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Advancements in protein nanoparticle vaccine platforms to combat infectious disease

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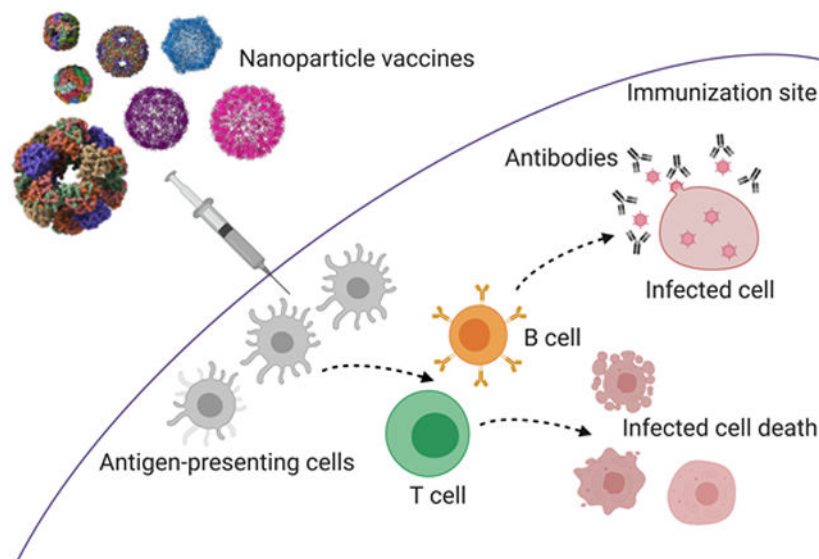
Abstract

Infectious diseases are a major threat to global human health, yet prophylactic treatment options can be limited, as safe and efficacious vaccines exist only for a fraction of all diseases. Notably, devastating diseases such as acquired immunodeficiency syndrome (AIDS) and coronavirus disease of 2019 (COVID-19) currently do not have vaccine therapies. Conventional vaccine platforms, such as live attenuated vaccines and whole inactivated vaccines, can be difficult to manufacture, may cause severe side effects, and can potentially induce severe infection. Subunit vaccines carry far fewer safety concerns due to their inability to cause vaccine-based infections. The applicability of protein nanoparticles (NPs) as vaccine scaffolds is promising to prevent infectious diseases, and they have been explored for a number of viral, bacterial, fungal, and parasitic diseases. Many types of protein NPs exist, including self-assembling NPs, bacteriophage-derived NPs, plant-derived NPs, and human virus-based vectors, and these particular categories will be covered in this review. These vaccines can elicit strong humoral and cellular immune responses against specific pathogens, as well as provide protection against infection in a number of animal models. Furthermore, published clinical trials demonstrate the promise of applying these NP vaccine platforms, which include bacteriophage-derived NPs, in addition to multiple viral vectors that are currently used in the clinic. The continued investigations of protein NP vaccine platforms are critical to generate safer alternatives to current vaccines, advance vaccines for diseases that currently lack effective prophylactic therapies, and prepare for the rapid development of new vaccines against emerging infectious diseases.

Graphical Abstract

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Various protein nanoparticles (NPs) have been utilized as scaffolds to display antigens for vaccination against infectious agents. NPs can stimulate humoral immune responses, cellular immune responses, or a combination of both.

Keywords

protein nanoparticles; antigen delivery; infectious disease vaccines; immune response; virus-like particles

1. Introduction

Bacteria, viruses, parasites, and fungi are the causative agents of infectious diseases [Lindahl & Grace, 2015], which vary in severity from the seasonal flu to fatal outbreaks of Ebola and endemic malaria [Fedson, 2016]. The most effective approach to prevent pathologies caused by these infectious organisms is vaccination [Omer et al., 2009][Delany et al., 2014]. Although drugs have been developed and approved to inhibit the growth or replication of these infectious organisms, these are only taken after an individual has contracted the organism. The power of vaccines lies in the fact that they can be administered long before patients come in contact with the organism, and they have the power to prevent disease pathology and systemic infection [Omer et al., 2009][Delany et al., 2014][Mbow et al., 2010]. Conventional vaccine approaches include live attenuated vaccines, inactivated whole organism vaccines, subunit vaccines, and toxoid vaccines [Kallerup & Foged, 2015]. The field has made significant progress in developing and bringing to market vaccines which protect against several diseases that pose serious threats to human health, including measles, mumps, rubella, smallpox, polio, hepatitis B (HBV), human papillomavirus (HPV), diphtheria, and influenza [Mailand & Frederiksen, 2017].

Despite the great benefit of vaccines, several have had documented issues with safety, which can have a serious impact on community trust, vaccination rates, and subsequently the loss

of control of an infectious agent in a population. For example, live attenuated vaccines, although highly effective and protective, can be lethal. Dengvaxia[®], the vaccine designed to prevent dengue disease, is a recent example of the dangers associated live attenuated vaccines; despite rigorous pre-clinical testing and clinical trials, the dangers of this vaccine were not realized until after it was deployed in a mass vaccination campaign of children in the Philippines [Cohen, 2019]. Historically and more recently, live attenuated oral polio vaccines have caused paralytic poliomyelitis [Desai et al., 2014][Alexander et al., 2004] [Mbaeyi et al., 2018]. Early rotavirus vaccines resulted in intussusception in children and negatively impacted the public's opinion and adoption of other vaccine candidates that were subsequently developed to protect against this highly contagious and transmissible diarrheal disease but which did not have these same safety concerns [Koch et al., 2017][Foster, 2007] [Soares-Weiser et al., 2012]. Incidents of post-immunization autoimmunity have also been linked to adverse responses to vaccination [Agmon-Levin et al., 2009][Cerpa-Cruz et al., 2013][Vadala et al., 2017]. These examples demonstrate the need to research and implement alternative vaccine approaches that are safer and elicit better antigen-specific immunogenicity.

Subunit vaccines are considered the safest design approach, but often struggle to induce robust immune responses to immunization [Vartak & Sucheck, 2016]. This can be overcome with the use of adjuvant platforms, as evidenced by the many subunit vaccines that have garnered FDA approval, including Shingrix, DPT (diphtheria, pertussis, tetanus), Prevnar 13[®], and Gardasil[®]9 [Shah, 2019][FDA, 2020b][Gupta et al., 1995][Sucher et al., 2011] [Manini & Montomoli, 2018]. Many adjuvants have achieved FDA approval and others are actively being studied for their ability to increase immunogenicity [Lee & Nguyen, 2015] [Sun et al., 2018][Jin et al., 2018]. Based on our experience, we believe nanoparticles (NPs) are an exciting adjuvant strategy that should be considered when novel subunit vaccines are being developed, especially for infectious diseases that still lack effective vaccines, including HIV-1, tuberculosis, and SARS-CoV2 (COVID-19) [Pati et al., 2018].

Nanoparticle (NP) scaffold platforms include inorganic NPs, polymeric NPs, liposomes, virus-like particles (VLPs), self-assembling NPs, and dendrimers [Pati et al., 2018][Al-Halifa et al., 2019]. NPs have already proven advantageous for the development of vaccines against cancer [Neek et al., 2019]. VLPs have virus-like nanostructures, but lack infectious genetic materials, providing safety while inducing humoral or cellular immune responses through the recognition of repetitive subunits on the surface of the particle by the immune system [Fuenmayor et al., 2017]. NPs with a size ranging from 20- to 200-nm can drain freely into lymph nodes and are presented to resident dendritic cells (DCs), making them ideal scaffolds for delivering antigens and immunostimulants [Manolova et al., 2008] [Bachmann et al., 2010]. The goal of vaccination is to elicit antigen-specific, high affinity antibody responses, especially IgG and IgA [den Hartog et al., 2020], against the immunizing antigens. As summarized in Figure 1, antigen-presenting cells (APCs) including DCs, macrophages, and B cells, take up extracellular proteins and degrade them into peptides for presentation to CD4⁺ helper T cells, which subsequently generate humoral immune responses. The CD4⁺ T cell and B cell interaction is crucial for the production of long-lived plasma cells which generate antigen-specific antibodies [Bachmann et al., 2010] [Look et al., 2010]. These antibodies can neutralize the infectious agent, thus preventing

infection, or mark it for phagocytosis by APCs where it is destroyed in lysosomes. In addition to interacting with CD4⁺ helper T cells, DCs can also trigger cytotoxic CD8⁺ T cell (CTL) cellular responses, resulting in the direct killing of infected cells [Look et al., 2010].

In this comprehensive review, we showcase the protein NPs that are being investigated for development into vaccine platforms for numerous infectious diseases, specifically focusing on self-assembling protein NPs, different types of VLPs (Fig. 2), and how they stimulate immune responses to vaccination. Furthermore, we discuss human-infecting viral vectors and their approved use as vaccine platforms for a number of infectious diseases [Sanchez et al., 2020][Ollmann, 2020]. Although others have also used these protein NP platforms as frameworks for drug and gene delivery systems [Lee et al., 2017][Ren et al., 2011][Molino et al., 2014], our main focus in this review is to highlight the investigations of protein NPs as structural scaffolds upon which to develop new vaccines. Examples of these vaccine NP scaffolds are shown in Figure 2. Furthermore, the humoral and cellular immune responses elicited from these NP vaccines, the adjuvants used, the route of administration, and other information will be covered.

2. Self-assembling Nanoparticles

In this section, studies on the development of infectious disease vaccines using non-viral self-assembling NPs as scaffolds will be reviewed. The NPs included in this section are small heat-shock protein, ferritin, and E2. We will discuss the structure of these NPs and the immune responses elicited after immunization with antigens presented on these NPs. A summary of this information is presented in Table 1.

2.1 Small heat-shock protein

Heat shock proteins (HSPs) are ubiquitous chaperones that bind to non-native or misfolded proteins, preventing aggregation. They are an evolutionarily conserved protein family and have evolved to exhibit a diverse range of structures symmetries [Abisambra et al., 2011]. For example, small heat shock protein 16.5 (sHsp16.5), derived from the primitive thermophilic archaeon *Methanocaldococcus jannaschii*, assembles into a 24-subunit caged nanoparticle. Within the archaea, sHsp16.5 sequesters non-native or misfolded proteins, preventing their aggregation [Shi et al., 2013]. Heat shock proteins have been examined as adjuvants in cancer vaccines [Segal et al., 2006].

In the area of infectious disease, Harmsen and colleagues have utilized the sHsp16.5 scaffold to elicit enhanced immune response [Richert et al., 2012][Richert et al., 2013]. Ovalbumin (OVA) was conjugated to sHsp16.5 (OVA-sHsp) following expression and purification from *E. coli* [Richert et al., 2012]. Mice were vaccinated with OVA-sHsp and then challenged with influenza, resulting in enhanced OVA-specific IgG levels in mice compared to vaccination with OVA. Because the OVA-sHSP NP does not itself contain antigens specific for influenza, this shows that it acted as a non-specific adjuvant, making the lung environment more conducive to a local antibody response. Pre-treatment with unconjugated sHSP induced local IgA response and accelerated rates of influenza-specific IgG production [Richert et al., 2012]. Treatment with the caged NPs OVA-sHsp enhanced alveolar macrophage and dendritic cell (DC) numbers in inducible bronchus-associated lymphoid

tissue and the tracheobronchial lymph node, which gave an accelerated response to the influenza challenge [Richert et al., 2013]. This migration pattern suggested augmented local immunity via increased phagocytic activity at the local site of infection. The vaccine promoted CD4⁺ T cell expansion and increased trafficking to the lungs, increasing accumulation in the tracheobronchial lymph node, inducible bronchus-associated lymphoid tissue, and spleen. Therefore, vaccination with sHSP increased immune activity both locally and systemically [Richert et al., 2013].

2.2 Ferritin

Ferritin is a spherical self-assembling protein NP that stores iron and is ubiquitous in living organisms. The main class of ferritin is 12-nm in diameter and contains 24 subunits, and antigens can be fused genetically to the N-terminus of a single ferritin subunit [Wang et al., 2019c] [Theil, 2013]. Georgiev et al. presented the possibility of placing two distinct antigens on one single ferritin while controlling the ratio and the geometric arrangement, and suggested that additional antigens can be attached, based on symmetry [Georgiev et al., 2018]. It was demonstrated that ferritin NPs were mostly recognized by DCs and macrophages expressing SIGNR1, and after subcutaneous injection in mice, fluorescently-labeled ferritin NPs were located abundantly in the medulla and interfollicular region [Wang et al., 2020].

Ferritin NP vaccines have been shown to induce humoral immune responses. For example, for vaccines in which Enterovirus 71 epitopes were displayed on the exterior of ferritin NPs, higher antibody titer responses were obtained [Wang et al., 2019c]. Serum from mice that were immunized with bacterioferritin fused to the receptor-binding domain (RBD) of Middle East Respiratory Syndrome (MERS-CoV) blocked the interaction between RBD and human receptor DPP4 (hDPP4) [Kim et al., 2018]. This is significant because the interaction between the RBD on MERS-CoV and the cellular receptor hDPP4 initiates MERS-CoV infection. The IgG1, IgG2a and IgG2b responses stimulated by RBD bacterioferritin were also stronger than RBD alone [Kim et al., 2018]. In another investigation, the display of multiple replicas of HIV-1 envelope glycoprotein trimers on ferritin resulted in greater neutralizing antibody titers and antibody responses against tier 1A viruses [Sliepen et al., 2015].

Ferritin has been examined in preclinical research as a vaccine scaffold for various viral infectious diseases; these include HIV-1; influenza (H1N1 strain); chronic hepatitis B (HBV); hepatitis C (HCV); Hand, Foot, and Mouth Disease (HFMD); rotavirus; Epstein-Barr; respiratory disease caused by respiratory syncytial virus, and Middle East Respiratory Syndrome (MERS-CoV) [He et al., 2016] [Swanson et al., 2020][Wang et al., 2019c] [Georgiev et al., 2018][Wang et al., 2020][Li et al., 2019][Sliepen et al., 2015][Kanekiyo et al., 2013][Kanekiyo et al., 2015][Kim et al., 2018][Yan et al., 2020]. Ferritin vaccines have also been investigated for bacterial infections, including gonorrhea [Wang et al., 2017]. Phase 1 clinical trials were completed in 2019 to examine the safety of a ferritin-hemagglutinin (HA) vaccine for influenza, and another study has an estimated completion date in 2021 [ClinicalTrials.gov, 2020b][ClinicalTrials.gov, 2020c][Kanekiyo et al., 2013] [Yassine et al., 2015].

2.3 E2

The pyruvate dehydrogenase complex of *Geobacillus stearothermophilus* includes a protein NP with icosahedral symmetry, formed from the E2 subunit of the complex. E2 is a 60-mer that assembles to form a hollow particle that is approximately 24-nm in diameter [Milne et al., 2002][Krebs et al., 2014]. It can be engineered with mutations to modify its properties while maintaining its dodecahedron structure and thermal stability; the internal, external, and inter-subunit interfaces can also be engineered to attach guest molecules [Dalmau et al., 2008][Neek et al., 2018] or modulate its pH-stability [Dalmau et al., 2009a][Dalmau et al., 2009b]. Foreign peptides and full-length proteins can be genetically attached to the N-terminus of each subunit, resulting in a display of up to 60 copies of the molecules [Domingo et al., 2001]. Multiple proteins can also be displayed on a single E2 NP, with the constraints being the protein expression of the protein-E2 fusion, its solubility, and steric hindrance [Caivano et al., 2010][Domingo et al., 2001]. This strategy has enabled the scaffold to be used as an antigen-delivery system for the development of cancer vaccines [Neek et al., 2018][Molino et al., 2016] and for infectious diseases that include HIV-1 [Krebs et al., 2014][Caivano et al., 2010][Jaworski et al., 2012][He et al., 2016], dengue fever [McBurney et al., 2016], and malaria [Domingo et al., 2001].

Domingo et al. showed that the surface of E2 can be simultaneously conjugated to green fluorescent protein and the MAL1 epitope from the circumsporozoite protein of *Plasmodium falciparum*, a parasite that can cause malaria [Domingo et al., 2001]. For HIV-1 vaccine studies, He et al. demonstrated that glycoprotein gp140 trimers can be stabilized on both E2 and ferritin scaffolds, and the resulting NPs are able to engage with B cells [He et al., 2016]. Caivano et al. demonstrated the possibility of genetically fusing the HIV-1 protein Gag(p17) on the N-terminus of the E2 while still resulting in the proper folding of the E2 scaffold [Caivano et al., 2010]. The IgG1 response was the dominant antibody isotype produced after the delivery of HIV-1 Gag(p17)-E2 NPs, indicating induction of Th2-type immune responses [Caivano et al., 2010]. In another study, by co-administering E2 presenting the HIV-1 envelope (Env) third hypervariable region and Env(gp160) DNA, the antibody response and neutralizing antibody titer were increased and developed more rapidly [Jaworski et al., 2012]. In addition, the percentage of IFN γ -secreting CD8⁺ T cells increased when the antigen-E2 scaffold and antigen DNA were co-administered, compared to the groups that received DNA and E2 with antigen separately [Jaworski et al., 2012]. This co-administration strategy of antigen-E2 with DNA was applied to a dengue vaccine that was tested in macaques [McBurney et al., 2016].

3. Virus-like Particles (VLPs): Bacteriophage-derived

Bacteriophages are viruses that infect bacteria, but not human cells [Hess et al., 2020]. Several types of bacteriophage exist, each with unique-sized shape and dimensions. In particular, the P22, Q β , λ , T4, MS2, and filamentous phage platforms have been developed for vaccination against numerous viral, bacterial, fungal, and parasitic infectious diseases (see Table 2). This section discusses recent advances in these platforms in infectious disease vaccines. While most vaccines are at the pre-clinical stage, an H1N1 Q β vaccine showed promise in a Phase I clinical trial [Low et al., 2014].

3.1 *Salmonella* virus P22 (P22)

The P22 phage is composed of a 60-nm icosahedral capsid that encapsulates double-stranded DNA (dsDNA), and a tail region which facilitates viral infiltration into *Salmonella typhimurium* bacteria [Wang et al., 2019b]. Researchers have recently utilized the P22 phage and its naturally-encapsulated DNA to develop influenza vaccines. In the study by Richert, et al., previously mentioned above, investigators evaluated the performance of sHSP conjugated to OVA antigen (OVA-sHsp) in an influenza murine model [Richert et al., 2012]. For some of the experiments, P22 phage was used as an adjuvant in combination with sHSP or OVA-sHsp. Pre-treatment of P22 prior to a single dose of OVA-sHsp enhanced OVA-specific serum IgG levels as early as 4 days after influenza A challenge. Since even non-modified P22 phages may be immunogenic, using P22 as a vaccine platform is promising to induce specific immune response. Others have also utilized a P22-based VLP by genetically fusing the conserved influenza A nucleoprotein antigen to the P22 capsid coat protein [Patterson et al., 2013]. Mice were vaccinated prior to lethal influenza challenge. The vaccine enhanced IgG antibody levels and antigen-specific CD8⁺ cytotoxic T cell activity, resulting in 80% survival compared to 0% of the unimmunized group. Another study developed a homologous influenza A virus (PR8)-specific P22 vaccine. Globular head domains of the hemagglutinin protein were fused onto the P22 platform through enzymatic conjugation. Mice were immunized then challenged with PR8 influenza A virus, leading to 100% survival compared to 50% for unvaccinated mice. Furthermore, the vaccine doubled antigen-specific IgG antibody levels compared to controls [Sharma et al., 2020].

The P22 platform has also been used in the development of a respiratory syncytial virus (RSV) vaccine. Two different RSV antigens, matrix (M) and matrix 2 (M2) proteins, were co-encapsulated through genetic fusion. Mice vaccinated with this formulation developed higher M- and M2-specific CD8⁺ and CD4⁺ T proliferation in the lungs, as well as more tissue-resident memory CD8⁺ T cells. Compared to PBS and un-conjugated P22 controls, treatment with P22 that co-encapsulated proteins M and M2 led to reduced viral titers in the lungs [Schwarz et al., 2016].

3.2 *Escherichia* virus Q-beta (Q β)

Q β is a bacteriophage that infects *E. coli* through the F-pilus protein, which is not present on mammalian cells, and therefore the virus is non-infectious to humans [Singleton et al., 2018]. Q β is found in abundance in sewage and animal feces. Its icosahedral coat proteins encapsulate interior single-stranded RNA (ssRNA), forming a 25-nm capsid cage. Q β has been used as a vaccine scaffold for both infectious disease as well as cancer vaccine applications [Hess et al., 2019][Speiser et al., 2010].

The use of Q β -based vaccines has been most extensively applied towards preventing viral diseases. One study developed a vaccine for foot and mouth disease virus (FMDV) by expressing and purifying Q β phages fused to FMDV viral antigen [Skamel et al., 2014]. Researchers performed multiple rounds of biopanning selection to allow for *in vitro* evolution of the phage vaccine, and the final selected FMDV antigen-displaying phages were used to vaccinate guinea pigs. Sera confirmed high antibody affinity for whole FMDV [Skamel et al., 2014]. These studies demonstrate that *in vitro* evolution is a feasible method

to develop a Q β -based vaccine with a higher diversity of presented antigen, recapitulating the plethora of viral antigen mutants that may be present in an animal infected with FMDV and thereby improving vaccine immune response. In another more recent study, several formulations of HIV-1 antigen fragment proteins displayed on the Q β scaffold were developed through genetic engineering or chemical conjugation. Formulation performances were compared in a six-month long rabbit immunization study. Priming with Q β VLPs followed by boost with full length HIV-1 antigen (subtype B JRFL gp120) resulted in superior neutralization antibody titers [Purwar et al., 2018]. Taken together, studies confirm that antigens can be displayed on Q β -based vaccines through a variety of biochemical approaches.

The Q β platform has also been used to develop influenza vaccines. Researchers have created an influenza A H1N1 viral vaccine using the Q β scaffold, a formulation showing high safety and immunogenicity in a Phase I clinical trial [Jegerlehner et al., 2013]. Initial studies used the Q β platform to develop vaccines against influenza B and multiple influenza A strains, including H1N1, H5N1, and H3N2. For each, the ectodomain of HA (gH) was conjugated to Q β , yielding strain-specific gH conjugation to Q β . Mice were vaccinated and lethally challenged with the specific influenza strain. Each formulation enhanced survival rates and antibody titers at multiple long-term time points, including up to 48 days post-challenge [Jegerlehner et al., 2013]. These results showed that vaccine design using the Q β scaffold provides strong protection in mice against H1N1 or other clinically relevant strains of influenza, depending on the antigen displayed.

Using the H1N1-specific vaccine (gH1-Q β) formulation, further *in vivo* studies evaluated efficacy of the vaccine adjuvanted with alum (Alhydrogel[®]), an FDA-approved adjuvant widely used to enhance vaccine performance. In both mice and ferrets, gH1-Q β with or without alum led to comparable levels of protection compared to the currently approved Panvax[®] H1N1 vaccine. However, the gH1-Q β vaccine treatment in mice induced superior Th1-biased CD4 T cell response compared to Panvax[®] [Skibinski et al., 2013]. In a Phase I clinical trial, the gH1-Q β vaccine resulted in 70.3% patient seroconversion by day 42 post-immunization, with no deaths or serious adverse events [Low et al., 2014]. The addition of a 2% Alhydrogel[®] adjuvant did not enhance seroconversion, perhaps because the natural ssRNA within the Q β scaffold is sufficiently immunogenic [Low et al., 2014]. In another study, peripheral blood mononuclear cells of the patients treated with the gH1-Q β vaccine were analyzed [Skibinski et al., 2018]. The gh1 peptide stimulation of samples collected post-vaccination had higher pro-inflammatory cytokine secretions levels including IFN- γ and IL17A than stimulated cells collected pre-treatment [Skibinski et al., 2018]. The Q β platform in influenza vaccine development serves as a pioneer for protein VLP vaccination in clinical trials. Its results in these human trials continue to draw interest, encouraging further investigation of Q β for other types of infectious diseases.

Q β -based vaccines are also being developed to address parasitic infections. Malaria, caused by *P. falciparum*, is transmitted by mosquitos. The malaria antigen circumsporozoite protein (CSP) was conjugated onto the Q β VLP [Khan et al., 2015]. Mice were vaccinated with the Q β -CSP vaccine and challenged with sporozoites. Following a two-dose schedule, vaccinated mice exhibited higher antibody titer levels specific for both the full length CSP

antigen as well as its central repeat epitope (termed NANP). NANP is similarly targeted by the currently approved malaria vaccine RTS,S, but in contrast the Q β -CSP may further enhance protection by displaying the full-length CSP antigen [Khan et al., 2015]. Interestingly, the surface display of polysaccharides on Q β can also serve as a vaccine. Moura et al. identified a trisaccharide epitope (α -Gal) found on the surface of leishmaniasis-causing parasites *Leishmania infantum* and *Leishmania amazonensis* [Moura et al., 2017]. These epitopes were displayed on the Q β scaffold, and the resulting VLPs were used to immunize mice in a leishmaniasis model. Immunized mice had at least doubled IgG serum antibody levels at 3 weeks post-immunization versus naive mice. Following challenge with *L. amazonensis* or *L. infantum*, vaccinated mice had diminished parasite loads [Moura et al., 2017]. These results demonstrate the versatility of the Q β phage platform in vaccine development, which is effective in both viral and parasitic models within preclinical research and preliminary clinical studies.

3.3 Escherichia virus lambda (λ)

λ phages can infect *E. coli* by entering the lytic or lysogenic replication pathways, which potentially allows for quiescent inheritance of phage by daughter cells. The dsDNA genetic material for phage replication is encapsulated in the icosahedron head, which is fused to a tail region that facilitates capsid ejection into *E. coli* cytoplasm [Fogg et al., 2010]. The λ phage platform has been investigated for development of a vaccine protective against hepatitis B virus (HBV). λ VLPs were engineered to also express hepatitis B antigen (HBsAg), and these were used to vaccinate mice and rabbits. HBsAg-specific antibody titers for both animal models over several weeks were enhanced compared to control treatment with HBsAg [March et al., 2004]. Given these promising results, further studies in rabbits compared the same formulation with the FDA-approved Engerix-B, a vaccine providing only limited protection to patients older than 60 years of age [Clark et al., 2011][Tohme et al., 2011]. Treatment with the λ phage vaccine led to consistently higher specific antibody titers compared to immunization with Engerix-B. These high antibody levels were maintained for more than 100 days in rabbits following multiple vaccinations with the λ vaccine [Clark et al., 2011]. It would be interesting to compare protection levels following challenge with HBV, especially since these results suggest potential advantages compared to an approved vaccine. However, we are not aware of more recent activity using the λ phage platform for infectious disease vaccines.

3.4 Escherichia virus T4 (T4)

The T4 phage consists of an icosahedron head with a tail region and is composed of 4 different proteins that form fibers, facilitating entry into *E. coli* [Bartual et al., 2010]. The outer capsid consists of a Hoc protein at the center of each major capsid and smaller Soc proteins [Shivachandra et al., 2006]. Compared to other phage-based vaccines, recombinant T4-based vaccines have been investigated mostly to protect against bacterial infection. Both of the following bacterial vaccines were developed by genetically fusing antigen to either the Hoc or Soc regions of the T4 phages.

Anthrax infection is caused by the Gram-positive bacterium *Bacillus anthracis*. Usually treated by antibiotics, inhalation of *B. anthracis* is more difficult to treat and can be fatal.

One study investigated an anthrax vaccine by fusing protective antigen (PA) from *B. anthracis* to the N-terminus of the T4 Hoc regions [Peachmann et al., 2012]. The T4-PA vaccine was compared to alternative PA formulations, including a PA-based emulsion and PA-encapsulated liposomes with adjuvants. Rabbits were vaccinated with the different formulations and challenged with anthrax. The T4-PA vaccine enhanced IgG antibody levels and survival rates, although survival was 70%-80% that of the other formulations [Peachmann et al., 2012]. The T4-PA VLPs were immunogenic in themselves, likely due to the naturally-packaged dsDNA, and performance would likely be further improved by the inclusion of adjuvant (such as alum adjuvant used in the previously-mentioned Q β studies).

More recently, a dual *B. anthracis* and plague (*Yersinia pestis*) vaccine was developed by genetically fusing antigens from both bacteria to the T4 Soc capsid proteins [Tao et al., 2018]. Following lethal doses of anthrax toxin and *Y. pestis* challenge in mice, rats, and rabbits, vaccination resulted in at least 80% survival, regardless of species. Neutralizing antibodies were also consistently higher in animals treated with recombinant T4 displaying antigen versus non-modified T4 [Tao et al., 2018]. As anthrax and plague are both potential biological weapons, the prospect of an effective dual vaccine has important clinical relevance.

3.5 Escherichia virus MS2 (MS2)

The MS2 phage consists of a 26-nm diameter icosahedron shell encapsulating ssRNA. Following infection of *E. coli*, virion assembly is instructed by multiple, specific RNA sequence sites [Stockley et al., 2016]. MS2-based vaccines are being developed for multiple anti-viral applications, including prevention of FMDV and human papillomavirus (HPV) infections.

To address FMDV, a coat protein epitope of a structural protein of FMDV (VP1) was genetically fused into the coat protein of MS2 [Dong et al., 2015]. Extensive *in vivo* studies were performed, including vaccination and FMDV challenge of mice, guinea pigs, and swine. Overall health, antibody levels, and endpoint survival were evaluated, revealing comparable performance to the standard inactivated vaccine that is commonly used in livestock husbandry [Dong et al., 2015]. FMDV is highly contagious, resulting in widespread livestock deaths and costing individual countries billions of dollars annually, yet no specific treatment currently exists [Knight-Jones & Rushton, 2013]. Therefore, it is especially promising that multiple phage platforms (including MS2 and Q β) may offer potential prevention options.

Tumban and colleagues have recently developed an effective MS2-based HPV vaccine by genetically engineering MS2 VLPs to express a tandem HPV peptide (amino acids 17-31) with an HPV consensus peptide (amino acids 69-86). While the former peptide offers strong antibody response against HPV16 [Tumban et al., 2012], the latter offers a lower yet more widespread antibody response against several other HPV types [Tyler et al., 2014]. Vaccination of the mice with MS2 displaying tandem and consensus peptides not only protected mice from multiple HPV types, but antibody levels were comparable to those of mice immunized with the approved HPV vaccine, Gardasil^{®9} [Zhai et al., 2017]. This group further evaluated vaccine performance combined with 2 mucosal adjuvants in mice,

including cholera toxin and the approved adjuvant monophosphoryl lipid A (MPLA). While the former study showed efficacy against only 6 types of HPV, these studies also showed efficacy against 8 additional HPV strains, including HPV16, which is associated with more than 70% of HPV-associated head and neck cancers. Notably, the MS2-based formulation provides protection against several additional HPV types associated with head and neck cancer versus the FDA-approved Gardasil[®]9 [Zhai et al., 2019]. These results suggest that a MS2-based vaccine may potentially protect against a more widespread range of HPV types than vaccines currently on the market.

3.6 Filamentous phage

While there have been a number of different filamentous phages used in antigen delivery (e.g., M13, fd, f1), all are rod-shaped with repetitive structure encapsulating ssDNA, which may be recognized by immune-activating toll-like receptor TLR9 [Henry et al., 2015]. Filamentous phage rod bodies consist of repeating units of pVIII coat protein, which are common sites engineered to express antigen peptides [Sartorius et al., 2019]. Filamentous vaccines, which others have previously reviewed [Henry et al., 2015], have been developed to vaccinate against bacterial, fungal, viral, and parasitic infections, and we highlight a few here.

Phage vaccine platforms have been examined for prevention of fungal infections. Wang et al. generated anti-fungal vaccines by genetically fusing 2 antigen epitopes from the fungus *Candida albicans* onto M13 bacteriophage [Wang et al., 2014]. Both epitopes were derived from the protein Sap2, a protease released by *C. albicans* known to degrade host proteins. Vaccination with the hybrid phage vaccine enhanced survival of mice challenged with *C. albicans*, in addition to higher levels of IgG antibody and pro-inflammatory cytokines, and reduced fungal levels in the spleen and kidney [Wang et al., 2014]. In another application, the prevention of fungal infection caused by *Sporothrix globosa*, a fungus present in soil and plant matter that causes sporotrichosis, has been investigated. Filamentous phages were genetically engineered with antigen epitopes of the fungus onto the major coat protein (pVIII). Mice vaccinated with the recombinant phage showed enhanced survival rates, even superior to those immunized with heat-killed *S. globosa* [Chen et al., 2017]. Because there are no currently approved anti-fungal vaccines [Nami et al., 2019], these examples of positive results are important for further evaluation.

Filamentous-based vaccines are also being developed to provide protection against parasitic infection. Researchers engineered fd filamentous phage to express highly immunodominant peptides trans-sialidase and amastigote surface protein-2, motifs accessible from the surface of the parasite *Trypanisoma cruzi* [Gomes-Neto et al., 2018]. Murine vaccination with either of the recombinant phage designs led to enhanced IgG antibody titer levels and superior survival rates following challenge with *T. cruzi*. Immunization was inhibited in TLR9 knockout mice, showing that immunogenicity of the filamentous phage-based vaccine requires receptor recognition of the naturally-encapsulated ssDNA adjuvant [Gomes-Neto et al., 2018]. Symptoms of Chagas disease, predominantly caused by transmission of *T. cruzi* from infected bugs, can be mediated by current anti-parasitic treatments. However, some individuals are chronically infected, and vertical transmission is of especially high concern

in poorer regions of Latin America [Moncayo, 2017]. Therefore, the possibility of a vaccine would offer an encouraging option and merits further research.

4. Virus-like Particles (VLPs): Plant virus-derived

Plant viruses can infect specific plant types but lack the mammalian-replicable genetic material necessary to infect humans. In tandem with this characteristic and the utilization of genetic fusion to conjugate peptide antigens from a multitude of pathogens, these VLPs have been explored as vaccine platforms for numerous infectious diseases (see Table 3). This section will specifically focus on vaccine formulations and immunization studies that have used plant virus-derived VLPs as vaccine platforms.

4.1 Cowpea Mosaic Virus (CPMV)

The 30-nm self-assembling icosahedral cowpea mosaic virus (CPMV) is one of the most studied plant-based platforms for protein nanoparticle vaccine development. It has been investigated as a vaccine platform against bacterial and viral pathogens, including *Staphylococcus aureus*, *Pseudomonas aeruginosa*, Group B *Streptococcus*, and HIV-1. In more recent years, CPMV has also been utilized in the development of cancer vaccines [Lizotte et al., 2016][Murray et al., 2018][Wang et al., 2019a]. Most formulations of CPMV utilize the technique of genetic fusion to load immunogenic epitope peptides from the proteins of pathogens.

CPMV fused to the D2 peptide of the fibronectin-binding protein B (FnBP) of *S. aureus* has elicited strong humoral immune responses. Immunizations of mice and rats with these CPMV-D2 formulations have yielded high titers of FnBP-specific antibodies, generated neutralizing antibodies capable of inhibiting the binding of fibronectin required for disease pathogenesis, produced antibodies that induced opsonization of *S. aureus*, protected animals from weight loss when intravenously administered with pre-opsonized *S. aureus*, and protected against endocarditis [Brennan et al., 1999a][Brennan et al., 1999b][Brennan et al., 1999c]. In a model of chronic pulmonary infection, a fusion protein of CPMV and peptide derived from the outer-membrane protein F of *P. aeruginosa* afforded protection upon challenge with two different immunotypes of the bacterium. Immunization generated high titers of antigen-specific opsonizing antibodies that marked the bacteria for phagocytosis by human neutrophils [Brennan et al., 1999b]. Against the pathogen group B *Streptococcus*, mice immunized with CPMV genetically fused to S9 peptide, a mimic of the group B streptococcal type III capsular polysaccharide, elicited a Th1 response. This response was characterized by the production of peptide-specific IgG2a antibodies, as well as the generation of IFN- γ by antigen-stimulated lymphocytes derived from treated mice [Pomwised et al., 2016].

In the application against a viral pathogen, formulations of CPMV fused with a gp41 peptide have been effective in inducing humoral and T cell-mediated immune responses against HIV-1. Immunization studies in mice have shown strong gp41-specific antibody responses, *in vitro* proliferation of T cells from splenocytes, and HIV-1 neutralization [McInerney et al., 1999]. Interestingly, intranasal immunization also elicited strong gp41 peptide-specific

antibody responses, supporting the viability of CPMV as an intranasal vaccine delivery system that may improve patient compliance to vaccination [Durrani et al., 1998].

4.2 Potato Virus X (PVX)

Potato virus X (PVX) is a self-assembling rod-structured plant virus that naturally infects potatoes. Through genetic engineering, PVX can be modified for pathogen antigen conjugation. Specifically, fusing antigens onto the nanoparticle scaffold allows high loading capacity of repeating antigens, which is favored for eliciting an immune response.

Utilizing genetic fusions, the antigens gp41 peptide, nucleoprotein peptide, and R9 peptide have been conjugated to PVX to induce immune response against HIV-1, influenza (H1N1), and hepatitis C virus (HCV), respectively. Immunization studies in mice have elicited humoral response against both HIV-1 and HCV, including antigen-specific and neutralizing antibodies [Marusic et al., 2001][Uhde-Holzem et al., 2010]. In the case of HIV-1, the sera of immunized mice were also able to inhibit virus-induced syncytium, a phenomenon in which infected cells express glycoproteins from the virus, allowing infected cells to adhere to healthy cells, which in turn leads to accelerated disease progression. For infectious diseases that reside extensively within cells, a vaccine-elicited cellular response characterized by antigen-specific CD8⁺ T cell activation is desired for the eradication of infected cells. This activation of CD8⁺ T cells has been observed using this PVX-derived platform; when immunized with the PVX-nucleoprotein peptide formulation, mice produced antigen-specific CD8⁺ T cells [Lico et al., 2009].

4.3 Tobacco Mosaic Virus (TMV)

Tobacco mosaic virus (TMV) is a 300-nm x 20-nm rod-shaped plant virus [Butler, 1984]. As discussed, most plant virus-derived nanoparticle platforms utilize genetic engineering to load antigens, but studies with TMV have employed both genetic fusion and chemical conjugation to load peptide antigens. Due to the symmetry of the virus, the number of antigens per TMV particle can be high.

TMV has been studied as a vaccine platform for bacterial and viral pathogens. Mice immunized with a TMV-peptide antigen vaccine produced peptide-specific opsonic antibodies that reacted to all seven strains of *P. aeruginosa*, as well as conferred protection against challenge with wild-type *P. aeruginosa* in a mouse model of chronic pulmonary infection [Staczek et al., 2000]. The TMV platform has also been engineered against the bacterial pathogen *Francisella tularensis*. Utilizing both genetic fusion and chemical conjugation, protein antigens including chaperone protein DnaK, OmpA-like protein (OmpA), SucB protein, and lipoprotein Tul4 have been conjugated onto TMV [Banik et al., 2015][Mansour et al., 2018][McCormick et al., 2018]. Immunization of mice with these formulations induced strong humoral responses predominated by IgG1 and IgG2 antibodies, as well as protection against lethal respiratory challenge of *F. tularensis* LVS (live vaccine strain) even after 163 days post-primary immunization [Banik et al., 2015][Mansour et al., 2018][McCormick et al., 2018]. In these studies, the magnitude and quality of immune responses were shown to vary based on immunization routes (subcutaneous or intranasal), adjuvant administration, and antigen selection. Against anthrax, immunization of mice with

genetically-fused peptides from the protective antigen (PA) protein toxin of anthrax has shown to partially neutralize this toxin when challenged with anthrax spores [McComb et al., 2015]. In the case of *Yersinia pestis*, LcrV and F1 protein antigens that were chemically conjugated to TMV induced an immune response capable of complete protection against pneumonic infection after a lethal dose of *Y. pestis* in mice [Arnaboldi et al., 2016].

For viral pathogens, TMV vaccines have generally elicited antigen-specific humoral responses. Interestingly, immunization of mice with a TMV-monomeric hemagglutinin protein (HA) vaccine not only elicited antibodies, but also better protected mice against H1N1 influenza virus challenge than the commonly-used trivalent inactivated H1N1 vaccine [Mallajosyula et al., 2014]. Against FMDV, a livestock disease, a TMV-based vaccine elicited peptide-specific antibodies, as well as protection against FMDV challenge in guinea pigs and swine [Jiang et al., 2006].

4.4 Cucumber Mosaic Virus (CMV)

Cucumber mosaic virus (CMV) is a plant virus with a 30-nm self-assembling icosahedral capsid. Immunization in rabbits with a fusion protein of CMV and peptide derived from the hepatitis C virus (HCV) envelope protein E2 (R9 mimotope) has led to increases in peptide-specific antibodies [Piazzolla et al., 2005]. In patients with chronic HCV infection, the CMV vaccine induced a significant release of IFN- γ , IL-12, and IL-15 by peripheral blood mononuclear cells, indicating a strong innate immune response to the peptide antigen [Piazzolla et al., 2005][Nuzzaci et al., 2007]. A CMV vaccine formulation using Zika virus envelope protein (EDIII) as its antigen induced high levels of antigen-specific antibodies after a single immunization; this was capable of neutralizing Zika virus without enhancing infection by dengue virus *in vitro* [Cabral-Miranda et al., 2019]. Unique from many plant-based VLPs, a CMV construct against the parasitic disease, malaria, has also been studied. Immunization with the chemically-conjugated malaria TRAP protein (from *Plasmodium vivax*) induced an increase in humoral responses and a strong T cell response, and it offered protection against challenge with recombinant *Plasmodium berghei* that expressed *P. vivax* TRAP antigen [Cabral-Miranda et al., 2019].

4.5 Alfalfa Mosaic Virus (AIMV)

Alfalfa mosaic virus (AIMV) particles have a bacilliform structure with a diameter of 19-nm and a length varying from 30- to 56-nm. As with other plant virus-derived VLP scaffolds, genetic fusion of pathogen antigens to AIMV has been utilized to construct infectious disease NP vaccines. AIMV has shown promise in inducing antigen-specific immune responses against both respiratory syncytial virus (RSV) and *P. falciparum*. Human DCs incubated with the AIMV-RSV formulation have been shown to stimulate vigorous CD4⁺ and CD8⁺ T cell responses [Yusibov et al., 2005]. Strong cellular and humoral RSV-specific immune responses have been generated in non-human primates [Yusibov et al., 2005]. In vaccine formulations against *P. falciparum*, AIMV has also shown promise in mice following immunization, where serum antibodies were observed with complete transmission blocking activity through a 6-month study period [Jones et al., 2013]. This formulation then entered a Phase 1 first-in-human study where a dose dependent antigen-specific antibody response was observed, as well as no evidence of dose-related toxicity. However, it was also noted

that the transmission-reducing activity of the generated antigen-specific antibodies was weak, pointing to the need for further optimization [Chicester et al., 2018]. AIMV's ability to elicit an immune response against a parasite using a peptide antigen gives this vaccine platform a unique position amongst the plant virus-based VLPs.

4.6 Papaya Mosaic Virus (PapMV)

This 500-nm x 13-nm rod-shaped virus was first reported in 1962 when studying diseased papaya plants in Florida [Conover, 1962]. In attempts to develop a universal influenza vaccine, PapMV fused with peptides from the influenza M2 ion channel protein have been shown to induce production of antigen-specific antibodies that can recognize infected cells and protect mice from influenza (H1N1) challenge [Therien et al., 2017][Carignan et al., 2015]. Combining this formulation with multimerized nucleoprotein nanoparticles protected mice from challenges by both H1N1 and H3N2 influenza strains [Bolduc et al., 2018]. More impressive was the ability of free (unconjugated) PapMV VLPs, used as adjuvants, to dramatically increase the immunogenicity of the PapMV-antigen vaccine. This increased immunogenicity led to 100% protection against a challenge with the WSN/33 influenza strain [Denis et al., 2008]. *In vitro* activation of B cells and expansion of antigen-specific T cells using PapMV-peptides derived from the nucleocapsid protein and M1 matrix protein of the influenza virus have been supported by immunizations of mice that exhibited antigen-specific CD8⁺ T cell responses [Babin et al., 2013][Leclerc et al., 2007][Hanafi et al., 2010] [Laliberte-Gagne et al., 2019].

PapMV formulations using peptide antigen fusions have also been examined for vaccines against hepatitis C virus (HCV) and lymphocytic choriomeningitis virus (LCMV). Mice immunized with the PapMV-E2 glycoprotein epitope vaccine showed a long-lasting humoral response (more than 120 days) against the HCV peptide antigen [Denis et al., 2007]. Against LCMV, the PapMV-p33 epitope (derived from LCMV's surface glycoprotein) vaccine activated large numbers of antigen-specific CTLs which rapidly expanded following LCMV challenge and protected vaccinated mice against LCMV infection in a dose-dependent manner [Lacasse et al., 2008].

5. Virus-like Particles (VLPs): Recombinant human-virus derived viral vectors (new section)

Viral vectors fall under a niche category of protein nanoparticle vaccine platforms. Unlike the other particles discussed in this review, these NPs have the ability to mimic infection in humans by entering host cells and inducing host-cell expression of foreign antigen using viral mechanisms; however, they lack viral extranuclear replication capabilities [Matthews et al., 2013][Ura & Shimada, 2014][Bull et al., 2019]. This in turn induces an immune response to those foreign, pathogenic antigens. The first published work in 1984 describing viral vectors as scaffolds used to deliver vaccine antigens was a recombinant vaccinia virus engineered to express hepatitis B antigen in host cells [Moss et al., 1984]. Since those initial studies, a plethora of viral vectors have been investigated, leading to several FDA-approved vaccines. While effective, safety concerns related to their innate pathogenic nature in humans have been raised [Rauch et al., 2018]. In this section, the use of human-tropism

virus vectors as vaccine platforms for infectious disease will be summarized and discussed. Due to many extensive reviews already written in this area [Ho et al., 2020][Nieto & Salvetti, 2014][Ura & Shimada, 2014][Bull et al., 2019], we highlight only a few platforms that have been studied in the clinic, giving some examples which have been approved for use.

5.1 Hepatitis B surface antigen (HBsAg)

Globally, hepatitis B is a common viral infectious disease, and its surface envelope protein (HBsAg) has the ability to present foreign antigens for chimeric protein delivery and has been used in preclinical and clinical vaccine development [Ho et al., 2020][RTS,S Clinical Trials Partnership, 2015]. RTS,S/AS01 (RTS,S) (Mosquirix) is a malaria subunit vaccine developed by fusing a portion of *Plasmodium falciparum*-derived circumsporozoite (CS) onto HBsAg [Choi & Chang, 2013][Ho et al., 2020]. Naturally-acquired immunity against malaria is mediated by IgG levels and can be gained with age and exposure, decreasing risk of morbidity and mortality in adults; however, because children and infants may lack such protection, vaccination is important in this patient group [Dobano et al., 2019b][Doolan et al., 2009]. In recent years, clinical trials utilizing RTS,S/AS01 have examined safety, efficacies, immunogenicity, as well as a multitude of unique immune responses [Asante et al., 2020][Sanchez 2020]. Immune responses including unique antibody responses after variable doses and unique avidities of antibodies have been well documented [Sanchez et al., 2020][Dobano et al., 2019a] [Dobano et al., 2019b]. RTS,S vaccination of young children in Africa results in higher specific IgG antibody titers, correlating with increased protection against clinical malaria [Dobano et al., 2019b]. However, there are limitations; only about 56% of children receiving the RTS,S vaccine are protected from naturally-occurring malaria infection, with around 31% protection rates in infants [Dobano et al., 2019b][Vrieze, 2019]. The WHO has now set up a pilot vaccination program in Africa to monitor safety and efficacy concerns as the vaccine is deployed.

5.2 Adenovirus (Ad) vectors

One of the most studied viral vectors are adenovirus (Ad) vectors. Ad vectors have been examined in clinical trials for vaccines against HIV, influenza, malaria, and tuberculosis (TB), and more recently, SARS-CoV-2 [Mathews et al., 2013][Choi & Chang, 2013][Ura & Shimada, 2014][BioRender, 2020][Zhu et al., 2020][Ramezanpour et al., 2016][van Zyl-Smit et al., 2017][Liebowitz et al., 2020][van Zyl-Smit et al., 2017]. Ad-based vaccines for HIV-1 and malaria have been shown to induce antigen-specific CD4⁺ and CD8⁺ T cell responses and antibody responses in clinical trial volunteers [Mathews et al., 2013]. An Ad-based vaccine for malaria has been demonstrated to induce similar specific humoral and cellular immune responses as the RTS,S/AS01 vaccine described previously [Choi & Chang, 2013][Ramezanpour et al., 2016]. During the Ebola virus outbreak from 2014-2016, several Ad-based Ebola virus vaccines saw development and clinical trials through phase 1 and 2 [Lauer et al., 2017][Rauch et al., 2018]. More recently, Ad-based vaccines are being investigated to combat the outbreak of SARS-CoV-2, the causative infectious agent of COVID-19. Currently, three Ad-based SARS-CoV-2 vaccines are undergoing clinical trials with two candidates in phase 2 and one in phase 3 [BioRender, 2020][Zhu et al., 2020].

5.3 Vesicular stomatitis virus (VSV)

Vesicular stomatitis virus (VSV), a member of the *Rhabdoviridae* family, is an enveloped virus containing a single-stranded, negative-sense RNA [Rauch et al., 2018]. Due to the lack of a DNA intermediate during viral infection, VSV has become an attractive vaccine platform [Rauch et al., 2018]. In particular, a VSV-based vaccine has been developed for Ebola in which the VSV envelope glycoprotein was replaced by the corresponding protein from the Ebola virus [Marzi et al., 2011][Ollmann, 2020]. It progressed in clinical trials during the Ebola virus outbreak of 2014-2016, received conditional marketing authorization by the WHO, and was approved by the FDA in 2019. This vaccine (Ervebo) has now been used to vaccinate over 280,000 people (at the time of writing this article) during the ongoing 2018-2020 outbreak in Africa [Rauch et al., 2018][Ollmann, 2020].

5.4 Other viral vector platforms

The use of adeno-associated virus (AAV) vectors as vaccine platforms have also been studied for infectious diseases including HIV-1, HPV, influenza (H1N1), and hepatitis C [Nieto & Salvetti, 2014][Ura & Shimada, 2014][Ramezanpour et al., 2016][Demminger et al., 2020][Zhu et al., 2019][Naso et al., 2017][Vardas et al., 2010][ClinicalTrials.gov, 2020a]. The rarity of AAV-based vaccines in clinical trials is due to its relatively weak ability to induce humoral and cellular immune response when compared to other viral vectors [Ura & Shimada, 2014]. Alphaviruses replicate in the cytoplasm of infected cells, which eliminate the concern for potential integration of viral genes into the host genome, and a number of alphavirus-based vaccine formulations have been shown to induce both antigen-specific humoral and cellular responses [Choi & Chang, 2013] [Ramezanpour et al., 2016]. Chimeras of two alphaviruses, Venezuelan equine encephalitis virus (VEE)-Sindbis virus (SIN), have been pre-clinically shown to induce high neutralizing antibody titers and cell-mediated responses against HIV-1 [Choi & Chang, 2013]. Poxviruses have large genomes making them advantageous for larger antigen gene insertions and have been investigated as the viral vectors for malaria, tuberculosis, small pox, influenza, and HIV-1 vaccines in clinical trials [Choi & Chang, 2013][Ura & Shimada, 2014][Pitisuttithum et al., 2020][Laher et al., 2020][Levy et al., 2020][Pittman et al., 2019][Krejtz et al., 2014] [Minhinnick et al., 2016][Wilkie et al., 2020][de Barra et al., 2014] .

6. Future outlook

Since Edward Jenner's first cowpox vaccine protected against smallpox, the field of vaccinology has made tremendous impacts in quality and length of life for untold millions of people around the world [Delany et al., 2014]. However, our work is not yet finished. Vaccines are still lacking for well-characterized infectious agents, including HIV-1 and tuberculosis, and new, emerging infectious agents, such as the 2009 H1N1 flu or SARS-COV2, will necessitate the rapid discovery and deployment of vaccine candidates to curb the threat of extended global pandemics [Pati et al., 2018][Lurie et al., 2020]. Although a live attenuated or inactivated whole organism approach may work for some of these organisms, subunit vaccines remain the safest design approach [Vartak & Sucheck, 2016]. These subunit vaccines will require adjuvants and alternative approaches to be efficacious; based on our

experience and the data reviewed here, we believe a protein nanoparticle-based approach should be considered in the design of these vaccines.

As of the writing of this review, 174 vaccines against SARS-CoV2 (the causative agent of COVID-19) are in development, with 51 of these in clinical trials [BioRender, 2020]. Several approaches are being taken, including inactivated SARS-CoV2 particles, a genetically modified chimpanzee virus expressing SARS-CoV2 proteins, and nucleic acid based vaccines [Callaway, 2020]. Subunit vaccines with antigens attached to nanoparticles are also being developed, including a self-assembled VLP, NSP-10, which is expected to generate desired immune response after displaying SARS-CoV2 spike protein [Carter et al., 2020]. Several factors are important for the development of COVID-19 vaccines; it is crucial to display the correct antigens in the correct conformations to induce antigen-specific, high affinity neutralizing antibodies. It is also critical to avoid antibody-dependent enhancement of viral uptake and induction of Th2 immune responses which have the potential to cause allergic inflammation and vaccine-associated enhanced respiratory disease [Graham, 2020] [Peeples, 2020][Vabret et al., 2020].

Great accomplishments have been achieved in developing nanoparticle vaccines for deployment in humans. However, these platforms still face a few challenges before they are ready for widespread inclusion in human vaccine formulations. The first limitation is the lack of in-depth understanding of how every permutation of each nanoparticle-based platform will perform *in vivo*. The high degree of customization inherent to NP design (factors including size, charge, and shape) is both a blessing and a curse; we have not yet established the ideal combination of each of these factors to achieve ideal targeting, biodistribution, and clearance for each pathogen of interest, but such information is important to advance the field. One can observe this challenge being surmounted as materials scientists, engineers, immunologists, and vaccinologists work together to generate vaccines using protein nanoparticle technologies.

The second challenge is maintaining reproducibility of synthesis as production scales up, which is a difficulty faced by every new therapeutic technology aiming to make the leap from pre-clinical development to clinical deployment. We know that this is not an insurmountable problem as multiple nanoparticle-based platforms have been reproducibly scaled for human clinical trials. Much like the first challenge, it is only a matter of time before the synthesis and large-scale production parameters are determined for each of these platforms.

Despite these challenges, the field is primed to see the results of clinical trials for vaccines using several of the platforms that are discussed in this review, and to envision where and how they will be applied next for the improvement of human health worldwide. As the collective field of vaccine research and design moves forward to meet the challenge of infectious and transmissible disease in the post-COVID-19 pandemic era, protein nanoparticle-based platforms have shown tremendous potential and should be considered as a possible strategy in vaccine development.

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Abbreviations

AAV	adeno-associated virus
Ab	antibody
Ad	adenovirus
AIDS	acquired immunodeficiency syndrome
AIMV	alfalfa mosaic virus
APC	antigen presenting cell
CD28	cluster of differentiation 28, co-stimulatory molecule expressed on T cells
CD80	cluster of differentiation 80, co-stimulatory molecule expressed on APCs
CMV	cucumber mosaic virus
COVID-19	coronavirus disease of 2019
CPMV	cowpea mosaic virus
CSP	circumsporozoite protein
CT	cholera toxin
CTL	cytotoxic T lymphocyte, CD8 ⁺ T cell
DC	dendritic cell
DNA	deoxyribonucleic acid
dsDNA	double-stranded DNA
ELISA	enzyme-linked immunosorbent assay
FCA	Freund's complete adjuvant
FIA	Freund's incomplete adjuvant
FDA	U.S. Food and Drug Administration

FMDV	foot and mouth disease virus
HA	hemagglutinin
HBV	hepatitis B virus
HCV	hepatitis C virus
HFMV	Hand, Foot, and Mouth Disease
HIV-1	human immunodeficiency virus 1
HPV	human papillomavirus
IL	interleukin
IM	intramuscular injection
IN	intranasal administration
IP	intraperitoneal injection
ISCOM	immune stimulating complex
IV	intravenous injection
LCMV	lymphocytic choriomeningitis virus
λ	Escherichia virus lambda
MERS	Middle East Respiratory Syndrome
MHC	major histocompatibility complex
MPLA	monophosphoryl lipid A
MS2	Escherichia virus MS2
MVA	Modified vaccinia virus Ankara
MΦ	macrophage
NP	nanoparticle
OVA	ovalbumin
P22	salmonella virus P22
PapMV	papaya mosaic virus
PVX	potato virus X
Qβ	Escherichia virus Q-beta
RNA	ribonucleic acid
RSV	respiratory syncytial virus

RTS,S	RTS,S/AS01 malaria vaccine (trade name Mosquirix)
SARS-CoV2	severe respiratory syndrome coronavirus 2
SC	subcutaneous injection
sHSP	small heat shock protein
SIN	Sindbis virus
ssRNA	single-stranded RNA
T4	Escherichia virus T4
TB	tuberculosis
TCI	transcutaneous immunization
TCR	T cell receptor
T_{FH}	T follicular helper cell, CD4 ⁺ T cell
Th1	T helper 1 cell, CD4 ⁺ T cell
Th17	T helper 17 cell, CD4 ⁺ T cell
TMV	tobacco mosaic virus
TNF-α	tumor necrosis factor alpha
VEE	Venezuelan equine encephalitis virus
VLP	virus-like particle
VSV	vesicular stomatitis virus
WHO	World Health Organization

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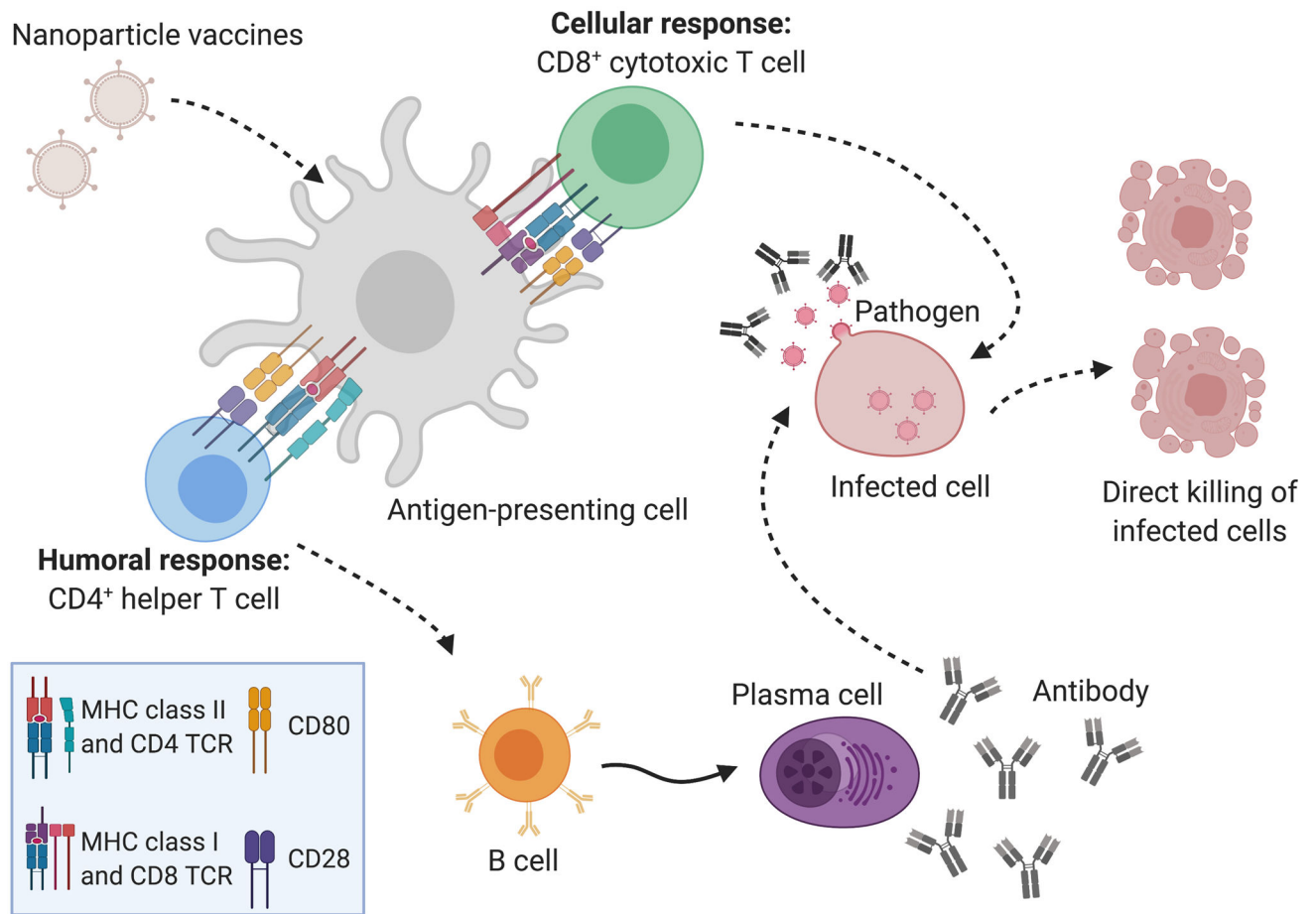


Figure 1. The interaction between the nanoparticle (NP) vaccines and the immune system results in immunological memory and protection against future infections.

NP platforms are capable of stimulating the host immune response to generate both cellular (CD8⁺ T cell) and humoral (antibody) responses to the pathogen antigen(s) included in the vaccine formulation. As detailed in this review, each unique NP platform can generate antigen-specific CD8⁺, CD4⁺ and/or antibody responses to immunization following uptake of the NP vaccine by local antigen-presenting cells at the immunization site. These antigen-presenting cells travel back to draining lymph tissues, where they process the nanoparticle vaccine platforms into protein peptides that are presented in MHC molecules to both CD8⁺ and CD4⁺ T cells. CD8⁺ T cells (green) are capable of directly killing infected cells (pink). CD4⁺ T cells (blue) interact with B cells (orange) to stimulate the maturation of the B cells into antibody-secreting plasma cells (purple). The antibodies are specific to the pathogen and can neutralize their ability to infect and proliferate. The antigen-specific cellular and/or humoral responses generated by the NP platforms will be maintained as memory cells against the pathogen, which can be quickly activated when the immunized individual encounters that specific pathogen in the future. Future infection with that specific pathogen results in a rapid and robust immune response that controls the proliferation and dissemination of the pathogen and significantly reduces disease pathology in the immunized individual.

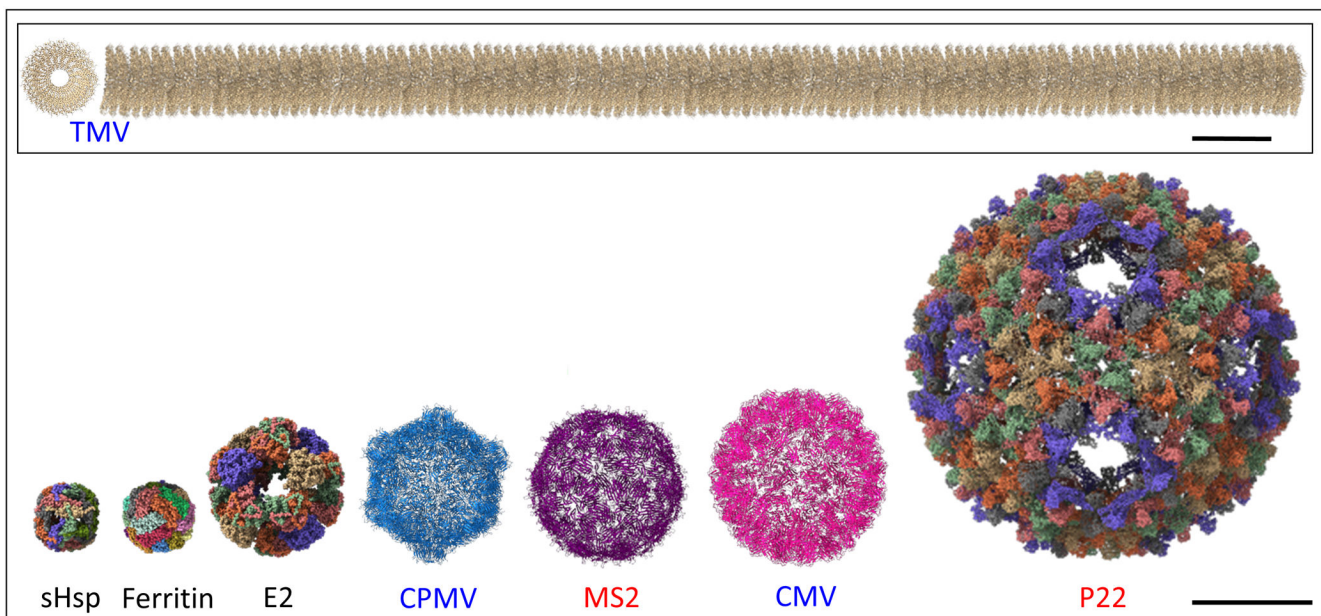


Figure 2. Crystal structures of several protein NP scaffolds investigated for vaccine development. Self-assembling, phage-derived, and plant-derived NPs are labeled in black, red, and blue, respectively. The inset shows TMV from the cross-sectional view (left) and side view (right). Scale bars denote 20-nm. NP structures were generated in ChimeraX using Protein Data Bank ID codes 1shs (sHsp), 1mfr (ferritin), 1b5s (E2), 5fmo (CPMV), 2ms2 (MS2), 1F15 (CMV), 3iyh (P22), and 6R7M (TMV).

Table 1.
Immune responses elicited by self-assembling protein nanoparticle vaccines.

Abbreviations: Ab (antibodies), CD4⁺ T cell (T helper cell), CD8⁺ T cell (cytotoxic T cell), CTB (bacterial cholera toxin B subunit), DC (dendritic cell), FCA (Freud's complete adjuvant), FIA (Freud's incomplete adjuvant), HFMD (Hand, Foot, and Mouth Disease), ID (intradermal injection), IFN- γ (interferon gamma), IgA (immunoglobulin A), IgG (immunoglobulin G), IM (intramuscular injection), IP (intraperitoneal injection), M Φ (macrophage), SC (subcutaneous injection), sHSP (small heat-shock protein), T_{FH} (T follicular helper), TNF- α (tumor necrosis factor alpha).

Scaffold Platform	Type of infectious disease	Disease	In vitro, In vivo	Route of Administration	Adjuvant co-administered	Elicited immune responses	Specific immune responses	References
sHSP	Viral	Influenza	In vivo	IN	sHsp, P22	Humoral	Antigen-specific Abs (IgG and IgA), protection against challenge	(Richert et al., 2012), (Richert et al., 2013)
						Cellular	T _{FH} and CD4 ⁺ T cell expansion, local DC and alveolar M Φ accumulation	
Ferritin	Viral	HIV-1	In vitro, In vivo	IM	Adjuvex, MPLA, ISCOMATRIX	Humoral	Neutralizing Abs	(Georgiev et al., 2018), (Slieden et al., 2015), (He et al., 2016)
		Influenza	In vitro, In vivo*	IM	Adjuvex	Humoral	Neutralizing Abs	
		Chronic hepatitis B	In vivo	SC	None	Humoral	IgG Abs, antigen-specific Abs, protection against challenge	(Wang et al., 2020)
		Hepatitis C	In vitro, In vivo	IP	Inject Alum	Humoral	Antigen-specific Abs, neutralizing Abs	(Yan et al., 2020)
		Rotavirus	In vivo	Oral	CTB	Humoral	IgA and IgG Abs, protection against challenge	(Li et al., 2019)
		HFMV	In vitro, In vivo	SC	FCA or FIA	Humoral	IgG Abs, neutralization titers, protection against challenge	(Wang et al., 2019c)
		Epstein-Barr virus infection	In vitro, In vivo	IM	SAS	Humoral	Neutralizing Abs, protection against challenge	(Kanekiyo et al., 2015)
		Respiratory syncytial virus infection	In vivo	IM	AF03	Humoral	Antigen-specific Abs, neutralizing Abs	(Swanson et al., 2020)

Scaffold Platform	Type of infectious disease	Disease	In vitro, In vivo	Route of Administration	Adjuvant co-administered	Elicited immune responses	Specific immune responses	References
		MERS	In vivo	IM	MF59	Humoral	IgG Abs	(Kim et al., 2018)
						Cellular	Expansion of CD4 ⁺ T cells that secreted IFN- γ and TNF- α	
E2	Viral	HIV-1	In vitro, In vivo	SC, IM, ID	IFA, FIA	Humoral	IgG Abs, neutralizing Abs,	(Caivano et al., 2010), (Krebs et al., 2014), (He et al., 2016)
						Cellular	CD8 ⁺ but not CD4 ⁺ T cells secreted IFN- γ , B cell activation	
	Dengue	In vitro, In vivo	IM, ID	Adjuplex	Humoral	Neutralizing Abs, protection against challenge	(McBurney et al., 2016)	
	Parasitic	Malaria	In vivo	Not specified	FCA or FIA	Humoral	Antigen-specific Abs	(Domingo et al., 2001)

* Asterisk indicates Phase I clinical trial.

Table 2.
Immune responses elicited by bacteriophage-derived protein nanoparticle vaccines.

Abbreviations: Abs (antibodies), CT (cholera toxin), CTL (cytotoxic T cell lymphocyte), FIA (Freud's incomplete adjuvant), FMDV (foot and mouth disease virus), IFN- γ (interferon gamma), IgG (immunoglobulin G), IL (interleukin), IM (intramuscular injection), IN (intranasal injection), IP (intraperitoneal injection), IT (intratracheal), IV (intravenous), MPLA (monophosphoryl lipid A), PO (perorally), respiratory syncytial virus (RSV), SC (subcutaneous), SE (stable emulsion of GLA-SE TLR-4 agonist), TCI (transcutaneous immunization), Th1 (T helper 1), Th17 (T helper 17).

Scaffold Platform	Type of infectious disease	Disease	In vitro, In vivo	Route of Administration	Adjuvant co-administered	Elicited immune responses	Specific immune responses	References
P22	Viral	Influenza	In vivo	IN, IT	None	Humoral	IgG Abs, protection against challenge	(Patterson et al., 2013), (Sharma et al., 2020)
						Cellular	CTL response	
Q β	Viral	FMDV	In vitro, in vivo	Not specified (guinea pigs)	None	Humoral	Antigen-specific Abs	(Skamel et al., 2014)
		HIV-1	In vitro, in vivo	IM	Adjuplex	Humoral	Antigen-specific Abs, neutralization Abs	(Purwar et al., 2018)
		Influenza	In vivo, in vitro	SC, IM	Alum	Humoral	IgG Abs, neutralization Abs, protection against challenge	(Jegerlehner et al., 2013), (Skibinski et al., 2013)
						Cellular	Specific Th1 response, cytokine release (IFN- γ)	(Skibinski et al., 2013)
		In vivo*	IM	Alum	Humoral	Antigen-specific antibodies, no serious adverse events	(Low et al., 2014)	
					Cellular	Cytokine release (IFN- γ , IL17A, IL-17F, IL-9, IL-5, IL-13, IL-21), specific CD4 ⁺ and CD8 ⁺ T cell proliferation	(Low et al., 2014), (Skibinski et al., 2018)	
		Parasitic	Malaria	In vivo	IP, SC	Alum, SE, Montanide	Humoral	IgG Abs, protection against challenge
Leishmaniasis	In vitro, in vivo		SC	None	Humoral	IgG Abs, protection against challenge	(Moura et al., 2017)	

Scaffold Platform	Type of infectious disease	Disease	In vitro, In vivo	Route of Administration	Adjuvant co-administered	Elicited immune responses	Specific immune responses	References
λ	Viral	Hepatitis B	In vivo, in vitro	IM	None	Humoral	Antigen-specific Abs, protection against challenge	(March et al., 2004) and (Clark et al., 2011)
						Cellular	Specific lymphocyte proliferation	(Clark et al., 2011)
T4	Bacterial	Anthrax	In vivo	IM, TCI	Alhydrogel, heat-labile enterotoxin, MPLA	Humoral	IgG Abs, protection against challenge	(Peachmann et al., 2012)
		Anthrax and plague	In vivo	IM	None	Humoral	IgG Abs, neutralization Abs, protection against challenge	(Tao et al., 2018)
MS2	Viral	FMDV	In vitro, in vivo	IM	FIA	Humoral	Neutralizing Abs, protection against challenge	(Dong et al., 2015)
		HPV	In vivo, in vitro	IM, PO	FIA, liquid CT, MPLA	Humoral	IgG Abs, neutralizing Abs, protection against challenge	(Tumban et al., 2012), (Tyler et al., 2014), and (Zhai et al., 2017)
		RSV	In vivo	IN	None	Humoral	Protection against challenge	(Schwarz et al., 2016)
Cellular	Specific splenocyte proliferation							
Filamentous (fd)	Fungal	<i>Candida albicans</i> (yeast infections)	In vivo, in vitro	Not specified	None	Humoral	IgG Abs	(Wang et al., 2014)
		Cellular	Specific splenocyte proliferation, cytokine release (IFN-γ, IL-2, IL-12, and IL-17)					
		Sporotrichosis	In vivo, in vitro	IV	None	Humoral	IgG Abs, protection against challenge Th1 and Th17 proliferation	(Chen et al., 2017)

* Asterisk indicates Phase I clinical trial.

Table 3.
Immune responses elicited by plant virus-derived protein nanoparticle vaccines.

Abbreviations: Abs (antibodies), AIMV (alfalfa mosaic virus), BE (back of ear), CD4⁺ T cell (helper T cell), CD8⁺ T cell (cytotoxic T cell), CMV (cucumber mosaic virus), CpG 1826 (Oligodeoxynucleotide 1826), CPMV (cowpea mosaic virus), CT (cholera toxin), CTL (cytotoxic T cell lymphocyte), DC (dendritic cells), dIC (Polyinosinic-polycytidilic acid), di-GMP (Cyclic diguanylate monophosphate), DOPS (dioleoyl phosphatidylserine), FCA (Freud's complete adjuvant), FIA (Freud's incomplete adjuvant), FMDV (foot and mouth disease virus), HCV (Hepatitis C virus), HIV-1 (human immunodeficiency virus 1), IFN- γ (interferon-gamma), IL (interleukin), IM (intramuscular injection), IN (intranasal), IP (intraperitoneal injection), ISCOM (immune stimulating complex), IV (intravenous), LCMV (lymphocytic choriomeningitis virus), MCT (microcrystalline tyrosine), M Φ (macrophage), PO (per-orally), PapMV (papaya mosaic virus), PVX (potato virus X), RSV (respiratory syncytial virus), SC (subcutaneous), TBSV (tomato bushy stunt virus), TMV (tomato mosaic virus).

Scaffold Platform	Type of infectious disease	Disease	In vitro, In vivo	Route of Administration	Adjuvant co-administered	Elicited immune responses	Specific immune response	References
CPMV	Bacterial	Staph infection	In vivo	IN, SC	ISCOM Matrix, FCA, FIA, QS-21	Humoral	Neutralizing Abs, opsonizing Abs	(Brannan et al., 1999a), (Brennan et al., 1999c), (Rennermalm et al., 2001)
		<i>Pseudomonas</i> infection	In vivo	SC	FCA, FIA, Alum	Humoral	Opsonizing Abs, Protection upon challenge	(Brennan et al., 1999b)
		Group B <i>Streptococcus</i>	In vivo	SC, IP	FCA, FIA	Humoral	Antigen-specific Abs	(Pomwised et al., 2016)
	Viral	HIV-1	In vivo	IN, PO, SC	Alum, FCA, Quil A, AdjuPrime, Ribi, CT	Humoral	Neutralizing Abs	(McInerney et al., 1999), (Durrani et al., 1998)
						Cellular	Cytokine release (IFN- γ)	(Pomwised et al., 2016)
							Cellular	Antigen-specific T cell proliferation
PVX	Viral	HIV-1	In vivo	IN, IP	None	Humoral	Neutralizing Abs	(Marusic et al., 2001)
		Influenza A (H1N1)	In vivo	SC	FIA	Cellular	Antigen-specific CD8 ⁺ T cell activation	(Lico et al., 2009)
		HCV	In vitro, In vivo	SC	GERBU adjuvant MM	Humoral	Antigen-specific Abs	(Uhde-Holzem et al., 2010)
TMV	Bacterial	<i>Pseudomonas</i> infection	In vivo	SC, IM	FCA, FIA, Aluminum hydroxide	Humoral	Opsonizing Abs, protection	(Staczek et al., 2000)

Scaffold Platform	Type of infectious disease	Disease	In vitro, In vivo	Route of Administration	Adjuvant co-administered	Elicited immune responses	Specific immune response	References
							against challenge	
		<i>Francisella tularensis</i>	In vivo, ex vivo	SC, IN	CpG 1826, Addavax, dIC, di-GMP	Humoral	Antigen-specific Abs, protection against challenge	(Banik et.al, 2015), (Mansour et.al, 2018), (McCormick et.al, 2018)
		Anthrax	In vivo	IP	None	Humoral	Neutralizing Abs	(McComb et. al, 2015)
		<i>Yersinia pestis</i>	In vivo	IN	None	Humoral	Antigen-specific Abs, protection against challenge	(Arnaboldi et. al, 2016)
	Viral	Influenza (H1N1)	In vivo	SC	Alum, Addavax	Humoral	Antigen-specific Abs, protection against challenge	(Mallajosyula et al., 2014)
		FMDV	In vivo	IP, BE	Aluminum hydroxide	Humoral	Antigen-specific Abs, protection against challenge	(Jiang et al., 2006)
CMV	Viral	HCV	In vitro, In vivo	SC, IM	FIA	Humoral	Antigen-specific Abs	(Piazzola et al., 2005), (Nuzzaci et al., 2007)
						Cellular	Cytokine release (IFN- γ , IL-12, and IL-15)	(Piazzola et al., 2005), (Nuzzaci et al., 2007)
		<i>Zika virus</i>	In vitro, In vivo	IM	DOPS	Humoral	Neutralizing Abs	(Cabral-Miranda et. al, 2019)
	Parasitic	Malaria	In vivo	IM	MCT, Alum	Humoral	Antigen-specific Abs, protection against challenge	(Cabral-Miranda et. al, 2017)
						Cellular	CD4 ⁺ and CD8 ⁺ T cell responses	(Cabral-Miranda et. al, 2017)
	AIMV	Viral	RSV	In vivo	IM	None	Humoral	Antigen-specific Abs
Cellular							CD4 ⁺ and CD8 ⁺ T cell responses	(Yusibov et al., 2005)
Parasitic		Malaria	In vivo *	IM	Alhydrogel	Humoral	Neutralizing Abs	(Jones et al., 2013) (Chichester et. al, 2018)
TBSV	Viral	HIV-1	In vitro, In vivo	SC	None	Humoral	Antigen-specific Abs	(Joelson et al., 1997)

Scaffold Platform	Type of infectious disease	Disease	In vitro, In vivo	Route of Administration	Adjuvant co-administered	Elicited immune responses	Specific immune response	References
PapMV	Viral	Influenza	In vitro, In vivo	SC, IP, IM	Alum	Humoral	Antigen-specific Abs, B cell expansion, protection against challenge	(Denis et al., 2008), (Hanafi et al., 2010), (Bolduc et al., 2018), (Carignan et al., 2015), (Therien et al., 2017)
						Cellular	Antigen-specific CD8 ⁺ T cell expansion and response	(Babin et al., 2013), (Leclerc et al., 2007), (Hanafi et al., 2010), (Laliberte-Gagne et al., 2019)
		HCV	In vivo	SC	None	Humoral	Antigen-specific Abs	(Denis et al., 2007)
		LCMV	In vivo	IV	None	Cellular	Antigen-specific CD8 ⁺ T cell activation, protection against challenge	(Lacasse et al., 2008)

* Asterisk indicates Phase I clinical trial.

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Table 4.
Clinical trial progress of recombinant viral vector vaccine platforms.

Abbreviations: AAV (adeno-associated virus vector), Ad (adenovirus vector), FDA (US Food and Drug Administration), Hepatitis B surface antigen (HBsAg), HIV (human immunodeficiency virus), HPV (human papillomavirus), RTS,S (RTS,S/AS01 malaria vaccine (trade name Mosquirix™)), TB (tuberculosis), VSV (vesicular stomatitis virus).

Scaffold Platform	Type of infectious disease	Disease	Clinical Status	References	
HBsAg	Parasitic	Malaria	Pre-clinical, Phase 1, Phase 2, Phase 3, FDA approved	(RTS,S Clinical Trials Partnership, 2015) (Dobano et al., 2019a) (Doolan et al., 2009) (Vrieze, 2019) (Asante et al., 2020) (Dobano et al., 2019b) (Sanchez et al., 2020)	
Ad	Bacterial	TB	Pre-clinical, Phase 1, Phase 2	(Matthews et al., 2013) (Ura & Shimada, 2014) (van Zyl-Smit et al., 2017)	
		Viral	HIV	Pre-clinical, Phase 1, Phase 2	(Choi & Chang, 2013) (Matthews et al., 2013) (Ura et al., 2014)
			Influenza	Pre-clinical, Phase 1, Phase 2	(Ura & Shimada, 2014) (Lauer et al., 2017) (Liebowitz et al., 2020)
			SARS-CoV-2	Pre-clinical, Phase 1, Phase 2, Phase 3	(BioRender, 2020) (Zhu et al., 2020)
	Ebola virus	Pre-clinical, Phase 1, Phase 2	(Lauer et al., 2017) (Rauch et al., 2018)		
Parasitic	Malaria	Pre-clinical, Phase 1, Phase 2	(Choi & Chang, 2013) (Matthews et al., 2013) (Ramezanzpour et al., 2016)		
AAV	Viral	HIV	Pre-clinical, Phase 1, Phase 2	(Ura & Shimada, 2014) (Nieto & Salvetti, 2014) (Ramezanzpour et al., 2016) (Vardas et al., 2010)	
		HPV	Pre-clinical, Phase 1	(Ura & Shimada, 2014) (Nieto & Salvetti, 2014) (ClinicalTrials.gov, 2020a)	
		Influenza (H1N1)	Pre-clinical	(Ura & Shimada, 2014) (Nieto & Salvetti, 2014) (Demminger et al., 2020)	
		Hepatitis C	Pre-clinical, Phase 1, Phase 2	(Zhu et al., 2019) (Naso et al., 2017)	
VSV	Viral	Ebola virus	Phase 1, Phase 2, Phase 3, FDA approved	(Rauch et al., 2018) (FDA, 2020a) (Ollmann, 2020)	
Alphavirus	Viral	HIV	Pre-clinical, Phase 1, Phase 2	(Choi & Chang, 2013) (Ramezanzpour et al., 2016)	
		Influenza	Pre-clinical, Phase 1, Phase 2	(Choi & Chang, 2013) (Ramezanzpour et al., 2016)	
Poxvirus	Viral	HIV	Pre-clinical, Phase 1, Phase 2, Phase 3	(Choi & Chang, 2013) (Ura & Shimada, 2014) (Pitisuttithum et al., 2020) (Laher et al., 2020) (Levy et al., 2020)	
		Smallpox	Pre-clinical, Phase 1, Phase 2, Phase 3	(Ura & Shimada, 2014) (Pitman et al., 2019)	
		Influenza	Pre-clinical, Phase 1, Phase 2	(Lauer et al., 2017) (Kreijtz et al., 2014)	
		TB	Pre-clinical, Phase 1	(Choi & Chang, 2013) (Minhinnick et al., 2016) (Wilkie et al., 2020)	
	Parasitic	Malaria	Pre-clinical, Phase 1, Phase 2	(Choi & Chang, 2013) (de Barra et al., 2014)	