

Combining HDAC inhibitors with oncolytic virotherapy for cancer therapy

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Abstract: Histone deacetylase (HDAC) enzymes play a critical role in the epigenetic regulation of cellular functions and signaling pathways in many cancers. HDAC inhibitors (HDACi) have been validated for single use or in combination with other drugs in oncologic therapeutics. An even more novel combination therapy with HDACi is to use them with an oncolytic virus. HDACi may lead to an amplification of tumor-specific lytic effects by facilitating increased cycles of viral replication, but there may also be direct anticancer effects of the drug by itself. Here, we review the molecular mechanisms of anti-cancer effects of the combination of oncolytic viruses with HDACi.

Keywords: epigenetics, glioma, oncolytic virus, HDAC inhibitor, HSV-1, cancer

Introduction

Significant genetic heterogeneity and instability are a hallmark of many cancers, and this contributes to treatment resistance. This is made even more complex by tumor epigenetic changes that also contribute to genomic instability and tumor plasticity.^{1,2} Epigenetic mechanisms include DNA methylation, post-translational modifications (PTMs) of histones and non-coding RNAs. One of the epigenetic modifiers attracting scientific attention as a potential therapeutic target comprises the histone deacetylases (HDACs). Advanced proteomic analyses with high-resolution mass spectrometry have enabled large-scale acetylome studies and subsequently have identified thousands of acetylation sites on the lysine moieties of more than 1,700 proteins in mammalian cells, including histones. For this reason, HDAC is also referred to as a lysine deacetylase.^{3,4} Pharmacological inhibitors of HDAC (HDACi) have been promisingly explored in therapeutic applications in cancer. In fact, pharmacological inhibitors, such as suberoylanilide hydroxamic acid (SAHA) and FK228, have been approved for clinical use to treat cancers by the Food and Drug Administration (FDA) in USA.

Oncolytic virotherapy was initially developed to eliminate cancer cells, with clinical treatment applications under development for over a decade.^{5,6} Oncolytic viruses (OVs) have also been applied to the area of cancer vaccination and immunotherapy. Genetically attenuated viruses are designed to selectively apply lytic effects against tumor cells (oncolysis), while simultaneously sparing non-tumor cells. However, as is the case with many therapeutic medications, OV therapy can be met with tumor resistance. This resistance may be due to less viral susceptibility and permissivity, attenuated apoptotic tumor death, and intrinsic/extrinsic antiviral immune responses. HDACi could address these resistance mechanisms to OVs, and are emerging as enhancers of oncolytic viro-

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therapy. In fact, recent work demonstrates that HDACi could improve OV therapy, reverse antiviral processes in tumor cells, and possibly promote OV-mediated tumor immunotherapy.

Oncolytic viruses

The concept behind OVs is for trying to find strains or mutants that are highly permissive and selective for infection and replication in malignant cells where they can produce their progeny for progressive oncolytic cycles throughout the tumor. In contrast to non-replicating viral vectors, OVs still maintain most of their viral genes intact, particularly those that encode genes required for the viral replicative life cycle, viral evasion mechanisms from host defense, and immune surveillance. OVs are engineered genetically or selected naturally from pathogenic viruses that include herpes simplex virus type 1 (HSV-1), adenovirus serotype 5 (Ad5), vascular stomatitis virus (VSV), measles virus, and many others. Strategies for acquiring and improving tumor selectivity in viruses range widely from genetic mutations and modification of transcriptional regulatory elements to structural modifications that alter infectivity and to arming with additional anticancer genes.^{7,8} OVs can also be combined with chemical drugs to obtain adjuvant and possibly synergistic anticancer effects both experimentally in animal models and clinically. The host still recognizes the OV as a foreign pathogen and will attempt to fight it via various immune responses that can diminish its oncolysis and lead to premature clearance of the OV. However, there exists a balance and this immune response can also lead to an anticancer effect. Immune activation in response to OV administration can thus also benefit treatment via a local cytokine storm in the tumor and recognition of tumor antigens. Understanding this balance between deleterious immunity that could lead to premature clearance of the OV and anticancer immunity that can lead to an effective anticancer therapy remains a target for scientific research and likely the objective to make OV therapy truly efficacious.^{9,10}

Several clinical trials using OVs have and are being conducted,^{7,11} but a recent phase III melanoma study using an HSV-based OV (oHSV1) is providing justification for approval by the US FDA as the first therapeutic viral oncolytic bioagent in the US (T-VEC; Amgen Inc. Thousand Oaks, CA, USA).¹² HSV-1, a member of alpha-herpes virus family, was the first virus that was genetically engineered to be oncolytic.⁶ It is one of the largest viruses by size (>150 kb) among OVs. The HSV-1 DNA genome is packed in the virion, which consists of three major layers, lipid envelope, tegument, and capsid. The kinetics of viral gene expression

consist of several immediate early (α) genes expressed right after infection, followed by early (β) genes expression. Viral DNA replication then triggers expression of late (γ) genes that encode viral structural components for progeny virions. In the design of oHSV1, appropriately placed mutations in one or several of these genes can attenuate lytic infection in normal tissues, while keeping infection of tumor tissues still to more permissive levels.⁸ Therapeutic success using OV is still a challenge since responsiveness and efficacy of OV may be different in different cancers and conditions due to different epigenetic states that affect viral permissiveness or susceptibility.

HDACs as modifiers of epigenetic silencing

PTM is an important biochemical process that occurs in both tumorigenesis and virus pathobiology. Acetylation and deacetylation of histones and non-histone substrates at lysine residues have emerged as crucial post-translational mechanisms. They regulate chromatin remodeling and accessibility of transcription factors that bind to specific sites in the regulatory regions of DNA to activate or suppress the transcription of specific genes by complexing with other proteins. Also, histone acetyltransferases (HATs) transfer acetyl groups to the ϵ -amino group of lysine on target substrates. Acetylation at lysine is a reversible process where HAT-mediated histone acetylation activates transcription, while the removal of the acetyl group causes chromatin condensation and transcriptional repression of genes. This deacetylation process is achieved by HDACs.

HDACs are classified into four groups based on similarity and function. Class I, II, and IV groups are classical zinc-dependent enzymes and are structurally distinct from the nicotinamide adenine dinucleotide (NAD⁺) dependent class III group, called sirtuins (SIRTs). Inhibitors of zinc-dependent HDACs have shown great potential in the treatment of epigenetic disorders including cancers. Many HDACi are currently in pre-clinical and clinical phases, and SAHA, FK228, and valproic acid (VPA), have been approved for clinical use by the FDA. Although SIRTs are considered to play key roles for physical stresses, metabolism, tumorigenesis, and aging, there is little information available regarding OV therapy with SIRTs and their inhibitors. Therefore, zinc-dependent HDACs and their inhibitors will be the focus of this review.

The class I group (HDAC1-3 and 8) is homologous to the yeast-reduced potassium dependency 3 (Rpd3) protein and is predominantly localized in the nucleus, but they also

localize in the cytoplasm to target cytosolic non-histone proteins. Class I HDACs are expressed ubiquitously in human tissues and global defects of this HDAC class in transgenic mice result in embryonic or perinatal lethality.^{13–16} HDACs often form complexes to deacetylase specific targets. HDAC1 and 2 have undergone little functional divergence from each other and can form homo- and hetero-dimer.^{17,18} HDAC1 and 2 are found in different transcriptional repressor complexes, including the Sin3 complex, the nucleosome remodeling and deacetylating complex, or the CoREST complex.^{19–21} HDAC3 is usually found in the complexes containing nuclear receptor corepressor 1 or the silencing mediator of retinoid and thyroid receptors.²²

The class II group (HDAC4-7, 9, and 10) is homologous to the yeast Histone deacetylase (Hda1) and localizes to the cytoplasm. This class II group is further subdivided into class IIa (HDAC4, 5, 7, and 9) and class IIb (HDAC6 and 10). The class IIa has a highly conserved deacetylase domain in its carboxyl-terminus and can be shuttled into the nucleus, based on the phosphorylation status of serine residues in the *N*-terminus, binding sites for the 14-3-3 proteins that regulate nuclear-cytoplasmic localization.²³ Since their intrinsic catalytic activities are relatively weak, class IIa HDACs usually interact with HDAC3 and other cofactors in order to increase the kinetics of deacetylation of target substrates.^{24,25} HDAC7 and 9 play key roles in lineage-determination in adaptive immunity.^{26–28} HDAC6 is unique among the HDAC family, in that it possesses two tandem catalytically active domains and an ubiquitin (Ub; and Ub-like modifiers) binding domain at its *C*-terminus. HDAC6 lacks *in vivo* deacetylase activity for nuclear histone proteins. Instead it is one of the major cytoplasmic deacetylases, targeting non-histone substrates, such as α -tubulin, Hsp90, and β -catenin.^{29–31} Biologically, HDAC6 functions in selective autophagy and in the aggresome processes of substrates that are conjugated to the moiety of Ub and Ub-like modifiers, and these processes also appear to impact the kinetics of viral infection/replication.^{32–35}

The class IV HDAC11 is the smallest one in the HDAC family. It contains mostly the catalytic domains and shares motifs with both class I and II HDACs. HDAC11 regulates the immunosuppressive cytokine molecule IL-10 in antigen-presenting cells.^{36,37}

HDAC1 and 2 as direct viral silencing factors

The HSV-1 genome is susceptible to host nuclear epigenetic mechanisms during infection, a process that allows the infected cell to regulate viral gene expression. HDAC1

and 2 are major enzymes that deacetylate the histone tails of nucleosomes containing HSV-1 genes and also play an important role in viral latency. Although HDACs themselves do not directly recognize and bind to DNA, HDAC1/2 are incorporated in the CoREST-REST-LSD1 repressor complex that binds repressor element 1 (RE-1) DNA sequences. This HDAC1/2 repressor complex primarily suppresses neuronal genes containing such RE-1s in their regulatory sequences in non-neuronal cells. The HSV-1 genome also contains multiple RE-1 sequences.^{38,39} HSV-1 infected cells that over-express wild-type REST were shown to preferentially lead to a latent program of viral life cycle, while HSV1 infected cells that express mutated REST (ie, lacking the binding capability to CoREST and HDAC1/2) preferentially lead to a lytic life cycle.^{40,41} Thus, HDAC1/2 and their repressor complex are critical factors in the decision between the lytic cycle or establishment of latency for HSV-1 in sensory neurons.⁴²

HSV-1 can interfere with HDAC1 and 2 mediated epigenetic silencing: the viral genes ICP0 and Us3 are known viral factors that inhibit HDACs. ICP0 is an immediate early (IE) protein, known as a viral Ub E3 ligase, and transfers Ub-chains to a target substrate for proteasomal degradation. Some intrinsic and innate antiviral factors including components of ND10 bodies (PML, SP100, and SUMO), p53 and the DNA sensor IFI16 are ICP0 substrates for degradation upon infection. ICP0 thus counteracts these antiviral host responses. ICP0's inhibition of HDAC1/2 seems to be independent of proteasomal degradation.³⁹ ICP0 instead disrupts CoREST-REST-LSD1 complex formation, dissociates HDAC1 from it and then translocates it to the cytoplasm.³⁹ In addition to HDAC1/2, ICP0 is also reported to counteract class II HDACs, implying multiple roles for ICP0 against diverse HDAC functions.⁴³ The Us3 gene products Us3 and Us3.5 kinase (hereafter, referred to as Us3-PKs) are also involved in the cytoplasmic distribution of HDAC1/2.^{44,45} In contrast to ICP0, Us3-PKs are not essential for viral replication in cultured cells, but are more critical for viral replication and neurotoxicity in mice. Us3-PKs show kinase activity against viral and cellular proteins and counteract apoptotic induction and innate immune responses led by type I interferons (hereafter referred to as IFN-I). To this end, HDAC1 and 2 also are subject to PTM from Us3-PKs.^{46–48} Although it has not been reported whether this phosphorylation of HDACs affects enzymatic activities, Us3-PKs overexpression changes HDAC1 localization into the cytoplasm.⁴⁴ Thus, ICP0 and Us3 independently interact with HDAC1/2 to displace them from viral DNA during infection.³⁹

The roles of ICP0 and Us3 in counteracting HDAC1 and 2 functions may not be critical in viral replication in cells that are not able to sustain the viral establishment of latency (ie, non-neuronal cells) and that preferentially sustain the lytic cycle. Viral gene expression of ICP0-null HSV-1 was temporally accelerated in cultured cells pretreated with sodium butyrate (a pan-HDACi), but the expression of viral genes from wild-type HSV-1 was not.⁴⁹ It is thus uncertain whether HDACi enhances viral replication and yields in these cells. An HSV-1 with a mutation in the Us3 gene is unlikely to exhibit altered replication in cultured cells when pretreated with HDACi (sodium butyrate or HDACi VII).⁴⁷ In proliferating or non-neuronal cultured cells, HDACs may not play a dominant role in suppression of the viral lytic cycle. Prior to ICP0 and Us3 gene activation, the major tegument protein VP16 regulates IE gene expression, required for the viral lytic cycle. Upon intracellular penetration, VP16 forms a complex with and recruits other histone modifiers, such as lysine-specific demethylase 1 (LSD1) and HAT (CLOCK) in addition to Oct1 and HCF1, to the promoters of IE genes.^{50,51} Thus, the viral VP16 complex competes with chromatin silencing mediated by the cellular HDACs-repressor complex. On the other hand, HDACs-repressor complex may play a more important role in preventing viral reactivation from the latent state where VP16, ICP0, Us3, and other viral proteins are absent. Experiments using *in vitro* and *in vivo* HSV-1 latency models indicate that latent HSV-1 in sensory neurons can be reactivated in response to sodium butyrate.^{52–54} The use of HDACi to reactivate human immunodeficiency virus (clinical trial in NCT01680094, NCT02513901, and NCT02336074) and Epstein–Barr virus (clinical trial in NCT01341834) from latency is used clinically as a means to flush out these hidden viruses from chronically infected patients.^{55,56} In summary, viral genes (ICP0 and Us3) associate with HDAC1/2 to suppress host repressor complexes and establish latency but HDACi may be more effective in cells that have already silenced the viral genome.

HDACi and OV in tumor

In cancer, hypoacetylation of histone tails and changing epigenetic signatures are hallmarks of cancer progression, metastasis, and therapeutic resistance, as well as tumor dormancy.^{57,58} OVs can be targets of genome silencing pathways that diminish oncolytic capability. Otsuki et al addressed the impact of oHSV1 in aggressive brain tumor type when treated with HDACi, mostly using VPA.⁵⁹ VPA is a widely used antiepileptic drug and is also used in migraine prophylaxis and bipolar disorders. The pharmacological actions of

VPA include multiple mechanisms to affect neuroprotection, but the finding that VPA can also act as a HDACi is relatively recent.^{59–61} Also, several clinical trials using VPA and vorinostat in high-grade glioblastoma patients are currently ongoing in combination with standard chemodrug temozolomide and/or radiation.⁶² When relatively high doses of VPA (5–30 mM) were administered to human glioma lines prior to oHSV1 infection, the cytotoxic effect of the virus was significantly more pronounced. This was also evidenced by using other pan-HDACi (trichostatin A [TSA], sodium butyrate, and APHA). VPA augmented oHSV1 replication in different glioma cells. This was correlated with higher expression of viral IE genes (ICP0 and ICP4) and late gene at early time points after infection. When VPA was administered to mice with an orthotopic xenograft glioma 1 day before oHSV1 intratumoral injection, there was a significant survival advantage over oHSV1-treated mice that were not treated with VPA.

This significant finding depended on the timing of VPA treatment. While treatment before viral infection (14 hours) significantly improved viral yields, concomitant treatment (8 hours) diminished this augmented OV effect.⁵⁹ Likely concomitant treatment caused a repression in viral DNA replication and reduction of late genes, whereas IE gene expression was not suppressed but enhanced. Since it is also known that prolonged treatment of HDACi causes cell arrest, pretreatment and cotreatment of HDACi may cause different changes in the cellular environment to affect oHSV1 infection. While VPA pretreatment could induce cell cycle arrest at G1 leading to efficient HSV-1 replication, the administration of HDACi along with the oHSV1 could dampen this benefit to support viral activity.⁶³ A similar negligible effect of TSA in viral replication was observed with oHSV1 infected glioma cells during TSA treatment.⁶⁴ In this instance, TSA treatment down-regulated cyclin D1 expression that is required for efficient HSV-1 replication. ICP0 facilitates the recruitment of cyclin D to viral DNA replication foci and stabilizes cyclin D proteins through degradation of Cdc34, an E2 Ub-conjugating enzyme.^{65,66} Cyclin D1 is also a critical driver of the G1-S phase transition and a tumorigenic factor for many cancers. Thus, despite little effect in viral replication, the synergistic cytotoxicity of TSA and oHSV1 was significant in glioma cells *in vitro* and *in vivo*.⁶⁴ In oral squamous cell carcinoma cells, TSA cotreatment led to up-regulation of oHSV1 replication at early time point (24 hours) and enhanced cytotoxicity of oHSV1.⁶⁷ This correlated with induction of cyclin-dependent kinase inhibitors p21, resulting in G0/G1 arrest in treated cells. In fact, the lack of HDAC1/2

can induce cell cycle arrest at the G1 phase through the up-regulation of cyclin-dependent kinase inhibitor proteins, p21 and p57, which are targets of Cullin 4-RING E3 ligases, an Ub E3 ligase complex.⁶⁸ CUL4B-containing Cullin 4-RING E3 ligases B and HDAC1/2 containing the Sin3A repressor complex are associated to deacetylase histone H3 and H4 on CDKN1A/p21 promoters, which can be blocked by TSA. Overall, HDACi can suppress tumor proliferation during oHSV1 infection. Whereas oHSV1 replication may be affected negatively or positively depending on the cellular context, HDACi can show enhanced cytotoxic effect with oHSV1 in glioma and squamous cell carcinoma.

The study by Otsuki et al also showed that HDACi can block type-I interferon pathway during oHSV1 infection in glioma cells.⁵⁹ IFN-I including IFN- α and IFN- β are crucial inflammatory cytokines for innate antiviral function in early phases of viral infection.⁶⁹ Extracellular IFN-Is bind to its receptors on the cellular surface and activates Janus tyrosine kinases followed by phosphorylation of STAT1 and STAT2 proteins.^{70–72} JAK/STAT pathways through IFN-I binding to its receptors stimulate transcription of genes containing IFN-stimulated gene response elements. While IFN- α is secreted from restricted cell types such as immune cells, IFN- β secretion occurs in nearly any types of cells in response to the stimulation. Administration of IFN- β protein is approved in the USA to treat multiple sclerosis and underway experimentally and clinically in glioma and other cancer therapy.^{73,74} In OV therapy, however, IFN treatment blocks the oncolytic effect and OV replication is greatly suppressed when IFN- β is added. Under this condition, VPA treatment improved oHSV1 oncolytic effects by neutralizing IFN-induced antiviral effects. At the molecular level, VPA suppressed up-regulation of STAT1 gene expression in response to IFN- β treatment. PKR and PML are involved in intrinsic antiviral functions against HSV-1. PKR is a kinase that phosphorylates translational initiation factor eIF-2 α to suppress viral protein synthesis and PML is one of the ND10 nuclear body proteins that associates with viral DNA early during lytic infection and represses viral replication. PKR and PML are ubiquitously expressed and their genes were up-regulated by IFN- β , whereas VPA suppressed them. The mechanism behind these findings implicates a non-canonical role of HDAC1/2 that functions as a transcriptional activator of a subset of genes that are induced by IFN-I.^{75–77} In addition to oHSV1, HDACi also inhibit IFN-induced antiviral responses that restrict oncolytic activities of VSV and vaccinia viruses in prostate cancer and melanoma cell lines, respectively.^{78,79} Thus, HDACi would be widely beneficial to

support OV replication in tumors where immune responses are intact and interfere with virus activity.

HDACi and immune cells in response to OV

HDACi may change intercellular antiviral immune responses. Natural killer (NK) cells are among the first responders against pathogens and they are recruited to pathogenic sites to attack virus and tumor cells. NK cells require expression of NK-cell specific recognition receptors on the recipient cells. NKG2D is one of the receptor molecules expressed on virus-infected cells and tumors. HDACi is known to induce immunosensitization of tumors by increasing NKG2D ligand expression and sensitize them to NK cytotoxicity that can be applied to cancer immunotherapy.⁸⁰ HSV-1 infection also increases susceptibility to NK lysis of the infected cells and natural cytotoxicity receptors NKp30 and NKp46 that are upregulated in response to HSV1 infection in gliomas.⁸¹ Therefore, early phases of the innate immune response can impact negatively on virotherapy and activation of both NK cells and macrophages in response to oHSV1 infection impedes viral replication and oncolysis. However, it is not clear if HDACi affects the cytotoxic properties and migration of NK cells in the setting of virotherapy of cancer. Thus, Alvarez-Breckenridge et al addressed the impact of HDACi against NK cells early during oHSV1 infection using a human glioma xenograft model.⁸² The injection of oHSV1 into the orthotopic glioma site of mice led to the infiltration of NK cells and IFN- γ product in that region, while VPA administration temporally suppressed them. When compared to cyclophosphamide that causes robust suppression of innate immune cells (macrophages, lymphocytes, and NK cells), VPA impaired the kinetics and abundance of recruited immune cells (NK and macrophage) after oHSV1 infection. At the cellular level, VPA attenuated IFN- γ production in NK cells in part through reduction in STAT5 phosphorylation and T-BET gene expression, a master regulator of IFN- γ gene expression.⁸³ In summary, HDACi can up-regulate NK ligand expression on tumor cell surface and suppress the inhibitory effect of NK cells against oHSV1 at the tumor site.⁸¹ Therefore, VPA pre-treatment may allow the oncolytic effect of oHSV1 in tumor bed to enhance NK-mediated tumor lysis at later time-point when the cytotoxic capability of NK cells recovers after clearance of VPA and viruses.

HDACi can assist OV-mediated immunotherapy. Multiple HDACs regulate the development process and maintain the lineage integrity of lymphocytes: hence, the utility of HDACi is recently expanding to include anti-inflammatory properties for autoimmune disease.^{26–28,84–87} Furthermore, the application

of HDACi in the immune system also boosts vaccinations to treat cancer and chronic infectious diseases as well.^{88–92} Thus, OV's are potential vaccine and immune-stimulatory agents to apply in tumor immunotherapy in this context.^{93,94} VSV is one of the oncolytic vaccine platforms that stimulates acute immunity along with immunogenic tumor cell death and boosts tumor-specific cytotoxic T lymphocytes when the virus encodes tumor-antigen to express in tumor. Thus, Bridle et al used HDACi to enhance this oncolytic vaccination strategy to target melanoma.⁹⁵ Oncolytic VSV encoding human dopachrome tautomerase gene (VSV-hDCT) was used to boost vaccination after priming with recombinant Ad expressing hDCT in naive mice bearing B16-F10 melanoma. When coadministered with MS-275, the VSV-reactive CD8+ response was rather attenuated, leading to delayed induction of antibody against the virus until the end of drug administration. However, prolonged VSV-induced lymphopenia was also observed by MS-275, associating with delayed reconstitution of peripheral CD4+, CD8+ T-cells, and B cells. As result, MS-275 increased tumor antigen (DCT)-specific CD8+ numbers with enhanced cytotoxic functionality in the context of TNF α and IFN- γ products at late time point while decreasing Foxp3+ Treg cell

number. Of interest, MS-275 coadministration suppressed this prime-boost induced autoimmune response against normal DCT+ melanocytes. This and other studies suggest that HDACi may suppress IFN-I products in diseases associated with autoimmune tissue damage and chronic infection, and thus can restore exhausted adaptive immunity.^{91,96–98}

Conclusion

There is an increasing body of evidence that dysregulated epigenetic factors are associated with chronic diseases such as cancer, inflammation, and infection. Targeting HDACs with pharmacological inhibitors can influence a broad range of transcriptomes. Acetyl-modification of nucleosome components and transcriptional accessories changes the immune and inflammatory signaling cascades, as well as lymphocyte lineages. In oncolytic therapy, HDACi could create a favorable environment for therapeutic viruses and we summarized currently published studies in Table 1 including those where HDACi-enhanced OV effects against tumors. These viruses could then function as self-productive, anti-tumor agents in the tumor bed, and promote tumor immunogenicity. There have not been clinical trials using combinations of HDACi

Table 1 A summary of published effects of combined OV and HDACi

Pathway	HDACi	Virus	Tumor type	Anti-tumor	Enhanced OV activity	Reference
Intracellular innate immunity						
Blocking IFN-I stimulating antiviral response	VPA	HSV-I (rQNestin34.5)	Glioma	Yes	Yes	59
Blocking IFN-I stimulating antiviral response	TSA	Vaccinia (VVdd-GFP)	Melanoma	Yes	Yes	79
Blocking broad innate antiviral responses	MS-275, SAHA	VSV (VSV- Δ 51-GFP)	Prostate cancer	Yes	Yes	78
NF- κ B activation	TSA	HSV-I (R849)	SCC	Yes	Yes	67
	SAHA	VSV (not specified)	Prostate cancer	Yes	Yes	101
Intercellular Immunity						
Blocking NK cells activation	VPA	HSV-I (rQNestin34.5)	Glioma	Yes	Yes	82
Enhanced Immunogenic vaccination	MS-275	VSV (VSV-hDCT)	Melanoma	Yes	–	95
Cell cycle						
Cyclin D1 down-regulation	TSA	HSV-I (G47 Δ)	Glioma	Yes	No	64
p21-induced G1/S arrest	TSA	HSV-I (R849)	SCC	Yes	No	67
p21-induced G1/S arrest	VPA	Ad (Ad5-Fiber-IRES- GFP)	Various cancer types	Adverse	No	102
	VPA	Ad (OBP-301)	Lung cancer	Yes	No	103
	SAHA	Ad (Δ 24-luc)	Lung cancer	Yes	Yes (in vitro), No (in vivo)	104
Viral receptor						
CAR gene up-regulation	FR901228, VPA	Ad (OBP-301)	Lung	Yes	Yes (FR901228)/ No (VPA)	105,103
	VPA	Ad (Ad5-Fiber-IRES-GFP)	Various cancer types	Adverse	No	102

Abbreviations: Ad, adenovirus serotype; HDACi, histone deacetylase inhibitors; hDCT, human dopachrome tautomerase; HSV-I, herpes simplex virus type I; IFN-I, type I interferons; NK, Natural killer; SAHA, suberoylanilide hydroxamic acid; SCC, squamous cell carcinoma; TSA, trichostatin A; OV, oncolytic virus; VPA, valproic acid; VSV, vesicular stomatitis virus; NF- κ B, nuclear factor kappa B; GFP, green fluorescence protein; CAR, Coxsackie virus and adenovirus receptor; IRES, internal ribosome entry site.

and virotherapy. At this point, it seems that all cancer indications may be worthwhile exploring in a possible clinical trial setting.

Although various HDAC inhibitors can be combined experimentally, there are mixed reports of therapeutic efficacy relative to their synergistic effects. This is because HDAC functions are complex, divergent, and redundant among classes. Therefore, the success of combined therapy may require careful evaluation of individual HDACs using a specific HDACi, whereas most studies used pan-HDACi (eg, VPA, TSA, and SAHA) in experimental OV virotherapy. Furthermore, the determination of the ideal time of HDACi administration is necessary for optimal OV therapy synergy with the host immune responses. Since host epigenetic roles in cancers and immune cell lineages are broad and also not restricted in lysine acetylation states, inhibition of other epigenetic modifiers (DNA/protein methylases and non-coding RNAs) are also needed to be evaluated in the setting of OV combination in the future.^{1,99,100}

Acknowledgment

This work was supported by NCI P01CA163205.

Disclosure

The authors report no conflicts of interest in this work.

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