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Outer Membrane Vesicles (OMVs) for Vaccination and Targeted Drug Delivery

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Abstract

Extracellular vesicles (EVs) are cell membrane-derived compartments that spontaneously secrete from a wide range of cells and tissues. Extracellular vesicles have shown to be the carriers in delivering drugs and siRNA. Among extracellular vesicles, bacterial outer membrane vesicles (OMVs) recently have gained the interest in vaccine development and targeted drug delivery. In this review, we summarize the current discoveries of OMVs and their functions. In particular, we focus on the biogenesis of OMVs and their functions in bacterial virulence and pathogenesis. Furthermore, we discuss the applications of OMVs in vaccination and targeted drug delivery.

Graphical abstract



Keywords

OMVs; Secretion pathway; Vaccines; OMVs-based drug delivery

Introduction

Despite the advances in engineering of synthetic nanoparticles and their surface bioconjugation for targeted drug delivery(1–7), reductionist's bio-functionalization of nanoparticles still remains insufficient in replicating complex intercellular interactions present in nature, thus impossibly avoiding exposure of exogenous features of synthetic

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nanoparticles to immune systems(8,9). Thus, it is urgent to develop new drug delivery platforms that possess intercellular interaction motifs in nature.

Intercellular communication is critical to maintain the homeostasis in a biological system. Recent studies have shown that cells secrete nanosized membrane vesicles (so called extracellular vesicles (EVs)) to transport signaling cargos between cells in a long distance (10). Many cell types secrete EVs (11–14). There are two types of EVs: exosomes and microvesicles. EVs are spherical membrane structures that are similar with synthetic liposomes, but EVs contain many membrane proteins that facilitate intercellular interactions.

EVs were first discovered in 1983, and it was found that reticulocytes released multivesicular bodies to the extracellular space (15). The secretion of EVs is strongly correlated to the pathogenesis of many diseases (such as cancer), therefore EVs have become biomarkers in diagnostics and early cancer detection (16). Interestingly, the recent studies have shown that EVs could serve as novel drug carriers in targeted drug delivery. To translate EVs, it is needed to develop new approaches to solve several issues, such as heterogeneity of EVs in composition and size, low production yield, inefficient drug loading and unlikely scalability. Nitrogen cavitation is exploited to generate cell membrane-derived nanovesicles since the mechanical force produced by nitrogen cavitation rapidly disrupts cells, and subsequently the cell membrane forms nanoscale vesicles (17,18). Their size and membrane composition are similar to those of EVs, thus the nanoscale vesicles become new drug delivery platforms to treat vascular diseases (17,18).

Except EVs derived from eukaryotes, prokaryotes (a unicellular organism that lacks a membrane-bound nucleus) also secret extracellular vesicles (EVs). For example, grampositive and gram-negative bacteria (13,19–21) have been reported to shed EVs. Grampositive bacteria have a single lipid membrane surrounded by a cell wall comprised of a thick layer of peptidoglycan and lipoteichoic acid, which is anchored to the cell membrane by diacylglycerol (22). Since gram-positive bacteria don't have "outer membrane" as compared with gram-negative bacteria, gram-positive bacteria release OMVs from the inner membrane and the released membrane vesicles go through the cell wall and form so-called "OMVs". Gram-positive bacteria-derived EVs were firstly discovered from *Staphylococcus aureus* by mass spectrometry (21). The size of EVs from gram-positive bacteria was reported to be ~20-100 nm in diameter, which was similar to EVs derived from gram-negative bacteria also spontaneously release extracellular vesicles, so-called outer membrane vesicles (OMVs). In this review, we will focus on OMVs derived from gram-negative bacteria.

A gram-negative bacterium possesses a unique membrane structure comprised of two membrane layers (outer membrane and inner membrane), a peptidoglycan layer, and a periplasm (23). Membrane proteins and lipopolysaccharide (LPS) distribute on the outer membrane while phospholipids mainly exist in the inner membrane. Periplasm is an oxidizing environment. A thin and rigid peptidoglycan layer in the periplasm adheres outer and inner membranes via proteins, such as OmpA. Several studies (24,25) have shown that peptidoglycan and outer membrane proteins (such as OmpA) are immunogenic, so they can be used in vaccination.

OMVs are a spherical structure with the size of 20-250nm, and they are comprised of bacterial cytoplasmic components that liberate during bacterial proliferation. OMVs play a vital role in pathogenesis, quorum signaling, nutrient acquisition and horizontal gene transfer (26). For instance, OMVs serve as vehicles to transfer toxins or various enzymes in bacterial infections. OMVs were first discovered in a cell culture medium of *Escherichia coli (E.coli)*, and subsequently were identified by an electron microscope, showing a small spherical structure with a single layer of membrane (27). It is found that other gram-negative bacteria also release similar membrane vesicles, implying an important role of OMVs in biological processes and revolution. Furthermore, OMVs can serve as novel vehicles in immunization and delivering therapeutics in the therapies of cancer and other diseases.

The focus of this review is to discuss the biogenesis of OMVs, and their pathogenesis for bacterial infections. We first review the status on vaccination development and targeted drug delivery using OMVs. In the end, we will summarize the advances in OMVs and discuss the future directions.

Biogenesis of OMVs

It is likely that OMVs are reserved among species. The generation of OMVs is spontaneous without the requirement of ATP. During their formation, OMVs are packed with proteins, lipids and DNAs. There are several steps to generate OMVs. Initially, proteins in a gramnegative bacterial envelop are homogenously distributed, and the outer membrane is linked to peptidoglycan. To start the vesicular formation of outer membrane, the binding between outer membrane and peptidoglycan is lost because the protein linkers detach, therefore the outer membrane liberates from a bacterium to form a vesicle. It was reported that antibiotics or autolysins are able to disrupt peptidoglycan, resulting in the outer membrane release to form OMVs (28). The mechanism of OMVs secretion is categorized based on their compositions and contents (Figure 1). During the formation of OMVs, the linking protein between outer membrane and peptidoglycan could release from the membrane and is incorporated inside OMVs. For example, the studies showed that OmpA, a linking protein was incorporated in OMVs (29-32). In some cases, the location in a bacterial envelope is enriched with vesicular-formation proteins that induce the OMVs generation. In this model, proteins concentrate in the inner membrane. When the adherence of peptidoglycan-outer membrane is removed, the dense proteins may participate in bulging of the bacterial outer membrane, leading to the formation of OMVs (33,34).

It has been an interest to understand how and why gram-negative bacteria secrete OMVs, and whether their secretion is spontaneous or random. For example, under stress, bacteria release OMVs for survival (35). The biogenesis of OMVs is an important topic in the future research.

It is noted that OMVs are complicated and heterogeneous. Analysis on their compositions and protein expression is very important to determine their biofunctions. *A. baumannii* release OMVs as a mechanism for the horizontal gene transfer, whereby carbapenem resistance genes are delivered to the neighbors (36). Additionally, proteomics is exploited to identify protein features in OMVs. Several studies have been performed to address whether

OMVs contain virulence factors of their sources. VacA, a virulence factor, found in *Helicobacter pylori* OMVs (37), indicates that virulence factors were incorporated into outer membrane vesicles during secretion. The adhesion molecules on OMVs (such as B-type flagellin and pili machinery: (PilA,F,Q,V,Y1)) that can serve as the virulence factors, were also identified in *P. aeruginosa* (38). Moreover, OMVs also contain outer membrane enzymes (lipases, peptidases, and ribonucleases) that are involved in pathogenesis. They include lipases, PagL (the most abundant protein in OMVs) (39), LipA (40), EstA, peptidases AaaA, PepA, PasP, MucD CtpA, Lon, IcmP and the M23 metaloprotease LasA (41–43). After OMVs were purified via ultracentrifugations, proteomics analysis showed that OMVs lacked the components of inner cell membrane and cytoplasm, but they shared higher similarities with outer membrane, including porins (OprB,C,D,E,H,O,Q), OstA (resistance to organic solvents and antibiotics) and OsmE (associated with the cell envelop integrity) (44,45). However, several studies on OMVs using SDS-PAGE are not always identical to those of the outer membrane of bacteria.

As mentioned above in Figure 1, OMVs not only contain the proteins from their source but also concentrate specific proteins compared with their parent bacteria during their formation. Studies on enterotoxigenic *Escherichia coli* and *P. aeruginosa* (49,50) show that several proteins are enriched in OMVs, such as LT and aminopeptidase, resulting in the increase of OMVs uptake by epithelial cells.

Proteomics has been applied to analyze the differences between bacteria and their derived vesicles, and to identify the abundance of proteins in OMVs. However, the proteomics results are still complicated. It is possible that the culture media and bacterial growth rates influence the heterogeneity of OMVs, leading to the inconsistence on proteomic analysis results. Another possibility is that proteins and virulence factors might mutate during the formation of OMVs.

Functions of OMVs in bacterial virulence and pathogenesis

OMVs play several roles in physiology and pathogenesis including horizontal gene transfer, quorum sensing, nutrient digestion, toxin secretion, misfolded protein secretion and immunomodulation (Figure 2). Here we focus on the role of OMVs in bacteria virulence and pathogenesis via the secretion of toxins and virulence factors.

Several studies have shown that OMVs directly regulate interactions between bacteria and the host via delivering of entrapped toxins, non-toxins, or bacterial virulence into host cells (55). For example, DNA packed in OMVs derived from *Neisseria gonorrhoeae* and *Borrelia burgdorferi* mediates the transfer of bacterial virulence to host cells (56). In addition to DNA packing, *P. aeruginosa* OMVs (38) have been shown to contain FliC, OprF and OprH/OprG and host-bacterium interaction proteins (EstA, FlgE, FlgK, etc) which participate in *P. aeruginosa* pathogenesis. Particularly, it was reported that EstA, a bacterial virulence factor, can induce nitric oxide and pro-inflammatory cytokines in macrophages. FlgE/K are flagellar proteins that stimulate innate immunity through Toll-like receptor 5 and play a role in the biofilm formation, an important virulence mechanism of *P. aeruginosa* (57,58).

Another example (50) is that enterogenic and uropathogenic *E.coli* ((ETEC) and (UPEC)) released the heat-labile enterotoxin into OMVs, and the endotoxins were transferred to host cells via OMVs. The mechanism further revealed that endotoxins were likely the ligands that mediated the binding of OMVs to lipid rafts of host cells, thus leading to the uptake of OMVs. Besides toxins and virulence factors, non-toxins including a variety of enzymes and proteins, have been shown to be entrapped in OMVs and delivered to host cells, therefore, affecting bacterium-host cell interactions. For example, *P. aeruginosa* OMVs released proteins associated with proteolysis, ion transport, and ion binding, which dysregulated host cells (38). The similar study (59) on *Treponema denticola* OMVs shows that they contain the necessary adhesins and proteolytic arsenal for the adherence to and the damage of eukaryotic cells.

OMVs can also affect bacterium-host cell interactions by entrapping cellular components and directly binding and even destroying host bacterial factors. For example, OMVs produced by *H.pylori*, presented Lewis antigens on their surface and were able to induce the host immune system activation (60). In this case, the OMVs directly bound to anti-Lewis antibodies in serum to decrease the self-defense ability of host cells, therefore, playing a very important role in *H. pylori* pathogenesis.

In summary, OMVs play a vital role in pathogenesis of bacterial infections since they contain many toxins from their source, which activate the host defense system. OMVs could be also applied to vaccine development.

OMVs in Vaccination

Infection can cause host immune responses, but sometimes excessive responses may lead to tissue damage, resulting in the death. OMVs play a central role in transporting toxins and virulence factors to host cells, and this transport mediates the host immune response. OMVs could be exploited to train the immune system to combat pathogens if they can be administered to the host in a control manner. Therefore, OMVs are a promising candidate for vaccine development against bacterial infections.

Studies have shown that OMVs could interact with epithelium cells, therefore, inducing the host immune response. The proteomics results from *Campylobacter jejuni* OMVs revealed that they contained many periplasmic and outer membrane-associated proteins (61). Several molecules are important in survival and pathogenesis, including the cytolethal distending toxin (CDT). Thus, OMVs could be an important alternative for the coordinated delivery of *C. jejuni* proteins into host cells. This idea was further confirmed by showing that *C. jejuni* OMVs possessed cytotoxic activity and induced a host immune response in T84 intestinal epithelial cells (IECs) (62). Similar studies in OMVs derived from various mucosal pathogens also showed that OMVs can interact with epithelial cells, resulting in the production of cytokines and chemokines that activated the pro-inflammatory response. For example, OMVs from all strains of *P. aeruginosa* elicited IL-8 secretion from lung epithelial cells to contribute the inflammation response (63). Moreover, *P. aeruginosa*-derived OMVs were shown to induce pulmonary inflammation via increasing chemokines and cytokines in the mouse lungs and mouse alveolar macrophages in a rodent model. Interestingly, OMVs

could induce inflammatory responses as compared with that of live bacteria (64), indicating that OMVs have the similar ability as live bacteria to induce innate immunity.

Interestingly, several studies in *Helicobacter pylori* showed that OMVs adhered to the epithelium to cause gastritis, rather than bacteria (65). These OMVs have also been shown to carry CayA and localize in the vicinity of cell-cell contact, therefore they may have an influence on host gene transcriptions, leading to infections and development of cancer (66).

OMVs also interact with various types of immune cells. It is shown that OMVs interact with innate immune cells. A study showed that N. meningitidis OMVs can stimulate human neutrophils, resulting in the production of TNF- α and IL-1 β and upregulation of CXCL8, CCL3 and CCL4 (67). L. pneumophila OMVs can generate pro-inflammatory cytokines from macrophages (68). Furthermore, Helicobacter pylori OMVs proteins can induce human eosinophil degranulation (69). Antigen presenting cells (dendritic cells), as a key connection between the innate and adaptive immunity, can be activated by OMVs. For example, OMVs from Salmonella spp induced the expression of CD86 and MHC class II molecules on dendritic cells and the production of TNF-a and IL-12, and promoted the development of protective B cell and T cell response in vivo (70). Similarly, OMVs derived from E.coli showed the increased uptake by dendritic cells and induced IL-6, IL-1 β production and antibodies production in vivo (71). OMVs can also interact with other host cells including endothelial cells and platelets cells. OMVs derived from E. coli OMVs up-regulate the expression of endothelial intercellular adhesion molecule-1 (ICAM-1), E-selectin and vascular cell adhesion molecule-1, and enhance the leukocyte binding on human microvascular endothelial cells (72). Moreover, Gingivalis OMVs (73) can enhance the platelet aggregation. In summary, OMVs strongly interact with the host via the activation of the innate and adaptive immune responses, therefore OMVs are an excellent candidate in vaccination.

We have discussed that OMVs are able to interact with a variety of cells. The uptake of OMVs have been investigated (74) and several internalization pathways have been found such as micropinocytosis(75), clathrin-mediated endocytosis (76), non calthrin-mediated endocytosis (lipid raft) (77), and membrane fusion (78). It is needed to address what receptors mediate the internalization of OMVs.

Adjuvants are required in most vaccine formulations to enhance the immune responses (79). Due to the unique features of OMVs, early efforts were focused on utilizing bacterial OMVs as adjuvants that were covalently complexed to antigenic proteins. *E. coli*-derived OMVs have been combined with malarial proteins in the development of the intranasal vaccine (80). In this work, they confirmed that OMVs can serve as adjuvants which can promote the immune response comparable to the cholera toxin adjuvant. Moreover, the authors suggested that OMVs can be applied as safe adjuvants and replace cholera toxin adjuvant because the cholera toxin adjuvant has the high toxicity and is difficult to be used in human clinical trials.

The studies showed that OMVs could not only be combined with proteins but also with other components. The study showed that *meningitides* derived OMVs can be complexed

with *Shigella*-specific lipopolysaccharides (LPS) to provide the immunity against *Shigella* keratoconjuctivitis (81). Another study showed that when combined with inactivated respiratory syncytial virus (iRVS), *Neisseria meningitidis* OMVs enhanced the protective immunity (82). The main mechanism was elucidated that hexa-acylated Lipid A moiety in the native LPS acted as a stimulator for a TLR4 receptor, therefore activating the innate immunity.

OMVs have been also used as vaccines to prevent bacterial infections. For example, *E.coli*derived OMVs efficiently prevented bacterium-induced lethality and OMVs-induced systemic inflammatory response syndrome via Th1 and Th17 cell responses (71,72). In this work, they performed immunization with E. coli-derived OMVs in a rodent model and demonstrated that *E. coli*-derived OMVs had the high protective effect as shown in Figure 3A. Moreover, they proved that this protective effectiveness can last 42 days after immunization (Figure 3B). It is shown that this protection was dependent on the induction of innate and adaptive immunity, including the production of anti-OMVs specific antibodies and T cell activation (Figure 3C&D). They also demonstrated the passive protection of OMVs via adoptive transfer of serum and splenocytes (data not shown). As mentioned above that immunization of OMVs mainly relies on the activation of innate and adaptive immunity, they also proved that the key antigen presenting cells (dendritic cells), functioning in the presenting of antigens of OMVs to the adaptive immunity-related cells, were fully activated (data not shown therefore). The activation of dendritic cells resulted in the production of immune-modulating cytokines (Th1- and Th17-polarizing cytokines), therefore, inducing the activation of T cells to OMVs. These cytokines included IFN- γ and IL-17, the key cytokines produced by Th1 and Th17 cells (data not shown). To understand whether this protective effect was Th-1 and Th-17-dependent and how such IFN-yand IL-17-dependent Th1 and Th 17 cell responses elicited the enhanced vaccine efficacy, IFN-γ, IL-17 and IL-4 knockout mice were used. The results showed that the knockout mice failed the survival compared to the wild-type mice immunized with OMVs (Figure 3E). In summary, they provide a comprehensive and new perspective on the immunological detail regarding OMVs being used alone in vaccination.

Similar studies showed that OMVs can serve as vaccines against *Edwardsiellosis* because the cytokines and chemokines were significantly increased after administration of OMVs (83). Moreover, *B. pertussis*-derived OMVs combined with alum adjuvants provided the protection from pertussis in a mouse model and this effect was comparable to the whole-cell formulation of vaccines (84). *B. parapertussis* derived OMVs also have been shown to have the cross-protection effect against both pertussis and parapertussis (85).

With the promising potential of OMVs in vaccination, OMVs-based vaccines have been tested in clinical trials (86). The *Meningitis type B* (MenB)-based vaccine has gained much attention. For several serogroups, conjugated vaccines consisting of capsular polysaccharide coupled to a carrier protein have already been in the market. *Meningitis type B* OMVs have been shown to have the efficacy range in 83%-85% (86,87) and have passed the phase I, phase II and multiple clinical studies (88–91). For instance, in clinical research, they showed that a four-dose schedule (three primary doses and one booster dose) for infants and a two-dose schedule for adolescents of the multi-component Men B vaccine (4CMenB) provided

the good result. These vaccines contained three surface-exposed recombinant proteins and New Zealand strain outer membrane vesicles (NZ OMVs) with PorA 1.4 antigenicity. The side effects are mainly associated with the injection site pain/tenderness and fever in infants. They are associated with the injection site pain, malaise and headache in adolescents. Thus, 4CMenB is conservatively estimated to provide 66-91% protection against B group of *meningococcal* strains worldwide (91). Moreover, Meningococcal-derived OMVs vaccines made of inactivation of lpxL1 gene, were also in the phase I trial (92,93).

With the promising perspectives of OMVs in vaccination, the major concern lies in the stability of OMVs when being administered in vivo. The kinetic biodistribution of OMVs in vivo demonstrated that OMVs were detected at the peak 3 hours after administration, then gradually decreased in most organs within 24 hours post injection (94), indicating that OMVs were likely to be eliminated during circulation. To address this concern, a report demonstrated that gold nanoparticles coated with OMVs can dramatically increase the stability of OMVs, therefore, resulting in higher immune activation as compared with OMVs administration alone (95). In their work, they incorporated gold NPs (AuNPs) into OMVs and proved that the membrane coating can effectively enhance gold NP stability in biological buffers, while gold NP cores stabilized OMVs (Figure 4A). With the higher stability, AuNPs demonstrated better efficacy in inducing B cell and T cell activation via producing higher IgG titer and cytokines (Figure 4B&C). Collectively, the future studies should be focused on increasing the stability of OMVs via membrane coating strategy to enhance OMVs efficacy for vaccination development.

Isolation and purification of OMVs

Purity of OMVs is an essential factor for their applications. Therefore, it is needed to develop novel methods to isolate and purify OMVs.

OMVs are directly obtained from cell suspensions. The most common isolation technique includes several steps of centrifugation. The first step is to separate cell debris through a 0.22 or 0.45 μ m filter, followed by the low speed centrifugation (2000-1000× g). To further purify and concentrate OMVs, the centrifugation is combined with the tangential flow microfiltration, such as using a filter of 50-100kDa (96) or a combination of 0.2 and 0.1 μ m filters(97). Finally, OMVs are collected after ultracentrifugation at 50,000-200,000× g. The ultra-centrifugation step can also be combined with the density gradient (98). Ammonium sulfate precipitation sometimes is applied as an alternative method for concentration of OMVs (99). In contrast of centrifugation approaches, OMVs are also isolated via detergents, such as DOC (Deoxycholic acid) (100).

However, some contaminants should be cautious during isolation and purification of OMVs. Bacterial components, such as pili, flagella and soluble components, may be mixed with OMVs since they cannot be separated by centrifugation approaches (101).

Bioengineered OMVs in vaccination

OMVs can also be engineered for vaccine development via incorporating of heterologous antigens. These heterologous antigens can be presented with or without surface exposure,

attached to the vesicles or non-attached and directly produced by the bacterium or combined in a later production stage. The generation of heterogenous OMVs vaccines mainly includes the following approaches:

1) Recombinant OMVs based on ClyA fusion protein

Antigens can be presented on the surface of OMVs with the exposure to the exterior side of the vesicles via a variety of fusion proteins, such as ClyA fusion (102–104). Cytolysin A (ClyA) is a transmembrane protein (a molecular weight at 34kDa) which is enriched in outer membrane vesicles (105). Genetic modification of OMVs is mainly focused on the fusion of antigens to the C terminus of ClyA which results in production of ClyA-antigen fusion proteins associated with OMVs. Chen's group (104) was the first to prove the concept via fusing of antigens with ClyA protein and express antigens on the cell surface. Their study showed that green fluorescent protein can be infused with ClyA of *E. coli*. The in vivo study showed that this recombinant GFP-OMVs can induce a strong production of GFP-specific antibodies without any adjuvants. Inspired by this work, ClyA was fused with antigens to generate anti-bacterial vaccines. Huang's group (102) fused Omp22 antigens from *Acinetobacter baumannii* into *E. coli* DH5α-derived OMVs, as shown in Figure 5, which showed the high protection in a murine sepsis model.

In this work (102), CytolysinA (ClyA) was successfully infused into *E. coli* membrane to generate chimeric ClyA fusion proteins, therefore, it can be used for the future engineering of infused Omp22-OMVs. It was confirmed that Omp22 antigen located on the surface of OMVs. This OMV-based vaccine lessened bacterial burdens in various tissues, and antiserum isolated from the mice demonstrated the bactericidal activity. Recombinant OMVs have also shown promising in creating anti-viral vaccines. Rosenthal's group (103,106) showed that the probiotic *E.coli* Nissle 1917 strain can be engineered to generate ClyA-GFP OMVs which elicited anti-GFP total IgG titers. The IgG titers were equivalently comparable with that of GFP adjuvanted with alum when the OMVs were administrated into the mice. Furthermore, they proved the protection against influenza infection using OMVs derived from Nissle 1917 strain fused with antigen protein M2e4xHet. Their results showed that bioengineered OMVs could elicit higher anti-M2e IgG2a antibodies which could facilitate the clearance of infected cells. The survival study demonstrated that bioengineered OMVs can save all mice when the mice were challenged with a lethal dose of influenza.

2) Recombinant OMVs based on other carrier fusion protein

Similarly, various fusion proteins are incorporated in OMVs. The fHbp was infused to express the Borrelial surface-exposed lipoprotein OspA on *Neisseria meningitidis* OMVs surface. The results showed that OMVs could able to elicit antigen specific antibodies as compared with that with luminal expression of OspA inside the cells (107). PspA or Ply fragments (108), ESAT6, Ag85B fragments, Rv2660c (109) and MOMP fragments (110) were able to be fused into fHbp fusion for vaccination.

Antigens can also be expressed in the outer membrane of OMVs. *P. aeruginosa* A104R antigen can be fused into outer membrane of *E.coli*-derived OMVs via OprI fusion, and through this way they can protect against African swine fever (111). Ail antigen can also be

fused into *E.coli*-derived OMVs (112). Many studies showed that antigens can be expressed in lumen of OMVs mainly via OmpA fusion. Kesty's study in 2004 was the first proof to concept to demonstrate the fusing of GFP into *E.coli*-derived OMVs via Tat signal (112). FLAG tag protein was fused in OMVs lumen (113). Several antigens have been infused with OmpA protein in order to localize them in OMVs lumen to target group A/B Streptococcus disease (109) and Chlamydia (114).

It is interesting to observe that lumen-fused OMVs vaccines only elicited minor specific antibody production. For example, the fusion PspA in *Salmonella enteriaca* in OMVs lumen showed lower antibody production and decreased protection. OMVs without any antigen fusion or purified fusion antigen showed neither the antibody production nor protective effect (115). One of possibilities is that the expressed antigens may change their conformations. The study observed the higher antibody titer when antigens were in their native structure(109). Additionally, when expressed on the surface of OMVs, bio-engineered OMVs would have higher antibody titers (115). However, the mechanism still remains unclear. It is also not clear whether the non-specific antibody response is dependent on strains. Therefore, more research should be focused on the specificity of OMVs responses.

Currently, most vaccines are generated from acellular organisms and their subunits (116). These vaccines are safer than utilization of live-attenuated or whole inactivated organisms. The vaccines usually don't contain the whole cell antigens, therefore, lacking of a broad protection compared to a whole inactivated organism. Moreover, the vaccines are also needed to combine with adjuvants to increase the efficacy. OMVs are derived from the bacterial membrane that contains a wide range of antigens required for immunization, thus they are potential to become novel vaccines (19). Since OMVs are nanoscale and can be engineered, they will increase the response of antigen-presenting cells (117,118).

While OMVs demonstrate the potential as a new platform for vaccination, there are several barriers when they are used in clinic. The heterogeneity of OMVs might cause the issues on reproducibility when they are largely scaled up. OMVs secretion strongly depends on the bacterial growth condition, therefore the preparation consistency is not guaranteed. Current techniques cannot scale up the production of OMVs required in clinic. Therefore, it is necessary to develop new approaches to generate not only the high quantity but also high purity of OMVs.

OMVs in drug delivery

Synthetic nanomaterials, such as liposomes, polymers, and metal-based nanoparticles have been broadly studied as drug carriers (119), but the simple bio-conjugation of synthetic nanoparticles is not efficient to replicate intercellular interactions that facilitate nanoparticle trafficking and delivery. Cell membrane derived nanovesicles possess the features of intercellular interactions, thus they are potential to become novel drug delivery platforms.

Similarly, OMVs have been reported to demonstrate a spherical structure which is the size range in 20-250 nm in Figure 6. OMVs contain a broad range of proteins that are derived

from their parent cells and can carry diverse cargos, therefore, they can serve as a new platform in targeted drug delivery. (120).

Enzyme degradation in serum is a problem to effectively deliver, so the incorporation of enzymes in OMVs may resolve this problem. Walper's group proposed an approach that phosphotriesterase (PTE) (EC 3.1.8.1) from *Brevundimonas diminuta*, containing a binuclear Zn/Zn active site, was selectively packaged within the OMVs (121,122). In this work, they used OmpA (outer membrane protein A) as an anchor to link PTE as described in Figure 7A. Furthermore, they demonstrated that the PTE-loaded OMVs exhibit native-like enzyme kinetics without changing enzyme activity of PTE (Figure 7B).

Extracellular vesicles are comprised of the adhesion molecules on vesicle surface and can bind target cells via the ligand-receptor recognition (17,18,123). Similarly, OMVs could target the cells of interest through the genetic and surface modification of OMVs for targeted drug delivery.

A pioneer work developed OMVs using bioengineered bacteria (*E.coli*) with the fusion of an anti-HER2 affibody to ClyA resulting in OMVs with the affibody displayed on the surface (124,125). OMVs could target and kill cancer cells in a cell-specific manner by electroporation loading with a small interfering RNA (siRNA) therapeutic targeting kinesin spindle protein mRNA (Figure 8 A). Although the loading efficiency was not high (data not shown), the amount of siRNA in the Affi_{HER2}OMVs was still sufficient to exert cytotoxic effects against the HER2-positive tumor cells, because siRNA loaded Affi_{HER2}OMVs were found selectively accumulate in the tumor sites after administration (Figure 8 B). Further study showed that this siRNA delivery strategy exhibited higher ability in tumor growth inhibition due to a significant reduction in KSP protein levels as compared with the free siRNA and non-targeted OMV^{siRNA} group (Figure 8 C). This genetically engineered OMVs were low toxic, inflammatory and immunological, thus they are considered as a safe platform for cancer therapy.

To enhance their functions, OMVs have been modified using nanotechnologies. The cells were decorated with synthetic nanocarriers to deliver drugs in a more controlled way. Gold nanoshells are nanoparticles which have been shown to successfully treat tumors in mice with tumor remission rate over 90% via being designed and fabricated to allow for the viability of the monocytes/macrophages during recruitment into the tumor (126). Additionally, the study showed that utilizing *E.coli* OMVs as a drug carrier to coat Au nanoparticles can sufficiently induce the activation and maturation of dendritic cells in the lymph nodes of the vaccinated mice, And these recombinant nanovesicles-induced antibodies production were durable and of higher avidity than OMVs only (95).

Conclusions and Future Perspectives

In Figure 9, we have summarized the overview of current research of OMVs. We have discussed the mechanisms of OMVs formation, and addressed how OMVs have been utilized in biomedical applications. We have demonstrated two major applications: vaccines and targeted drug delivery platforms.

We have shown that OMVs bulge from bacterium outer membrane via three major mechanisms. During their formation, OMVs can entrap periplasmic proteins. Understanding the mechanism by which OMVs are formed is critical for the medical applications of OMVs, but their biogenesis remains unclear. For example, how and why OMVs are produced? Is the formation of OMVs spontaneous or regulated? If it is regulated, what signaling pathways are involved? If we have a complete picture about their biogenesis, we may resolve the issues on their low production and complex composition.

We believe that we need novel approaches to address these questions. It is shown that secretion of OMVs is a ubiquitous process and OMVs have widely diverse functions than it is currently appreciated. Utilizing proteomics may enable to quantitatively analyze the composition of OMVs and how the composition is associated with their biological functions. Proteomics generates the huge and complex data, so it is needed to develop mathematical approaches to analyze them for the understanding of the OMVs biofunctions. The biogenesis of OMVs is important to develop effective OMVs-based vaccines.

OMVs are exploited as a new drug delivery platform as they are in the nanoscale range. OMVs can be recognized by dendritic cells, thus they may activate the innate and adaptive immune responses. However, the mechanism of OMVs internalization by host cells remains unclear. OMVs have been shown to contain a variety of virulence factors, including LPS and virulent proteins. Therefore, the safety issues should be cautious when OMVs are utilized as drug delivery carriers since LPS could cause innate immune response. This immunotoxicity of OMVs will be an interesting topic in drug delivery applications.

In summary, we have demonstrated that OMVs may be a new drug delivery system used in vaccination and targeted drug delivery. While the research on OMVs is in the early stage, their unique nanosized structure and biofunctions may be a promising platform in nanomedicine.

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Figure 1. Biogenesis of OMVs

Step1: Gram-negative bacteria cell envelop. In this stage, envelop proteins are homogenously distributed. Outer membrane is linked with peptidoglycan. Step2: Vesiculation initiation. The linking between outer membrane and peptidoglycan is lost through the movement of linking proteins or breaking the connection of outer membrane with peptidoglycan directly. Model A, B and C demonstrate three ways for OMVs production. Model A indicates the basal OMV production. Model B refers to the OMV production with enriched periplasma cargos. Model C shows the formation OMVs is located at specific proteins on the outer surface, and the dense proteins could induce the additional budding of OMV from gram-negative bacteria cell envelop.

Outer membrane Peptidoglycan

Inner membrane

Quorum sensing

Misfolded proteins secretion



Horizontal DNA transfer



Toxins secretion

Figure 2. Biofunctions of OMVs

3 LPS

OMVs release from bacteria outer membrane layer and automatically entrap various cellular molecules. Biofunctions of OMVs are categorized and described as above, including horizontal DNA transformation, quorum sensing, toxin secretion, nutrient digestion and misfolded protein secretion.

Outer membrane proteins

Immunomodulation

Nutrients digestion

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Figure 3. Immunization with *Escherichia coli* outer membrane vesicles protects bacteria-Induced lethality via Th1 and Th17 cell responses

A. Survival rates of OMV- and sham-immunized mice challenged with *E. coli*; **B.** Survival rates of OMV- and sham immunized mice challenged with *E. coli* 42 d after immunization; **C.** Serum levels of OMV-reactive IgG; **D.** OMV-specific production of IFN-g, IL-17,IL-4, and IL-10 from splenic T cells; **E.** Survival rates of wild-type, IFN-g^{-/-}, IL-17^{-/-}, and IL-4^{-/-} mice after the *E. coli* injection. (Reprinted with permission from Ref. 71. Copyright © 2013 by The American Association of Immunologists, Inc)

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Figure 4. Modulating antibacterial immunity via bacterial membrane-coated nanoparticles A. Stability of extruded OMVs and BM-AuNPs with time; **B.** BM-AuNPs eliciting strong bacterium-specific antibody responses in vivo; **C.** BM-AuNPs inducing pronounced bacterium-specific T cell activation in vivo. (Reprinted with permission from Ref 95. Copyright © 2015 American Chemical Society)

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Figure 5. OMVs biomodification in Acinetobacter baumannii vaccines development. Schematic diagram of the construction of recombinant Omp22-OMVs (Reprinted with permission from Ref. 102. Copyright © 2016, Springer Nature)



Figure 6. Cryo-TEM visualization OMVs derived from *B. pseudomallei* outer membrane vesicles (120)

Cryo-transmission electron micrograph of purified OMVs prepared from a late logarithmic culture of *B. pseuodomallei* strain 1026b. Bar indicates 100 nm.

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Figure 7. Bacterial nanobioreactors–directing enzyme packaging into bacterial outer membrane vesicles

A. Crystal structures for the proteins utilized in the biorthogonal membrane conjugation of PTE for packaging into outer membrane vesicles; **B.** PTE kinetic data fit to the standard Michaelis–Menten enzyme kinetics equation for NAI and CAI UC pellet, CAI and CA UC supernatant (Left); A Lineweaver–Burk analysis used for determining KM and kcat/KM (Right). (Reprinted with permission from Ref.122 Copyright © 2015 American Chemical Society)



Figure 8. Bioengineered bacterial outer membrane vesicles as cell-specific drug-delivery vehicles for cancer therapy

A. Schematic representation of OMVs expressing HER2-specific affibody(Affi_{HER2}OMV) and the application of Affi_{HER2}OMV in cancer therapy; **B.** Tumor-specific retention and accumulation of delivered siRNA in major vital organs; **C.**Tumor growth inhibition (TGI) after delivery of siRNA (Reprinted with permission from Ref.125 Copyright © 2014 American Chemical Society)



Figure 9. OMVs functionality in vaccination and drug delivery

Left: A. Natural derived OMVs vaccine formulas, including OMVs alone, OMVs combined with adjuvants, OMVs combined with bacterial antigens and AuNPs loaded OMVs; B. Bioengineered OMVs vaccine formula. Bacterial antigens are expressed as fusion proteins with outer membrane proteins located on the outer membrane surface of OMVs; Right: A. Directing enzyme packing delivery systems for packing PTE enzymes in OMVs; B. siRNA loaded bioengineered OMVs in cancer therapy.

Table 1

OMV-associated Virulence proteins

OMV-associated proteins	Activity	References
Vacuolating toxins (VacA)	Immunolocalization	(37)
Porin proteins (OprF, L,etc)	Membrane proteins, transport small molecules	(44,51,52)
OstA	Organic solvent tolerance protein OstA precursor	(44)
OsmE	Membrane proteins, cell envelop integrity	(45)
FilC	Flagellin type B, motility and attachment	(53)
Pili machinery (pilA)	Type 4 fimbrial precursor pilA	(44)
PagL	Lipid A 3-O-deacylase	(54)
EstA	Secreted factors (enzymes), metabolism	(54)
aaaA	Amino acid biogenesis and metabolism	(54)