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Vorinostat in autophagic cell death: a critical insight into autophagy-mediated, -associated and -dependent cell death for cancer prevention

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Abstract

Histone deacetylases (HDACs) inhibit the acetylation of crucial autophagy genes, thereby deregulating autophagy and autophagic cell death (ACD) and facilitating cancer cell survival. Vorinostat, a broad-spectrum pan-HDAC inhibitor, inhibits the deacetylation of key autophagic markers and thus interferes with ACD. Vorinostat-regulated ACD can have an autophagy-mediated, -associated or -dependent mechanism depending on the involvement of apoptosis. Molecular insights reveal that hyperactivation of the PIK3C3/VPS34–BECN1 complex increased lysosomal disparity and enhanced mitophagy. These changes were followed by reduced mitochondrial biogenesis and by secondary signals that enabled superactivated, nonselective or bulk autophagy, leading to ACD. Although the evidence is limited, this review focuses on molecular insights into vorinostat-regulated ACD and describes critical concepts for clinical translation.

Teaser:

Vorinostat activates excessive autophagy through hyperactivation of the PIK3C3/VPS34–BECN1 complex, resulting in disrupted lysosomal homeostasis, altering the coordination of mitophagy-mitochondrial biogenesis and leading to superactivated, nonselective/bulk autophagy that could act as an alternative cell death mechanism in potential cancer therapy.

Keywords

autophagy; cancer; cell death; histone deacetylases; vorinostat

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Conflict of interests

The authors declare that they have no conflicts of interest.

Introduction

Disruption of the ‘epigenetic code’ as the result of legitimate epimutations and post-translational modifications drives the aberrant gene expression responsible for malignant transformation, cancer initiation and progression. Histone deacetylation, an antagonistic post-translational epigenetic co-repressor mechanism, promotes carcinogenesis and tumor progression through deacetylation of tumor suppressor genes and transcription factors.^{1, 2} By removing acetylated lysine residues, histone deacetylases (HDACs) favor a non-permissive chromatin conformation that aids transcriptional repression. HDACs mediate transcriptional repression that provokes the proliferation of cancer cells, cell cycle progression and inhibition of apoptosis and autophagy.² HDAC inhibitors (HDACis) reverse the HDAC-mediated transcriptional repression, and thus activate tumor-suppressor and cell-death-associated genes.³ In apoptosis-deficient cancer cells specifically, HDACis target autophagy-associated molecular markers to mediate autophagic cell death (ACD; formerly called type II programmed cell death on the basis of its morphological characteristics) and thus these inhibitors possess immense potential for cancer prevention.⁴

Autophagy, a catabolic cellular self-degradative mechanism, has been under investigation in recent times as an alternative cell death mechanism. Under normal conditions, autophagy serves as a cytoprotective quality control mechanism that eliminates redundant cargos and damaged cellular organelles. Under stress conditions, autophagy provides a continuous source of energy required for cell survival and energy homeostasis.⁵ However, the Nomenclature Committee on Cell Death has denoted ACD as a separate ‘cell death subroutine’, with a mechanism that is distinct from that of apoptosis.⁶ In certain circumstances, autophagy without the involvement of alternative cell death pathways can be lethal, as a result of failed or excessive cargo degradation.⁷ ACD has been established as a genuine cell death mechanism that has distinct autophagic morphologies, along with the participation of unique molecular components and excessive autophagic flux.⁸ In fact, ACD can have an autophagy-mediated, -associated or -dependent mechanism, depending on its direct role in the cell death process and its association with apoptosis.⁹ Several ACD-mediating drugs have been under clinical investigation, but the HDACis have garnered more interest in recent times because of the principal involvement of HDACs in cancer progression.

Among all of the broad spectrum and selective HDACis, vorinostat, a hydroxamate-class Zn²⁺-dependent pan-HDACi, has been studied extensively for its involvement in the induction of ACD.¹⁰ With limited reports available, however, this review focuses on how vorinostat mediates ACD, with particular emphasis on autophagy-mediated, -associated and -dependent cell death. In addition, we emphasize the molecular mechanisms that are associated with vorinostat-induced ACD, such as superactivated autophagy, mitophagy, disrupted lysosomal homeostasis, and the involvement of secondary signals. We try to focus especially on the clinical landscape for vorinostat in different cancer subtypes, looking at autophagy as the principal regulatory mechanism that determines the pharmacokinetics and pharmacodynamics of the drug.

HDACs, HDACis and autophagy: an intricate interplay

HDACs mediate the transcriptional regulation of several proteins and play a key role in cancer initiation and progression. Through dynamic post-translational modifications and non-permissive chromatin conformations, HDACs target key proteins and transcription factors that are associated with cell proliferation, cell differentiation, cell survival, cell cycle progression, apoptosis and autophagy.^{1, 2} The deacetylation of crucial molecular components that are associated with tumor suppressor genes and apoptotic genes promotes the survival of cancer cells. Conversely, HDACis oppose the survival and proliferation of cancer cells by causing dynamic alterations in the acetylation and deacetylation status of key target proteins and transcription factors.¹ For example, HDACis target the acetylation of apoptotic genes for cancer regulation. In apoptosis-deficient cells, however, HDACis target autophagy regulatory genes to induce ACD.¹⁰ Autophagy, a catabolic cellular self-degradative pathway, has been under investigation in recent times as an alternative mechanism of cell death.⁵ Several reports have indicated that HDACs deregulate key autophagic genes to sustain cancer cell survival.³ In this context, HDACis are effective yet selective cytostatic agents that target autophagy to inhibit cancer cell survival.

HDACs mediate the deregulation of cytoprotective autophagy in cancer progression, and the HDAC inhibitor vorinostat acts as an autophagy modulator

Deacetylation of key autophagic genes promotes cytoprotective autophagy that encourages cancer cell proliferation, survival and drug resistance.¹¹ The application of broad-spectrum multipharmacological HDACi in cancer therapy has emerged as a prominent therapeutic strategy in cancer treatment.^{3, 12} Hence, understanding the molecular mechanisms and critical molecules that are associated with cancer cell survival will provide insight into the best opportunities for therapeutic applications.

Deacetylation of key molecular markers during the stage-specific progression of autophagy for cancer progression

Cancer cells preferentially activate cytoprotective autophagy as a survival mechanism to overcome apoptosis-mediated cell death.⁵ From this perspective, regulation of the deacetylation status of selective autophagy regulatory genes drives cytoprotective autophagy and promotes cancer cell survival and progression.¹ Cancer cell survival and progression, in lieu of the deacetylation of stage-specific autophagic regulatory genes, are described below.

Deacetylation of autophagic marker proteins during autophagy initiation—In MCF-7 breast cancer cells, downregulation of ESR1/estrogen receptor α is transcriptionally regulated by HDAC1 and RPS6KB/p70S6K. The phosphorylation of HDAC1 and activation of the PI3K–MTOR signaling pathway is regulated by RPS6KB/p70S6K, which recruits HDAC1 to the *ESR1* promoter and enhances drug resistance.¹³ In triple-negative breast cancer cells, activators of HDACs and MTORC1 enhance the expression of EIF4EBP1 to promote cell viability.¹⁴ In endometrial cancer, HDAC6 supports cancer cell proliferation, metastasis and invasion by modulating the PTEN–AKT–MTOR signaling pathway.^{15, 16} Although the direct involvement of autophagy was not reported in the study by Zheng *et al.*,¹⁵ the dynamic regulation of AKT–MTOR is expected to play a cytoprotective role in

nullifying a *MIR206*-mediated carcinostatic effect. In human multiple myeloma cell lines, dynamic co-regulation of HDAC and MTOR promotes the stability of the oncogenic protein MYC, which promotes cancer cell survival and progression.¹⁷

Deacetylation of autophagic marker proteins during phagophore membrane expansion—In hepatocellular carcinoma cells, HDAC1 overexpression regulates enhanced LC3-II lipidation to induce cytoprotective autophagy and thus to sustain hepatocarcinogenesis.^{18, 19} In Waldenstrom macroglobulinemia cells, HDAC4 and HDAC5 induce cytoprotective autophagy by enhancing the expression of LC3-II and RAB7, which abrogates apoptosis regulated by LBH589 (a specific HDAC4 and HDAC5 inhibitor) and *MIR9* (microRNA 9).^{20, 21} In human cervical cancer (HeLa) cells, HDAC6 deacetylates LC3-II, promoting autophagosome formation.²² Tubacin, a specific HDAC6 inhibitor, results in partial acetylation of LC3-II, suggesting the involvement of other HDACs in the deacetylation process. Moreover, tubacin-induced LC3-II acetylation further impairs the autophagic flux by promoting the accumulation of SQSTM1/p62.²³ In gastric cancer cells, class III HDACs deacetylate BECN1, ATG5, ATG7 and Atg8-family proteins, which preferentially support cell survival, proliferation, drug resistance, invasion and metastasis. Moreover, deacetylation of BECN1 by class III HDACs promotes epithelial-to-mesenchymal transition (EMT) in hepatocellular carcinoma via autophagic degradation of CDH1/E-cadherin.²⁴

Deacetylation of autophagic marker proteins during flux in autophagy—In metastatic prostate cancer (DU145) cells, a dynamic co-regulation of SQSTM1 and HDAC6 directs the acetylation of TUBA/ α -tubulin, thereby reducing the stability and motility of this complex. The less stable and motile acetylated form of TUBA/ α -tubulin impairs the autophagy flux, and subsequently promotes invasion, metastasis and EMT.²⁵ In MYC/v-Myc myelocytomatosis-overexpressing neuroblastoma cells (BE[2]-C, Kelly and IMR32 cells), HDAC10 facilitates the onset of cytoprotective autophagy against doxorubicin-induced cytotoxicity via deacetylation of HSPA/heat shock protein 70 and HSPA8/HSC70. Mechanistically, HDAC10-mediated deacetylation of HSPA8 results in the selective transport of cargos to lysosomes in a LAMP2-dependent molecular cascade, whereas HSPA maintains lysosomal membrane integrity for sustained autophagic flux. Hence, the autophagosome formation that results from HDAC10-mediated deacetylation induces cytoprotective autophagy in neuroblastoma cells.¹¹

Vorinostat as a broad-spectrum pan-HDACi

As HDACs promote the survival and progression of cancer cells, there is a dire need for strategies to inhibit HDAC and thus prevent cancer.³ Several HDACi, such as vorinostat, panobinostat, romidepsin, Trichostatin A, tubacin and tubastatin, mediate ACD in several cancer subtypes. Among all the pan-HDACi, vorinostat and romidepsin have been approved by the US Food and Drug Administration (FDA). Panobinostat has been used experimentally in several solid cancers and has shown the ability to induce ACD at a lower dosage than vorinostat. Panobinostat has been under clinical trial in recent times.^{1, 26} In the ACD context, vorinostat (suberanilohydroxamic acid (SAHA)), a member of the hydroxamate class of broad-spectrum pan-HDACi, has been studied extensively and approved by the

FDA for the treatment of cutaneous T-cell lymphoma. Vorinostat inhibits class I, II and IV HDACs that display Zn²⁺-dependent deacetylase activity. Chemically, vorinostat binds to the Zn²⁺ ion present in the active site of HDACs, thereby restricting their enzymatic activity.²⁷ Several reports have implied the potential of vorinostat as an anticancer drug in various cancer subtypes because of its abilities to induce cell cycle arrest, apoptosis, and DNA-damage repair, thereby inhibiting metastasis and angiogenesis, and because of its autophagy-modulating activities.²⁷ From a cancer treatment perspective, the ability of cancer cells to escape apoptosis suggests autophagy as an alternative cell death mechanism that could be modulated by therapeutics.^{5, 8} Various reports suggest that vorinostat-mediated autophagy can function in this capacity, and this broad-spectrum pan-HDACi has become highly prevalent in cancer therapeutics. Moreover, vorinostat can act on multipharmacological targets (EGFR, RTKs and synergistic agents), making it a more effective drug against heterogeneous tumors.²⁸

Modes of ACD induced by vorinostat

The Nomenclature Committee on Cell Death has denoted ACD as a separate ‘cell death subroutine’ with distinct morphological characteristics, including rounded cell structure, detachment from the substratum, deposition of small vesicles, excessive accumulation of cytoplasmic vacuoles, and double-membrane compartments that resemble autophagosomes containing damaged cellular aggregates, damaged organelles and cytoplasmic cargos.^{6, 8} ACD can further be divided into autophagy-mediated (autophagic facilitation of apoptotic cell death), autophagy-associated (co-existence of autophagic and apoptotic cell death) and autophagy-dependent (autophagy as an explicit cell death mechanism that is independent of apoptosis) modes.⁹ With limited evidence available, we summarize all three modes of ACD in the context of nomenclature, mode and molecular mechanism of action (Figure 1). Table 1 lists the modes of ACD seen after vorinostat treatment in several *in vivo* and *in vitro* models of numerous cancer subtypes.

Where and how does ACD occur after vorinostat treatment

How autophagy, the catabolic process that helps in cell survival under stress conditions, becomes lethal by over-consuming cellular components is not fully understood because of the lack of molecular evidence. Nevertheless, recent investigations have provided multiple indications of autophagy being lethal in the context of cellular dependency.⁸ A few of the suggested molecular mechanisms are discussed below (Figure 2).

Hyperactivation of the PIK3C3/VPS34–BECN1 complex induces superactivated autophagy

The PIK3C3/VPS34–BECN1 complex plays a significant role in the onset of hyperactivated autophagy, which drives autophagic cell death.⁸ Previous reports have demonstrated that the activation of BECN1 after vorinostat treatment induces ACD in several cancer subtypes.²² Moreover, it has also been reported that vorinostat downregulates BCL2 and BCL2L1/Bcl-xL protein expression during autophagy associated with cell death, with subsequent activation of BECN1.²² In a molecular context, DAPK1 mediates the kinase cascades that are involved in the phosphorylation of protein kinase D (PRKD), which then phosphorylates and activates PIK3C3/VPS34.²⁹ Second, the phosphorylation of BECN1 is mediated by

DAPK1, which subsequently dissociates BECN1 from BCL2 and promotes its association with PIK3C3/VPS34. In acute myeloid leukemia, vorinostat, in synergism with tyrosine kinase inhibitors and FLT3 (fms-related receptor tyrosine kinase 3) inhibitors, mediates the derepression of DAPK1.³⁰ Furthermore, a cyclic synergistic administration of sorafenib and vorinostat in acute myeloid leukemia mediates DAPK1 derepression.³¹ Although the reports are limited, it appears that the functional association of vorinostat and DAPK1 activation might drive hyperactivation of the PIK3C3/VPS34–BECN1 complex and the subsequent induction of ACD. Interestingly, another pan-HDACi panobinostat, LBH589, induces DAPK1-dependent ACD via enhanced LC3 lipidation. In autophagy-deficient cells, LBH589 promotes DAPK1-independent apoptotic cell death.³²

Sustained autophagic flux to restrain bulk autophagy for autophagic cellular morphologies

Progressive autophagic flux overwhelms the cellular cytoplasm with the accumulation of non-cargo-containing autophagic vacuoles and empty autophagosomes. In the later stages of bulk autophagy, the autophagic vacuoles fill the cytoplasmic volume, which is presumably devoid of intracellular organelles such as mitochondria, Golgi and endoplasmic reticulum (ER).^{8, 33} In addition to overconsumption of intracellular organelles, rerouting of cellular membranes to facilitate the production of autophagosomes disrupts cellular membrane homeostasis.⁸ The mechanistic observations in TAMR/MCF-7, MDA-MB-231 and 4T1 cells, as well as those in *apaf1*-knockout mouse embryonic fibroblasts (MEFs) and BCL2L1-overexpressing HeLa cells, have suggested the onset of ACD with prominent autophagic cellular morphologies following vorinostat treatment.^{10, 34}

Death from disrupted lysosomal homeostasis

Lysosomal homeostasis acts as a key component for maintaining bulk autophagy for subsequent activation of autophagic cell death. Conservation of the lysosomal pool during hyperactivated autophagy induction promotes enhanced autophagic flux.³⁵ In this context, it has been reported that vorinostat treatment improves lysosomal function (lysosomal acidification), number and enzymatic activity; particularly proteolytic including CTSB (cathepsin B), CTSD (cathepsin D) and CTSL (cathepsin L).³⁶ Moreover, vorinostat activates lysosomal biogenesis through acetylation of TFEB and its nuclear translocation. Besides TFEB, vorinostat enhances the expression of TFEB target genes such as UVRAG and LAMP1, which are important during the stage-specific progression of autophagy.³⁶ By contrast, a recent finding from a drug screening using the lysosomal-METRIQ probe has demonstrated vorinostat functions as a downregulator of lysosomal membrane trafficking and activity.³⁷ It has been expected that vorinostat-mediated downregulation of lysosomal activity might inhibit autophagic flux leading to autophagosome accumulation and subsequent activation of autophagy-mediated cell death.

Although Transcription Factor EB (TFEB) acts as a master regulator of autophagy and lysosomal biogenesis, selective cargo sorting and degradation via chaperone-mediated autophagy with the help of HSP90 and LAMP2 also play a key roles in ACD.³⁸ A combination of vorinostat with doxorubicin and paclitaxel enhances the acetylation of HSP90 and reduces the ectopic expression of HSP90 client proteins. This reduction in the expression of HSP90 client proteins, together with reduced expression of LAMP2A,

prevents the lysosomal degradation of CDKN2A, which in turn elicits ACD.^{38, 39} Moreover, acetylated microtubules facilitate autophagosome–lysosome fusion, which is critical for hyperactivated autophagy.⁴⁰ Acetylation of tubulin (at Lys40) is enhanced in both the labile and stable populations of microtubules, and results in the recruitment of dyneins and KIF1 that deliver autophagosomes centripetally towards lysosomes along microtubules.^{40, 41} In this context, vorinostat-mediated microtubule acetylation might expedite the enhanced autophagosome–lysosome fusion required for ACD.

Death from excessive mitophagy followed by compromised mitochondrial biogenesis

Excessive removal of mitochondria through autophagy creates a problem for mitochondria-depleted cells in generating the energy required for cellular homeostasis. The association of autophagy markers with mitochondria leads to lethal autophagy of these organelles.⁴² A mechanistic investigation revealed the role of smARF/CDKN2A (CDKN2A is the translational precursor of smARF) in mitophagic cell death. smARF/CDKN2A overexpression leads to autophagy-dependent cell death independent of caspase-mediated apoptosis.^{8, 43} Moreover, under hypoxia or external stimuli, enhanced expression of smARF/CDKN2A sustains bulk autophagy via SQSTM1 degradation. Subsequently, smARF/CDKN2A deregulation induces PINK1–PRKN/Parkin mitophagy.^{8, 44} In cutaneous T-cell lymphoma, the deregulated expression of CDKN2A that is induced by vorinostat treatment reduces tumor proliferation, in a process that might involve autophagy-dependent cell death resulting from excessive mitophagy.⁴⁵

In another investigation, mitochondrial translocation of orphan nuclear receptor NR4A1/TR3/Nur77 increased mitochondrial membrane permeability through the permeability transition pore ANT1–VDAC1, leading to mitophagic cell death. Mechanistically, the mitochondrial translocation of NR4A1/TR3/Nur77 occurs via the interaction of BNIP3L/Nix with the outer mitochondrial membrane proteins TOMM20 and TOMM70.⁴⁶ It has been shown that vorinostat upregulates the expression of NR4A1/TR3/Nur77 and its associated signaling mediators such as CDKN1A/p21.⁴⁷ Limited evidence suggests that the dynamic interaction of vorinostat and NR4A1/TR3/Nur77 might be associated with cell death as the result of mitophagy. In addition to NR4A1/TR3/Nur77, BNIP3 also mediates mitophagy-dependent cell death. In the context of autophagy-associated cell death, a synergistic combination of several drugs with vorinostat regulates the expression of BNIP3, which might follow the induction of mitophagic cell death in close coordination with apoptosis.⁴⁸ Moreover, the extensive interplay between DAPK1 and BNIP3 illustrates the possibility of mitophagic cell death.⁸

Mitochondrial energy homeostasis post mitophagy is thought to be maintained by mitochondrial biogenesis. PPARGC1A/PGC-1 α is a critical regulator of mitochondrial biogenesis, working alongside several transcriptional and signaling cascades.⁴⁹ In osteosarcoma cells, vorinostat downregulates the expression of PPARGC1A, which might be responsible for diminishing the metastatic behavior of these cancer cells.⁵⁰ In addition to this, vorinostat also suppresses the MYC/c-MYC-regulated expression of NFE2L2/NRF2 (a transcription factor associated with PPARGC1A), and restores the function of KEAP1, which might promote mitochondrial biogenesis.⁵¹ Moreover, acetylated TP53/p53

inhibits the expression of AMPK, a critical regulator of cellular energy metabolism via mitochondrial biogenesis. Vorinostat treatment results in the acetylation of TP53 and, subsequently, AMPK, potentially inhibiting mitochondrial biogenesis.⁵²

Secondary signals and autophagy-associated signaling cascades elicit cell death

Ceramides are critical for the induction of apoptosis and necrosis-independent cell death.⁸ Furthermore, several reports have demonstrated the essential involvement of ceramides in the induction of autophagy-dependent cell death. In gastrointestinal tumor cells, vorinostat in synergism with sorafenib resulted in enhanced ceramide generation, which in turn mediates PPP2/PP2A-DAPK1-dependent cell death.⁵³ Furthermore, vorinostat directly causes the acetylation of the HIF1A/HIF1- α -associated chaperone HSP90, subsequently hindering the nuclear translocation of HSP90 and consequently inhibiting the transcriptional activity that might regulate HIF1A-dependent ACD.⁵⁴ In addition to this, vorinostat-mediated acetylation deregulates the expression of oncogenic RAS, leading to PMAIP1/Noxa- and BECN1-dependent ACD.⁵⁵ More importantly, the acetylation of cytoplasmic LC3, which might be achieved through vorinostat therapy, is critically involved in the regulation of superactivated autophagy that leads to cell death.⁵⁶

Vorinostat-mediated regulation of autophagy for cancer therapy

The clinical efficacy of vorinostat in terms of regulating autophagy, specifically the expression of key autophagic markers in tumors, has been demonstrated in several cancer subtypes. In addition to its ability to regulate autophagy, however, the clinical pharmacology (i.e., pharmacokinetics and pharmacodynamics) of vorinostat, used either as a mono-chemopreventive agent or in combination with autophagy activators or inhibitors, is immensely important.⁵⁷⁻⁵⁹ A number of pharmacological parameters have been estimated in different phases of randomized and non-randomized clinical trials (Table 2). These include the dose and periodicity of treatment, drug synergism, drug efficacy, drug-related adverse toxicity, pharmacodynamics targets and biomarkers of primary tumor tissues, pharmacokinetic drift, the area under curve, median half-life, and drug concentrations in tumor, blood and plasma.⁵⁷ With limited clinical trials involving an assessment of autophagy regulation available, we have summarized clinical pharmacology results for various vorinostat combinations in Table 2.

Challenges for clinical translation and scope for future expansion

After the approval of vorinostat by the FDA in 2006 for the treatment of cutaneous T-cell lymphoma, several clinical trials in different phases have looked at the use of vorinostat as a mono-therapeutic agent or in combination with other drugs as a treatment for hematological and solid tumors.²⁷ The clinical outcomes achieved by vorinostat as a mono-therapeutic agent against solid tumors have been disappointing. The restricted pharmacokinetic properties of the drug, which affect its absorption, distribution, excretion, metabolism, targeted drug delivery, and short half-life, hinder the clinical translation of vorinostat. Moreover, adverse side effects and toxicity during early-phase clinical trials have further restricted its clinical translation.⁵⁷ In addition to this, the complexity of the autophagic response, which ranges from pro-survival to lethal for cancer cells, makes

the use of vorinostat more challenging. The drug synergism of vorinostat with specific autophagy mediators may provide a way to overcome these challenges for translation, ultimately delivering valuable outcomes in personalized and precision cancer therapy. The combination of chloroquine (CQ) or hydroxychloroquine (HCQ) with vorinostat has shown promising clinical results in several cancer subtypes.^{57, 58, 60} Hence, the design of stage-specific autophagy-targeting drugs and drug combinations will offer improved clinical efficacy over that provided by vorinostat alone, owing to their improved pharmacokinetics and pharmacodynamics. Importantly, the identification of drug targets and an understanding of the mechanistic regulation of ACD will make it easier to use vorinostat to provide an enhanced carcinostatic effect in apoptosis-deficient or -resistant cancer cells.

Conclusions

Deacetylation of key molecular markers that are involved in apoptosis and autophagy plays a crucial role in the initiation and progression of cancer. With the dire need to identify and establish autophagy as an alternative cell death mechanism, apoptosis-resistant and -deficient cancer cells deregulate autophagy-related genes through deacetylation. By reversing such deregulation, HDACis impart a critical response in cancer cells and facilitate ACD. Vorinostat, a broad-spectrum HDACi, enables autophagy-mediated, -associated and -dependent death of cancer cells. By acetylation of several key components of the autophagy machinery, vorinostat promotes superactivated autophagy through hyperactivation of PIK3C3/VPS34-BECN1. In addition, vorinostat mediates nonselective autophagy via lysosomal disparity and mitophagy, with subsequent inhibition of mitochondrial biogenesis, that expedites ACD. Notably, an understanding of the clinical pharmacology of ACD regulation allows vorinostat to be implemented as a mono-therapeutic or as a synergistic drug for cancer prevention. More investigation is warranted to delineate the mechanistic involvement of vorinostat-mediated inhibition of acetylation in ACD induction. Further elucidation of the vorinostat specificity of novel molecular ACD targets will afford enhanced therapeutic value for personalized cancer therapy.

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Highlights

1. HDACs mediate deregulation of cytosolic autophagy for cancer progression.
2. Vorinostat mediates autophagy-mediated, associated and dependent cell death.
3. Hyperactivation of PIK3C3/VPS34-BECN1 complex aids vorinostat regulated cell death.
4. Lysosomal disparity; biogenesis and functionality facilitates cell death.
5. Excessive mitophagy followed by compact mitochondrial biogenesis provokes cell death.

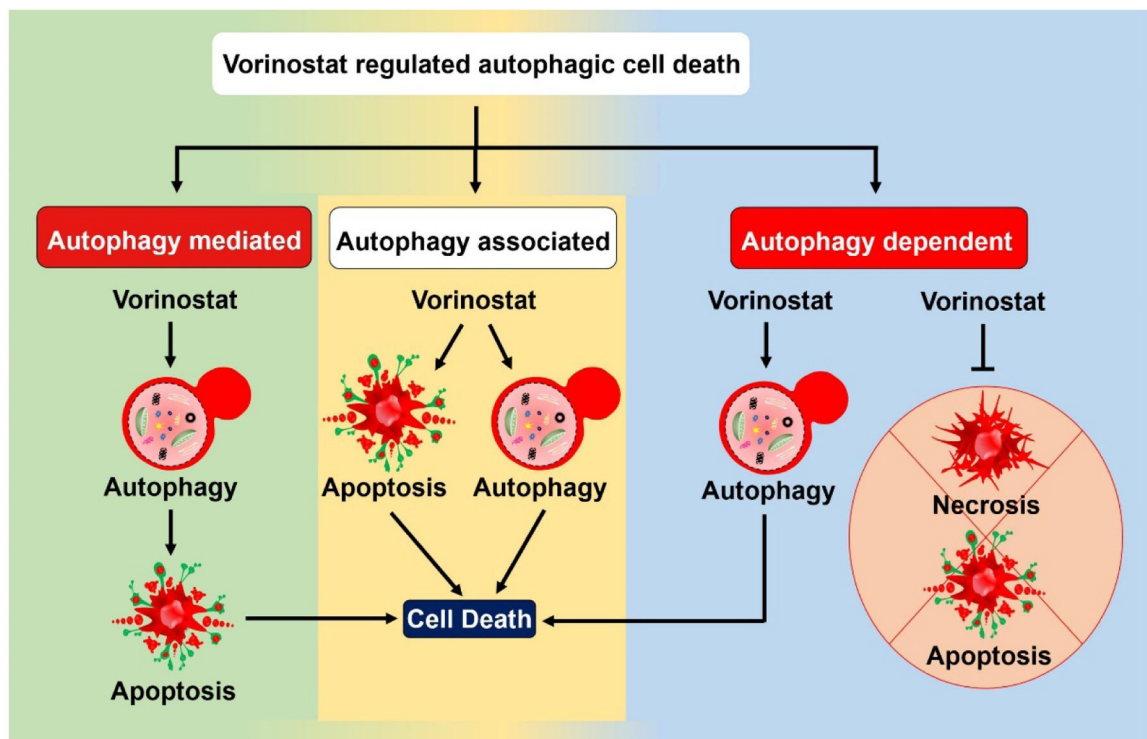


Figure 1. Intricate role of vorinostat-regulated autophagy in cell death. Vorinostat-regulated autophagic cell death (ACD) can be grouped into autophagy-mediated, -associated and -dependent modes depending on the involvement of apoptosis. In autophagy-mediated death (autophagic facilitation of apoptotic cell death) cell, the autophagy pathway activates the apoptotic cell death pathways. In autophagy-associated cell death (in which autophagic and apoptotic cell death co-exist), the induction of autophagy accompanies apoptotic cell death. Autophagy-dependent (autophagy as an explicit cell death mechanism) cell death occurs independently of the apoptotic and necrotic cell-death pathways.

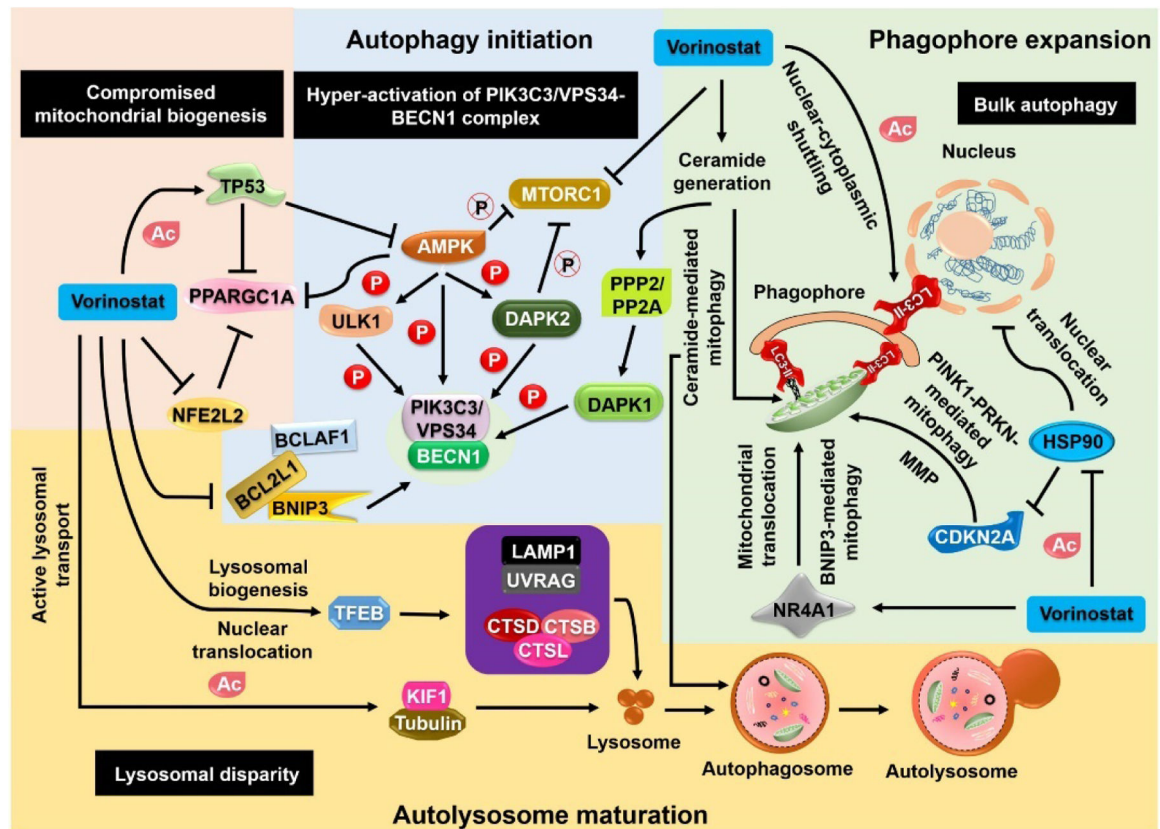


Figure 2. Mechanisms through which vorinostat-regulated autophagy promotes cell death. Lethal autophagy can be introduced via hyperactivation of the PIK3C3/VPS34–BECN1 complex in the autophagy initiation stage. Apart from direct inhibition of mTOR (MTORC1), vorinostat activates AMP-activated protein kinase (AMPK) and causes the phosphorylation of particular substrates (ULK1, DAPK2), which inhibits mTOR expression. Further modulation of the phosphorylation status of BECN1 by PPP2/PP2A–DAPK1 inhibits its interaction with BCL2 homology domain proteins, leading to the sequestration of BNIP3 and BCLAF1. Despite bypassing canonical initiation of autophagy, the noncanonical pathways drive the formation of autophagosomes. Secondary signals, such as ceramide generation, that recruit mitochondria to the phagophore membranes initiate mitophagy by interacting with LC3-II. Moreover, vorinostat regulates the acetylation of LC3-II to mediate nuclear–cytoplasmic shuttling, which has a substantial role in enhancing autophagosome formation. In addition, vorinostat inhibits the acetylation of HSP90, preventing its nuclear translocation. The inhibition of HSP90 translocation inhibits CDKN2A degradation and facilitates PINK1–PRKN-mediated mitophagy. Furthermore, vorinostat mediates the mitochondrial translocation of NR4A1/Nur77, thereby aiding BNIP3-mediated mitophagy. During the autolysosome maturation stage, vorinostat causes nuclear translocation of Transcription Factor EB (TFEB), leading to further activation of downstream lysosomal targets such as LAMP1, UVRAG, CTSD, CTSB, and CTSL and to enhanced lysosomal biogenesis. In addition to lysosomal biogenesis, vorinostat mediates the acetylation of tubulin and KIF1, resulting in enhanced lysosomal transport. During the compromised mitochondrial biogenesis, vorinostat acetylates TP53, which in turn inhibits

AMPK and PPARGC1A. Moreover, vorinostat-mediated inhibition of NFE2L2 also inhibits PPARGC1A-mediated mitochondrial biogenesis.

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Table 1:

Different modes of vorinostat regulated autophagic cell death (ACD).

Types of ACD	Drug entity	Cancer type or model	Molecular mechanisms of action	Reference
Autophagy-mediated cell death	Synergism of temozolomide and vorinostat with chloroquine	Glioblastoma (C6 and U251MG) cells	Temozolomide induces cyto-protective autophagy via sustained upregulation of AMPK–ULK1 and inhibition of AKT–mTOR. Vorinostat inhibits mTOR signaling to aid cyto-protective autophagy. Inhibition of temozolomide-vorinostat regulated cyto-protective autophagy by chloroquine facilitates the onset of apoptotic cell death pathways.	61
		GL261 glioma tumor-bearing C57BL/6 mice	Temozolomide (10 mg/kg dose, every 3 days for 10 days) vorinostat (i.p. 25 mg/kg/day for 10 days) and chloroquine (pH 7; 20 mg/kg/day for 10 days) inhibits cyto-protective autophagy to aid apoptosis.	
	Vorinostat in synergism with chloroquine	Glioblastoma (U87MG) cells	Autophagic flux inhibition by chloroquine facilitates vorinostat-mediated apoptosis via hyperaccumulation of autophagosomes, prevention of the elimination of damaged mitochondria and increased lysosomal pH with an eventual generation of reactive oxygen species (ROS).	60
	Vorinostat in combination with quinacrine	T-cell ALL (Jurkat and Molt-4) cells	Quinacrine in synergism with vorinostat facilitates the inhibition of mitophagy, enhances the K-63 associated ubiquitination of mitochondrial proteins, enhances the accumulation of mitochondrial aggregates, enhances ROS production, decreases mitochondrial membrane potential, enhances the expression of Bax and downregulates the expression of Bcl-2.	62
		Jurkat-cell-bearing SCID mice	Vorinostat (i.p. 70 mg/kg/d for 12 d) in synergism with quinacrine (i.p. 50 mg/kg/d for 12 d) mediates the onset of apoptosis, resulting in reduced tumor cell proliferation.	
	Vorinostat in synergism with chloroquine	Acute myeloid leukemia (AML) t(8;21) cells	Inhibition of vorinostat-mediated pro-survival autophagy by chloroquine results in excessive accumulation of ubiquitinated proteins (AML1-ETO) and aggregates, leading to cell death.	63
	Synergism of gefitinib-vorinostat with metformin and spautin-1	EGFR-TKI resistant PC-9GR and H1975 non small cell lung cancer (NSCLC)	Metformin and spautin-1 reduce the lipidation of LC3, reduces the phosphorylation of Beclin-1 and enhances the expression of P62, leading to chemosensitization of the resistant NSCLC towards apoptosis (evidenced by upregulation in the expression of BIM and Bax and downregulation of antiapoptotic proteins Bcl-2, Bcl-XL and Mcl-1).	64
	Vorinostat	Neuroblastoma (SK-N-SH and QDDQ-NM)	Decreased LC3 lipidation and decreased expression of ATG5 and Beclin-1, leading to chemosensitization of neuroblastoma cells towards apoptosis.	65
	Vorinostat in synergism with simvastatin	Triple negative breast cancer (TNBC) (MDA-MB-468, MDA-MB-231 and MDA-MB-453) cells	Inhibition of autophagic flux, leading to accumulation of LC3II, P62, NDP52 and NBR1, reduced the prenylation of Rab7 that is required for autolysosome formation. Eventual onset of apoptosis through caspase-dependent signaling pathway.	66
	Synergism of bortezomib and vorinostat with chloroquine	Colon cancer (HT-29 and HCT-8) cell	Disruption of autophagic flux, bulk generation of superoxide as chief regulator of ubiquitinated protein accumulation, and subsequent onset of apoptosis.	67
Vorinostat in synergism with chloroquine	HCT-8 colon cancer-bearing female BALB/c xenograft mouse model	Chloroquine (60mg/kg/d for 21 d) and vorinostat (100mg/kg/d for 21 d) treatment results in the inhibition of autophagy, leading to ubiquitinated protein accumulation, greater superoxide generation and the subsequent onset of apoptosis.	68	
		Vorinostat in combination with sorafenib		Hepatoma (HepG2, Hep3B and PLC/PRF/5) cells

Types of ACD	Drug entity	Cancer type or model	Molecular mechanisms of action	Reference
	Vorinostat and pharmacological inhibitors of autophagy	Glioblastoma stem cells (GSCs) and GSC-bearing mouse xenograft model	Inhibition of autophagy by pharmacological agents (3MA, wortmannin, BafA1 and chloroquine) and genetic agents (ShLC3, shBeclin-1 and shATG5) results in the onset of apoptotic cell death (evidenced by cleaved Caspase 3 and PARP).	69
Autophagy-associated cell death	Vorinostat in combination with olaparib	TNBC (MDA-MB-231 and MDA-MB-2468) cells	Vorinostat mediates the onset of autophagy-associated cell death, which in turn enhances the cytotoxic effect of olaparib (a PARP inhibitor) through PTEN modulation to sustain tumor cell proliferation and to induce apoptotic cell death.	70
	Vorinostat in combination with BIBW2992/WZ4002	EGFR-TKI NSCLC (PC-9 and NCI-H1975) cells	Enhanced expression of LC3II and Beclin-1 and downregulation of p62 dynamically regulates the onset of caspase-dependent intrinsic apoptosis.	71
		NCI-H1975-bearing mouse xenograft model	Vorinostat (25mg/kg) in combination with BIBW2992 (10mg/kg) and WZ4002 (25mg/kg) induces autophagy (enhanced LC3 lipidation and Beclin-1 expression) and apoptosis (evidenced through cleaved PARP and Caspase 3, and by downregulation of Bcl-XL).	
	Vorinostat in combination with tamoxifen	TAMR/MCF-7 and TAMR/T47D cells	Induces both apoptotic cells death (evidenced by cleaved Caspase 3, 7, 9 and PARP, upregulation of Bax, and downregulation of Bcl-2 together with ER-associated gene transactivation) and autophagic cell death (evidenced by enhanced LC3 lipidation and expression of Beclin-1, ATG 5 and ATG 7).	34
	Vorinostat in combination with ionizing radiation	MDA-MB-231/4T1 cells and 4T1-bearing Balb/c <i>in vivo</i> mouse model	Co-existence of apoptotic cell death (evidenced by annexin-v-positive cells, nuclear pyknosis and chromatin condensation) and autophagic cell death (evidenced by enhanced expression of LC3II and Beclin-1).	72
	Vorinostat in combination with decitabine	Ovarian cancer (SKOv3 and Hey) cells	Co-existence of apoptotic (evidenced by enhanced expression of Caspase 3/7) and autophagic cell death (evidenced by enhanced LC3 lipidation and overexpression of ARH1 and PEG3).	73
Autophagy-dependent cell death	Vorinostat in combination with BIBW2992/WZ4002	EGFR-TKI NSCLC (PC-9) cells	Induction of autophagic cell death independent of apoptosis: treatment with the pan caspase inhibitor Z-VED-FMK does not abrogate cell death.	71
	Vorinostat	TAMR/MCF-7-bearing athymic mouse xenograft model	Administration of vorinostat (i.p. 50mg/kg, every 2 d for 21 d for 10 treatment cycles) induces autophagic cell death (evidenced through enhanced expression of LC3, Beclin-1, ATG5 and ATG7).	34
	Vorinostat	Bcl-XL overexpressing cervical cancer (HeLa) cells	Induction of non-apoptotic cell death despite inhibition of cytoplasmic shuttling of cytochrome C and caspase activation with distinct autophagic cellular morphologies.	10
	Vorinostat in combination with pazopanib	Fibrosarcoma (HT1080), uterine sarcoma (MES) and osteosarcoma (Saos-2) cells	Autophagic cell death with an evident increase in LC3 lipidation and Beclin-1 expression. Inhibition of the mTOR-p70S6K pathway for the induction of lethal autophagy.	74
	Vorinostat and pharmacological inhibitors of autophagy	Gastric cancer (AGS and KATO-III) cells	Induction of autophagy (enhanced lipidation of LC3 and decreased expression of p62) followed by autophagy flux inhibition facilitates autophagy-dependent cell death.	75
	Vorinostat	Hepatocellular carcinoma (Huh7, Hep3B and HepG2)	Enhances LC3 lipidation, Beclin-1 expression, downregulation of p62 and inhibition of the PI3K-AKT-mTOR signaling axis. Phosphorylation of ER stress regulates markers such as PERK and eIF2 α , leading to autophagy-dependent cell death.	76
	Vorinostat	Endometrial stromal sarcoma (ESS-1) cells	Downregulation of p-mTOR, mTOR and pS6K expression, leading to induced autophagy-dependent cell death that was independent of activation of Caspase 3.	77

Table 2:

Detailed evaluation of the clinical pharmacology of vorinostat in the regulation of autophagy.

Details of clinical trial	Drug entity	Cancer type	Clinical pharmacology properties and their relevance	Autophagy-associated molecular mechanisms of action	Reference
Phase I clinical trial (NCT01023737 [*])	Hydroxychloroquine (HCQ) (400–1000) and vorinostat (300) mg/m ² /day	Patients with advanced or malignant solid tumors (colorectal and renal)	Evaluation of safety, tolerability, drug response rate, progression-free survival and anti-tumor activity.	Inhibition of autophagy by HCQ in combination with vorinostat critically regulated autophagy, resulting in inhibition of cancer progression with minimal toxicity.	–
Phase I clinical trial	HCQ and oral ingestion of vorinostat (400 mg/m ² , once a day for 21 d)	Colorectal, soft tissue sarcoma, ovarian cancer, melanoma, NSCLC, renal cell carcinoma, prostate, breast and bladder cancers	Evaluation of minimal dose-related (MDR) toxicity at maximum tolerated dose (MTD), C _{max} of 768 and 786 µg/L, AUC of 3387 and 2410 µg × hr/L, median half time of 2.06 h and 1.3 h and median Vd/f of 304–309 L.	Enhanced expression of cathepsin D (CTSD), which is responsible for the onset of HCQ-vorinostat-regulated autophagy-mediated apoptosis. Accumulation of autophagic vacuoles (AVs). Enhanced expression of CDKN1A, SQSTM1 (p62) and MAP1LC3B.	57
Early phase clinical trial	HCQ (600) and oral ingestion of vorinostat (400) mg/m ² , once a day for 21 d)	Advanced and refractory metastatic colorectal cancer	Evaluation of median overall survival (6.7 months), median progression-free survival (2.8 months) and minimal drug-related adverse toxicity.	Enhanced CTSD and p62 level.	58
Phase II randomized, therapeutic clinical trial (NCT02316340 [*])	HCQ (600 mg/m ²) and oral ingestion of vorinostat (400 mg/m ² , once a day for 28 days)	Chemorefractory metastatic colorectal cancer	Evaluation of enhanced anti-tumor efficacy, overall survival and median progression-free survival	Autophagy inhibition leading to enhanced tumor immunity (decreased expression of IL-1b, IL-6, IL-10, IL-2, IFN-γ and TNF-α).	–
Phase II clinical trial (NCT00882206 [*])	Vorinostat (230 mg/m ² , oral gavage, twice a day), intravenous injection of decitabine (15 mg/m ²), prednisone (40 mg/m ² , twice a day for 8 days), vincristine (1.5 mg/m ²), doxorubicin (60 mg/m ²), Pegasparginase (2,500 IU/m ²) and intrathecal cytarabine (70 mg for 14 d)	Relapsed and refractory acute lymphoblastic leukemia (ALL)	Evaluation of enhanced chemosensitization and identification of 32 differentially methylated genes in the responding groups as compared to non-responding ALL patients	Hypomethylation of cell death (apoptosis and autophagy) and transcription-regulating genes, such as SQSTM1, MAP3K5, SIRT1, DNABJ6, BTA1 and UB2E1.	–

* The NCT reference codes identify the clinical trials, and details of these clinical trials can be found in [ClinicalTrials.gov](https://clinicaltrials.gov).