trigger apoptosis to set cancer cells on the road to ruin. Multiple mechanisms have been implicated in apoptosis inducing action of platycodin D, including up-regulation of Fas/FasL, mitochondrial dysfunction, Bcl-2 family protein modulation, ROS generation, inhibition of inhibitors of apoptosis, induction of mitotic arrest, Mitogen activated protein kinase (MAPK) pathway activation, and suppression of telomerase activity and pro-survival pathways such as AKT. Platycodin D-induced apoptosis through multiple mechanisms has been shown in Figure 2.

Targeting cancer cells by extrinsic apoptosis

The extrinsic apoptotic pathway is triggered by the activation of death receptors (DR) by their respective ligands [1]. Theses DRs such as fibroblast-associated antigen (Fas), tumour necrosis factor receptor-1, DR3, DR4, DR5 and DR6 are transmembrane receptors containing a cystein-rich extracellular domain for ligand binding as well as a cytoplasmic domain of about 80 amino acids called death domain which plays a critical role in transmitting the death signals from the cell surface to the intracellular signalling proteins [60, 61, 69]. Upon ligand binding, the DRs such as Fas and DR4/5 interact with Fas-associated death domain (FADD) and procaspase-8 to form death-inducing signalling complex (DISC) within the cell. Within this complex, FADD recruits procaspase-8 through its death effector domain which becomes activated by auto-proteolysis and dissociates



Fig. 1 (A) Chemical structure and natural source of platycodin D. (B), Platycodin D inhibits cancer progression and development by inhibiting proliferation, survival, angiogenesis, invasion, metastasis and cell cycle progression, and ultimately inducing apoptosis and autophagy by modulating expressions of various genes.

from the DISC to activate apoptotic machinery by directly activating the downstream effector caspase-3, resulting in cell death *via* the type I extrinsic apoptotic pathway, or cleaves Bid, a pro-apoptotic member of the Bcl-2 family, leading to activation of the type II extrinsic apoptotic pathway [1, 70].

Killing cancer by targeting DRs in cancer cells has been considered as a useful and promising approach by researchers [61, 70, 71]. Recently, platycodin D has been investigated for its potential value as a source of novel anti-cancer drug in various cancer cell lines. It has been found to induce extrinsic apoptosis in different human cancer cells including AGS gastric adenocarcinoma. MCF-7 breast adenocarcinoma, and PC3 prostate cancer and HaCaT cells. In PC3 cells, platycodin D has been shown to induce activation of caspase-8 [45], while in MCF-7 cells it has been shown to induce activation of caspase-8 and Bid [72]. In HaCaT cells, platycodin D induced extrinsic apoptosis by up-regulating the expression of Fas and FasL both at mRNA as well as at protein levels and inducing proteolytic cleavage of caspase-8 and -3 in a time-dependent manner [57]. The induction of Fas and FasL appears to come from platycodin D-induced NF-KB activation in HaCaT cells. Pre-treatment of cells with 100 μM TPCK (NF-κB inhibitor) inhibited the translocation of NF- κ B into nucleus and abrogated

platycodin D-induced apoptosis in HaCaT cells indicating the potential involvement of NF- κ B in platycodin D-induced apoptosis. Further results from this study revealed that platycodin D induces activation of NF- κ B by stimulating I κ B kinase (IKK) activity and I κ B α degradation and this inducible NF- κ B is made up of p50 and p65 sub-units as evident by the reduction in NF- κ B DNA binding activity in the presence of anti-p50 and anti-p65 antibodies.

To further confirm the functional link between NF- κB activation and Fas receptor up-regulation by platycodin D, HaCaT cells were transfected with luciferase construct pFLF1 containing a promoter region between -1435 and +236 of the Fas receptor gene with the NF- κB binding site for 18 hrs, followed by a dose-dependent treatment of platycodin D (10–50 μM) for 3 hrs and luciferase activity was determined. Platycodin D increased luciferase activity dose-dependently indicating that NF- κB is involved in positive regulation of Fas receptor promoter.

In AGS gastric cancer cells, platycodin D has been reported to induce type I as well as type II extrinsic apoptosis by inducing the expression of FasL and proteolytic cleavage of caspase-8 Bid, caspase-3, caspase-9 and poly (ADP-ribose) polymerase (PARP) in a dose- and time-dependent manner [73, 74]. Type II extrinsic apopto-

Fig. 2 A schematic model of platycodin Dinduced apoptosis in cancer cells. Platycodin D induces ROS generation and increases the expression of transcriptional factor FOXO3a, ROS activates ASK-1 by reducing the expression of thioredoxin. Once activated, ASK-1 activates p38 and JNK by phosphorylation. Transcriptional factor FOXO3a and activated p38 and JNK triggered both extrinsic and extrinsic apoptosis by inducing the transcription of apoptosis related genes such as FasL, Bim, c-Jun. Moreover, platycodin D induces intrinsic apoptosis by modulating the expression of Bcl-2 family proteins and mitochondrial dysfunction, inducing endoplasmic reticulum stress, increasing the expressions of Eqr-1 and NAG-1 which results in caspase-3 activation. Caspase-3 then executes apoptotic cell death by cleaving PARP and activating CAD.



sis was mediated by translocation of tBid into mitochondrial membrane thereby modulating the expressions of Bcl-2 family proteins and ultimately releasing cytochrome c into cytosol which is a hallmark of mitochondrial apoptosis. Both type I and type II extrinsic pathways converge at caspase-3, which is the main executioner of apoptotic cell death [74].

These effects were mediated by platycodin D-induced activation of p38/AP-1 (activator protein 1) signalling. Pre-treatment of cells with SB203580 (p38 inhibitor) completely reversed platycodin Dinduced apoptosis and reduced the up-regulation of FasL and Bax/ Bcl-2 ratio and suppressed the activation of AP-1, caspase-8, -9, -3 and PARP. These effects were further confirmed by knockdown of p38 expression through siRNA. Moreover, pre-treatment of cells with parthenolide (AP-1 inhibitor) inhibited the expressions of pro-apoptotic proteins Bim and c-Jun in response to platycodin D. Parthenolidemediated inhibition of AP-1 activation was also verified by EMSA. In addition to p38, platycodin D also increased the phosphorylation of JNK and decreased the phosphorylation of ERK in AGS cells. Platycodin D-induced apoptosis was not affected by SP600125 (JNK inhibitor), however, U0126 (ERK inhibitor) potentiated the platycodin Dmediated apoptosis indicating anti-apoptotic effect of ERK activation in AGS cells. Taken together, these sets of data provided clear evidence that platycodin D induces apoptosis in AGS mainly through p38/AP-1 signallings [74].

Contrary to the NF- κ B activation in HaCaT cells [57], platycodin D inhibited the activation of NF- κ B in AGS cells [74]. NF- κ B activation in HaCaT cells in response to platycodin D treatment appears as apoptosis promoter while platycodin D-induced apoptosis in AGS cells seems to occur independent of NF- κ B activation. A similar NF- κ B inhibitory effect of platycodin D has been reported in MDA-MB-231 breast cancer cells [75] and LPS-stimulated macrophages [41]. This indicates that platycodin D modulates the expression of NF- κ B depending upon the type of cells. Based on available data, it can be concluded that platycodin D inhibits NF- κ B activation in cancer cells while induces NF- κ B expression in normal cells.

Targeting cancer cells by intrinsic apoptosis

Unlike extrinsic apoptosis, the intrinsic apoptotic pathway is initiated by a diverse array of non-receptor mediated stimuli that produce intracellular signals that act directly on the target within the cell [60]. All of these stimuli alter the functioning of mitochondria. Mitochondria being an important component of the apoptosis execution machinery have been extensively reviewed [1, 61]. As the activation of mitochondria has been considered as 'point of no return' in apoptotic process, the manipulation of mitochondria activation to initiate apoptotic cell death has been considered as a potential therapeutic approach.

Bcl-2 family proteins have been envisaged the main mediators of mitochondrial apoptosis [76–78]. Modulation of Bcl-2 family proteins results in opening of the mitochondrial permeability transition pore, dissipation of mitochondrial membrane potential and subsequent release of many pro-apoptotic proteins such as cytochrome c, second mitochondrial activator of caspases (Smac/DIABLO), apoptosis inducing factor (AIF), Endonuclease G (Endo G) and caspase-activated DNase (CAD) from mitochondrial inter-membrane space into cytosol [79]. Cytochrome c, once released into the cytosol, interacts with apoptotic protease activating factor-1, leading to the activation of caspase-9 [80, 81]. Active caspase-9 then activates caspase-3, which in turn cleaves substrates such as inhibitor of caspase-activated DNase (ICAD) and PARP, leading to nucelosomal DNA fragmentation and thus apoptosis. Smac/DIABLO promotes caspases activation through neutralizing the inhibitory effects of inhibitor of apoptosis

proteins such as xiap and survivin [82, 83], while AIF and Endo G induce caspase-independent apoptosis by directly inducing DNA damage and chromatin condensation reffered as stage 1 condensation. ICAD is then released from mitochondria and translocated into nucleus where, after cleavage by caspase-3, it induces oligonucleosomal DNA fragmentation and more pronounced and advanced chromatin condensation termed as stage II condensation [60, 79].

The apoptotic effects of platycodin D has been investigated in various cancer cell lines. It has been shown to trigger intrinsic apoptosis mainly by modulating the expressions of Bcl-2 family proteins in different cancer cell lines of human origin including gastric cancer [73, 74]. breast cancer [72], liver cancer [56], brain cancer [84] and prostate cancer [45]. Platycodin D has significantly reduced the Bcl-2/Bax ratio and increased the expression of cleaved caspase-9, caspase-3 and PARP in PC-3, AGS, HepG2 and MCF-7 cancer cells. In U251 human glioma cells, it has been reported to reduce Bcl-2/Bax ratio and increase DNA fragmentation and caspase-3 activation. In addition to suppressing Bcl-2/Bax ratio, platycodin D has been reported to induce the expression of Bim and c-Jun through AP-1 transcriptional activity in AGS gastric cancer cells [74]. The anti-cancer activity of platycodin D has also been validated in PC-3 prostate cancer xenograft model [45]. The data demonstrated that 2.5 mg/kg dose of platycodin D suppressed about 56% growth over a period of 4 weeks.

Based on available data, it appears that platycodin D holds the promise to induce intrinsic apoptosis by interacting with Bcl-2 family proteins. As Bcl-2 family proteins appear to be an attractive drug target for cancer therapy, it is therefore imperative to explore platycodin D for its potential to target Bcl-2 family protein in a wide variety of cancer cells both *in vitro* and *in vivo* cancer models to develop it into a promising chemotherapeutic drug.

Targeting cancer cells by ROS-mediated apoptosis

Reactive oxygen species are oxygen containing reactive chemical entities which have been shown to play a vital role in various cellular processes including proliferation, cell survival, differentiation, gene expression, and regulating the activity of enzymes and eliminating pathogens and foreign particles [85, 86]. Compelling evidence suggests that cancer cells exhibit high oxidative stress which plays important role in cancer cell survival, proliferation, disruption of cell death signalling, angiogenesis, metastasis and drug resistance [87–89]. Recent studies have shown that in contrast to tumour promoting effect of increased ROS, this biochemical property of cancer cells can be exploited for therapeutic benefits. Exploiting the vulnerability of cancer cells with intrinsic oxidative stress to further ROS insult above a toxic threshold level by exogenous ROS generating phytochemicals to selectively kill cancer cells have been shown to be feasible in various *in vitro* and *in vivo* experimental models [82, 87–91].

Platycodin D has recently been reported to induce apoptosis in U937 leukaemia [92] and MCF-7 breast cancer cells [72, 93] by targeting ROS metabolism through different mechanisms. In U937 human leukaemia cells, platycodin D (15 μ M) have been shown to

induce ROS-dependent apoptosis. Treatment of cells with platycodin D (15 μ M) for 48 hrs has induced apoptosis *via* mitochondrial membrane potential dissipation, and increased expression of transcription factor early growth response-1 (Egr-1), non-steroidal anti-inflammatory drug (NSAID)-activated gene-1 (NAG-1), cleaved caspase-3 and PARP and DNA fragmentation. Pre-treatment of cells with NAC (ROS inhibitor) completely reversed while broad-spectrum caspase inhibitor (z-VAD-fmk) partially inhibited the apoptotic effects of platycodin D in U937 cells [92]. As platycodin D treatment started to increase ROS generation within 30 min. while induced the expression of EGR-1 and NAG-1 after 1 and 2 hrs respectively, it could be concluded that ROS act as an upstream signalling molecule to induce NAG-1 expression through EGR-1 activation.

In MCF-7 breast cancer cells, the apoptotic effect of platycodin D has been found to be associated with ROS generation, mitochondrial dysfunction, Bcl-2 family protein modulation, caspase-9, -8, -3 and PARP cleavage, stress-activated MAPK proteins activation and ER stress. A reversal of all these ROS-mediated events was observed with supplementation of ROS scavenger NAC. Pre-treatment of cells with SP600125 (JNK inhibitor) and SB203580 (p38 inhibitor) has partially reversed the apoptosis indicating that involvement of pathways other than MAPK [72]. Platycodin D-mediated apoptosis via ROS signalling has been shown in Figure 2. Further detailed investigations demonstrated that platycodin D induced activation of ASK-1 by downregulating the expression of thioredoxin and phosphorylating ASK-1 at Threonine 845 and dephosphorylating at Serine 967. Activated ASK-1 leads to the activation (phosphorylation) of p38 and JNK. Moreover, platycodin D exerted ER stress in MCF-7 cells by phosphorylating PERK (PKR-like endoplasmic reticulum kinase) and eukaryotic initiation factor 2 α and glucose-regulated protein 78/binding immunoglobulin protein and CHOP/GADD153 up-regulation, and caspase-4 activation. The induction of ER stress-related signals by platycodin D treatment was decreased by the antioxidant NAC, suggesting that ROS generation by platycodin D is upstream of ER stress [93].

Targeting cancer cells by cell cycle-mediated apoptosis

Cell cycle arrest at a particular checkpoint is one of the major causes of cell death in cancer cells [94]. The accurately orchestrated sequence of events that lead to cell division is regulated by a coordinated interaction of a variety of cyclins with their respective cyclindependent kinases at various checkpoints [94, 95]. Natural compounds such as taxol and vinca alkaloid that arrest cell cycle at G2/M checkpoint by targeting microtubules have proven to be one of the best classes of cancer chemotherapeutic drugs in both haematopoietic and solid tumours [96, 97]. Microtubules represent the best cancer target identified so far [98, 99]. Platycodin D has been shown to arrest cell cycle at G2/M checkpoint in various cancer cell lines. In human hepatoma HepG2 cells, it induced G2/M arrest in a dosedependent manner [100]. In U937 leukaemia cells, platycodin D appears to arrest cell cycle at G2 phase by inhibition of cyclin B1/ cdk1 complex *via* Wee1 expression and M phase by promoting tubulin polymerization [101]. Taken together, the data suggest that platycodin D induced endoreduplication (END) through mitotic arrest as evident by increased expression of cdk2 and mitosis markers such as histone H3 serine 10 phosphorylation, BubR1 and MPM-2 and this END ultimately leads to mitochondrial apoptosis. Unlike eteposide and merbarone which induce END *via* Topo II inhibition, platycodin-induced END is not associated with Topo II activity.

Platycodin D has also been found to induce G2/M phase arrest in PC3 cells in a dose-dependent manner [45]. It has been shown to inhibit the expressions of E2F1, MDM2, Cyclins (B1 & D1) and CDKs (1, 2, 4 & 6). These effects were associated with up-regulation of tumour suppressor protein FOXO3a (Forkhead box transcription factor) and its direct downstream target genes p21 and p27 in PC3 cells. Platycodin D-induced expression of FOXO3a appears to come from suppressive effects of platycodin D on Pl3K/AKT and ERK pathways as shown in Figure 3.

Targeting cancer cells by non-specified apoptosis

In addition to aforementioned apoptotic cell death processes which can be initiated either by activation and ligation of DRs to their respective ligands or by mitochondrial activation, platycodin D has been shown to induce apoptotic cell death in U937 leukaemia cells by inducing DNA fragmentation, LDH release and cleavage of caspase-3 and PARP in a dose-dependent manner through a non-specified apoptotic process [102]. The apoptotic cell death found to be linked with reduced activity of telomerase via down-regulation of hTERT (Telomerase reverse transcriptase) expression. Platvcodin D treatment decreased the protein levels of c-Myc and Sp1 (Regulators of hTERT transcription) and reduced their DNA binding activity. In addition to hTERT transcription inhibition, platycodin D treatment also suppressed the activity of telomerase by inhibiting the phosphorylation and nuclear translocation of hTERT via inhibition of AKT activation. The set of data demonstrate that cytotoxic effects of platycodin D in U937 leukaemia cells seems to result from suppression of telomerase activity through transcriptional and post-transcriptional inhibition of hTERT activity [102]. Platycodin D has also been shown to inhibit growth and DNA synthesis and promote apoptosis and caspase-3 activity in MDA-MB-231 cells in a dose-dependent manner [103].

Targeting autophagy in cancer cells

Autophagy is an evolutionary highly conserved catabolic process that plays a vital role in degradation of misfolded proteins and damaged



Fig. 3 A schematic representation of platycodin D-induced G2/M phase cell cycle arrest in cancer cells. Platycodin D inhibits G2-M phase transition by increasing the expression of transcription factor FOXO3a, p53, p21 and Weel, and decreasing the expression of MDM2 and induces M phase arrest by sustainable tubulin polymerization. Binding of ligand such as growth factor (EGF) to the growth factor receptor (EGFR) promotes the activation of downstream pro-survival signalling pathways such as PI3K/AKT and MAPK/ERK pathways. Activated AKT and ERK translocate to the nucleus and inhibit the transcription of apoptosis promoting (Bim, FasL, TRAIL) and cell cycle regulatory genes (p21, p27) by phosphorylating FOXO3a and exporting it out of nucleus. ERK regulates the expression of MDM2 that ubiquitinates FOXO3a in response to ERK pathway activation leading to proteasomal degradation. Platycodin D induces G2/M phase arrest and apoptosis by increasing the expression of FOXO3a in nucleus by inhibiting AKT and ERK.

organelles [104, 105]. It plays a very prominent role in various physiological and pathological conditions such as cancer [104, 106]. A substantial amount of research evidence exists to support autophagy as a bonafide tumour suppressor mechanism. Autophagic cell death or type II programmed cell death is an alternative mechanism of cancer cell death in apoptosis deficient/resistant cells. Accumulating evidence from various studies has shown that natural compounds induce autophagic cell death in apoptosis defective cells [106-108]. The connection between apoptosis and autophagy is complex and is currently being investigated for therapeutic benefits [107]. For example, anti-apoptotic Bcl-2 protein also inhibits autophagy by inhibiting beclin; an autophagy gene initially identified as tumour suppressor. Inhibition of anti-apoptotic Bcl-2 protein has been shown to induce apoptosis as well as autophagy in cancer cells [108]. Inhibition of protective autophagy has been shown to induce apoptosis and increase the therapeutic efficacy of cisplatin and 5-fluorouracil in human cancers. Other studies showed that autophagy switches to apoptosis and inhibition of autophagy delays apoptosis [106]. Thus, identification and characterization of the type of autophagy and interaction between apoptosis and autophagy may provide a unique strategy for the development of highly selective and more effective anti-cancer drugs for various carcinomas.

LC3 is a hallmark of autophagy and conversion of LC3-I to LC3-II shows autophagy induction [109, 110]. Recently, platycodin D has been explored for its potential to induce autophagy. The compound was found to be potent inducer of autophagy at a concentration range of 1.25-30 µM in a wide range of cancer cells including HepG2, Hep3B, MCF-7, MDA-MB-231, A549, 95D and NCI-H460 cells [111]. In Hep3B, MCF-7, MDA-MB-231, A549 and 95D cells, platycodin D triggered autophagy in a dose- and time-dependent manner by increasing the expression of LC3-II. In HepG2 cells, an increased level of LC3-II expression was observed when cells were treated with a combination of platycodin D and late-stage autophagy inhibitors (CQ and BAF) compared to platycodin D treatment alone. Combine treatment of cells with platycodin D and autophagy inhibitors enhanced the apoptotic effect in HepG2 cells indicating that platycodin D induced protective autophagy in HepG2 cells. Moreover, platycodin D increased phosphorylation of ERK and U0126 (MEK inhibitor) reduced the phosphorylation of ERK and attenuated platycodin D-induced expression of LC3-II indicating that ERK activation is involved in platycodin D-mediated autophagy in HepG2 cells [111].

In A549 and NCI-H460 human non-small cell lung carcinoma cells, platycodin D induced autophagy by increasing the conversion of LC3-I to LC3-II and accumulation of Atg-3, Atg-7 and Beclin in a dose- and time-dependent manner. Platycodin D treatment reduced the phosphorylation of several members of the Pl3K/Akt/mTOR signalling pathway such as p-Akt (Ser473), p-p70S6K (Thr389) and p-4EBP1 (Thr37/46), and increased the expression of phosphorylated p38 and JNK in A549 and NCI-H460 cells [112]. Platycodin D co-treatment with LY294002 (Pl3K inhibitor) or rapamycin increased the expression of LC3-II in a synergistic fashion. Moreover, platycodin D reversed insulin-mediated activation of Pl3K/Akt/mTOR signalling and suppression of autophagy in these cancer cell lines. Platycodin D also increased the phosphorylation of p38 MAPK and JNK. Pre-treatment of cells with SB203580 (p38 inhibitor) and SP600125 (JNK inhibitor)

decreased the ratio of LC3-II/LC3-I as compared to platycodin D treatment alone. Taken together, platycodin D trigger autophagy *via* inhibiting PI3K/Akt/mTOR signalling pathway and activating JNK and p38MAPK signalling pathway in A549 and NCI-H460 cells. The effect of platycodin D on autophagy induction has been represented in Figure 4.

Targeting tumour cell development and progression with platycodin D

Apart from type I and type II programmed cell death, platycodin D has been shown to interfere with tumour progression by interacting with multiple targets.

Targeting proliferation of cancer cells

In normal tissues, cell proliferation is controlled by irreversible entry into post-mitotic terminal differentiation [113]. Cancer cells escape this terminal differentiation by up-regulating the expression of c-Myc which is a potent inhibitor of differentiation in many cell lineages [113, 114]. Myc has been found deregulated in nearly half of the human solid tumours and leukaemia and appears frequently associated with cancer progression [115]. Besides c-Myc, overexpression of cylin D1 [116] and phospholipase $C\gamma 1$ (PLC $\gamma 1$) [117] has been linked with higher rates of cellular proliferation in many cancers.



Fig. 4 Molecular mechanism of platycodin D-induced autophagy in cancer cells: Platycodin D induces autophagy through Pl3K/AKT and MAPK pathways. Platycodin D converted LC3-I to LC3-II by increasing the expression of Beclin-1, Atg3 and Atg 7. Inhibition of Pl3K/AKT and activation of MAPK signalling pathways by platycodin D resulted in autophagy by accumulating LC3-II in cancer cells (\vdash Inhibition; \leftarrow Activation).

Platycodin D inhibited proliferation in several human cancer cells in a time- and dose-dependent manner [41, 45, 56, 72–75, 84, 92, 100–102, 118]. Apart from its anti-proliferative effects determined by MTT assay, platycodin D has been shown to inhibit the expressions of c-Myc in U937 leukaemia cells [102], phosphorylated PLC γ 1 in HUVECs [118] and cyclin D1 in PC3 prostate [45] and AGS gastric [73] cancer cells. Platycodin D also inhibited tumour growth and expression of Ki-67, a well known marker of cell proliferation in MDA-MB-231 xeno-graft tumours in athymic nude mice at a dose of 5 mg/kg [75].

Targeting pro-survival pathways in cancer cells

One of the major characteristic features of cancer cells is their ability to avoid apoptotic cell death by activating pro-survival signalling. PI3K/AKT [1] and NF- κ B [119, 120] signalling pathways are frequently overexpressed in a variety of human cancers. Several antiapoptotic proteins such as Bcl-2 and survivin which are known to be crucial for cancer cell survival are the downstream target genes of AKT and NF- κ B. In MDA-MB-231 breast cancer cells, platycodin D inhibits both constitutive and EGF-induced activation of AKT and mTOR [75]. It also inhibits constitutive and LPS-stimulated activation of NF- κ B in MDA-MB-231 [75] and RAW 264.7 [41] cells respectively. Platycodin D has also been reported to inhibit Pl3K/AKT signalling pathway in HUVECs [118], HepG2 [100], U251 [84], NCI-H460, A549 [112], U937 [102] and AGS [73] cells.

In addition to AKT and NF- κ B signalling, many studies have demonstrated that platycodin D is a potent inhibitor of anti-apoptotic protein Bcl-2 and survivin. It down-regulated Bcl-2 in MCF-7 [72], AGS [73], HepG2 [100], PC3 [45] and U251 [84], and survivin in HepG2 [56] and AGS [73] cells.

Targeting angiogenesis in cancer cells

Angiogenesis is a complex physiological process by which new blood vessels are formed from pre-existing structures [121]. Angiogenesis plays a vital role in the growth and survival of solid neoplasm and is considered as a hallmark of cancer progression. Although angiogenesis is regulated by multiple signalling pathways, it is now universally accepted that VEGF and its cognate receptor VEGFR-2 are the most prominent regulators [122]. VEGF signalling stimulates cellular pathways that promote rapid tumour growth and facilitate metastatic potential of cancer by forming new blood vessels for adequate blood supply [122]. The development of highly selective drugs that target VEGF such as Bevacizumab or VEGFR-2 such as Sorafenib has improved the prognosis of patients with metastatic cancer but fail to achieve the desire outcome because of activation of other compensatory mechanisms such as fibroblast growth factor and COX-2/ PGE2. Thus, simultaneous targeting of VEGF and other compensatory pathways may provide a useful strategy for the development of most effective drugs for metastatic tumours.

Platycodin D has recently been investigated for its anti-angiogenic effects. Luan *et al.* [118] showed that platycodin D inhibits the growth, migration and tube formation in HUVEC in a dose-dependent

manner at a very low concentration (0.3-30 µM). It also inhibits angiogenesis in chick embryo chorioallantoic membrane. The antiangiogenic effect of platycodin D in tumour was validated using HCT-15 xenograft mouse model. Platycodin D inhibited tumour growth, decreased tumour microvessel density and increased apoptosis in tumour tissue at a dose of 6 mg/kg/day for 14 days without any toxic effects to mice. The anti-angiogenic activity of platycodin D appeared to result from its suppressive effect on VEGF-induced VEGFR-2 phosphorylation and its downstream targets such as AKT, JAK-2, Src and PLCy1 [118]. Platycodin D has decreased the expression of proangiogenic factor VEGF and COX-2 in a dose-dependent manner (10-20 µM) in AGS cells [73]. In RAW 264.7 cells, platycodin D downregulates the LPS-induced expression of COX-2 and iNOS at both transcriptional and translational levels. Platycodin D also inhibited the LPS-induced synthesis of prostaglandin E2 (PGE2) the enzymatic product of COX-2 in RAW 264.7 cells [41]. Collectively, platycodin D exerts angio-suppressive activity by regulating the expressions of VEGF, VEGFR-2, iNOS, COX-2 and PGE2 in different cells.

Targeting invasion and metastasis

Tumour invasion and metastasis are complex, multi-step biochemical processes and are considered as one of the hallmarks of cancer biology. Invasion is the direct migration and penetration of cancer cells into neighbouring tissues while metastasis is the ability of cancer cells to penetrate into lymphatic or blood vessels, circulate through the bloodstream and then invade in normal tissues elsewhere in the body [123]. Degradation of basement membrane and remodelling of extracellular matrix (ECM) are the characteristic features of cancer invasion and metastasis. Matrix metalloproteinases (MMPs) especially MMP-2 and -9 are aberrantly expressed in cancers and are believed to play vital role in cancer invasion and metastasis [124, 125]. Metastasis has been reported to responsible for about 90% of cancer patient deaths and is considered as the most challenging obstacle in successful treatment of cancer [125].

Platycodin D has been shown to inhibit migration and invasion of MDA-MB-231 cells in a dose-dependent manner (5–15 μ M) as seen from wound healing assay and matrigel transwell chamber assay respectively. Treatment of cells with 15 μ M platycodin D inhibited the adhesion of cells to fibronectin, collagen type I, laminin and fibrinogen by 42%, 54%, 49% and 50% respectively indicating that suppressive effect of the compound on MDA-MB-231 cells invasion is associated with decreased adhesion of cells to ECM proteins. Zymographic analysis revealed that platycodin D markedly inhibited the activity of MMP-9, while MMP-2 activity was slightly decreased. Platycodin D also down-regulates the mRNA expression of MMP-9 as evident from real-time PCR analysis [75]. Platycodin D's anti-invasion and anti-metastatic activity may be associated to its ability to inhibit proteolytic activity and mRNA expression of MMP-9.

In HepG2 cells, platycodin D inhibited colony formation and suppressed adhesion to matrigel dose-dependently ($1.25-10 \mu M$). Platycodin D also inhibits 12-O-tetradecanoylphorbol 13-acetate (TPA)-induced migration and invasion of HepG2 cells as seen from transwell chamber assay [56]. Platycodin D has also decreased the

expression and phosphorylation of FAK, a component of cell substratum adhesion, in AGS [74] and HUVEC [118]. Among other players of invasion and metastasis, platycodin D reduced VEGF-induced phosphorylation of JAK2 and Src in human umbilical vein endothelial cells in a dose-dependent fashion [118].

In vivo anti-tumour effects of platycodin D

Apart from *in vitro* anti-cancer activity of platycodin D, various tumour models and drug administration routes have been tested for platycodin D *in vivo*. Both intraperitoneal as well as oral administration of platycodin D has been shown to exhibit significant anti-tumour activity. Intraperitoneal administration of platycodin D at a dose of 5 mg/kg bodyweight significantly reduced the volume and size of MDA-MB-231 xenograft tumour. Immunohistochemical analysis of tumour tissues shows that platycodin D inhibited the expression of Ki-67 and EGFR [75]. EGFR inhibition seems to be major mechanism of platycod in D's *in vivo* anti-tumour efficacy as demonstrated by *in vitro* cell studies.

In PC3 prostate cancer xenograft model, intraperitoneal administration of 2.5 mg/kg platycodin D achieved 56% tumour growth inhibition. At a very low dose of 1 mg/kg, platycodin D also reduced tumour size but not significantly. Immunobloting analysis of tumour tissues showed that platycodin D enhanced the expression of FOXO3a, p21, p27, Bax and cleaved PARP, while decreased the expression of MDM2, Bcl-2, cyclin B1 and D1 and CDK-1, -2, -4 and -6. FOXO3a seems to hold the promise of platycodin D-mediated anticancer activity [45].

A recent study by Park *et al.* [44] has shown anti-tumour effect of platycodin D in H520 cell xenograft Balb/c *nu-nu* nude mice. Oral administration of 50, 100 and 200 mg/kg platycodin D, once a day for 5 weeks from 15 days after tumour cells inoculation has significantly reduced the tumour size and weight with remarkable immune-stimulatory and anti-cachexia effects in a dose-dependent manner. Gemcitabine (160 mg/kg) has been used as positive control in this study. Gemcitabine also exerted potent anti-tumour effects but aggravated the cancer-related immune suppressions and cachexia in this experiment.

More recently, Lee *et al.* [103] has shown that platycodin D inhibits breast cancer-induced bone destruction by inhibiting osteoclastogenesis and inducing apoptosis in MDA-MB-231 breast cancer cells in Balb/c *nu/nu* mice. Oral administration of 2 mg/kg platycodin D has significantly inhibited MDA-MB-231 cell-induced osteolysis and tumour growth in bone marrow in intratibial mouse model. Collective data of this study demonstrate that platycodin D inhibits recombinant mouse receptor activator of nuclear factor kappa-B ligand-induced osteoclast formation by inhibiting the expression and nuclear translocation of NFATc1 and c-Fos in bone marrow-derived macrophages and ultimately reduces osteoclast-mediated bone resorption.

In addition to anti-tumour effects, platycodin D has also been explored for its toxic effects on general body organ using male and

female ICR mice. A single oral dose of platycodin D up to 2000 mg/kg could be safely administered without causing treatment related mortality, abnormal clinical signs, remarkable changes in body and organ weights and significant histopathological changes in 14 principal organs till 14 days of post-treatment in both male and female mice [43]. The Korean Food and Drug Administration [43] guidelines indicate that the maximum platycodin D dosage recommendation is 2000 mg/kg.

Platycodin D has also been demonstrated to ameliorate cisplatininduced nephrotoxicity in ICR mice by attenuating oxidative stress and cell death in kidney [58]. It should also be noted that platycodin D was well tolerated and no mortality or any sign of pharmacotoxicity was observed in aforementioned tumour bearing nude mice during all experimental periods.

Platycodin D in combination therapy

The combined application of saponins with other anti-tumour drugs has drawn a considerable attention in cancer chemotherapy because of huge structural diversity and adjuvant activity of saponins. Because of excellent anti-tumour effect of platycodin D in various in vitro and in vivo cancer models, it has recently been investigated for its combined application with other anti-tumour drugs in various cell lines of breast cancer. In MCF-7 (oestrogen receptor positive) and MDA-MB-231 (oestrogen receptor negative) cell lines of breast cancer, platycodin D has been shown to enhance the anti-cancer activity of doxorubicin. While treatment with either platycodin D (10 μ M) or Doxorubicin (2.5 μ M) in vitro resulted in apoptotic cell death, a combine treatment of both demonstrated a significant growth suppressive effect and PARP cleavage in both cell lines compared to treatment with either agent alone [126]. As platycodin D increased the intracellular accumulation of doxorubicin in given concentration only in MDA-MB-231 cells, the equally increased growth suppressive effects of combine treatment in both cell lines could not be associated with enhanced accumulation of doxorubicin in cells.

Platycodin D has also been shown to exhibit greater growth inhibitory and anti-invasive effects in combination with osthol in MDA-MB-231 and 4T1 breast carcinoma cells [42]. Platycodin D (75 μ M) in combination with osthol (15 μ M) synergistically inhibits the growth and invasion of both cancer cell lines. Both these compounds have been shown to exert varying effects on the expression of TBRII. Smad2 and Smad3: however, a combine treatment of both agents effectively reduced the mRNA and protein expressions of TBRII, Smad2 and Smad3 in both cell lines. Moreover, combine treatment effectively decreased the TGF (transforming growth factor)-B-induced phosphorylation of Smad2 and Smad3. Taken together, the data demonstrate that platycodin D-osthol combination inhibits proliferation and blocks invasion of MDA-MB-231 and 4T1 breast cancer cells by inhibiting TGF-β/Smad signalling. Thus, platycodin D may be developed into a potential lead compound for treating breast cancer in combination with other chemotherapeutic agents.

Concluding remarks and future perspectives

In this review, we have highlighted the recent progress of platycodin D in various *in vitro* and *in vivo* cancer models. Collective data from multitudinous studies has provided persuasive evidence for promising role of platycodin D in treatment of many type of cancers. It has been shown to interact with multiple mechanisms to kill cancer cells in a wide variety of tumours. The most exciting possibility to develop platycodin D into a lead is its low toxicity to normal cells and good adjuvant property. The major drawback to develop saponins into a successful chemotherapeutic drug is their severe toxicity to normal cells. Contrary to most of the saponins, platycodin D has been shown to exert less toxic effects on normal cells; however, the exact mechanism still remains unexplored. As platycodin D has been shown to exhibit excellent anti-cancer activity by interfering with multiple mechanisms which are the hallmark of cancer, further detailed studies regarding its structure–activity relationship for haemolysis and

adjuvant property should be conducted with special focus on the role of sugar chains. As evident by study of Chun *et al.*[73], deglycosylation of platycodin D at C-3 and C-28 reduces its cytotoxicity while *o*acetylation at rahmnose C-2 or C-3 as well as dehydroxylation at C-24 of platycodin D has been shown to increase its cytotoxicity against various cancer cells. Medicinal chemistry studies may be needed to improve the anti-cancer and adjuvant activity and reduce the haemolytic ability of platycodin D to develop it into a successful chemotherapeutic drug either alone or in synergistic combination with other drugs.

It is hoped that the data collected here will provide the researchers with valuable information and establish the basis for the future development of platycodin D into a promising anti-cancer drug in the years to come.

Conflicts of interest

The authors confirm that there are no conflicts of interest.

16.

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