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Advancements in the delivery of epigenetic drugs

Samantha A. Cramer^{1,2}, Isaac M. Adjei¹, and Vinod Labhasetwar^{1,3,*}

¹Department of Biomedical Engineering, Lerner Research Institute, Cleveland Clinic, Cleveland, OH 44195

²Department of Chemical and Biomedical Engineering, Cleveland State University, Cleveland, OH 44115

³Taussig Cancer Institute, Cleveland Clinic, Cleveland, OH 44195

Abstract

Introduction—Advancements in epigenetic treatments are not only coming from new drugs but from modifications or encapsulation of the existing drugs into different formulations leading to greater stability and enhanced delivery to the target site. The epigenome is highly regulated and complex; therefore it is important that off-target effects of epigenetic drugs be minimized. The step from *in vitro* to *in vivo* treatment of these drugs often requires development of a method of effective delivery for clinical translation.

Areas covered—This review covers epigenetic mechanisms such as DNA methylation, chromatin remodeling and small RNA mediated gene regulation. There is a section in the review with examples of diseases where epigenetic alterations lead to impaired pathways, with an emphasis on cancer. Epigenetic drugs, their targets and clinical status are presented. Advantages of using a delivery method for epigenetic drugs as well as examples of current advancements and challenges are also discussed.

Expert opinion—Epigenetic drugs have the potential to be very effective therapy against a number of diseases, especially cancers and neurological disorders. As with many chemotherapeutics, undesired side effects need to be minimized. Finding a suitable delivery method means reducing side effects and achieving a higher therapeutic index. Each drug may require a unique delivery method exploiting the drug's chemistry or other physical characteristic requiring interdisciplinary participation and would benefit from a better understanding of the mechanisms of action.

Keywords

Epigenetics; Epidrugs; Drug delivery; Cancer; Nanoparticles; Prodrugs; RNA

1. Introduction

Epigenetic mechanisms are known to play a critical role in cancer initiation and development and are implicated in other diseases as well such as multiple sclerosis,

*Author for correspondence: Vinod Labhasetwar, Ph.D., Department of Biomedical Engineering/ND20, Cleveland Clinic, 9500 Euclid Avenue, Cleveland, OH 44195, Tel: 216/445-9364, Fax 216/444-9198, labhasv@ccf.org.

neurological disorders, asthma, and depression [1-4]. By exploiting epigenetic regulation, the expression of genes can be controlled or regulated without the risks associated with genetic changes in the DNA sequence itself. There are several epigenetic drugs approved or in clinic trials including 5-azacytidine, 5-aza-2'-deoxycytidine (decitabine), Suberoylanilide hydroxamic acid (SAHA), valproic acid, and entinostat [5-14].

Several methods of enhancing delivery include a delivery system such as different nanocarriers, altering the chemical structure of the drug (pro-drugs), and administration in combination with another drug are aimed at increasing effectiveness of these drugs [15]. Choosing a form of delivery vector includes taking into account properties of the drug or agent to be encapsulated such as the polarity of the drug (some popular epigenetic drugs such as decitabine and 5-azacytidine are hydrophilic), molecular weight of the drug, pharmacokinetics and biodistribution, and effective dose and time of treatment needed at the diseased site [16-18]. Other factors come into play as well such as the target area's pH, mode of cellular uptake, and the desired intracellular release location [18]. Surface modifications can be used to alter or enhance the intracellular interactions or interactions occurring prior to cellular uptake possibly with the blood when given intravenously or gut tissue and fluids when given orally [19].

This review focuses on epigenetic targets where there are drugs in the clinical or preclinical phase targeting their action (i.e. DNMT and HDAC) and the methods of delivering these drugs including delivery systems, altering the chemical structure of the drugs themselves (prodrugs) and using combination therapies to increase efficacy.

2. Epigenetics

Epigenetics is the study of changes in gene activity that do not involve changes in the nucleotide sequence of the gene. Instead of changing nucleotide sequence, gene expression is controlled by modifying nucleotide residues, chromatin structure or degrading the translated messenger ribonucleic acid (RNA). Three distinct mechanisms for epigenetic regulation have been identified. These include DNA methylation, chromatin remodeling (histone modification), small RNA (siRNA) and micro RNA (miRNA) mediated gene regulation [20-22].

Methylation of cytosine residues on DNA is a reversible or irreversible epigenetic mark that regulates several biological processes including gene silencing, imprinting and X-chromosome inactivation [23]. In mammals, DNA methylation is mediated by DNA methyl transferases (DNMT) which add methyl groups to nucleotides in CG context mostly as clusters near gene promoters (termed CpG islands) where they control the expression of the genes they are associated with [24, 25].

The remodeling of chromatin involves post-translational modifications to nucleosomes, particularly to the histone core proteins which allow or restrict access of transcription machinery to the bound DNA. Post-translational modifications of histones generally include acetylation and methylation although modifications such as ubiquitination can also occur [26]. In this section, the most common histone modifications, acetylation and methylation will be discussed in further detail.

Acetylation of histones by histone acetyltransferases (HAT) using acetyl co-A as acetyl donor decreases the positive charge on histones which decreases their affinity to DNA double helix in the nucleosome. This action makes the DNA accessible to transcription factors for gene expression. The added acetyl groups can be removed by histone deacetylases (HDAC) which leads to transcription repression due to chromatin compaction [27].

Methylation of histones occurs on the basic amino acids arginine, lysine and histidine with each experiencing different degrees of methylation [21, 28]. Methyl transferases, which use S-adenosyl methionine as methyl donor, methylate histones to generate methylation profiles that can encourage or prevent gene expression. The methylases are substrate specific and some are sensitive to the degree of methylation [23-25]. Histone methylation can be stable or removed by demethylases.

Unlike the other epigenetic mechanisms that act on just the chromatin, small RNA mediated gene regulation occurs in both the nucleus and cytoplasm. The most common example of RNA mediated gene regulation in the nucleus is the X-inactive specific transcript (Xist) mediated X-chromosome inactivation [29] which results in the random inactivation of one X-chromosome in each cell in females [30]. In the cytoplasm, gene regulation is through the activity of miRNA which are small non-coding single stranded RNAs about 20 to 25 nucleotides in length that regulate a wide range of cellular processes including cellular proliferation, cell death, and developmental processes [31, 32]. Only a small section of the human genome encodes for miRNAs but they may regulate as many as 60% of human genes due to the nature of miRNA interaction with RNA, which enable each miRNA to have several potential targets [33].

3. Diseases with epigenetic mechanisms

The different epigenetic mechanisms modulate high order DNA structure and ultimately gene expression. Stable changes in the epigenetic status of a gene or its associated histones without genetic changes or mutations can lead to diseases. In most diseases however, etiology of a disease is a combination of both epigenetic changes and genetic mutations [34]. Several diseases are implicated where malfunction in epigenetic mechanisms occur (Table 1). In this review we discuss in more detail the role of epigenetic mechanisms in cancer.

3.1 Cancer

Changes in different epigenetic mechanisms have been associated with cancer. Of these, the methylation status of cancer is the most studied. In general, there is an alteration in the methylation status of the promoters of several tumor suppressor genes [35, 36]. Hypermethylation associated with cancer development occurs in the CpG islands of tumor suppressor genes such as retinoblastoma tumor-suppressor gene (*Rb*) and breast cancer type 1 susceptibility protein (*BRCA1*), and in genes involved in regulation of the cell cycle, DNA repair and induction of apoptosis. On the other hand, genes that prevent apoptosis and enhance cell survival are hypomethylated [37, 38]. Aberrant changes in activities of HAT and HDAC can lead to cancer.

There are widespread changes in the expression of miRNA in cancer compared to adjacent normal tissues with miRNA expression profiling of tumors being used in the diagnosis, staging and monitoring of progression and response to treatments [39, 40]. MicroRNAs that have been shown to be down regulated in cancer include the *let-7* family of miRNAs which target *RAS* oncogenes [41, 42]. The reduced expression of these miRNAs results in increased tumor growth and decreased survival [43, 44].

4. Drugs with epigenetic mechanisms

Several classes of drugs that target epigenetic mechanisms are in preclinical or clinical trials (Table 2). Most of these drugs target methylation (inhibit DNMT) or deacetylation (inhibit HDAC). Listed below are the activities of some of the common epigenetic inhibitors.

4.1 Demethylating agents

5-azacytidine (also known as azacytidine) is an analog to the nucleoside cytidine. Azacytidine is incorporated into DNA and RNA and reversibly inhibits DNA methyl transferase. Because azacytidine is incorporated into both DNA and RNA, it inhibits synthesis of DNA, RNA and proteins [45]. 5-Aza-2'-deoxycytidine (decitabine, DAC)) is another analog of cytidine. Decitabine is incorporated only into DNA, unlike 5-azacytidine which is incorporated into both DNA and RNA. Decitabine incorporated into DNA irreversibly bind to DNMTs leading to rapid depletion of the enzyme and subsequent hypomethylation of DNA [46].

Despite the success of epigenetic drugs such as 5-Aza-2'-deoxycytidine in clinical trials against hematologic malignancies and its antiproliferative effect *in vitro*, its efficacy in solid tumors has been disappointing [47]. This is due to the difference in the pharmacokinetics and pharmacodynamics of 5-Aza-2'-deoxycytidine in between solid and hematologic malignancies. 5-Aza-2'-deoxycytidine has a very high clearance rate *in vivo* (the half-life is 10-35 min) [48]. In addition, 5-Aza-2'-deoxycytidine rapidly degrades in acidic conditions, which most solid tumors exhibit [48, 49]. The discrepancy between the *in vitro* efficacy of epigenetic drugs, including 5-Aza-2'-deoxycytidine, and their *in vivo* efficacy in solid tumors has been attributed to the slowly dividing nature of cancer cells *in vivo* versus *in vitro* [50, 51]. Since 5-Aza-2'-deoxycytidine is S-phase specific, its activity in the tumor, when injected intravenously, may not last long enough for all the tumor cells to pass through the S-phase and thus for 5-Aza-2'-deoxycytidine to be effective. It is estimated that it takes approximately 5-15 days, depending on the tumor type and growth rate, for all the cancer cells in solid tumors to pass through S-phase [51-53]. Since 5-Aza-2'-deoxycytidine targets rapidly dividing cells such as bone marrow stem cells, its repeated high dosing to achieve therapeutic levels in solid tumors was found to have serious side effects, such as chronic myelosuppression and leukopenia [50, 54]. The toxicity of 5-Aza-2'-deoxycytidine at higher doses is due to its DNA-damaging activity rather than DNA-demethylation activity [55].

4.2 HDAC inhibitors

Inhibitors of HDAC arrest growth and induce differentiation of cancer cells. These inhibitors have also been reported to induce apoptosis in cancer cells by changing the expression of

genes associated with this process. Common HDAC inhibitors include valproic acid and vorinostat. HDAC inhibitors alter the expression profiles of 2-5% of genes [27, 56, 57].

Valproic acid is an 8-carbon fatty acid that has previously been used in treating epilepsy and bipolar disorder. Valproic acid inhibits HDAC at physiological concentrations used for treating these neurological conditions. As an inhibitor of HDAC, valproic acid acts as a competitive inhibitor to acetyl groups on histone N-terminal tails by binding to the catalytic site of the enzyme [58].

Vorinostat, also known as suberoylanilide hydroxamic acid (SAHA), is an inhibitor of class I and II HDACs [59]. SAHA inhibits cell growth and is known to be better tolerated by normal cells than cancer cells [60]. As a chelator, SAHA acts by binding to the co-factor zinc at the active site of the enzyme making it unavailable for catalysis [61]. Entinostat, also known as MS-275 or SNDX-275, is a synthetic benzamide derivative that inhibits class I HDACs [62, 63]. It is being evaluated in phase I/II clinical trials for Hodgkin's lymphoma and phase III clinical trials for metastatic lung cancer [13, 14]. Entinostat has also shown promise as a pretreatment that could be used to resensitize resistant cells to chemotherapies [64].

5. Delivery of epigenetic drugs

A method of delivering these epigenetic drugs is often necessary due to fast degradation by enzymes *in vivo*. For instance, decitabine is rapidly degraded by cytidine deaminase which is present in high amount in tissues like liver, gut, etc [48, 65]. In addition, since it does not bind to proteins, its excretion is quite rapid, thus requiring drug to be continuously administered as an infusion [66]. Because of this, systemic drug levels drops rapidly; reaching to almost undetectable levels within 30 min once the infusion is stopped. Additional advantages of nanoparticle-based drug delivery systems may include the ability to target cancer (active, passive, and enhanced permeability and retention [EPR] effect), greater drug stability and lower drug concentrations during administration. All of these advantages have been found to result in lower systemic toxicity and a higher therapeutic index [15, 67, 68]. Targeting has been exploited in cancer therapies and can be passive or active. Passive targeting takes advantage of the EPR effect. This property of the blood vessel walls is caused by abnormal angiogenesis around a primary tumor. The abnormal and rapid formation of the vessels causes them to be 'leaky'. Nanoparticles, liposomes, and other delivery vectors take advantage of this property and the poor lymphatic drainage characteristic of tumors to achieve greater accumulation in tumor tissue than elsewhere in the body [69, 70]. Active targeting seeks to exploit an extracellular or intracellular mechanism or the increased acidity of a tumor environment (due to increased levels of lactic acid as a product of glycolysis) [15]. Certain proteins or receptors are overexpressed in cancer cells and these can be exploited by conjugating a ligand to the nanoparticle or other delivery vector [71]. Cellular uptake of the delivery system could be by several receptor-mediated endocytotic pathways including clathrin-dependent, clathrin-independent macropinocytosis, and caveole-mediated [15, 18]. The mechanism of uptake will depend on the size, shape, and surface characteristics such as charge and hydrophobicity of the vector [72].

There are several nanocarriers for drug delivery including polymeric nanoparticles, liposomes, dendrimers, nanogels, and biological vectors (Fig 1). Other ways to enhance the delivery of epigenetic drugs are prodrugs and using a drug combination for synergistic effect [15].

5.1 Nanoparticles

Nanoparticles (NPs) for drug delivery applications are usually less than 300 nm in diameter [68]. They can be prepared using natural or synthetic polymers. Polymers commonly used in NP formulations include Polyethyleneglycol (PEG), Polylactic acid (PLA), Polyglycolic acid (PGA), Poly(lactic-co-glycolic acid) (PLGA), dextran, and chitosan. NPs are often broken down into monomers or small molecular weight fragments which are removed metabolically through preexisting biological pathways or excreted. For example, PLGA is broken down into lactic acid and glycolic acid. Lactic acid goes through the tricarboxylic acid cycle and is eliminated as carbon dioxide and water [73]. Glycolic acid can be excreted through the kidney or may go through the tricarboxylic acid cycle and be eliminated as carbon dioxide and water [74]. By altering the formulation and manufacturing parameters, tight control is possible over the properties influencing degradation and release kinetics such as molecular weight, composition (lactide to glycolide ratio), hydrophobicity, and crystallinity. The release kinetics may be a first-order release or a triggered release [68]. NPs can be prone to phagocytosis resulting in decreased circulation time and decreased tumor accumulation [75]. Surface modifications (commonly PEG) and/or targeting ligands can be engineered onto the nanoparticles to increase cellular uptake, blood circulation half-life, and impact biodistribution [68].

5.2 Liposomes

Liposomes are self-assembling bilayers of lipids with an aqueous core. Their characteristics such as surface charge, functionality, and size can be controlled by altering the type of lipid or the ratio of the lipids [68]. They are often surface-modified with polyethylene glycol (PEG) to avoid their uptake and clearance by the reticuloendothelial system (RES) and to improve stability *in vivo* [76]. One advantage of liposomes is that they can be used to deliver both hydrophilic and hydrophobic drugs. The hydrophilic drugs get encapsulated in the inner space of the liposomes, while the hydrophobic drugs are enclosed between the two layers of lipids. The loading for liposomes tends to be less than that of other delivery methods. They also have shown instability in the bloodstream and have a rapid burst release. LiPlasome Pharma has developed non-targeted liposomes with lipids that can be degraded by phospholipase A2 (PLA2) which is up-regulated in the tumor environment. There has also been research into using amphoteric liposomes to deliver nucleic acids [68]. This could include delivering nucleic acids such as RNA sequences which work through epigenetic mechanisms. A novel formulation involving enriched liposomal carriers with encapsulated anacardic acid in the liposomal bilayer with a vitamin C gradient, loaded with mitoxantrone has been developed [77]. The epigenetic agent is the anacardic acid which has been shown to have anticancer properties. This novel formulation has multiple facets. The cytotoxicity of mitoxantrone was shown to be enhanced by the anacardic acid and the vitamin C when used on human melanoma cell lines A375 and Hs294T. However, these same two compounds –

anacardic acid and vitamin C – were cytoprotective when used on a normal human fibroblast line [77].

5.3 Dendrimers

Dendrimers are globular macromolecules, about 5-10 nm across. By controlling their properties, the polymer chains can be designed to be degraded for tailored drug release and lengthened circulation time. A dendrimer is built around a core of functional monomers and layers of multifunctional monomeric units are added outwards in a stepwise fashion [78]. Amphiphilic dendrimers can self-assemble to form micelles with hydrophilic surface groups. Delivery of many therapeutics, imaging agents, and targeting agents is possible by using different functionalized monomers [68]. Dendrimers can aid in the delivery of drugs, diagnostic agents, and targeting molecules [68]. CALAA-01, a targeted transferrin-cyclodextrin-siRNA nanoparticle, targets the M2 subunit of ribonucleotide reductase. This product may be effective against solid tumors and is now in phase 1 clinical trials [79]. More studies could be done to elucidate dendrimer biodistribution so that tissue localization can be predicted and improved. The steric hinderance associated with the structure of dendrimers could be a disadvantage [80]. But, it also provides a mechanism for a cascade release, which could be useful in some circumstances such as sequential treatment with an epigenetic drug followed by a chemotherapeutic.

5.4 Nanogels

Nanogels can be formulated so that they release their encapsulated drug under specific conditions or a stimulus such as temperature, pH, magnetic fields, or biomolecule recognition [81]. Vijayaraghavalu and Labhasetwar in their study developed N-isopropylacrylamide (NIPAM)-based biodegradable nanogels which can be loaded with epigenetic drugs (Fig 2). It is known that DNA Methyltransferase 1 (DNMT1) promotes DNA methylation to maintain cancer drug resistance. Decitabine is a potent hypomethylating agent, but its effect is transient because of its short half-life *in vivo* (10-35 min), and accelerated degradation in acidic conditions such as those found at a tumor site [48].

The efficacy of decitabine -loaded nanogels was shown in doxorubicin-resistant breast cancer cells (Fig 3), decitabine-resistant melanoma cells, and leukemia cells. The data demonstrated that decitabine in nanogel sustained DNMT1 depletion, prolonged cell arrest in the G2/M cell-cycle phase, and significantly enhanced antiproliferative effect of decitabine [82]. Decitabine-loaded nanogels potentially could be explored for treating solid-tumor because of better drug effect than with decitabine in solution which is cleared rapidly from the circulation following intravenous administration.

5.5 Biological

Biological methods of delivery include the “Nanocell” – a bacteria (~300nm) void of DNA thereby preventing mutations and replication. This has been loaded with molecules of different solubility and charge such as doxorubicin, paclitaxel, and siRNAs. There is a potential for an immune response due to the lipopolysaccharides [83].

5.6 Prodrugs

Prodrugs are drugs that have been modified so they are inactive until some event *in vivo* which activates them. Epigenetic drugs may be modified to increase their stability leading to slower degradation *in vivo* and/or the modification can better target the drug to the delivery site (intracellular or extracellular). This is often done by converting the parent drug to an ester, or manipulating the pH [15]. By converting to an ester, the variable expression of esterases *in vivo* can be used to cleave the prodrug activating the drug. The pH can be manipulated so that the drug is activated in the acidic conditions of tumor tissue [15]. In some cancers, folic acid receptors are overexpressed. In these cases, attaching a folic acid to the drug can enhance cellular uptake in the target cells [15, 84].

S110, a dinucleotide containing decitabine, has been shown to have decreased depletion by cytidine deaminase [65]. This is attributed to the specificity of cytidine deaminase. The S110 and decitabine were similar in their ability to inhibit DNA methylation, induce expression of the *p16* tumor suppressor gene, inhibit tumor cell growth, stability in aqueous solution, and cytotoxicity. The decreased deamination by cytidine deaminase could increase bioavailability. A lower dose would lead to fewer side effects [65].

CP-4200, an elaidic acid derivative of azacytidine has been used in mouse models to demonstrate better therapeutic efficacy than azacytidine alone. This is partly due to this chemical modification causing less dependence on nucleoside transporters [85]. Rather than encapsulating the drug in a liposome, a lipophilic element such as a cholesterol or fatty acid can be attached to the drug to aid in cellular uptake.

HDAC inhibitors – psammaplin A and FK228 (Depsipeptide) – are naturally occurring prodrugs. They work epigenetically as antitumor agents and their disulfide bond is cleaved by glutathione to form an active thiol [86, 87].

5.7 Drug combinations

Enhancing the delivery could also be done by using a combination of drugs. For instance, using tetrahydrouridine (THU) in combination with decitabine to treat β -thalassemia. In this case, THU acts as a competitive inhibitor of cytidine deaminase, which is normally responsible for the rapid metabolism of decitabine in the in the intestines and liver [88]. Low doses of decitabine have been shown to reactivate γ -globin gene (HBG) expression to treat β -thalassemia [89, 90]. Vijayaraghavalu et al have shown that sequential treatment of decitabine and doxorubicin is highly synergistic in inducing cytotoxic effect in drug resistant breast cancer (MCF-7/ADR) cells than simultaneous treatment. The combination treatment could provide an effective therapeutic strategy to overcome drug resistance that can potentially minimize doxorubicin-induced cardiotoxicity because lower doses of doxorubicin may be needed to achieve tumor regression [91].

UVI5008 is a novel epigenetic therapy that independently inhibits three epigenetic targets thereby becoming a combination therapy within itself. These targets are DNMTs, HDACs, and sirtuins (a class of NAD⁺-dependent HDACs) [92]. *In vivo* experiments have shown UVI5008 is cancer cell-selective. It is also not reliant on p53, Bcl-2 modifying factor, and/or TNF-related apoptosis-inducing ligand (TRAIL), which increases the range of cancers it

could be used to treat [92, 93]. Using one drug to inhibit multiple targets, rather than two or three, is beneficial as it increases ease of manufacturing and research into ADMET profiles, and conceivably reduces costs.

5.8 Other delivery methods of note

Belinostat, a histone deacetylase inhibitor, has been used in clinical trials both intravenously and through oral administration. Oral administration of a drug may be suitable to strengthen the effects on the target and to increase patient convenience. The results show that the oral route is well tolerated for high doses but further studies are necessary to determine dosing and scheduling [94].

To test the potency of an epigenetic drug there is now a cell-based assay called EPISSAY. This uses a non-malignant human breast cancer cell line – MCF10A – with a silenced triple-mutated bacterial nitroreductase (TMnfsB) fused with Red-fluorescent Protein (RFP). The RFP expression can be easily observed and is indicative of the expression of RFP-TMnfsB. The EPISSAY was used to test the potency of decitabine with and without PEGylated liposomal encapsulation. A 50% higher potency was observed when the decitabine was encapsulated in the PEGylated liposomes. The EPISSAY was also shown to be able to indicate the potency of HDAC inhibitors such as Vorinostat [95]. The promoter used in this system – a CMV promoter – had no major changes in methylation before and after treatment with decitabine. Though decitabine is usually solely referred to as a demethylating agent, there have been studies that show it can also work as an HDAC inhibitor [96-98].

6. Conclusions

This is an overview of the current epigenetic drugs and delivery methods being used in coordination. Delivery methods have been broadly defined as carriers such as nanoparticles, liposomes, dendrimers, and nanogels as well as prodrugs and drug combinations. Both epigenetic drugs and delivery methods are in relatively early stages of development and the research with them together is limited. Targeting and sustaining the effect of epigenetic drugs in combination with other drugs could be an effective strategy. There are many epigenetic targets and enzymes not mentioned in this review because no drug has been developed to target them [15]. Efforts are being made to identify these targets and to develop inhibitors/activators [99].

7. Expert Opinion

Nanoparticles, liposomes, prodrugs and other delivery methods hold the key to increasing the efficacy and lowering systemic toxicities of drugs, notably chemotherapeutics. Exploitation of target characteristics such as an acidic microenvironment and/or the altered expression of receptors on the cell surface to influence biodistribution thereby reducing off-target toxicities should. Sometimes the target is an overexpressed receptor. Ideally, in a clinical setting, the presence of these overexpressed receptors in a particular patient would be confirmed. A formulation that could successfully target an overexpressed receptor could theoretically treat metastasis as it could seek out the cancer without the aid of the EPR effect.

Successful targeting has to be paired with an increased retention time for improvement in the efficacy of the drug. For instance, a chemotherapeutic epigenetic drug needs to reach the cancer cells, be taken up by the cells, be released from their encapsulation (if there is one), and get to their intracellular target possibly by a method of endosomal escape.

The majority of epigenetic drugs are hydrophilic and therefore more difficult than other drugs to load into nanoparticles which are mostly hydrophobic. Thus it would require modification of the original method or developing totally new method depending on the original techniques to encapsulate epigenetic drugs. Other factor to consider while developing formulations is their stability in an aqueous environment.

A better understanding of epigenetic mechanisms would provide a larger base for finding novel epigenetic therapeutics. Currently, there are far more delivery methods than there are epigenetic drugs. Epigenetic drugs have potential, especially as a chemotherapy demonstrated by *in vitro* studies. Some epigenetic drugs are unstable and, like other chemotherapeutics, have a high systemic toxicity. This is why epigenetic drugs in particular could benefit from a delivery system.

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Article highlights

- Recent advances in epigenetic treatments are often coming not from new drugs but from modifications to these drugs or encapsulation of the drugs leading to enhanced delivery to the target site.
- Epigenetic drugs have the potential to be a powerful tool against a number of diseases, especially cancers. The epigenetic target may have different roles and expression throughout the body. This is why a delivery method which decreases side effects by targeting and therefore increases the therapeutic index is beneficial.
- Each drug may require a delivery method exploiting the drug's chemistry or another characteristic requiring interdisciplinary collaboration and would benefit from a better understanding of the mechanisms of action.

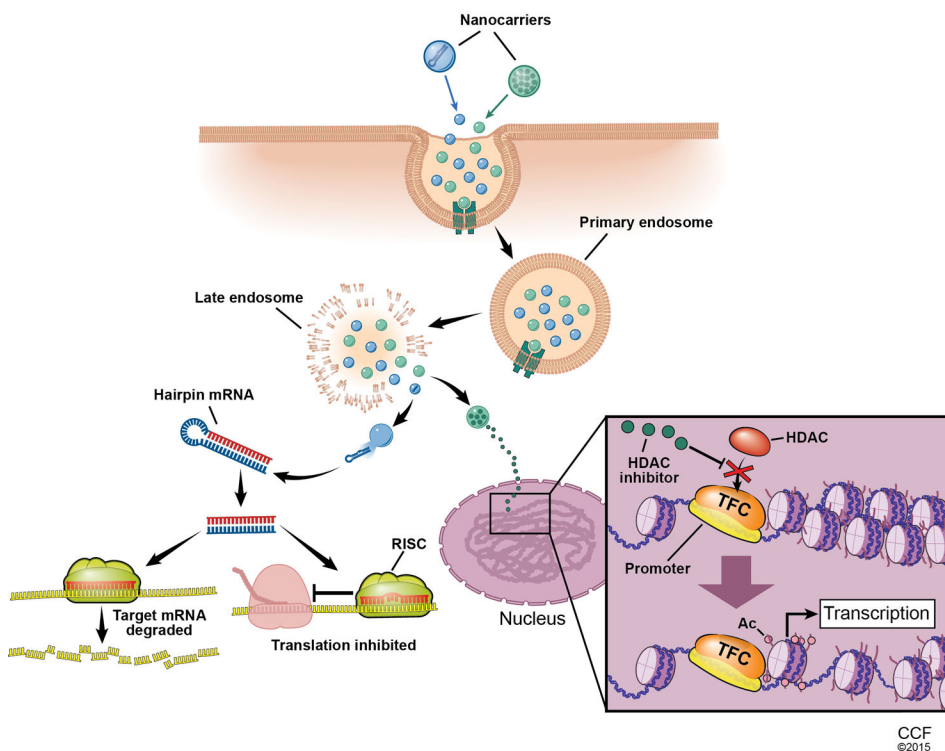


Fig. 1. Example of delivery of epigenetic therapeutics using nanoparticles

Schematic showing two possible mechanisms of epigenetic drugs being delivered via nanoparticles. The nanoparticles are taken into the cell by pinocytosis and enter an endosome. After endosomal escape two different paths are illustrated. The nanoparticle loaded with pre-miR releases its contents in the cytoplasm and is processed by Dicer. Next, it interacts with RISC to either degrade the target mRNA or inhibit its translation. The nanoparticle loaded with HDAC inhibitor drug enters the nucleus and releases the drug. There it blocks the action of HDAC causing an increase in transcription. *Abbreviations:* pre-miR, precursor miRNA; RISC, RNA-induced silencing complex; HDAC, histone deacetylase inhibitor; TFC, transcription factor complex.

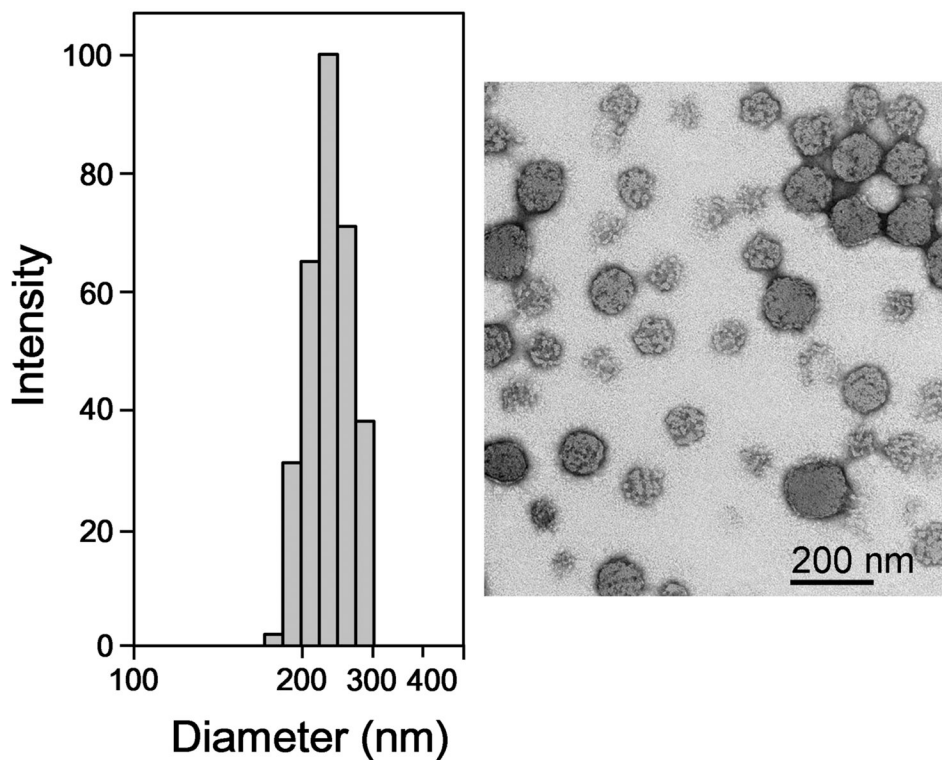


Fig. 2. Hydrodynamic diameter and particle size distribution, and transmission electron microscopic analysis of the decitabine-loaded nanogels formulation. Nanogels are synthesized using a combination of N-isopropylacrylamide (NIPAM), vinyl pyrrolidone (VP), and PEG–maleic anhydride (PEG-MA). NG-70 contains 70% NIPAM, 20% VP and 10% PEG-MA. The ratios of these three polymers were varied and nanogels formed were tested for their physical characteristics and drug loading. Composition of other nanogels is given in original publication. Figure reproduced with permission from Elsevier through RightsLink Copyright Clearance Center [82].

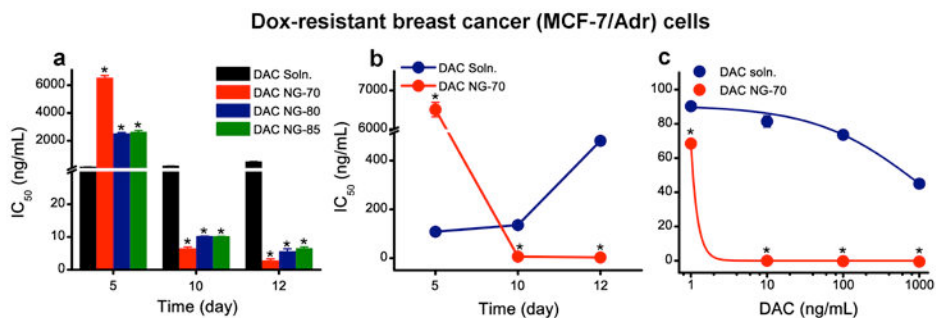


Fig. 3. Antiproliferative effect of decitabine-loaded nanogel in doxorubicin-resistant (MCF-7/Adr) breast cancer cells. **a)** Comparison of IC₅₀ of decitabine in solution vs. in different nanogel formulations over time. Cells treated with decitabine-nanogel formulations demonstrated sustained antiproliferative effect compared with cells treated with decitabine in solution. Efficacy of DAC nanogel depends on the nanogel composition, with NG-70 showing a more sustained antiproliferative effect than other formulations of nanogels. **b)** Comparison of IC₅₀ of decitabine in solution vs. decitabine loaded in nanogel (NG-70), depicting the transient antiproliferative effect of decitabine in solution vs. the sustained effect with DAC nanogel. **c)** Dose-response curves showing the difference in efficacy of decitabine in solution and decitabine in nanogel (NG-70) at 12 d post treatment. Data are expressed as mean \pm s.e.m, $n = 6$ * $p < 0.0005$, # $p < 0.005$ DAC solution vs. DAC nanogel. Figure reproduced with permission from Elsevier through RightsLink Copyright Clearance Center [82].

Table 1

Epigenetic diseases and impaired pathways.*

Disease	Epigenetic alteration	Impaired pathway
Alpha-Thalassemia X-Linked Intellectual Disability (ATRX) Syndrome	Hypomethylation	Chromatin and transcriptional deregulation [100].
Immunodeficiency, centromere instability and Facial anomalies (ICF) syndrome	Hypomethylation	Loss of DNMT3b activity at multiple immunological and developmental genes [101].
Multiple sclerosis	Hypermethylation	Silencing of <i>FOXP3</i> [2].
	Hypomethylation	Increased expression of PAD2 alters myelin processing [3].
Asthma	Histone acetylation	Decreased inhibition of NF-κB via histone acetylation [4].
Depression	Hypermethylation	Decreased expression of glucocorticoids[1].
Beckwith–Wiedemann syndrome	Hypomethylation	Increased activity of <i>KCNQ1OT1</i> , a long non-translated RNA [102].

* Examples of some diseases where an epigenetic malfunction is implicated. *Abbreviations:* DNMT3b, DNA-methyltransferase 3 beta; FOXP3, scurf; PAD2, peptidylarginine deiminase 2; NF-κB, nuclear factor kappa-light-chain-enhancer of activated B cells.

Table 2

Epigenetic drugs, target, and clinical status.*

Drug class	Example	Target	Clinical status
Aza-nucleosides	5-Azacytidine	DNMT	<p>a. Approved for myelodysplastic syndrome (MDS) [103].</p> <p>b. Relapsed acute myeloid leukemia (AML) non-responsive to chemotherapy (off-label use)</p>
	5-Aza-2'-deoxycytidine (Decitabine)	DNMT	<p>a. Approved for MDS [104].</p> <p>b. AML – off-label use in elderly AML patients</p> <p>c. Phase III – AML [105].</p>
Hydroxamic acids	Suberoylanilide hydroxamic acid (SAHA)	HDAC	<p>a. Approved for cutaneous T-cell lymphoma [106].</p> <p>b. Phase I/II – Recurrent glioblastoma, metastatic gastric cancer [10, 11].</p> <p>c. Phase II – Metastatic melanoma [107].</p>
	Belinostat	HDAC	<p>a. Approved for refractive peripheral T-cell lymphoma [108].</p> <p>b. Phase I – Solid tumors or lymphoma [109].</p>
	Panobinostat	HDAC	<p>a. Phase II – Metastatic thyroid cancer [110].</p> <p>b. Phase III – Hodgkin's lymphoma [111].</p>
Depsiptides	Romidepsin	HDAC	<p>a. Approved for cutaneous T-cell lymphoma [112].</p> <p>b. Approved for peripheral T-cell lymphoma.</p> <p>c. Phase I/II – Recurrent high grade glioma [113].</p>
Short-chain fatty acids	Valproic acid	HDAC	<p>a. Phase II – Advanced thyroid cancers of follicular origin [114].</p> <p>b. Status epilepticus – off label use for refractory disease.</p>
Benzamide	Entinostat	HDAC	<p>a. Phase I/II – Hodgkin's lymphoma [13].</p> <p>b. Phase III – Metastatic lung cancer [14].</p>

* Drugs with epigenetic mechanisms including their drug class, target, and clinical status. *Abbreviations:* DNMT, DNA-methyltransferase; HDAC, histone deacetylase.