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Cucurbitacins as Multifunctional Anticancer Agents: Chemistry, Molecular Mechanisms, Translational Challenges, and Emerging Therapeutic Opportunities

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ABSTRACT: Cancer continues to be a significant health challenge worldwide and one of the primary contributors to morbidity and mortality. Despite significant progress in the field of immunotherapy, targeted therapy and molecular oncology, the need for improved and safer anti-cancer drugs remains. For a long time, natural ingredients have been a significant source of novel medications. Among these bioactive substances, Cucurbitacins diverse pharmacological properties have made them very promising as madidates. Cucurbitacins represent a class of extremely oxygenated tetracyclic triterpenoids that are predominantly isolated from Cucurbitaceae plants but also isolated from many other plants and fungi. They have been shown to have potent anti tumour properties in various experiments using different types of cancers, including breast cancer, lung cancer, colon cancer, pancreatic cancer, hepatocellular carcinoma, ovarian cancer, prostate cancer, and skin cancer. The updated discussion builds on the idea of the Cucurbitacins as single cytotoxic agents, but incorporates a more comprehensive perspective on multitarget pharmacology, tumour signalling plasticity and translational oncology. When appropriate, the emphasis changes from descriptive reporting to mechanistic interpretation, comparison to other studies and identification of unanswered questions useful for clinical development.

In this review paper, the chemistry, biosynthesis, pharmacological properties and anticancer activity of cucurbitacins have been briefly discussed, with emphasis on the four main members of the group, namely cucurbitacin Q, I, E and D; and cucurbitacin B. These substances have anticancer effect thru a variety of methods, such as suppression of the JAK/STAT3, PI3K/Akt/mTOR, and NF- κ B pathways; activation of autophagy and apoptosis; suppression of angiogenesis; disruption of the cytoskeleton; and control of EMT and cancer stem cells. Overall, cucurbitacins are potential and natural compounds to develop anticancer drugs. Furthermore, recent advances in NDDS, combination strategies with conventional chemotherapeutic agents, pharmacokinetic limitations, safety concerns, and future clinical prospects are discussed. Although substantial preclinical evidence supports the therapeutic value of Cucurbitacins, their clinical development remains constrained by challenges such as poor bioavailability, potential systemic toxicity, and limited clinical evaluation. Taken together, A diverse family of naturally occurring substances with great potential for cancer treatment is cucurbitacins. Continued research integrating medicinal chemistry, molecular pharmacology, nanotechnology, and clinical oncology may facilitate the development of cucurbitacin-based treatments for precision cancer medicine.

1. INTRODUCTION

Cancer is a complex and diverse disease with the following characteristics: Abnormal cell proliferation, resistance to programmed cell death, persistent angiogenesis, invasion of adjacent tissue, metastatic dissemination, and emergence of treatment resistance (1,2). Worldwide health surveys report cancer remains one of the leading causes of mortality and a significant burden on health care budgets worldwide (2–4). Many patients today have improved life expectancy due to advances in surgery, chemotherapy, radiotherapy, immunotherapy and targeted therapy, but there are still problems with treatment, such as side effects, resistance to the drugs, recurrence of the disease and poor tumour specificity (2,5). Natural products have been a significant source for developing anticancer drugs. These days, there are numerous approved anti-cancer drugs of natural origin. A large number of the currently approved anti-cancer agents are derived from nature or are natural products (6,7). Examples include paclitaxel, vincristine, drugs based on calprotectin, and anthracycline antibiotics. In the past few years, the plant-derived phytochemicals with multi-targeted anticancer activities have gained more attention, and the cucurbitacins have always been identified as showing a high therapeutic potential (7).

Cucumber, pumpkin, watermelon, bitter melon, squash and gourd species of the Cucurbitaceae family are the primary source of the highly oxygenated tetracyclic terpene class called cucurbitacins. (8,9,11). These secondary metabolites give many cucurbit plants a very bitter flavor and help to provide natural insect protection, herbivore protection, and microbiological disease protection (9,11). To date, over 200 cucurbitacin-related compounds and derivatives have been found and classified in a few structural groups (8,11).

Cucurbitacins have several pharmacological properties such as hepatoprotective, antidiabetic, antioxidant, antibacterial and antiviral properties in addition to their ecological properties. (9,12,13). Their anticancer activity is particularly remarkable as they may simultaneously regulate multiple cellular signalling pathways related to tumour genesis, tumour development, tumour metastasis and treatment resistance (8,10,14). Cucurbitacins are appealing candidates for overcoming tumour heterogeneity and lowering the risk of treatment resistance due to their capacity to target a variety of molecular processes (10,14,15).

The anticancer effects of cucurbitacins are believed to be mediated by their effects on various cellular and molecular pathways that play a role in tumour formation and progression (8,10,14). These chemicals are likely to induce programmed cell death (apoptosis), halt the cell cycle, block angiogenesis, and prevent cancer invasion and metastasis, according to all data available (5,12,13). Moreover, it has been demonstrated that cucurbitacins are able to affect different processes, such as inducing autophagy, disturbing the cytoskeletal architecture and enhancing the production of reactive oxygen species (ROS), all of which cause Mayer cell death (13,14). They are also known to inhibit important cancer-promoting signalling pathways, such as JAK/STAT3, PI3K/Akt/mTOR and NF- κ B (13,14,16). Furthermore, Cucurbitacins have been shown to inhibit epithelial–mesenchymal transition (EMT) and modulate the behaviour of Mayer stem cells, both of which are crucial in tumour progression, metastasis, and resistance to treatment (10,15).

Although promising in vitro and in vivo results have been obtained, the clinical development of the cucurbitacins is still hindered (8,9,14). Their medicinal use has been hampered by a number of factors, including as low water solubility, limited bioavailability, quick metabolic breakdown, and possible systemic toxicity (9,17). But recent advances in pharmacological and nanotechnological methods have opened up new avenues for improvement of their therapeutic potential. Methods such as prodrug, targeted drug delivery systems, nanoparticle formulation, and structural modification have consistently demonstrated their ability to enhance drug stability, bioavailability and therapeutic efficacy, and decrease undesirable effects (17).

2. CHEMISTRY AND STRUCTURAL CHARACTERISTICS OF CUCURBITACINS

2.1 General Chemical Structure

Cucurbitacins are tetracyclic triterpenoids that emerge from squalene cyclisation and are biosynthesized via the mevalonate route. They are produced from the cucurite skeleton, which consists of thirty carbon atoms organized in a highly oxygenated framework (8,9,11).

The basic cucurbitane skeleton contains:

The cucurbitane skeleton represents the characteristic tetracyclic triterpenoid framework of Cucurbitacins and is composed of four fused rings (A–D) (8,11). Multiple hydroxyl groups, ketone moieties, double bonds, and an unsaturated side chain

functionalize this core structure, all of which are crucial in determining the physicochemical and biological characteristics of distinct analogues (11,12). The presence and specific arrangement of these structural elements are essential for the biological activity of Cucurbitacins, as they influence molecular interactions with biological targets (8,9,11). Consequently, these features contribute to the diverse pharmacological properties of Cucurbitacins, including their anticancer, anti-inflammatory, antioxidant, and other therapeutic activities (9,12)

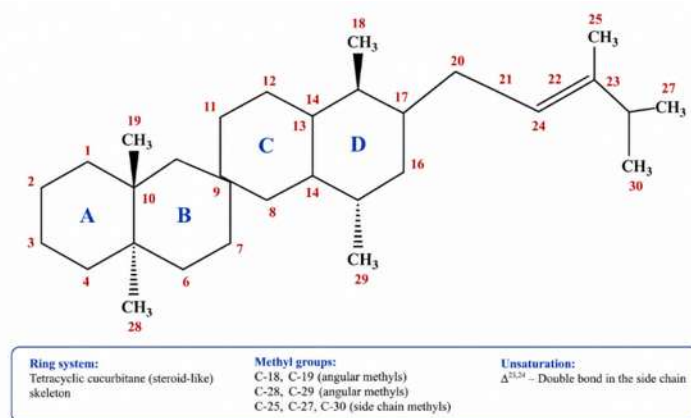


Figure1. Generalized Cucurbitacin Skeleton

IUPAC NAME

(8R,9S,10R, 13R, 14S, 17R)-17-[(2E)-6,6,10-trimethylundeca-2,4-dien-1-yl]-1,2,3,4,4,5,6,7,8,9,10,11,12,13,14,15,16-tetradecahydrocyclopenta[a]phenanthrene

2.2 Major Types of Cucurbitacins

Sr. No.	Cucurbitacin Type	Major Biological Activities	Reference
1	Cucurbitacin B	STAT3 inhibition, apoptosis induction	(13,18,19)
2	Cucurbitacin D	Anti-metastatic activity	(20,21)
3	Cucurbitacin E	Cytoskeletal disruption	(22,23)
4	Cucurbitacin I	JAK/STAT inhibition	(18,22)
5	Cucurbitacin Q	Anti-proliferative effects	(8,24)
6	Cucurbitacin R	ROS generation	(8,24)

Table 1: Major Cucurbitacin Analogues and Their Major Biological Activities

Cucurbitacins are classified into more than 12 main classes. The two that have been examined the most are cucurbitacin B and cucurbitacin I.

2.3 Structure–Activity Relationship (SAR)

The structural properties of cucurbitacins have a significant impact on their biological action, and even minor modifications within the cucurbitane framework can markedly alter their pharmacological properties (11,12,18). The α , β -unsaturated carbonyl group is considered the principal pharmacophore responsible for the biological activity of many cucurbitacin analogues (12,17,18). This electrophilic moiety can undergo Michael addition reactions with nucleophilic amino acid residues, particularly cysteine residues present in signalling proteins, leading to modulation of pathways such as STAT3, NF- κ B, and PI3K/Akt (8,17,18,22). The number, position, and stereochemical orientation of hydroxyl groups substantially influence

cytotoxic activity, aqueous solubility, and target selectivity (11,12,18). Increased hydroxylation generally enhances hydrogen-bonding interactions with biological macromolecules and may improve affinity for molecular targets (12,25). Acetylation of hydroxyl groups increases lipophilicity and can facilitate membrane permeability and intracellular accumulation of cucurbitacin derivatives (17,25). The position and geometry of carbon–carbon double bonds also contribute to molecular conformation and biological activity, and the presence of a double bond between C-23 and C-24 has been associated with enhanced potency in several naturally occurring analogues (11,12,25). Furthermore, modifications of the side chain, including changes in chain length, oxidation state, and functional group substitution, can significantly affect cytotoxicity, selectivity, and pharmacokinetic behavior (20,25). Collectively, structure–activity relationship studies indicate that the combined effects of hydroxylation pattern, acetylation status, electrophilic carbonyl functionality, double-bond configuration, and side-chain modifications determine the potency and therapeutic potential of cucurbitacins (6,11,12,16,17,25).

Important Structure–Activity Relationship (SAR)

Observations

Studies on the structure–activity relationship of Cucurbitacins have demonstrated that specific structural modifications can markedly influence their biological and pharmacological properties (1–4). In general, an increased degree of hydroxylation is associated with enhanced cytotoxic activity, likely due to improved hydrogen-bonding interactions with biological macromolecules and greater affinity for molecular targets (12,22,25). Acetylation of hydroxyl groups has been reported to increase the lipophilicity of cucurbitacin derivatives, which may facilitate membrane permeability and improve cellular uptake (3,5,6). Furthermore, the presence of a double bond between C-23 and C-24 in the side chain is considered an important structural feature that contributes to the biological potency of several cucurbitacin analogues (2,7,8). Another critical determinant of activity is the carbonyl functionality, particularly when it forms part of an α , β -unsaturated carbonyl (neon) system. This electrophilic moiety readily undergoes Michael addition reactions with nucleophilic residues, especially cysteine thiol groups in cellular proteins, leading to the modulation of key signaling pathways involved in cell proliferation, apoptosis, and inflammation (3,6,9–11). Collectively, these structural characteristics play a pivotal role in defining the potency, selectivity, and therapeutic potential of Cucurbitacins (2,6,8,10,12,13).

3. NATURAL SOURCES AND BIOSYNTHESIS

3.1 Plant Sources

Cucurbitacins are naturally occurring, highly oxygenated tetracyclic triterpenoids that are mostly found in plants belonging to the Cucurbitaceae family.(14,15). Members of this family, including *Citrullus colocynthis*, *Cucumis sativus*, *Bryonia dioica*, *Momordica charantia*, and *Trichosanthes cucumerina*, represent the major natural reservoirs of different cucurbitacin analogues (8,14). Individual plant species differ considerably in both the type and concentration of cucurbitacins they produce owing to genetic, environmental, and developmental factors (8,15). Among the naturally occurring analogues, cucurbitacin B, C, D, E, and I are the most extensively investigated because of their remarkable pharmacological and anticancer activities (2,16).

Important Sources

Sr. No	Plant Species	Major Cucurbitacins	References
1.	<i>Citrullus colocynthis</i>	Cucurbitacin E	(2,14)
2.	<i>Cucumis sativus</i>	Cucurbitacin C	(14,15)
3.	<i>Bryonia dioica</i>	Cucurbitacin D	(2,8)
4.	<i>Momordica charantia</i>	Cucurbitacin B	(8,17)
5.	<i>Trichosanthes cucumerina</i>	Cucurbitacin I	(2,17)

Table 2: Major Plant Sources of Cucurbitacins and Their Predominant Cucurbitacin Analogues

Although cucurbitacins are predominantly associated with higher plants, cucurbitacin-like triterpenoids and structurally related metabolites have also been reported in certain fungi and marine organisms, indicating a broader natural distribution of these bioactive compounds (16,18).

3.2 Biosynthetic Pathway

The biosynthesis of cucurbitacins originates from the mevalonate (MVA) pathway, which produces the universal isoprenoid precursor 2,3-oxidosqualene through a series of enzymatic reactions (14,18). Cyclization of 2,3-oxidosqualene by oxidosqualene cyclase's generates the characteristic cucurbitadienol skeleton, which serves as the primary precursor for the biosynthesis of all cucurbitacin analogues (18,19). Subsequent oxidation, hydroxylation, and rearrangement reactions catalyzed primarily by cytochrome P450 monooxygenases introduce multiple oxygen-containing functional groups, giving rise to structurally diverse cucurbitacins (14,19,20). The cucurbitane skeleton consists of four fused rings (A–D) and is subsequently modified by hydroxylation, ketone formation, acetylation, glycosylation, and side-chain oxidation to produce different cucurbitacin derivatives (14,19). Variations in the number, position, and stereochemistry of these functional groups are responsible for the remarkable structural diversity and biological activities of cucurbitacins (14,16,19). These structural modifications significantly influence their pharmacological properties, including anticancer, anti-inflammatory, antioxidant, antimicrobial, hepatoprotective, antiviral, and antidiabetic activities (2,17,20). Cytochrome P450 enzymes play critical roles in catalyzing oxidation reactions that generate structurally diverse cucurbitacin analogues and regulate the accumulation of these metabolites in different plant species (14,20).

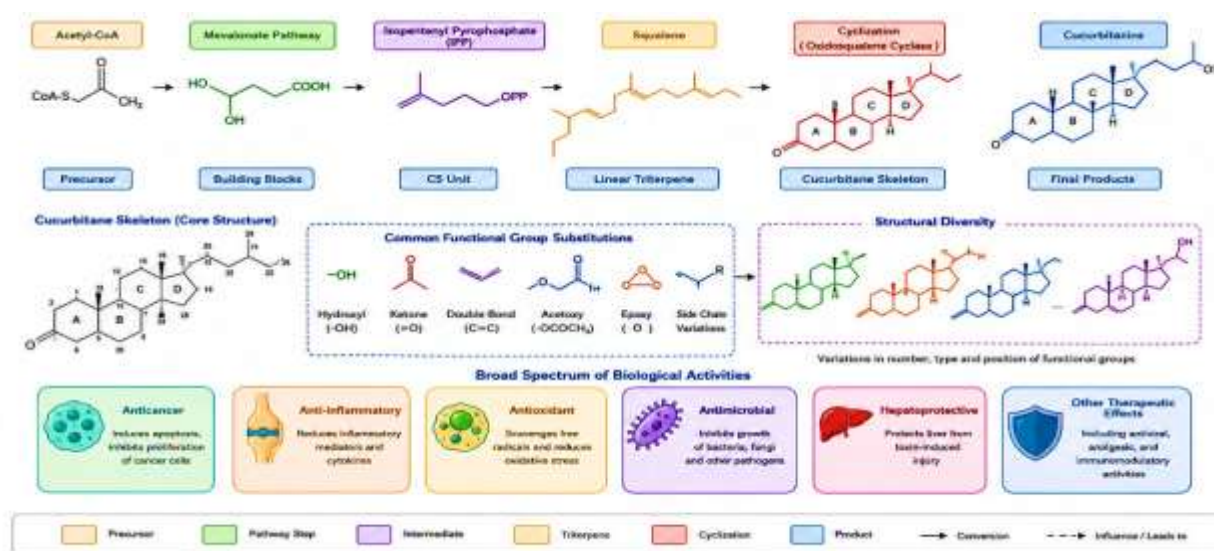


Figure 2. Simplified Biosynthetic Pathway

Cytochrome P450 enzymes play critical roles in catalysing oxidation reactions that generate structurally diverse cucurbitacin analogues and regulate the accumulation of these metabolites in different plant species (14,20).

4. PHARMACOLOGICAL PROPERTIES OF CUCURBITACINS

In addition to their well-documented anticancer potential, Cucurbitacins possess a broad spectrum of pharmacological activities that contribute to their therapeutic significance (3,8,9). These naturally occurring triterpenoids have hepatoprotective, immunomodulatory, antiviral, anti-inflammatory, antioxidant, and antibacterial properties (3,8,9,17). The capacity of cucurbitacins to influence many cellular signalling pathways involved in inflammation, oxidative stress, cell proliferation, and apoptosis is responsible for their wide range of biological actions (2,3,9).

4.1 Anti-Inflammatory Activity

Through the modulation of various important mediators and signal transduction pathways implicated in the inflammation process, cucurbitacins exhibit potent anti-inflammatory properties (3,9). This results in inhibition of inflammatory enzymes such as COX-2 and iNOS as well as reduction in the release of various pro-inflammatory cytokines such as IL-6 and TNF- α .

(2,3,9). Cucurbitacins also inhibit the NF- κ B signalling pathway that acts as an important regulatory pathway for inflammatory genes (3,21,22). In light of the fact that inflammation has been shown to have an important role in the initiation and progression of cancers, anti-inflammatory properties of cucurbitacins have been considered one of the major mechanisms underlying their chemoprotective effects (22,27).

4.2 Antioxidant Activity

Cucurbitacins have an essential function in regulating cellular oxidative stress by lowering the production of ROS, enhancing the inherent antioxidant defenses of the body and maintaining mitochondrial health (2,17,23). The above actions assist in maintaining cellular redox balance and protecting healthy cells from oxidative stress (17, 23). It is important to observe that the activity of cucurbitacins results in a controlled increase in the level of intracellular ROS in cancer cells, resulting in oxidative stress that is beyond the antioxidant capacity of the cell and thus causes apoptosis (2,17,24). Excess ROS activity leads to the loss of the mitochondrial membrane potential, activates the intrinsic apoptotic pathway and causes caspase-mediated cell death (17,24,25). Cucurbitacins not only induce ROS activity but also regulate antioxidant signalling pathways such as Nrf2/Keap1 pathway, affecting the synthesis of antioxidant enzymes such as GPx, CAT and SOD (2,25). Cucurbitacins not only induce ROS activity but also regulate antioxidant signalling pathways such as Nrf2/Keap1 pathway, affecting the synthesis of antioxidant enzymes such as GPx, CAT and SOD (2,25).

4.3 Antimicrobial and Antiviral Activity

The interest in the pharmacological properties of cucurbitacins has grown significantly owing to their wide-spectrum antibacterial and antiviral activity as well as anticancer activity (2,8,17). The inhibitory effects of cucurbitacins on the development of various kinds of pathogenic fungi, Gram-positive and Gram-negative bacteria have been proven in scientific experiments (2,8). Such antimicrobial effects are believed to be related to the interference with the key intracellular signalling pathways, metabolic suppression in cells and destruction of membranes in microbes (2,17). Moreover, a number of studies prove that cucurbitacins possess antiviral activity against various DNA and RNA viruses due to the suppression of virus replication, the suppression of interactions between virus and host and the modulation of host immunity (8,9). In addition, the suppression of JAK/STAT and NF- κ B signalling pathways has been identified in various studies as the factor promoting the antiviral activity of cucurbitacins by limiting the virus proliferation and inflammation induced by the virus (9,17). Despite the promising results in this field, additional studies in vivo and clinical trials are required to identify the effectiveness and safety of using cucurbitacins as the antiviral and antibacterial agents (2,8,9,17).

4.4 Immunomodulatory Activity

It has been demonstrated that cucurbitacins possess immunomodulatory properties by affecting the activation, differentiation and cytokine production of immune cells (9,10,17). Cucurbitacins affect the secretion of proinflammatory cytokines and chemokines; therefore, they help to maintain the balance of immune responses and regulation of inflammatory responses (9,17). In addition, cucurbitacins can regulate the signalling pathways including JAK/STAT3 and NF- κ B involved in immune cell function, inflammatory signalling and tumour immune escape mechanisms (9,10,17). By inhibiting these pathways, cucurbitacins could increase antitumour immunity, reduce chronic inflammation, and change the tumour microenvironment for immune-mediated destruction of cancer cells (10,17). Thus, cucurbitacins become promising compounds for the development of combination treatments using immunotherapy and targeted anticancer therapy (10,17,26). However, further investigation is needed to understand the immunomodulatory activities of cucurbitacins and apply these compounds in clinical practice (9,10,17,26).

5. MOLECULAR MECHANISMS OF ANTICANCER ACTIVITY

5.1 Induction of Apoptosis

By eliminating dysfunctional cells, either through damage, malfunctions, or unnecessary cells, apoptosis – which is a highly regulated and energy-requiring type of programmed cell death – acts as a significant factor in maintaining tissue homeostasis (27, 28). Contrary to necrosis, apoptosis allows for maintenance of tissue integrity, since the process occurs systematically and without causing inflammation (12, 28). Malignant cell resistance to apoptosis forms part of the cancer hallmarks which causes uncontrolled growth, formation of tumours, and resistance to conventional cancer therapies (5, 12, 28). Therefore, reinstating apoptotic signalling has formed a key therapeutic approach in cancer treatment (5, 12). Various in vitro studies have

demonstrated that cucurbitacins induce apoptosis in several human cancer cell lines by triggering the intrinsic and extrinsic apoptotic pathways (24, 29-31).

Intrinsic (Mitochondrial) Pathway

Intracellular stress signals, including DNA damage, oxidative stress, hypoxia, endoplasmic reticulum stress, and chemotherapeutic drug exposure, are examples of some stimuli that activate the intrinsic apoptotic pathway (24, 27). Mitochondrial outer membrane permeabilization followed by the release of cytochrome c into the cytosol is a direct consequence of injury caused by the above stimuli on the integrity of the mitochondrial membrane (28, 29). Once cytochrome c connects with the apoptotic protease-activating factor-1 (Apaf-1) to form the apoptosome, it activates caspase-9. Executioner caspases, which include caspase-3 and caspase-7, follow (27, 28, 30). Proteins of the Bcl-2 family strictly regulate the mitochondrial pathway, where anti-apoptotic proteins, such as Bcl-2 and Bcl-xL, prevent the permeabilization of the mitochondrial membrane while the pro-apoptotic proteins, including Bax and Bak, promote it (5,12). Cucurbitacins induce mitochondrial apoptosis through the activation of the mitochondrial pathway. Proteins of the Bcl-2 family strictly regulate the mitochondrial pathway, where anti-apoptotic proteins, such as Bcl-2 and Bcl-xL, prevent the permeabilization of the mitochondrial membrane while the pro-apoptotic proteins, including Bax and Bak, promote it (5,12). Cucurbitacins induce mitochondrial apoptosis by activating the caspase cascade, upregulating Bax expression, downregulating Bcl-2, and promoting cytochrome c release, all of which result in apoptotic cell death, according to several studies (24, 29–32).

Extrinsic (Death Receptor) Pathway

Death ligands present outside the cell, such as FasL and tumour necrosis factor-related apoptosis-inducing ligand (TRAIL), bind to death receptors in the cell surface to initiate the extrinsic apoptotic pathway (12,27). The activation of initiator caspase-8 occurs via death-inducing signalling complex (DISC) formation, which happens after the interaction of ligand and death receptors (5,28). Programmed cell death results from the cleavage and activation of executioner caspases by activated caspase-8, primarily caspase-3 (5,27). Moreover, through mitochondrial membrane permeabilization, activated caspase-8 cleaves Bid to truncated Bid (tBid), thus linking both intrinsic and extrinsic apoptotic pathways (12,28). It has been proven experimentally that cucurbitacins have increased apoptotic signalling in cancer cells, which enhances death receptor-mediated apoptosis via caspase-8 activation, DISC formation, and Bid cleavage (29–32).

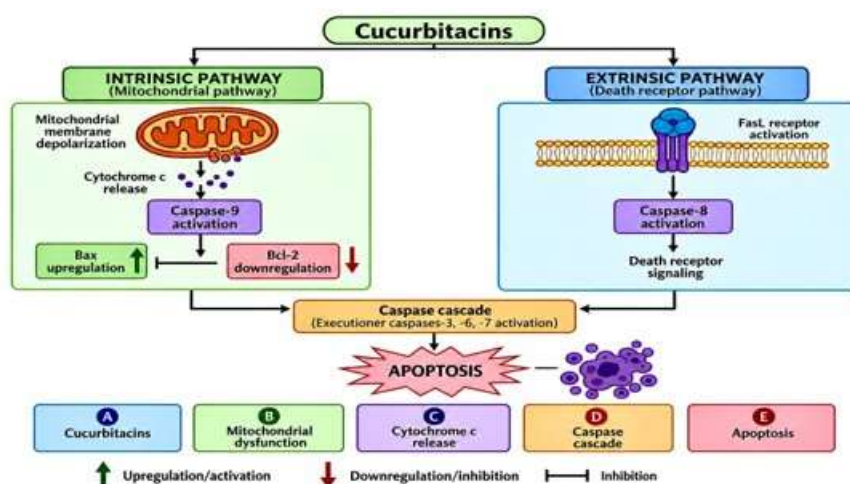


Figure 3: Molecular Mechanisms of Cucurbitacin-Induced Apoptosis in Cancer Cells

5.2 Cell Cycle Arrest

The tightly regulated process of cell cycle development ensures precise DNA replication and cell division (30, 33). Cancer cells often evade these regulatory checkpoints, resulting in unregulated proliferation (34, 35). One of cucurbitacins' primary anticancer actions is their ability to induce cell cycle arrest, which prevents malignant cells from multiplying and dividing (30, 36). Depending on the kind of cancer and the specific cucurbitacin analogue, cell cycle arrest can happen during the G1, S, or G2/M phase (33, 37). This impact is produced via modulating cyclins, cyclin-dependent kinases (CDKs), and cell cycle

regulatory proteins such as p21, p27, and p53 (37, 38). Cucurbitacins inhibit the growth of cancer cells and encourage apoptosis by interfering with the advancement of the cell cycle (30, 38).

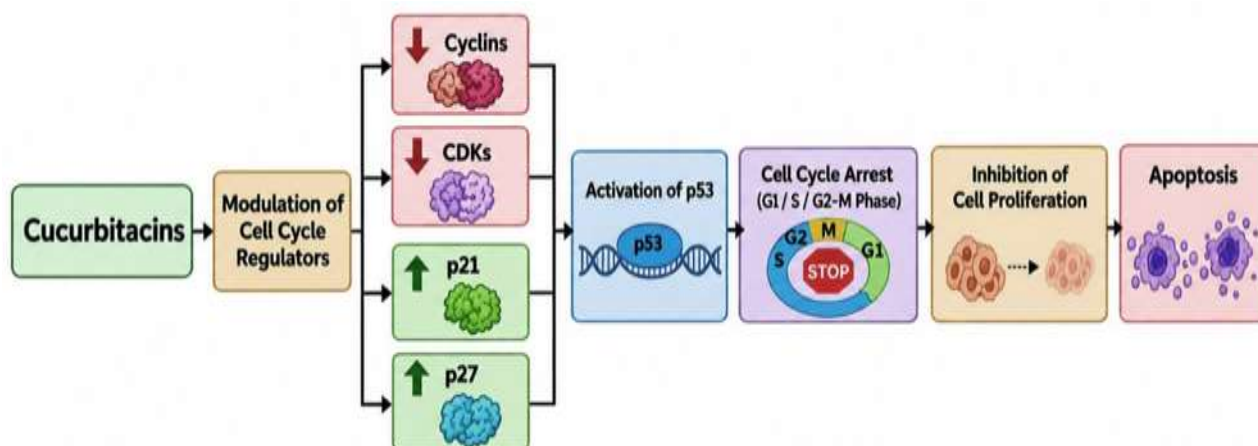


Figure 4: Mechanism of Cucurbitacin-Induced Cell Cycle Arrest in Cancer Cells

5.3 Inhibition of JAK/STAT3 Signalling

Angiogenesis, immunological evasion, cell proliferation, survival, and metastasis are all regulated by the Janus kinase/signal transducer and activator of transcription 3 (JAK/STAT3) signalling pathway (40,41). In a number of human malignancies, persistent STAT3 activation is linked to treatment resistance and a poor prognosis (42, 43). Thus, blocking the JAK/STAT3 pathway has become an essential therapeutic approach for cancer therapy (40,44). By preventing STAT3 phosphorylation and nuclear translocation, cucurbitacins, especially Cucurbitacin I, have shown significant inhibitory efficacy against STAT3 signalling (45,46). This inhibition increases the anticancer effects of cucurbitacins by lowering the expression of genes involved in tumour growth, survival, and metastasis (45, 47).

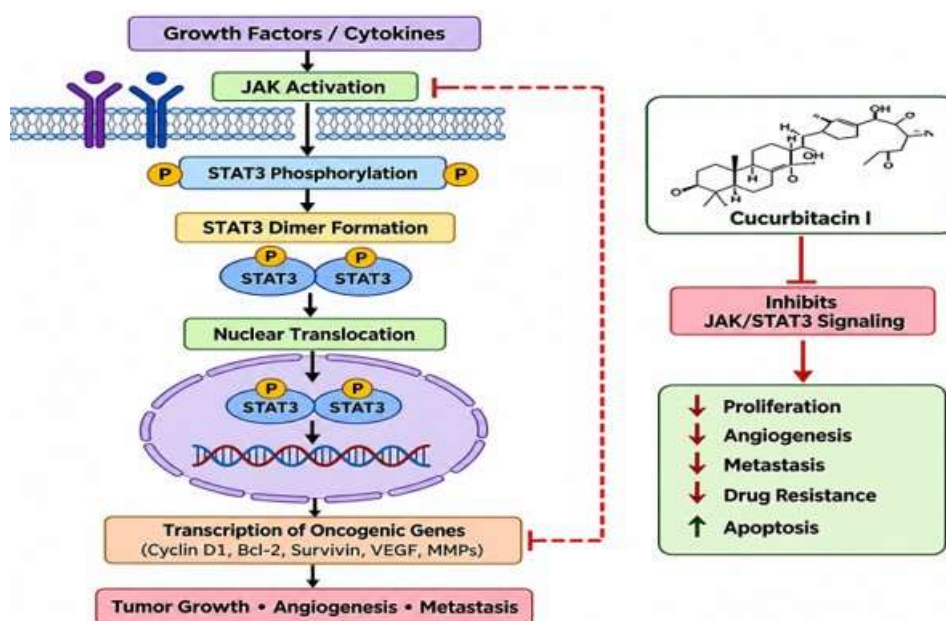


Figure 5: Inhibition of the JAK/STAT3 Signalling Pathway by Cucurbitacin I

5.4 PI3K/Akt/mTOR Pathway Suppression

One of the main intracellular systems that controls cell growth, proliferation, survival, metabolism, and protein synthesis is the phosphatidylinositol 3-kinase (PI3K)/Akt/mammalian target of rapamycin (mTOR) signalling pathway (48, 49). Many human malignancies have this system improperly active, which promotes tumour growth, metastasis, and treatment resistance

(50,51). Consequently, blocking the PI3K/Akt/mTOR pathway has become a crucial treatment approach for cancer (48,52). According to research, cucurbitacins block downstream signalling by blocking the phosphorylation and activation of PI3K, Akt, and mTOR (53,54). This results in an inhibition of tumour growth, an increase in apoptosis, and a decrease in the proliferation of cancer cells (53,55).

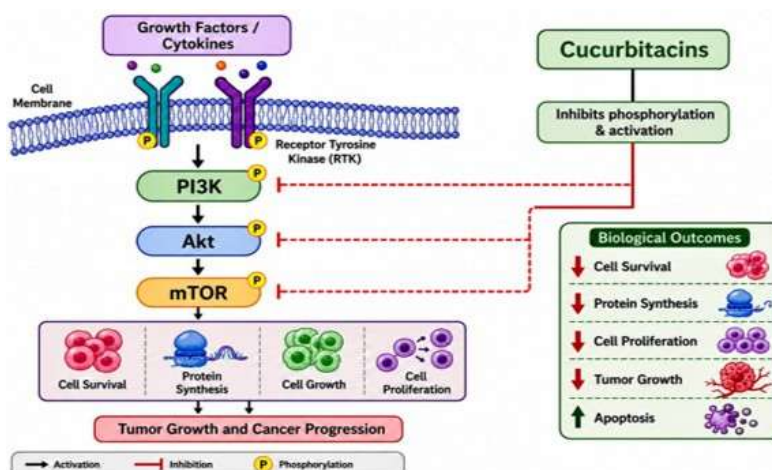


Figure 6: Suppression of the PI3K/Akt/mTOR signalling pathway by Cucurbitacins

5.5 NF- κ B Signalling Inhibition

Nuclear factor-kappa B (NF- κ B) is a pivotal transcription factor that regulates inflammation, cell proliferation, apoptosis, angiogenesis, immune responses, and tumour progression (56,57). Constitutive activation of NF- κ B has been reported in numerous human malignancies and is closely associated with chronic inflammation, enhanced tumour cell survival, metastatic progression, and resistance to chemotherapy (57–59). Under resting conditions, NF- κ B remains sequestered in the cytoplasm through its interaction with inhibitor of κ B (I κ B) proteins (56,59). Following stimulation by inflammatory cytokines, growth factors, or oxidative stress, I κ B undergoes phosphorylation by the I κ B kinase (IKK) complex, leading to its ubiquitination and proteasomal degradation, thereby permitting NF- κ B nuclear translocation (57,58). Activated NF- κ B subsequently induces the transcription of genes involved in inflammation, cell survival, angiogenesis, invasion, metastasis, and anti-apoptotic signalling (59,60). Experimental evidence indicates that cucurbitacins suppress NF- κ B signalling by inhibiting IKK activation, preventing I κ B degradation, blocking nuclear translocation of the p65 subunit, and reducing the expression of NF- κ B-regulated genes (61–63). Consequently, cucurbitacins decrease the production of pro-inflammatory cytokines and anti-apoptotic proteins while enhancing apoptosis and suppressing tumour progression, highlighting NF- κ B inhibition as an important mechanism underlying their anticancer activity (60–63).

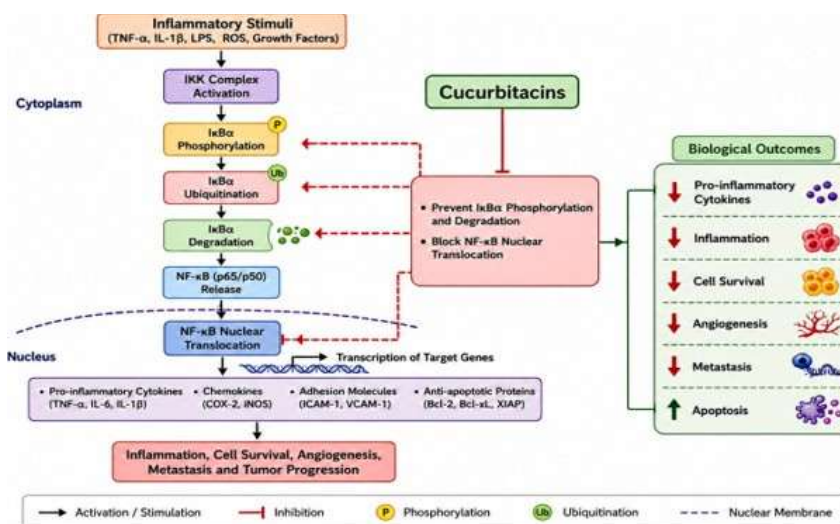


Figure 7. Inhibition of the NF- κ B Signalling Pathway by Cucurbitacins

5.6 Reactive Oxygen Species (ROS) Generation

Reactive oxygen species (ROS) are highly reactive oxygen-containing molecules that function as important mediators of cellular signalling, proliferation, differentiation, and survival under physiological conditions. However, excessive ROS production disturbs intracellular redox homeostasis, resulting in oxidative stress that damages DNA, proteins, lipids, and other cellular components (64,65). Cancer cells generally maintain higher basal ROS levels than normal cells because of their increased metabolic activity and mitochondrial dysfunction, making them more susceptible to further oxidative stress induced by anticancer agents (65,66). Several studies

have demonstrated that cucurbitacins markedly increase intracellular ROS accumulation, leading to mitochondrial membrane depolarization, oxidative DNA damage, endoplasmic reticulum (ER) stress, and activation of apoptotic signalling pathways (24,67,68). ROS generated by cucurbitacins also promotes cytochrome *c* release, caspase activation, and mitochondrial dysfunction, thereby enhancing intrinsic apoptosis in cancer cells (67,68). Furthermore, excessive ROS production contributes to inhibition of tumour cell proliferation and sensitizes cancer cells to chemotherapy by disrupting cellular antioxidant defence mechanisms (24,69). Collectively, ROS-mediated oxidative stress represents one of the major mechanisms through which cucurbitacins exert their anticancer effects (24,67–69).

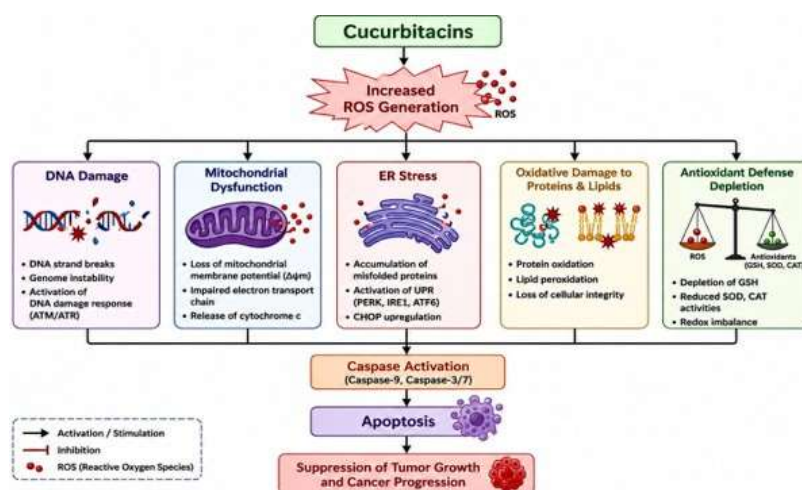


Figure 8: ROS-Mediated Anticancer Mechanism of Cucurbitacins

5.7 Autophagy Modulation

Autophagy is a highly conserved lysosome-dependent intracellular degradation process that maintains cellular homeostasis by eliminating damaged organelles, protein aggregates, and dysfunctional macromolecules (70,71). Under physiological conditions, autophagy serves as a cytoprotective mechanism that enables cells to survive nutrient deprivation, oxidative stress, and other adverse conditions (70,71). In cancer, however, autophagy exhibits a dual role by either promoting tumour cell survival or facilitating programmed cell death depending on the tumour type, genetic background, and stage of disease (72,73). Increasing evidence indicates that cucurbitacins regulate autophagy in a dose-, time-, and cell type-dependent manner through modulation of multiple signalling pathways, including PI3K/Akt/mTOR and AMPK signalling (69,74,75). Experimental studies have demonstrated that cucurbitacins alter the expression of autophagy-related proteins such as microtubule-associated protein light chain 3 (LC3-II), Beclin-1, p62/SQSTM1, and ATG proteins, thereby influencing autophagic flux and cellular fate (69,74–77). Furthermore, autophagy induced by cucurbitacins frequently interacts with apoptotic signalling, and the balance between these two processes determines the overall therapeutic response of cancer cells (75–77). Therefore, modulation of autophagy represents an important mechanism contributing to the anticancer activity of cucurbitacins and may provide opportunities for combination therapies targeting both apoptosis and autophagy (69,74–77).

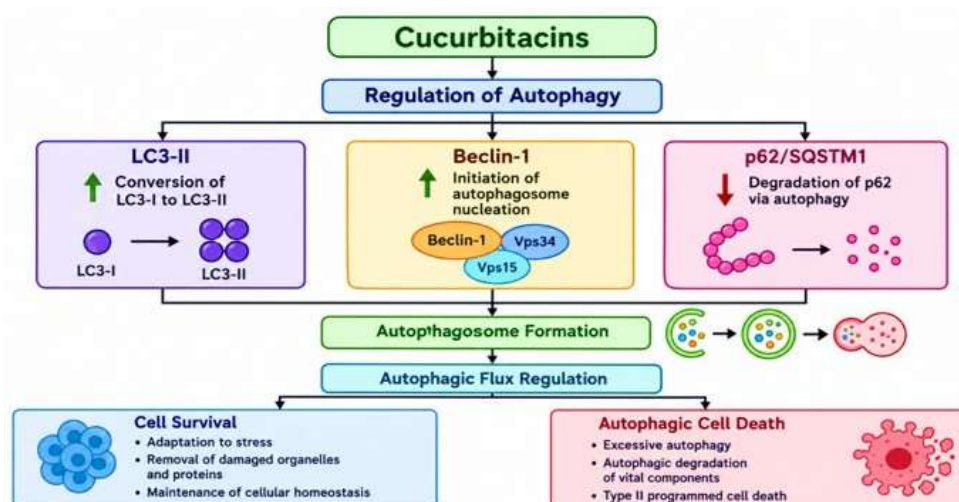


Figure 9: Modulation of Autophagy by Cucurbitacin

5.8 Cytoskeletal Disruption

Maintaining cell shape, intracellular transport, cell division, migration, and invasion all depend on the cytoskeleton, a highly dynamic intracellular network made up of actin filaments, microtubules, and intermediate filaments (78,79). Remodelling of the cytoskeleton is a hallmark of malignant transformation and plays a critical role in tumour progression and metastasis (78,80). Among naturally occurring triterpenoids, cucurbitacins are unique because they exert profound effects on the actin cytoskeleton, resulting in rapid morphological alterations and inhibition of cancer cell motility (8,81). Experimental studies have demonstrated that cucurbitacins induce depolymerization, aggregation, and abnormal reorganization of filamentous actin (F-actin), leading to loss of cell polarity, disruption of focal adhesions, and impairment of migratory capacity (8,81,82). In addition to actin remodelling, certain cucurbitacin analogues have been shown to interfere with microtubule dynamics and mitotic spindle organization, thereby contributing to G2/M cell-cycle arrest and inhibition of tumour cell proliferation (82,83). Recent mechanistic studies further suggest that cucurbitacins alter actin dynamics indirectly through interactions with actin-regulatory proteins, including cofilin, rather than by directly binding polymerized actin filaments (81,84). Collectively, disruption of cytoskeletal organization suppresses cancer cell migration, invasion, and metastatic dissemination while promoting apoptotic cell death, representing an important component of the multifaceted anticancer activity of cucurbitacins (8,81–84).

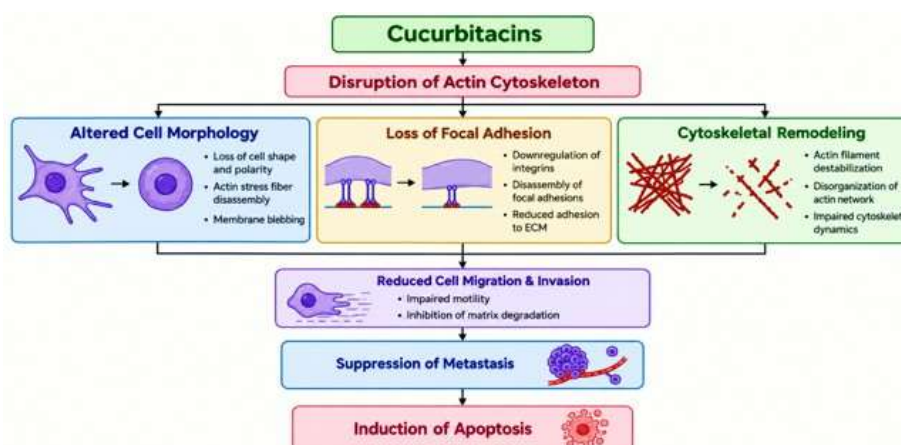


Figure 10: Cucurbitacin-Induced Cytoskeletal Disruption and Inhibition of Cancer Cell Migration

6. Hallmarks of Cancer Targeted by Cucurbitacins

Tumour start, development, metastasis, and treatment resistance are all made possible by a set of biological abilities known as the hallmarks of cancer (12,85). Sustained proliferative signalling, growth suppressor evasion, resistance to apoptosis, replicative immortality, angiogenesis stimulation, invasion and metastasis activation, metabolic reprogramming, immunological evasion, and genomic instability are some of these characteristics (85,86). Unlike conventional chemotherapeutic agents that often target a single molecular pathway, cucurbitacins exert broad-spectrum anticancer activity by simultaneously modulating multiple hallmarks of cancer through regulation of interconnected signalling networks (17,87). Experimental evidence has demonstrated that cucurbitacins suppress tumour cell proliferation, induce apoptosis, inhibit angiogenesis, interfere with epithelial–mesenchymal transition (EMT), reduce metastatic potential, modulate oxidative stress, and alter the tumour microenvironment (39,87–89). Moreover, inhibition of key oncogenic pathways, including JAK/STAT3, PI3K/Akt/mTOR, and NF- κ B, enables cucurbitacins to overcome several mechanisms responsible for tumour progression and therapeutic resistance (39,88–90). Owing to their multitarget mode of action, cucurbitacins have emerged as promising natural compounds for the development of novel anticancer therapies capable of targeting multiple cancer hallmarks simultaneously (17,39,87–89,91).

6.1 Inhibition of Angiogenesis

Because it provides oxygen and nutrients to rapidly reproducing cancer cells, angiogenesis—the physiological process by which new blood vessels are formed from pre-existing vasculature—is crucial for tumour progression beyond 1-2 mm³ (92,93). Vascular endothelial growth factor (VEGF), hypoxia-inducible factor-1 α (HIF-1 α), fibroblast growth factors (FGFs), platelet-derived growth factor (PDGF), and other pro-inflammatory cytokines are the main drivers of tumour angiogenesis (93,94). Aggressive tumour development, metastatic spread, and a poor clinical prognosis are often linked to overexpression of these angiogenic mediators (94,95). Numerous experimental studies have demonstrated that cucurbitacins inhibit angiogenesis by suppressing VEGF expression, reducing HIF-1 α accumulation, inhibiting endothelial cell proliferation and migration, and interfering with STAT3-mediated angiogenic signalling (30,91,96,97). In addition, cucurbitacins decrease micro vessel density within tumours and impair the formation of capillary-like structures, thereby limiting tumour vascularization and nutrient supply (30,91,97). Through simultaneous inhibition of angiogenic signalling pathways and endothelial cell function, cucurbitacins effectively suppress tumour progression and metastatic potential, highlighting their promise as antiangiogenic agents in cancer therapy (30,91,96,97).

Sr. No.	Target	Effect of Cucurbitacins	Biological Outcome	References
1.	VEGF	Down Regulation	Reduced angiogenesis	(30,96)
2.	HIF- 1 α	Inhibition	Reduced VEGF expression	(30,96)
3.	STAT 3	Suppression	Decreased angiogenic signalling	(30,97)
4.	Endothelial cell proliferation	Inhibited	Reduced vessel formation	(91,97)
5.	Tube formation	Suppressed	Impaired neovascularization	(30,91)
6.	Tumour micro vessel density	Decreased	Reduced tumour growth	(96,97)

Table 3: Anti-angiogenic Mechanisms of Cucurbitacins

6.2 Anti-Angiogenic Effects

Angiogenesis, the physiological process by which new blood vessels are formed from pre-existing vasculature, is crucial for tumour development beyond 1-2 mm³ because it provides oxygen and nutrients to rapidly proliferating cancer cells (92,93). Vascular endothelial growth factor (VEGF), hypoxia-inducible factor-1 α (HIF-1 α), fibroblast growth factors (FGFs), platelet-

derived growth factor (PDGF), and other pro-inflammatory cytokines are the main drivers of tumour angiogenesis (93,94). Furthermore, by preventing endothelial cell migration, proliferation, and capillary tube development, cucurbitacins obstruct the production of new blood vessels required for the progression of cancer (99,101). produced from pre-existing vasculature—is essential for the growth of tumours larger than 1–2 mm³ (92,93). Tumour angiogenesis is primarily driven by vascular endothelial growth factor (VEGF), hypoxia-inducible factor-1 α (HIF-1 α), fibroblast growth factors (FGFs), platelet-derived growth factor (PDGF), and other pro-inflammatory cytokines (93,94). Furthermore, cucurbitacins impede the establishment of new blood vessels necessary for cancer growth by inhibiting endothelial cell migration, proliferation, and capillary tube formation (99,101). Cucurbitacins have also been shown in several studies to interfere with angiogenesis-associated signalling molecules, such as matrix metalloproteinases (MMPs) and focal adhesion kinase (FAK), which results in decreased neovascularization and poor tumour vascularization (100,102). Through inhibition of tumour angiogenesis, Cucurbitacins restrict tumour growth, reduce metastatic potential, and enhance their overall anticancer efficacy (99–102).

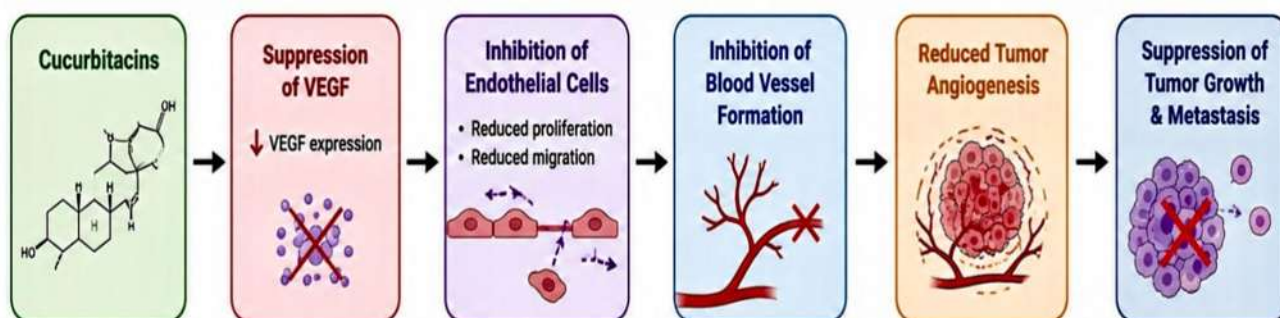


Figure 11: Anti-Angiogenic Mechanism of Cucurbitacins

6.3 Anti-Metastatic Effects

Cancer cells separate from the original tumour, infiltrate adjacent tissues, intravasate into the circulation, survive in the bloodstream, extravasate into distant organs, and form additional tumours during the intricate, multi-step process of metastasis (103,104). One of the main obstacles to cancer treatment is metastatic spread, which accounts for around 90% of cancer-related fatalities (103,105). By focusing on many signalling pathways involved in tumour invasion and migration, experimental research has shown that cucurbitacins have strong anti-metastatic effect (100,106). By controlling EMT-associated transcription factors and re-establishing epithelial traits in cancer cells, cucurbitacins inhibit the epithelium–mesenchymal transition (EMT). Additionally, these substances reduce extracellular matrix breakdown and tumour invasion by inhibiting the production and activity of matrix metalloproteinases (MMP-2 and MMP-9) (100,107). Additionally, cucurbitacins interfere with focal adhesion dynamics and actin cytoskeletal architecture, which impairs cell migration and lowers the potential for metastasis (106,107). Collectively, these mechanisms suppress cancer cell invasion, migration, and distant metastasis, highlighting Cucurbitacins as promising anti-metastatic agents for cancer therapy (100,105–107).

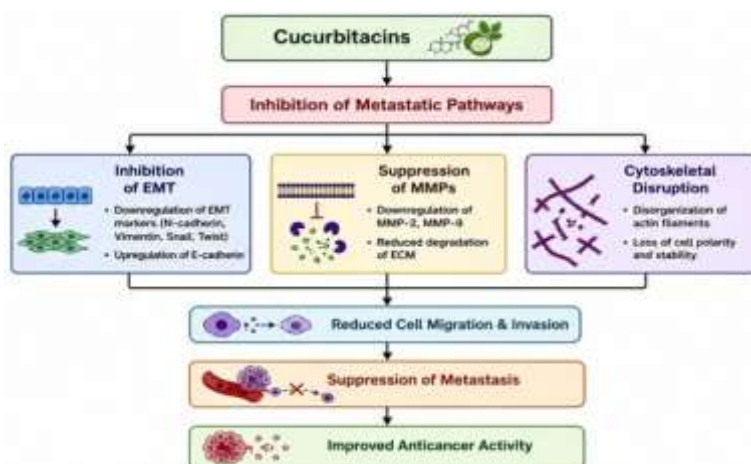


Figure 12: Molecular Mechanisms Underlying the Anti-Metastatic Effects of Cucurbitacins

6.4 Targeting of Cancer Stem Cells

Tumour recurrence, metastasis, and treatment resistance are all significantly influenced by cancer stem cells (CSCs), a tiny subset of tumour cells distinguished by their ability to self-renew, differentiate, and initiate tumours (108,109). A growing body of research indicates that CSCs are inherently resistant to traditional chemotherapy and radiation, which may contribute to disease recurrence after treatment (109,110). According to recent research, Cucurbitacins efficiently target CSCs by inhibiting stemness-associated signalling pathways such as Notch, JAK/STAT3, and Wnt/ β -catenin (106,111). Additionally, stem cell markers such CD44, ALDH1, Nanog, Oct4, and Sox2 have been shown to be downregulated by cucurbitacins, which reduces their ability to self-renew and initiate tumours (106,110). Moreover, Cucurbitacins' suppression of CSC-associated signalling increases chemosensitivity and lowers the risk of cancer recurrence and metastatic spread (99,106). Cucurbitacins are attractive therapeutic possibilities for overcoming drug resistance and enhancing long-term clinical outcomes in cancer treatment because they target CSC maintenance and survival (99,106,110,111).

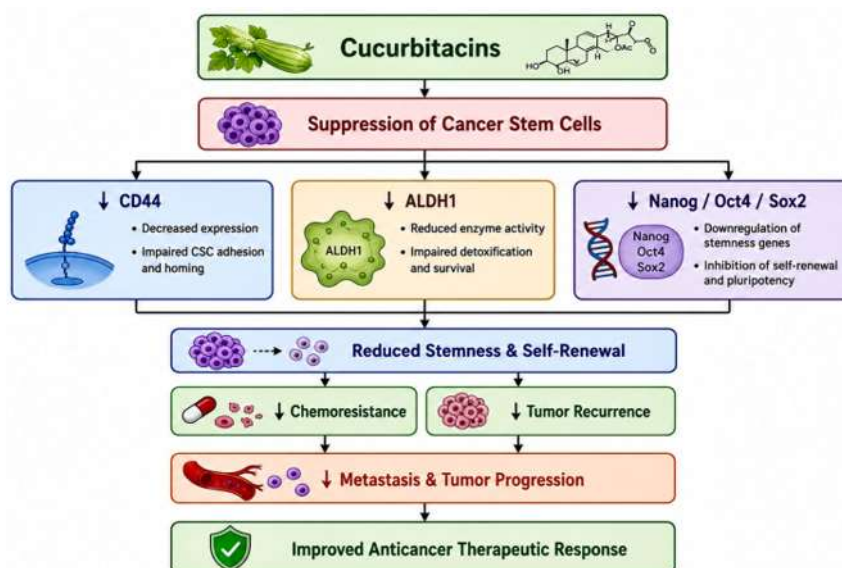


Figure 13: Modulation of Cancer Stem Cell Signalling by Cucurbitacins

7. Cucurbitacins in Specific Cancer Types

Cucurbitacins' primary molecular targets and treatment efficiency fluctuate depending on the biological traits and genetic landscape of various cancer types, despite the fact that the basic molecular pathways underlying their anticancer action are mostly consistent (11,112). Numerous preclinical studies have shown that different cucurbitacin analogues have strong antitumour activity against a wide range of solid and haematological malignancies by regulating several signalling pathways related to cell proliferation, apoptosis, angiogenesis, metastasis, oxidative stress, autophagy, and therapeutic resistance (1,11,112). Among the different analogues, Cucurbitacins B, D, E, I, and Q have shown particularly promising anticancer activity against breast cancer, lung cancer, colorectal cancer, pancreatic cancer, leukaemia, glioblastoma, and several other malignancies through inhibition of oncogenic pathways such as JAK/STAT3, PI3K/Akt/mTOR, NF- κ B, MAPK, VEGFR, and Wnt/ β -catenin signalling (1,88,112). In addition to their direct cytotoxic effects, increasing evidence indicates that cucurbitacins can enhance the efficacy of conventional chemotherapeutic agents, suppress cancer stem cell populations, inhibit tumour invasion and metastasis, and partially overcome multidrug resistance, highlighting their potential as

multifunctional adjuncts in cancer therapy (1,11,88). Consequently, substantial research efforts have focused on evaluating the therapeutic potential of cucurbitacins across different cancer types, providing a strong rationale for the development of novel targeted treatment strategies and future clinical translation (1,11,88,112).

Sr. No.	Cancer Type	Predominant Cucurbitacin Analogue(s)	Major Molecular Targets	References
1	Breast Cancer	B, D, I	STAT3, Akt, NF- κ B	#####
2	Lung Cancer	E, I	EGFR, JAK/STAT3, PI3K/Akt	#####
3	Colorectal cancer	D, E	Wnt/ β -catenin, PI3K/Akt	(7,117–119)
4	Pancreatic cancer	B	KRAS, JAK/STAT3	(113,120–122)
5	Leukaemia	I	JAK/STAT3, Caspases	(10,123–127)

Table 4: Major Cucurbitacin Analogues Investigated in Different Cancer Types and Their Primary Molecular Targets

7.1 Breast Cancer

Despite significant advancements in early diagnosis and focused therapy, breast cancer remains one of the top causes of cancer-related death and is the most often diagnosed disease among women globally (17,113). Treatment resistance, tumour recurrence, and metastatic progression remain significant therapeutic problems, especially for patients with advanced-stage illness and triple-negative breast cancer (TNBC) (11,113). By inhibiting several oncogenic signalling pathways involved in cell survival, proliferation, invasion, and metastasis, Cucurbitacin B, Cucurbitacin D, and Cucurbitacin I have shown notable antiproliferative activity against breast cancer cells among the different cucurbitacin analogues (11,17,114,115). According to experimental research, cucurbitacins suppress constitutive STAT3, Akt, and NF- κ B signalling, which lowers the production of anti-apoptotic proteins, inhibits the growth of cancer cells, and triggers caspase-dependent apoptosis (11,114,115). Additionally, cucurbitacins reduce the potential for metastasis by increasing intracellular reactive oxygen species (ROS), inducing mitochondrial dysfunction, suppressing the epithelial–mesenchymal transition (EMT), and impairing the invasive and migratory capabilities of breast cancer cells (17,114,115). Crucially, preclinical studies have shown that Cucurbitacin D reverses STAT3- and NF- κ B-mediated chemoresistance in drug-resistant breast cancer cells, increasing the cytotoxic efficiency of doxorubicin, indicating its possible use as an adjuvant treatment agent (115). Due to their capacity to concurrently control apoptosis, oxidative stress, oncogenic signalling pathways, and treatment resistance, these results collectively suggest that cucurbitacins constitute prospective multifunctional agents for breast cancer therapy (11,17,113–115).

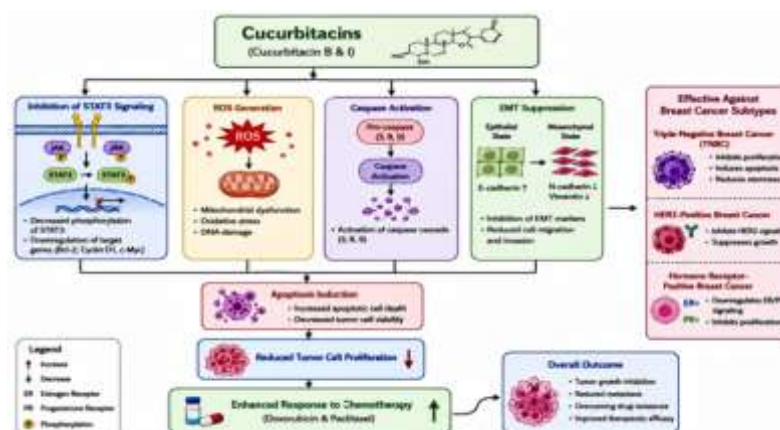


Figure 14: Molecular mechanisms underlying the anticancer effects of Cucurbitacins in breast cancer

7.2 Lung Cancer

Non-small cell lung cancer (NSCLC) accounts for approximately 85% of all lung cancer cases and remains one of the leading causes of cancer-related mortality worldwide despite remarkable advances in targeted therapy and immunotherapy (113,116). The development of therapeutic resistance, persistent activation of oncogenic signalling pathways, and metastatic progression continue to limit long-term clinical outcomes in patients with advanced NSCLC ((113,117). Among the various cucurbitacin analogues, Cucurbitacin B, Cucurbitacin E, and Cucurbitacin I have demonstrated potent antiproliferative activity against NSCLC through modulation of multiple signalling pathways involved in cell growth, apoptosis, and metastasis (7,117–119). Experimental studies have shown that Cucurbitacin E suppresses EGFR/MAPK signalling, inhibits cell-cycle progression, and activates caspase-dependent apoptosis in A549 lung cancer cells (118). Similarly, Cucurbitacin B inhibits STAT3 and PI3K/mTOR signalling while activating AMPK-dependent pathways, resulting in reduced proliferation and enhanced apoptotic cell death in both EGFR-wild-type and EGFR-mutant NSCLC cells (7,117). Furthermore, Cucurbitacin I effectively suppresses STAT3 activation in CD133-positive lung cancer stem cells, thereby reducing tumorigenicity, inhibiting self-renewal, and enhancing the sensitivity of NSCLC cells to chemotherapy and radiotherapy (119). Collectively, these findings indicate that cucurbitacins exert multifaceted anticancer effects against lung cancer by targeting EGFR, STAT3, PI3K/Akt/mTOR, and cancer stem cell-associated signalling pathways, highlighting their potential as promising adjunctive therapeutic agents for NSCLC (7,117–119).

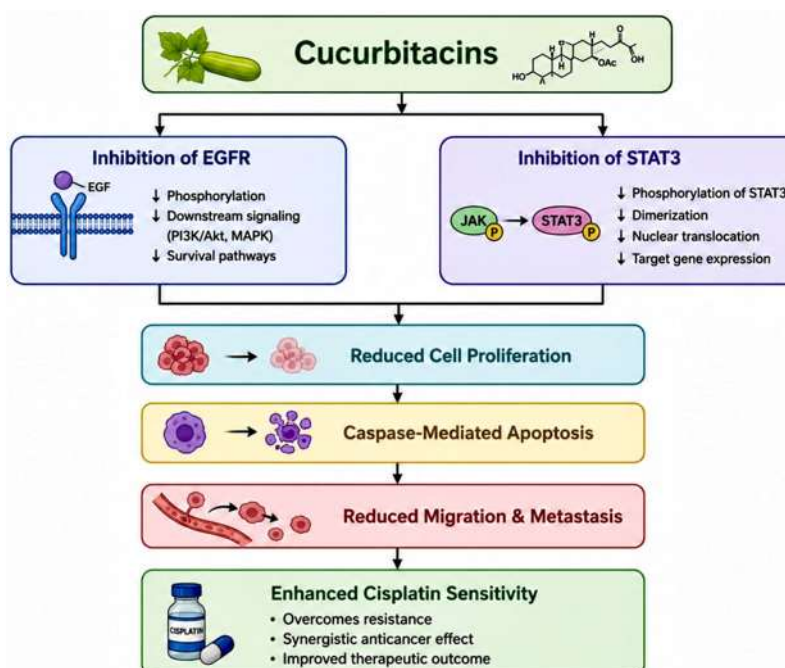


Figure 15: Molecular Mechanisms of Cucurbitacins in Lung Cancer (NSCLC)

7.3 Colorectal Cancer

Colorectal cancer (CRC) is one of the most common gastrointestinal malignancies and remains a leading cause of cancer-related mortality worldwide despite considerable advances in screening, surgical management, and systemic therapies (113,120). Aberrant activation of the Wnt/ β -catenin, PI3K/Akt, and EGFR signalling pathways plays a pivotal role in colorectal tumour initiation, progression, and resistance to chemotherapy (120,121). Recent experimental studies have demonstrated that several cucurbitacin analogues, particularly Cucurbitacin D and Cucurbitacin E, exert potent antiproliferative activity against colorectal cancer cells by simultaneously targeting multiple oncogenic signalling pathways (121,122,128). Cucurbitacins suppress Wnt/ β -catenin signalling, thereby reducing the nuclear accumulation of β -catenin and inhibiting the transcription of genes associated with cell proliferation, invasion, and tumour progression (121,122). In addition, these compounds induce cell-cycle arrest, activate mitochondrial-mediated apoptosis through regulation of Bax, Bcl-2, and caspase-3, and suppress the proliferative capacity of colorectal cancer cells (122,128). Furthermore, inhibition of PI3K/Akt signalling and modulation of oxidative stress contribute to the antitumor activity of cucurbitacins, ultimately reducing tumour growth and enhancing apoptotic cell death (128,129). Collectively, these findings suggest that cucurbitacins represent promising multitarget therapeutic agents for colorectal cancer owing to their ability to inhibit oncogenic signalling, induce apoptosis, and suppress tumour progression (121,122,128,129).

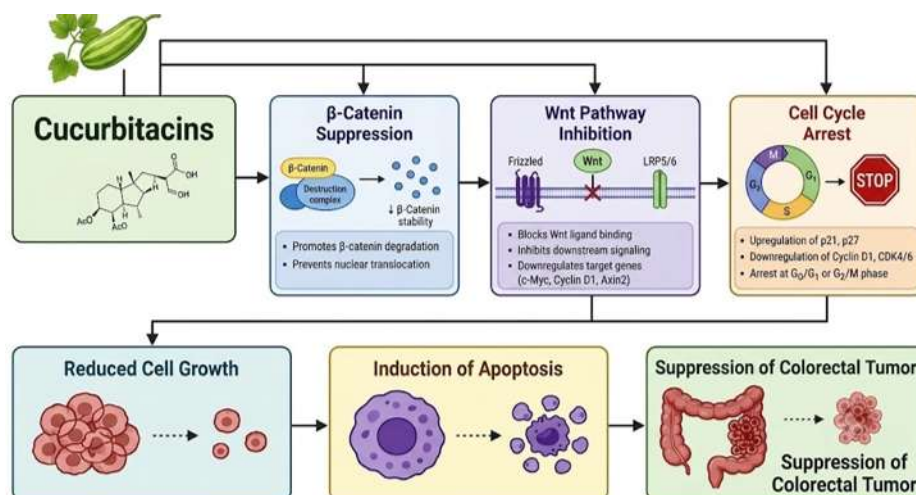


Figure 16: Molecular Mechanisms of Cucurbitacins in Colorectal Cancer

7.4 Pancreatic Cancer

Pancreatic ductal adenocarcinoma (PDAC) is one of the most aggressive and lethal malignancies, with a five-year survival rate of less than 10% owing to its late diagnosis, rapid metastatic progression, and poor response to conventional chemotherapy (123,130). Constitutive activation of oncogenic pathways, particularly KRAS, JAK/STAT3, PI3K/Akt, and NF- κ B, plays a central role in pancreatic tumour initiation, progression, chemoresistance, and maintenance of cancer stem cell populations (130,131). Among the different cucurbitacin analogues, Cucurbitacin B has demonstrated significant antitumor activity against pancreatic cancer by suppressing multiple signalling pathways associated with tumour growth, invasion, and therapeutic resistance (131–134). Experimental studies have shown that Cucurbitacin B inhibits JAK/STAT3 signalling, resulting in reduced STAT3 phosphorylation, downregulation of anti-apoptotic proteins, suppression of tumour cell proliferation, and induction of caspase-dependent apoptosis (131,132). In addition, Cucurbitacin B interferes with KRAS-associated downstream signalling and inhibits PI3K/Akt activation, thereby reducing cell survival, impairing metastatic potential, and enhancing mitochondrial-mediated apoptotic cell death (132,133). Recent evidence further suggests that cucurbitacins suppress the expression of cancer stem cell-associated markers, including CD44, CD133, and ALDH1, thereby reducing self-renewal capacity and increasing the sensitivity of pancreatic cancer cells to conventional chemotherapeutic agents such as gemcitabine (133,134). Collectively, these findings indicate that cucurbitacins represent promising multitarget therapeutic

agents for pancreatic cancer by simultaneously inhibiting oncogenic signalling pathways, inducing apoptosis, suppressing cancer stem cell characteristics, and overcoming chemoresistance (131–134).

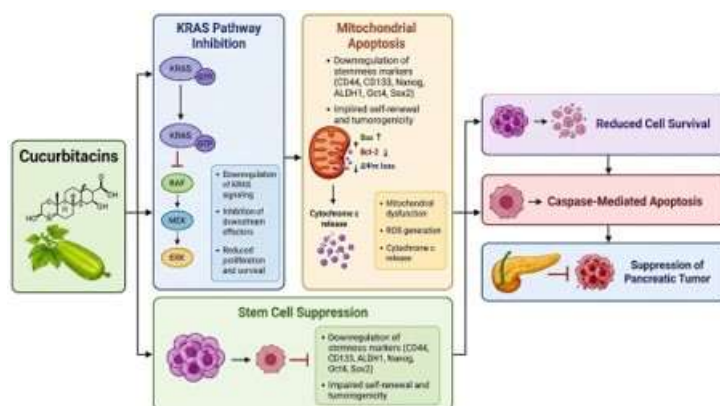


Figure 17: Molecular Mechanisms of Cucurbitacins in Pancreatic Cancer

7.5 Leukemia

Leukaemia comprises a heterogeneous group of haematological malignancies characterized by the uncontrolled proliferation and accumulation of abnormal hematopoietic cells in the bone marrow and peripheral blood. Despite substantial advances in chemotherapy, targeted therapy, and hematopoietic stem cell transplantation, treatment failure and disease relapse remain major clinical challenges, particularly in acute and drug-resistant Leukaemia (124,135). Persistent activation of oncogenic signalling pathways, especially the JAK/STAT3, PI3K/Akt, and NF- κ B pathways, contributes to leukemic cell survival, uncontrolled proliferation, resistance to apoptosis, and therapeutic failure (10,124). Among the various cucurbitacin analogues, Cucurbitacin I has demonstrated remarkable antileukemic activity by targeting multiple signalling pathways involved in leukemic cell growth and survival (10,125–127). Experimental studies have shown that Cucurbitacin I potently inhibits STAT3 phosphorylation and its nuclear translocation, resulting in suppression of STAT3-dependent transcription, downregulation of anti-apoptotic proteins, and induction of caspase-mediated apoptosis in Leukaemia cells (10,125). Furthermore, Cucurbitacin I induces cell-cycle arrest, disrupts mitochondrial membrane potential, and activates intrinsic apoptotic pathways through modulation of Bax, Bcl-2, and caspase-3, thereby significantly reducing leukemic cell viability (125,126). Recent investigations also indicate that cucurbitacins may enhance the sensitivity of leukemia cells to conventional chemotherapeutic agents and represent promising candidates for combination therapy to overcome drug resistance (126,127). Collectively, these findings suggest that cucurbitacins, particularly Cucurbitacin I, possess considerable therapeutic potential for Leukaemia through simultaneous inhibition of oncogenic signalling pathways, induction of apoptosis, and suppression of leukemic cell proliferation (10,125–127).

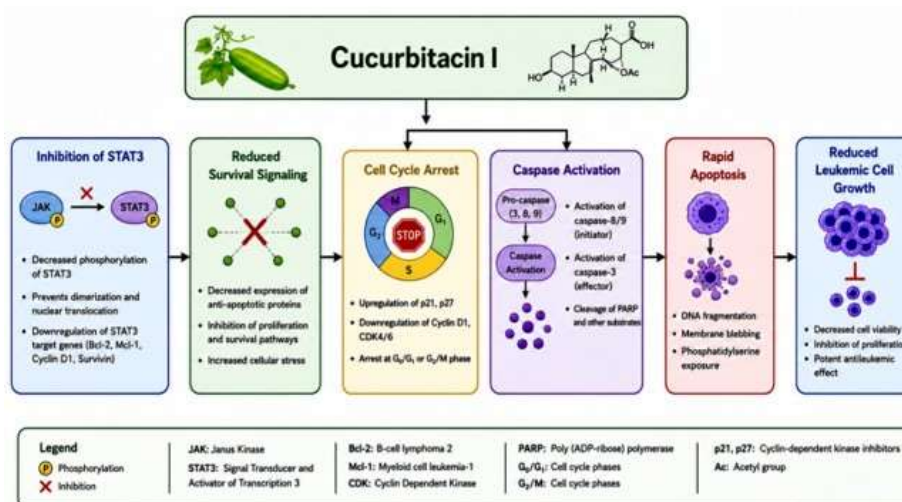


Figure 18: Molecular Mechanisms of Cucurbitacin I in Leukemia

8. Synergistic Combination Therapy

Combination therapy has emerged as an effective therapeutic strategy for cancer treatment by combining agents with complementary mechanisms of action to enhance antitumour efficacy, reduce systemic toxicity, and overcome drug resistance (136,137). Growing evidence indicates that cucurbitacins can sensitize cancer cells to conventional chemotherapeutic agents and targeted therapies through simultaneous modulation of multiple oncogenic signalling pathways involved in cell proliferation, apoptosis, metastasis, and therapeutic resistance (17,138). Experimental studies have demonstrated that cucurbitacins exhibit synergistic interactions with several anticancer drugs, including doxorubicin, cisplatin, paclitaxel, and gemcitabine, resulting in enhanced apoptosis, inhibition of survival signalling, increased oxidative stress, and improved therapeutic response in various malignancies (11,138,139). Furthermore, the ability of cucurbitacins to inhibit key signalling pathways such as JAK/STAT3, PI3K/Akt/mTOR, NF- κ B, and EGFR highlights their potential as adjuvant agents capable of improving treatment efficacy and overcoming acquired drug resistance (17,138,139). Collectively, these findings suggest that combination therapy involving cucurbitacins represents a promising approach for improving clinical outcomes and may facilitate the development of more effective multimodal anticancer treatment strategies (11,17,136–139).

8.1 Combination with Chemotherapy

Combination chemotherapy remains a cornerstone of cancer treatment; however, the development of multidrug resistance, dose-limiting toxicity, and nonspecific systemic effects often compromise therapeutic efficacy (140,141). Consequently, considerable efforts have been directed toward identifying naturally occurring compounds that can enhance the activity of conventional chemotherapeutic agents while minimizing adverse effects (131,141). Accumulating preclinical evidence indicates that cucurbitacins exhibit significant synergistic interactions with several clinically used chemotherapeutic drugs, including doxorubicin, cisplatin, paclitaxel, and gemcitabine (24,141,142). These synergistic effects are primarily mediated through suppression of JAK/STAT3, PI3K/Akt, and NF- κ B signalling pathways, increased intracellular reactive oxygen species (ROS) generation, activation of mitochondrial-dependent apoptosis, and reversal of drug resistance mechanisms (24,131,142,143). Among the various analogues, Cucurbitacin B has been reported to enhance the cytotoxic activity of doxorubicin and gemcitabine, whereas Cucurbitacin E and Cucurbitacin I have demonstrated synergistic effects with cisplatin and paclitaxel, respectively, resulting in greater inhibition of tumour cell proliferation and induction of apoptosis than either agent alone (24,142,143). Overall, these findings suggest that cucurbitacin-based combination therapy represents a promising strategy to improve chemotherapy responsiveness, reduce therapeutic resistance, and achieve enhanced anticancer efficacy while potentially lowering the required doses of conventional chemotherapeutic agents (24,131,142,143).

Examples

Sr. No	Chemotherapeutic Agent	Combined Cucurbitacin	Therapeutic Outcome	References
1.	Doxorubicin	Cucurbitacin B	Enhanced apoptosis	(24)
2.	Cisplatin	Cucurbitacin E	Reduced resistance	(142)
3.	Paclitaxel	Cucurbitacin I	Synergistic cytotoxicity	(143)
4.	Gemcitabine	Cucurbitacin B	Improved pancreatic cancer response	(131)

Table 4: Representative Combinations of Cucurbitacins with Conventional Chemotherapeutic Agents and Their Therapeutic Outcomes

8.2 Combination with Targeted Therapy

Targeted therapies have significantly improved cancer treatment by selectively inhibiting molecular pathways involved in tumour growth and progression. However, the emergence of acquired resistance and activation of compensatory signalling pathways frequently limit their long-term clinical efficacy (140,144). Recent preclinical studies suggest that cucurbitacins can enhance the therapeutic efficacy of targeted anticancer agents through simultaneous inhibition of multiple oncogenic signalling pathways, including JAK/STAT3, EGFR, PI3K/Akt/mTOR, and VEGF signalling (1,11,17). By targeting these

interconnected pathways, cucurbitacins suppress tumour cell proliferation, promote apoptosis, inhibit angiogenesis, and reduce metastatic potential, thereby improving the overall antitumour response (11,17). In addition, cucurbitacins have been reported to overcome resistance to targeted therapies by blocking compensatory survival pathways and downregulating the expression of proteins associated with drug resistance and tumour progression (1,11). Although most evidence has been obtained from in vitro and animal studies, these findings indicate that cucurbitacins may serve as promising adjuvant agents in combination with molecularly targeted therapies. Nevertheless, further preclinical investigations and well-designed clinical trials are required to validate their efficacy and safety before clinical application (1,145).

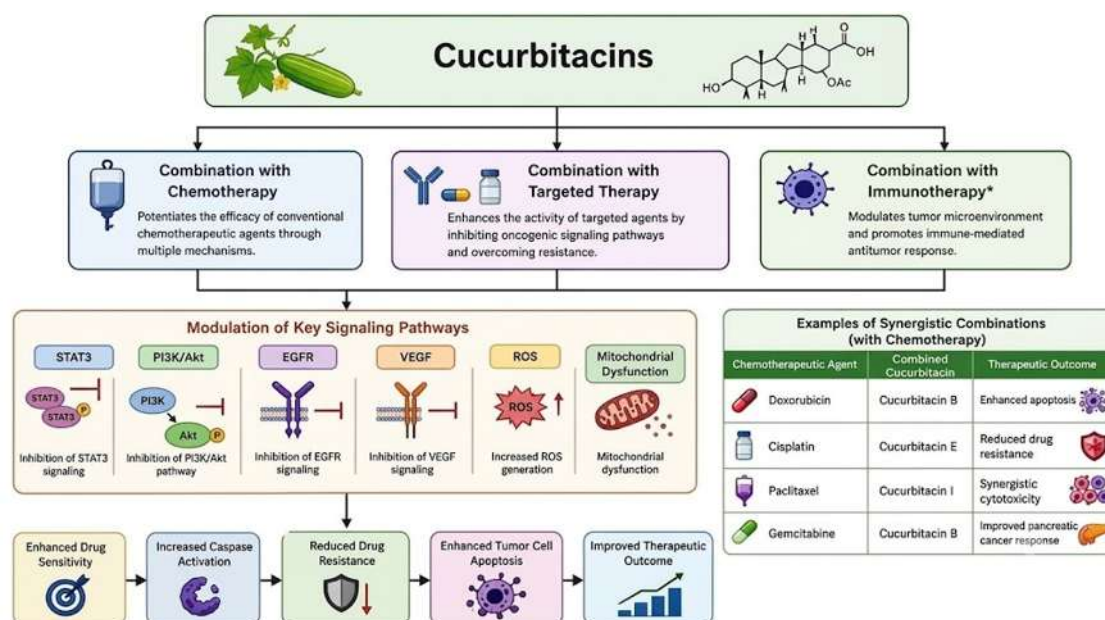


Figure 19 : Proposed mechanisms underlying the synergistic anticancer effects of cucurbitacins in combination with chemotherapy and targeted therapy

8.3 Combination with Immunotherapy

Cancer immunotherapy has revolutionized oncology by harnessing the host immune system to recognize and eliminate malignant cells. Immune checkpoint inhibitors targeting programmed cell death protein-1 (PD-1), programmed death-ligand 1 (PD-L1), and cytotoxic T-lymphocyte-associated protein-4 (CTLA-4) have demonstrated remarkable clinical success in several malignancies. (146). However, only a subset of patients achieves durable responses because tumour-associated immunosuppression, immune evasion, and the immunosuppressive tumour microenvironment often limit therapeutic efficacy (146,147). Recent preclinical studies suggest that cucurbitacins possess immunomodulatory properties that may complement current immunotherapeutic strategies by regulating inflammatory signalling pathways and modifying the tumour immune microenvironment (26,148). Inhibition of JAK/STAT3 and NF- κ B signalling by cucurbitacins reduces the production of immunosuppressive cytokines while promoting antitumour immune responses (26,47,148). Experimental evidence further indicates that cucurbitacins can enhance the activation and cytotoxic function of CD8⁺ T lymphocytes, reduce the accumulation of immunosuppressive cells within the tumour microenvironment, and improve immune-mediated tumour cell killing (26,47). These immunomodulatory effects may increase the responsiveness of tumours to immune checkpoint blockade and other immunotherapeutic approaches. Although the available findings are encouraging, current evidence is derived predominantly from preclinical investigations, and direct clinical evidence remains limited. Therefore, further mechanistic studies and well-designed clinical trials are required to establish the safety and therapeutic efficacy of cucurbitacins in combination with cancer immunotherapy (47,149).

9. Translational Challenges and Emerging Therapeutic Strategies

Despite the remarkable anticancer potential demonstrated by cucurbitacins in numerous preclinical studies, their successful clinical translation remains limited because of several pharmacokinetic, pharmaceutical, and toxicological challenges (150,151). The clinical development of cucurbitacin-based therapeutics is hindered by poor aqueous solubility, low oral bioavailability,

rapid metabolic degradation, short systemic half-life, nonspecific tissue distribution, and dose-dependent toxicity, all of which reduce their therapeutic efficacy and limit clinical applicability (151–153). To overcome these limitations, considerable research efforts have focused on the development of advanced drug delivery systems, structural modification of cucurbitacin molecules, targeted delivery approaches, and rational combination therapies (152–154). Nanotechnology-based formulations, including liposomes, polymeric nanoparticles, micelles,

and solid lipid nanoparticles, have demonstrated the ability to improve drug solubility, prolong circulation time, enhance tumour-specific accumulation through the enhanced permeability and retention (EPR) effect, and reduce systemic toxicity (153–155). Furthermore, structural optimization of cucurbitacins and the synthesis of novel derivatives have been investigated to improve pharmacokinetic properties while preserving or enhancing their biological activity (154,156). Combination strategies involving conventional chemotherapeutic agents, molecularly targeted therapies, and immunotherapeutic approaches have also shown promising synergistic effects in preclinical studies by enhancing apoptosis, overcoming drug resistance, and suppressing multiple oncogenic signalling pathways simultaneously (155–157). Although these emerging therapeutic strategies have significantly improved the translational potential of cucurbitacins, most investigations remain at the in vitro or animal-study stage. Therefore, comprehensive pharmacokinetic evaluation, toxicity assessment, standardized formulation development, and well-designed clinical trials are essential to establish the safety and efficacy of cucurbitacin-based therapies for cancer treatment (156–158).

9.1 Translational Challenges

Despite the remarkable anticancer efficacy of cucurbitacins demonstrated in numerous in vitro and in vivo studies, their successful translation into clinical oncology remains limited owing to several pharmaceutical, pharmacokinetic, and toxicological challenges (17,139). One of the major limitations is their poor aqueous solubility, which restricts dissolution in biological fluids and significantly reduces oral absorption and systemic bioavailability (139,151). Furthermore, cucurbitacins undergo rapid metabolic degradation and extensive first-pass metabolism, resulting in a short plasma half-life and limited drug exposure at tumour sites (151,152). Another important challenge is the dose-dependent toxicity associated with cucurbitacins. Although these compounds exhibit selective cytotoxicity toward cancer cells at appropriate concentrations, higher doses have been associated with hepatotoxicity, gastrointestinal toxicity, nephrotoxicity, and other systemic adverse effects in experimental models (152,153). The narrow therapeutic window and limited safety data remain significant barriers to their clinical application. In addition, the absence of standardized pharmaceutical formulations, poor tumour-specific accumulation, and variability in pharmacokinetic behavior further hinder the clinical development of cucurbitacin-based therapeutics (153,155). These limitations highlight the need for innovative drug delivery strategies, structural optimization, and rational combination therapies to improve therapeutic efficacy while minimizing systemic toxicity (150,155). Overall, overcoming these translational barriers is essential for advancing cucurbitacins from promising preclinical anticancer compounds to clinically viable therapeutic agents (17,139,150–153,155).

9.2 Clinical Studies and Translational Potential

Despite extensive evidence demonstrating the anticancer activity of cucurbitacins in cellular and animal models, their clinical translation remains at an early stage (11,17). To date, most investigations have focused on elucidating the molecular mechanisms underlying their antitumour activity, including inhibition of JAK/STAT3, PI3K/Akt/mTOR, NF- κ B, and Wnt/ β -catenin signalling pathways, induction of apoptosis, suppression of angiogenesis, and inhibition of metastasis (2,11). However, these encouraging preclinical findings have not yet been translated into substantial clinical evidence. The limited clinical advancement of cucurbitacins is primarily attributed to their unfavorable pharmacokinetic properties, dose-dependent toxicity, poor aqueous solubility, and the absence of standardized pharmaceutical formulations suitable for human administration (2,87,151). Furthermore, the lack of comprehensive pharmacokinetic studies, long-term safety evaluations, and well-designed randomized clinical trials has hindered regulatory approval and routine clinical application (87,151). Recent advances in formulation science, nanotechnology-based drug delivery systems, and rational combination therapies have substantially improved the translational potential of cucurbitacins by enhancing bioavailability, reducing systemic toxicity, and increasing tumour-selective drug accumulation (151,152). These developments provide a promising foundation for future clinical investigations and support the continued evaluation of cucurbitacins as potential adjuvant or targeted anticancer agents (152). Future clinical studies should focus on establishing optimal dosage regimens, evaluating long-term safety, identifying predictive biomarkers of therapeutic response, and assessing the efficacy of cucurbitacin-based formulations in

combination with existing chemotherapeutic, targeted, and immunotherapeutic agents (150,152). Such investigations will be essential for bridging the gap between preclinical discoveries and clinical implementation (153).

9.3 Emerging Therapeutic Strategies

Emerging therapeutic strategies are being actively investigated to enhance the clinical applicability of cucurbitacins and overcome the pharmacokinetic and toxicological limitations associated with these natural compounds (150,151). Recent advances have focused on improving drug delivery, increasing tumour specificity, enhancing bioavailability, and minimizing systemic toxicity through innovative pharmaceutical and molecular approaches (151,155). Nanotechnology-based drug delivery systems, including liposomes, polymeric nanoparticles, micelles, and solid lipid nanoparticles, have shown considerable promise in improving the physicochemical properties of cucurbitacins. These delivery platforms enhance aqueous solubility, prolong systemic circulation, facilitate controlled drug release, and promote preferential accumulation within tumour tissues via the enhanced permeability and retention (EPR) effect. Consequently, nanocarrier-based formulations have demonstrated improved therapeutic efficacy and reduced off-target toxicity in preclinical studies (151,154,155). Another promising strategy involves the structural modification of cucurbitacin molecules to generate semisynthetic derivatives with improved pharmacokinetic properties and enhanced anticancer activity (2,154). Rational chemical modifications may improve metabolic stability, reduce toxicity, and optimize interactions with molecular targets while preserving the characteristic biological activity of the parent compounds (154). In addition, advances in targeted drug delivery have enabled the development of ligand-conjugated nanoparticles capable of selectively delivering cucurbitacins to tumour tissues by recognizing cancer-specific receptors, thereby improving therapeutic selectivity and minimizing damage to normal tissues (156,157). Combination with conventional chemotherapy, molecularly targeted agents, and immunotherapeutic approaches also represents a promising strategy for enhancing antitumour efficacy while overcoming therapeutic resistance (152,157). Collectively, these emerging therapeutic strategies have substantially improved the translational potential of cucurbitacins and provide a strong foundation for future clinical development. Nevertheless, additional pharmacokinetic investigations, toxicity studies, and randomized clinical trials are required before these approaches can be successfully translated into routine oncology practice (152,158).

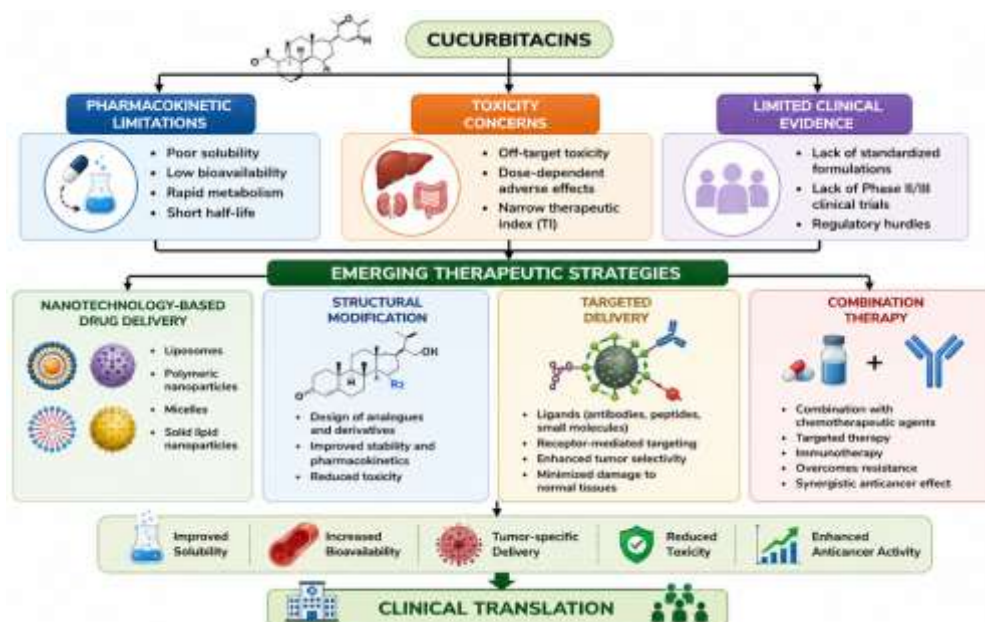


Figure 20 : Translational Challenges and Emerging Therapeutic Strategies for Clinical Development of Cucurbitacins

Summary of the molecular mechanisms, translational challenges, and emerging therapeutic strategies of cucurbitacins in cancer therapy. The figure illustrates the principal anticancer mechanisms of cucurbitacins, major barriers limiting clinical translation, and current strategies aimed at improving their therapeutic efficacy and clinical applicability (159,160).

10. Future Perspectives

The future of cucurbitacin research is highly promising because of the growing understanding of its molecular mechanisms and broad-spectrum anticancer activity. Numerous preclinical studies have demonstrated that cucurbitacins effectively inhibit tumour cell proliferation, induce apoptosis, suppress angiogenesis, and interfere with multiple oncogenic signalling pathways, including JAK/STAT3, PI3K/Akt/mTOR, NF- κ B, Wnt/ β -catenin, and MAPK. However, despite these encouraging findings, several challenges such as poor aqueous solubility, limited oral bioavailability, rapid metabolism, and systemic toxicity continue to hinder their successful clinical application. Future investigations should therefore focus on improving the pharmacokinetic profile, therapeutic selectivity, and safety of cucurbitacin-based therapies to facilitate their transition from laboratory research to clinical oncology.

One of the most important future directions is the development of advanced nanotechnology-based drug delivery systems. Nanocarriers, including liposomes, polymeric nanoparticles, solid lipid nanoparticles, dendrimers, and micelles, have demonstrated considerable potential to enhance drug solubility, prolong systemic circulation, improve tumour accumulation through the enhanced permeability and retention (EPR) effect, and minimize off-target toxicity. Surface modification with tumour-specific ligands such as antibodies, peptides, folic acid, and aptamers may further improve active targeting and increase therapeutic efficacy while reducing adverse effects.

Structural modification of naturally occurring cucurbitacins also represents an important strategy for future drug development. Medicinal chemistry approaches can generate semisynthetic derivatives with improved physicochemical properties, enhanced pharmacological potency, reduced toxicity, and greater selectivity toward malignant cells. The development of prodrug formulations and chemically modified analogs may further overcome limitations associated with poor stability and extensive first-pass metabolism, thereby improving bioavailability and therapeutic performance.

Future clinical research is expected to emphasize combination therapy, where cucurbitacins are administered alongside conventional chemotherapeutic agents, molecularly targeted drugs, immunotherapeutic agents, or radiotherapy. Such combination strategies may produce synergistic anticancer effects, overcome multidrug resistance, reduce the required therapeutic dose of individual agents, and decrease treatment-related toxicity. Additionally, the immunomodulatory properties of cucurbitacins may enhance antitumour immune responses and improve the effectiveness of immune checkpoint inhibitors.

The integration of biomarker-guided precision medicine is another promising direction. Identification of predictive biomarkers, including STAT3 activation status, PI3K/Akt pathway alterations, epithelial-to-mesenchymal transition (EMT) markers, inflammatory mediators, and reactive oxygen species (ROS)-associated signatures, may facilitate personalized patient selection and optimize therapeutic outcomes. Such biomarker-based approaches could enable clinicians to identify patients who are most likely to benefit from cucurbitacin-based treatment while minimizing unnecessary toxicity.

Artificial intelligence (AI), machine learning, and computational drug discovery are expected to play an increasingly significant role in accelerating cucurbitacin research. AI-assisted molecular docking, virtual screening, quantitative structure–activity relationship (QSAR) analysis, pharmacokinetic prediction, toxicity modeling, and de novo drug design can substantially reduce drug development time and

improve candidate selection. These computational approaches may facilitate the identification of novel cucurbitacin derivatives with enhanced efficacy and improved safety profiles. Future translational research should also focus on conducting large-scale, randomized, multicenter clinical trials to establish the long-term safety, pharmacokinetics, optimal dosing regimens, and clinical efficacy of cucurbitacin formulations across different cancer types. Standardization of extraction procedures, purification methods, formulation techniques, and quality-control protocols will be essential to ensure reproducibility and regulatory acceptance for clinical use.

Finally, successful clinical translation of cucurbitacins will require close collaboration among medicinal chemists, pharmacologists, oncologists, nanotechnology researchers, molecular biologists, bioinformaticians, and regulatory scientists. Such multidisciplinary efforts will accelerate the development of safe, targeted, and personalized cucurbitacin-based therapeutics. With continued advances in nanomedicine, biomarker-guided therapy, artificial intelligence, and precision oncology, cucurbitacins have significant potential to emerge as an important class of next-generation anticancer agents for improving patient outcomes.

11. Research Gaps

Although substantial progress has been made in elucidating the anticancer mechanisms of cucurbitacins, several critical research gaps continue to impede their successful clinical translation. Most available evidence is derived from in vitro experiments and animal models, whereas robust clinical investigations evaluating the safety, pharmacokinetics, and therapeutic efficacy of cucurbitacins in human patients remain extremely limited. Consequently, the clinical relevance of many promising preclinical findings has yet to be established.

Another major limitation is the lack of comprehensive pharmacokinetic and toxicological studies. Poor aqueous solubility, low oral bioavailability, rapid metabolism, and dose-dependent toxicity remain significant barriers to clinical application. Furthermore, standardized pharmaceutical formulations, optimized dosing regimens, and long-term safety evaluations have not been sufficiently investigated, limiting reproducibility and regulatory advancement.

The molecular mechanisms underlying cucurbitacin activity also require further investigation. Although multiple signalling pathways, including JAK/STAT3, PI3K/Akt/mTOR, NF- κ B, Wnt/ β -catenin, and MAPK, have been identified as therapeutic targets, the precise interactions between these pathways, tumour heterogeneity, and resistance mechanisms remain incompletely understood. In addition, limited information is available regarding predictive biomarkers that could identify patients most likely to benefit from cucurbitacin-based therapies.

Another important research gap is the limited exploration of structure–activity relationships (SAR) and rational structural modification of cucurbitacins. A better understanding of how specific structural features influence biological activity, selectivity, and toxicity would facilitate the development of safer and more effective derivatives with improved pharmacological properties.

Finally, although nanotechnology-based delivery systems and combination therapies have demonstrated encouraging preclinical outcomes, their scalability, manufacturing reproducibility, regulatory feasibility, and clinical applicability require further systematic evaluation. Addressing these research gaps through multidisciplinary collaboration will be essential for advancing cucurbitacins from promising experimental compounds to clinically effective anticancer therapeutics.

12. Conclusion

Cucurbitacins represent a unique class of naturally occurring tetracyclic triterpenoids that have attracted considerable attention because of their broad-spectrum anticancer activities and ability to modulate multiple oncogenic signalling pathways simultaneously. Extensive preclinical investigations have demonstrated that cucurbitacins inhibit tumour growth through diverse molecular mechanisms, including induction of apoptosis, cell cycle arrest, suppression of JAK/STAT3, PI3K/Akt/mTOR, and NF- κ B signalling, inhibition of angiogenesis and metastasis, modulation of autophagy, generation of reactive oxygen species, and disruption of cytoskeletal organization. These multifaceted biological activities highlight the potential of cucurbitacins as promising candidates for the treatment of a wide range of human malignancies.

In addition to their direct anticancer effects, cucurbitacins have demonstrated encouraging therapeutic potential when combined with conventional chemotherapy, molecularly targeted agents, and emerging immunotherapeutic strategies. Such combination approaches may enhance therapeutic efficacy, overcome drug resistance, and reduce the doses of standard anticancer drugs, thereby minimizing treatment-associated toxicity. Furthermore, advances in nanotechnology-based drug delivery systems and structural optimization have provided effective strategies for improving the pharmacokinetic properties, bioavailability, and tumour selectivity of cucurbitacins, thereby addressing several limitations that have hindered their clinical development.

Despite these promising findings, the translation of cucurbitacins from experimental models to routine clinical oncology remains challenging because of poor aqueous solubility, limited oral bioavailability, rapid metabolism, dose-dependent toxicity, and the scarcity of well-designed clinical studies. Addressing these limitations through optimized formulations, targeted delivery systems, comprehensive pharmacokinetic evaluation, and rigorous safety assessment will be essential for successful clinical translation.

Future research should focus on the development of standardized pharmaceutical formulations, identification of predictive biomarkers, elucidation of molecular targets using advanced omics technologies, and validation of therapeutic efficacy through large-scale clinical trials. The integration of artificial intelligence-assisted drug discovery, precision oncology, and multifunctional nanocarriers may further accelerate the development of cucurbitacin-based therapeutics and facilitate personalized cancer treatment strategies.

Overall, the available evidence indicates that cucurbitacins possess significant potential as multifunctional anticancer agents capable of targeting numerous hallmarks of cancer through diverse molecular mechanisms. Continued multidisciplinary research integrating medicinal chemistry, pharmaceutical sciences, molecular oncology, and clinical medicine will be crucial for translating these promising natural compounds into safe, effective, and clinically applicable anticancer therapy.

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AUTHOR CONTRIBUTIONS

- **Dr. Rao V. Javvaji** : Conceptualization, Investigation and review
- **Yash N. Nandan** : Review of Literature and Preparation of skeleton.
- **Suraj S. Ingale** : Data Curation and Preparation of Diagrams
- **Pratik K. Patil** : Study of Literature and Supervision
- **Pratiksha V. Sangale** : Writing and Review.
- **Dev T. Desale** : Skeleton Drafting & Tabulation of Data
- **Gargi V. Pawar** : Collection of literature, Editing
- **Varsha S. Pillai** : Supervision, Curation of Data & Editing

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