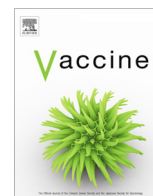




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## Platform technologies for modern vaccine manufacturing



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### ABSTRACT

Improved understanding of antigenic components and their interaction with the immune system, as supported by computational tools, permits a sophisticated approach to modern vaccine design. Vaccine platforms provide an effective tool by which strategically designed peptide and protein antigens are modularized to enhance their immunogenicity. These modular vaccine platforms can overcome issues faced by traditional vaccine manufacturing and have the potential to generate safe vaccines, rapidly and at a low cost. This review introduces two promising platforms based on virus-like particle and liposome, and discusses the methodologies and challenges.

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### 1. Advancing from traditional vaccine production

Vaccination continues to be a leading defense strategy against infectious pathogens. Traditional vaccines that employ whole-cell antigens to raise an immune response have been irrefutably successful in the control or localized eradication of diseases such as poliomyelitis, measles, mumps, rubella, influenza and hepatitis A and B [1–3]. Eradication of smallpox was declared in 1980 after a global immunization effort by WHO [4]. Rinderpest was the second disease globally eradicated by traditional vaccine means as declared by the World Organization for Animal Health in 2011 [5]. Despite this success, live attenuated and inactivated vaccines possess several major drawbacks. Both live attenuated and inactivated vaccines require the production of large volumes of pathogens in the form of viruses and bacteria. This lengthy culturing process contributes to a considerable lag time between antigen production and vaccine delivery. Furthermore, it demands specialized containment facilities and poses considerable risk to the operators and environment due to the infectious nature of the material [6,7]. Despite adequate passaging to diminish virulence, live attenuated pathogens are capable of reverting to virulent strains as evidenced with simian immunodeficiency virus [8], African horse sickness [9] and infectious bronchitis virus vaccines [10]. The genuine threat of vaccine-derived polio associated with Sabin's oral polio vaccine has hindered immunization programs worldwide [11,12]. Inactivated polio vaccine has less of a biosafety risk to vaccine recipients as inactivated poliovirus is incapable of replication, thereby eliminating the possibility of vaccine-derived polio.

However, inactivation of microorganisms can compromise the native conformation of antigenic epitopes resulting in reduced immunogenicity [13]. Pathogens that display high levels of antigenicity owing to high mutation rates (e.g. RNA viruses such as influenza and human immunodeficiency virus [14,15]) or existing as multiple genotypes and serotypes (e.g. rotavirus [16,17], enterovirus [18] and the Group A Streptococcus [19]) present a challenge for developing efficacious vaccines. While this is an important consideration for all vaccine manufacturing platforms, the current timescale of traditional vaccine manufacturing highlights their inadequacy.

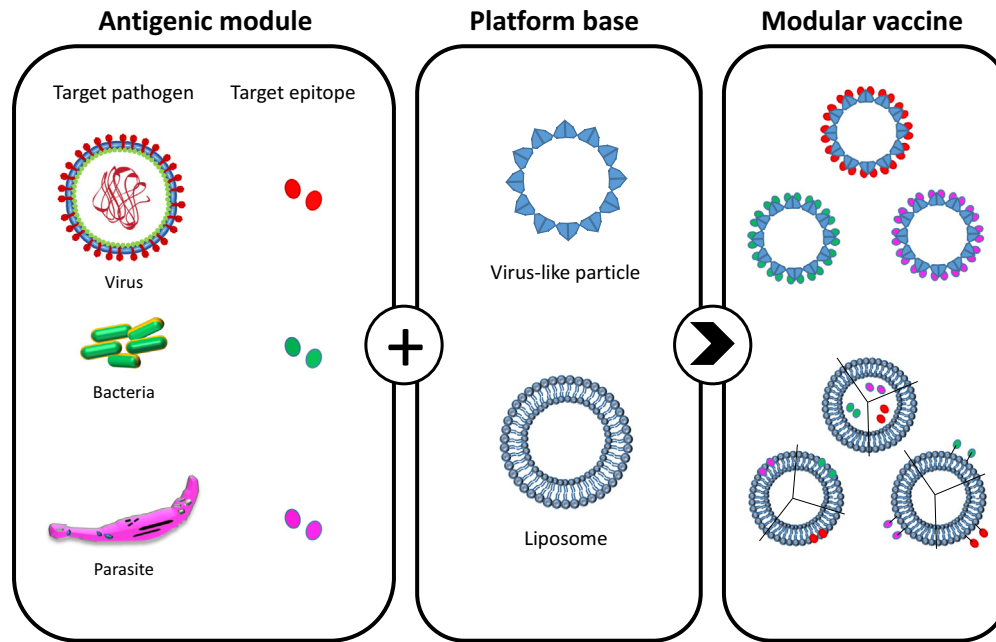
Outbreaks of H1N1 influenza, Middle East Respiratory Syndrome, Ebola and Zika over the last decade, are timely reminders that improved modern vaccine technology is necessary to shorten the developmental and production time of vaccines. Vaccine platform technologies, the formulation of antigens of choice with a pre-defined platform base, have the potential to address vaccine manufacturing challenges such as speed, safety and efficacy. Platforms based on virus-like particle (VLP) and liposomes are discussed, with a focus on the challenges and opportunities offered by these vaccine platform technologies.

### 2. Modular vaccine approach

A tailorable platform that supports safe and simple manufacture of target antigens at high capacity has the potential to rapidly respond to an emerging disease. Most vaccine platform technologies consist of a platform base carrier (Fig. 1) that is amendable to modularization with target antigenic components of pathogens (known as modules). Independently, these components exhibit weak immunogenicity and poor stability. To harness the

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**Fig. 1.** Modularization of target epitopes onto VLP and liposome vaccine platforms. Antigenic modules from a variety of microorganisms may be modularized onto the surface of VLPs through electrostatic interaction, chemical conjugation or genetic fusion. In liposomes, these antigenic modules may be encapsulated into the aqueous core, adsorbed into the lipid bilayer or conjugated (both covalently or non-covalently) to the vesicle surface.

immunostimulatory properties of such antigens, platform carriers are engineered and developed to enhance the antigenicity but without the infectious trait of pathogens. Such engineering also allows the production of novel vaccine candidates that cannot be obtained through traditional methods (attenuation and inactivation). Basic research to determine suitable modules with antigenic potential is a prerequisite of this modular approach, yet the use of generic platforms supports streamlined and standardized vaccine development, potentially reducing the cost of development.

A well-exploited platform is based on VLP technology. VLPs are highly ordered structures, with varying degrees of complexity, which stimulate both innate and adaptive immune responses [20,21]. These intrinsic properties contributed to the commercialization of VLP-based vaccines against human papillomavirus (HPV), hepatitis B and E [22–24]. The self-adjuvanting properties of VLPs, due to their particulate structure and optimal size for uptake by antigen presenting cells [20,25], makes them an attractive tool for increasing the immunogenicity of antigens. Antigens encapsulated within VLPs can also be used as vectors for drug delivery [26]. Well reported platforms based on self-assembling proteins include HPV L1 [27], Hepatitis B core [28] or surface antigen [29,30], murine polyomavirus VP1 [31,32] and bacteriophages MS2 [33], AP205 [34,35] and Q $\beta$  [36]. High antigen-specific antibody titers and protective efficacies have been demonstrated across a range of peptide epitopes and protein domains modularized onto these VLP platforms. As reported, a pre-existing immunity against the VLP proteins from previous exposure to the platform does not diminish the immune response against the antigenic modules [37,38]. Mosquirix™ (RTS,S/AS01, GlaxoSmithKline), a protein-based malaria vaccine comprising circumsporozoite protein and Hepatitis B surface antigen, has demonstrated safety and protection in children and infants in a Phase III trial [39], and WHO has recently announced the first pilot studies in sub-Saharan Africa [40].

Liposomes are another favorable vaccine platform owing to their natural ability to induce an immune response [41]. Composed of an aqueous core and a uni- or multilamellar phospholipid

bilayer, these lipid-based vesicles have immense adaptability and parameters with relation to size, charge, lipid, adjuvant composition and antigen presentation are manipulable [42]. As a result of this versatility, liposomal-based platforms are less well-defined than VLP-based platforms. Surface charge of the vesicle is reported to be an important factor that influences the immune response [42–44]. Cationic formulations are considered the most effective tools in liposomal antigen delivery due to their ability to bind antigen presenting cells through electrostatic interactions and form antigen depots at the site of injection [45,46]. The combination of positively charged dimethyloctadecylammonium (DDA) with the immunostimulant, trehalose-6,6-dibehenate (TDB) was engineered for the delivery of the tuberculosis antigen, Ag85B-ESAT-6 [45] and is possibly the best characterized. DDA:TDB is also considered as a potential platform for *Chlamydia* vaccines [47].

### 3. Vaccine design

The strategy for modularizing antigenic peptide or protein module onto the platform base is the key driver for inducing the protective immune response. Maintaining both the native conformational structure of the antigenic module post modularization and the integrity of the immunostimulating platform base are of equal importance. The rules to guide vaccine design are still limited. Although computational simulation tools and structure-based vaccine design are still in their infancy, they offer alternative possibilities to traditional empirical vaccine development [48,49].

Modularization of chosen antigens onto VLPs is achieved through electrostatic interaction [50], chemical conjugation or genetic fusion [51]. Electrostatic interaction requires minimal processing but these non-covalent interactions can be weak and stability is questionable. A variety of linkage chemistries suitable for chemical conjugation result in a more permanent interaction albeit this requires more complex manufacturing processes under potentially harsh conditions that may alter protein structure. Permanent and regular module display is afforded through genetic fusion, eliminating downstream processing yet insertion sites for modules

**Table 1**  
Platform manufacturing technologies for modularization.

Mechanism of Modularization	Advantages and Challenges	Platform	Disease	References
VLP – Molecular insertion	Simple molecular cloning Co-production of platform and module Reproducible module display Identification of insertion site Determination of suitable linkers Limitations on module size Steric hindrance with large modules	Bacteriophage AP205	Influenza (M2)	[84]
		Cucumber Mosaic Virus	Alzheimer's disease (Amyloid $\beta$ )	[85]
		Hepatitis B Core	Newcastle disease virus	[86]
			Malaria (Circumsporozite)	[28]
			Dengue virus type 2 (Envelope domain III)	[87]
		Human Papillomavirus L1 Capsid	Influenza (M2e)	[88]
			Tuberculosis (CFP-10)	[89]
		Murine Polyomavirus	Human respiratory syncytial virus	[27]
			Influenza (M2e)	[32]
			Group A Streptococcus (J8)	[31]
Tobacco mosaic virus	Rotavirus (VP8*)	[38]		
	Poliovirus (type 3)	[90]		
	Foot-and-mouth disease	[91]		
VLP – Conjugation	Conjugation of large modules without affecting VLP assembly Range of conjugation chemistries Quantification of conjugation efficiency Removal of unconjugated material Location of module dependent upon method of conjugation Harsh conditions alter epitope structure	Bacteriophage AP250	Malaria (Circumsporozite)	[34]
		Bacteriophage Q $\beta$	Malaria (Pfs25 / VAR2CSA), Tuberculosis (Ag58A)	[59]
			Malaria (Pfs25 / CIDR)	[58]
		Hepatitis B Core	Influenza (Hemagglutinin)	[92]
		Rabbit Haemorrhagic Disease Virus	Influenza A (M2e)	[93]
		Cationic liposome	Human papillomavirus type 16 (E6)	[94]
			Leishmania	[61] [65]
Liposome – Encapsulated	Module protected from proteases Longer circulation time Low encapsulation efficiency	Cationic liposome	Hepatitis E	[62]
			Duck Tembusu virus	[70]
			Human papillomavirus type 16 (E7)	[75]
Liposome – Surface conjugation	Modularization possible on pre-formed liposomes Range of conjugation chemistries Harsh conditions alter epitope structure Determination of suitable linkers Removal of unconjugated material	DMPC-DMPG-cholesterol-MPL <sup>a</sup>	Human immunodeficiency virus type 1 (gp41)	[95]
		Metallochelating liposome	<i>Candida albicans</i> (Heat shock protein 90)	[66]
		Neutral liposome	Group A Streptococcus	[76]
		Oleoyl liposome	Hepatitis C virus	[96]
		Cationic liposome	Tuberculosis (Ag85B-ESAT-6)	[97]
Influenza (Hemagglutinin)	[98]			
Liposome – Adsorbed	Minimal preparation Lacks control of module orientation or display	Cationic and neutral liposomes		

<sup>a</sup> DMPC, 1,2-dimyristoyl-sn-glycero-3-phosphocholine; DMPG, 1,2-dimyristoyl-sn-glycero-3-phosphorylglycerol; MPL, monophosphoryl lipid A.

can place limitations on antigen size and may be incompatible with VLP assembly. Peptides are more amenable to VLP surface display than large protein domains although conformational structure can be compromised, ultimately affecting the quality of the immune response [52–54]. Displaying large protein domains has the added benefit of presenting multiple epitopes in the correct structure which may increase immunogenicity. However, expression of large genetically-fused antigens is a challenge owing to protein folding errors or compromised VLP formation through steric hindrance [55,56]. To overcome these issues, strategies such as linker designs [56], antigen titration [38,56,57], split-intein conjugation [34,58,59] and a tandem core fusion strategy [60] are implemented to enable ease of large antigen modularization.

For liposomal vaccine platform, antigens can be encapsulated into the hydrophilic aqueous core [61,62], intercalated into the lipid bilayer or surface attached [63]. Successful modularization of antigens up to 150 kDa have been reported [64–66], larger than those described for VLPs. Modularization with surface attached antigens often elicit superior immune responses in comparison to encapsulated antigens perhaps owing to intracellular processing which is possible for the latter [67]. Despite this, encapsulation protects antigens from protease degradation, facilitates longer circulation time and can generate effective immune responses [68–70]. Low encapsulation efficiency is common due to antigen loss from the vesicle during the manufacturing process which involves film extrusion and high shear methods [71]. Incubating antigens with pre-formed liposomes in the presence of 30% v/v ethanol improves encapsulation efficiency [71,72] and may aid a more

streamlined manufacturing process whereby peptides can be encapsulated post-production. Unlike VLP technology, modules cannot be genetically fused to the carrier thus surface exposed antigens rely heavily upon bioconjugate technologies such as covalent conjugation (i.e. palmitoylation). Lipidation can compromise peptide conformation potentially resulting in altered immune responses [73]. Incorporating appropriate linkers between the module and the fatty acid to create spatial separation can address this [74–76]. As demonstrated by Lipotek Pty Ltd [77] and others [66,78], the use of nitrilotriacetic acid (NTA) - histidine conjugation is promising, yet this remains a relatively unexplored area of liposome technology. NTA conjugation offers the opportunity of assembling entire protein domains [66] onto pre-formed liposomes whilst removing costly purification processes. Novel liposomal platforms encapsulate immunostimulants (including diphtheria toxoid and TLR9 agonists) independent of surface attached target antigen [76,79]. This spatial segregation of antigenic components (with the immunostimulant exposed only upon intracellular processing) has been shown to enhance target specific immune responses. Table 1 summarizes manufacturing technologies for modularization.

#### 4. Platform-based vaccine manufacturing

The long and complex vaccine development process (development, testing, regulatory) requires a huge investment of resources which includes time, facilities and money. Vaccine manufacturing

processes are often customized and conducted in dedicated facilities for separate vaccines due to the characteristics of vaccine antigens and safety issues. These factors pose barriers for a fast response that is critical for controlling modern-day disease outbreaks that spread rapidly, as observed for H1N1 influenza in 2009 [80] and most recently Zika [81]. A platform approach for vaccine manufacturing ideally streamline the bioprocess, and shorten vaccine product development and delivery (time to market).

Platform technologies allow the standardization of upstream and downstream processes, given that the platform base remains unchanged. Certainly, processes will need optimization with modularization of different antigenic modules, but vaccine platform technologies provide flexibility and possibility for multi-product facilities. Prior knowledge, experience and production facility set-up is immensely beneficial. Merck Research Laboratories used their prior knowledge and know-hows from developing hepatitis B VLP vaccine (Recombivax) as the key decision factor when choosing to use the same host (*Saccharomyces cerevisiae*) for the production of HPV VLP vaccine (Gardasil) [82]. Similarly, the decision on the choice of adjuvant to formulate HPV VLP was made based on Recombivax.

The desire to lower cost of goods, thus leading to cheaper vaccines in the market has been well discussed and debated in papers and at conferences. The largest vaccine market is in developing countries, where vaccines would have a significant impact on public health, but these low-income countries face vaccine accessibility and affordability challenges. In combination with modular single-use technologies [83], modern vaccine manufacturing based on platform technologies may potentially lower capital and operating costs, resulting in affordable vaccines.

Another benefit of platform technologies is the potential reduction of regulatory burden. The level of proof and documentation required for new antigenic module on the generic platform may lessen as regulatory authorities are well informed by regulatory track records on the platform base. In the scenario of a disease outbreak, a close collaboration with regulatory authorities may lead to fast-track development of a safe and effective vaccine for the public, against an emerging pathogen.

The benefits of VLP and liposome platform technologies are many but perhaps the most significant is their potential to generate multivalent vaccines. Vaccines designed for immunization against multiple strains of an antigenically diverse pathogen are possible through display of different modules on a single platform or formulation of multiple platform products. Future work is expected to optimize the methodologies by which modules are incorporated into each platform to ensure the success of modern vaccines.

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## References

- [1] Cutts FT, Lessler J, Metcalf CJ. Measles elimination: progress, challenges and implications for rubella control. *Expert Rev Vaccines* 2013;12:917–32.
- [2] Roush SWM, Trudy V. Historical comparisons of morbidity and mortality for vaccine-preventable diseases in the United States. *J Am Med Assoc* 2007;298:2155–63.
- [3] Van Panhuis WGB, Shawn, Zadorozhny Vladimir, Lee Bruce Y, Eng Heather, Cross Anne, et al. Contagious diseases in the United States from 18 88 to the present. *New Engl J Med* 2013;369:2153. 6.
- [4] Henderson DA. The eradication of smallpox—an overview of the past, present, and future. *Vaccine* 2011;29(Suppl 4):D7–9.
- [5] Mariner JH, House JA, Mebus CA, Sollod AE, Chibeu D, Jones BA, et al. Rinderpest eradication: appropriate technology and social innovations. *Sci Transl Med* 2012;337:1309–12.
- [6] Uddowla S, Hollister J, Pacheco JM, Rodriguez LL, Rieder E. A safe foot-and-mouth disease vaccine platform with two negative markers for differentiating infected from vaccinated animals. *J Virol* 2012;86:11675–85.
- [7] Steel J, Lowen AC, Pena L, Angel M, Solorzano A, Albrecht R, et al. Live attenuated influenza viruses containing NS1 truncations as vaccine candidates against H5N1 highly pathogenic avian influenza. *J Virol* 2009;83:1742–53.
- [8] Whatmore AC, Cook N, Hall GA, Sharpe S, Rud EW, Cranage MP. Repair and evolution of nef in vivo modulates simian immunodeficiency virus virulence. *J Virol* 1995;69:5117.
- [9] Weyer CT, Grewar JD, Burger P, Rossouw E, Lourens C, Joone C, et al. African horse sickness caused by genome reassortment and reversion to virulence of live, attenuated vaccine viruses, South Africa, 2004–2014. *Emerg Infect Dis* 2016;22:2087–96.
- [10] Zhang Y, Wang HN, Wang T, Fan WQ, Zhang AY, Wei K, et al. Complete genome sequence and recombination analysis of infectious bronchitis virus attenuated vaccine strain H120. *Virus Genes* 2010;41:377–88.
- [11] Nathanson N, Kew OM. From emergence to eradication: the epidemiology of poliomyelitis deconstructed. *Am J Epidemiol* 2010;172:1213–29.
- [12] Bandyopadhyay AG, Garon J, Seib K, Orenstein Wa. Polio vaccination: past, present and future. *Future Microbiol* 2015;10:791–808.
- [13] Fan YC, Chiu HC, Chen LK, Chang GJ, Chiou SS. Formalin inactivation of Japanese encephalitis virus vaccine alters the antigenicity and immunogenicity of a neutralization epitope in envelope protein domain III. *PLoS Negl Trop Dis* 2015;9:e0004167.
- [14] Treanor J. Influenza vaccine — outmaneuvering antigenic shift and drift. *New Engl J Med* 2004;350:218–20.
- [15] Lipsitch M, O'Hagan JJ. Patterns of antigenic diversity and the mechanisms that maintain them. *J R Soc Interface* 2007;4:787–802.
- [16] Miles MG, Lewis KD, Kang G, Parashar UD, Steele AD. A systematic review of rotavirus strain diversity in India, Bangladesh, and Pakistan. *Vaccine* 2012;30 (Suppl 1):A131–9.
- [17] Chung JY, Kim MS, Jung TW, Kim SJ, Kang JH, Han SB, et al. Detection of rotavirus genotypes in Korea 5 years after the introduction of rotavirus vaccines. *J Korean Med Sci* 2015;30:1471–5.
- [18] Xu M, Su L, Cao L, Zhong H, Dong N, Dong Z, et al. Genotypes of the enterovirus causing hand foot and mouth disease in Shanghai, China, 2012–2013. *PLoS ONE* 2015;10:e0138514.
- [19] Steer AC, Carapetis JR, Dale JB, Fraser JD, Good MF, Guilherme L, et al. Status of research and development of vaccines for *Streptococcus pyogenes*. *Vaccine* 2016;34:2953–8.
- [20] Keller SAB, Bauer Monika, Manolova Vania, Muntwiler Simone, Saudan Philippe, Bachmann Martin F. Cutting edge: limited specialization of dendritic cell subsets for MHC class II-associated presentation of viral particles. *J Immunol* 2010;184:26–9.
- [21] Boisgerault F, Moron G, Leclerc C. Virus-like particles: a new family of delivery systems. *Expert Rev Vaccines* 2002;1:101–9.
- [22] Kushnir N, Streatfield SJ, Yusibov V. Virus-like particles as a highly efficient vaccine platform: diversity of targets and production systems and advances in clinical development. *Vaccine* 2012;31:58–83.
- [23] Pils S, Joura EA. From the monovalent to the nine-valent HPV vaccine. *Clin Microbiol Infect* 2015;21:827–33.
- [24] Lua LH, Connors NK, Sainsbury F, Chuan YP, Wibowo N, Middelberg AP. Bioengineering virus-like particles as vaccines. *Biotechnol Bioeng* 2014;111:425–40.
- [25] Manolova VF, Flace Anna, Bauer Monika, Schwarz Katrin, Saudan Philippe, Bachmann Martin F. Nanoparticles target distinct dendritic cell populations according to their size. *Eur J Immunol* 2008;38:1404–13.
- [26] Zdanowicz M, Chroboczek J. Virus-like particles as drug delivery vectors. *Acta Biochim Pol* 2016;63:469–73.
- [27] Murata Y, Lightfoote PM, Rose RC, Walsh EE. Antigenic presentation of heterologous epitopes engineered into the outer surface-exposed helix 4 loop region of human papillomavirus L1 capsomeres. *Virology* 2009;6:81.
- [28] Sällberg M, Hughes J, Jones J, Phillips TR, Milich DR. A malaria vaccine candidate based on a Hepatitis B virus core platform. *Intervirology* 2003;45:350–61.
- [29] Shchelkunov SN, Salyaev RK, Pozdnyakov SG, Rekoslavskaya NI, Nesterov AE, Ryzhova TS, et al. Immunogenicity of a novel, bivalent, plant-based oral vaccine against hepatitis B and human immunodeficiency viruses. *Biotech Lett* 2006;28:959–67.
- [30] Ballou WR. The development of the RTS, S malaria vaccine candidate: challenges and lessons. *Parasite Immunol* 2009;31:492–500.
- [31] Middelberg AP, Rivera-Hernandez T, Wibowo N, Lua LH, Fan Y, Magor G, et al. A microbial platform for rapid and low-cost virus-like particle and capsomere vaccines. *Vaccine* 2011;29:7154–62.
- [32] Wibowo N, Chuan YP, Lua LHL, Middelberg APJ. Modular engineering of a microbially-produced viral capsomere vaccine for influenza. *Chem Eng Sci* 2013;103:12–20.
- [33] Fu Y, Li J. A novel delivery platform based on Bacteriophage MS2 virus-like particles. *Virus Res* 2016;211:9–16.
- [34] Janitzek CM, Matondo S, Thrane S, Nielsen MA, Kavishe R, Mwakalinga SB, et al. Bacterial superglue generates a full-length circumsporozoite protein virus-like particle vaccine capable of inducing high and durable antibody responses. *Malar J* 2016;15:545.
- [35] Pastori C, Tudor D, Diomede L, Drillet AS, Jegerlehner A, Rohn TA, et al. Virus like particle based strategy to elicit HIV-protective antibodies to the alpha-helic regions of gp41. *Virology* 2012;431:1–11.



- [36] Bessa JS, Schmitz Nicole, Hinton Heather J, Schwarz Katrin, Jegerlehner Andrea, Bachmann Martin F. Efficient induction of mucosal and systemic immune responses by virus-like particles administered intranasally: implications for vaccine design. *Eur J Immunol* 2008;38:114–26.
- [37] Chuan YP, Rivera-Hernandez T, Wibowo N, Connors NK, Wu Y, Hughes FK, et al. Effects of pre-existing anti-carrier immunity and antigenic element multiplicity on efficacy of a modular virus-like particle vaccine. *Biotechnol Bioeng* 2013;110:2343–51.
- [38] Tekewe A, Fan Y, Tan E, Middelberg AP, Lua LH. Integrated molecular and bioprocess engineering for bacterially produced immunogenic modular virus-like particle vaccine displaying 18 kDa rotavirus antigen. *Biotechnol Bioeng* 2017;114:397–406.
- [39] Penny MA, Verity R, Bever CA, Sauboin C, Galactionova K, Flasche S, et al. Public health impact and cost-effectiveness of the RTS, S/AS01 malaria vaccine: a systematic comparison of predictions from four mathematical models. *Lancet* 2016;387:367–75.
- [40] WHO. <[http://www.who.int/immunization/research/development/malaria\\_vaccine\\_qa/en/](http://www.who.int/immunization/research/development/malaria_vaccine_qa/en/)>; November 2016.
- [41] Allison AC, Gregoriadis G. Liposomes as immunological adjuvants. *Nature* 1974;252:252.
- [42] Watson DS, Endsley AN, Huang L. Design considerations for liposomal vaccines: influence of formulation parameters on antibody and cell-mediated immune responses to liposome associated antigens. *Vaccine* 2012;30:2256–72.
- [43] Kraaijeveld CAS, Schilham M, Jansen J, Benaissa-Trouw B, Harmsen M, Van Houte AJ, et al. The effect of liposomal charge on the neutralizing antibody response against inactivated encephalomyocarditis and Semliki Forest viruses. *Clin Exp Immunol* 1984;56:509–14.
- [44] Nakanishi TK, Kunisawa Jun, Hayashi Akira, Tsutsumi Yasuo, Kubo Kazuyoshi, Nakagawa Shinsaku, et al. Positively charged liposome functions as an efficient immunoadjuvant in inducing cell-mediated immune response to soluble proteins. *J Control Release* 1999;61:233–40.
- [45] Christensen D, Agger EM, Andreasen LV, Kirby D, Andersen P, Perrie Y. Liposome-based cationic adjuvant formulations (CAF): past, present, and future. *J Liposome Res* 2009;19:2–11.
- [46] Tandrup Schmidt S, Foged C, Korsholm KS, Rades T, Christensen D. Liposome-based adjuvants for subunit vaccines: formulation strategies for subunit antigens and immunostimulators. *Pharmaceutics* 2016;8.
- [47] Yu H, Jiang X, Shen C, Karunakaran KP, Jiang J, Rosin NL, et al. Chlamydia muridarum T-cell antigens formulated with the adjuvant DDA/TDB induce immunity against infection that correlates with a high frequency of gamma interferon (IFN-gamma)/tumor necrosis factor alpha and IFN-gamma/interleukin-17 double-positive CD4+ T cells. *Infect Immun* 2010;78:2272–82.
- [48] He L, Zhu J. Computational tools for epitope vaccine design and evaluation. *Curr Opin Virol* 2015;11:103–12.
- [49] Park MS, Park SY, Miller KR, Collins EJ, Lee HY. Accurate structure prediction of peptide-MHC complexes for identifying highly immunogenic antigens. *Mol Immunol* 2013;56:81–90.
- [50] Gleiter SS, Stubenrauch Kay, Lilie Hauke. Changing the surface of a virus shell fusion of an enzyme to polyoma VP1. *Protein Sci* 1999;8:2562–9.
- [51] Peacey M, Wilson S, Baird MA, Ward VK. Versatile RHDV virus-like particles: incorporation of antigens by genetic modification and chemical conjugation. *Biotechnol Bioeng* 2007;98:968–77.
- [52] Anggraeni MR, Connors NK, Wu Y, Chuan YP, Lua LH, Middelberg AP. Sensitivity of immune response quality to influenza helix 190 antigen structure displayed on a modular virus-like particle. *Vaccine* 2013;31:4428–35.
- [53] Scarselli MA, Aricò Beatrice, Brunelli Brunella, Savino Silvana, Di Marcello Federica, Palumbo Emmanuelle, et al. Rational design of a meningococcal antigen inducing broad protective immunity. *Sci Transl Med* 2011;3:91ra62.
- [54] Dormitzer PR, Grandi C, Rappuoli R. Structural vaccinology starts to deliver. *Nat Rev Microbiol* 2012;10:807–13.
- [55] Chakerian B. Virus-like particles: flexible platforms for vaccine development. *Expert Rev Vaccines* 2007;6:381–90.
- [56] Lua LH, Fan Y, Chang C, Connors NK, Middelberg AP. Synthetic biology design to display an 18 kDa rotavirus large antigen on a modular virus-like particle. *Vaccine* 2015;33:5937–44.
- [57] Cohen J, Nussenzweig V, Vekemans J, Leach A. From the circumsporozoite protein to the RTS,S/AS candidate vaccine. *Human Vaccines* 2010;6:90–6.
- [58] Brune KD, Leneghan DB, Brian JJ, Ishizuka AS, Bachmann MF, Draper SJ, et al. Plug-and-Display: decoration of Virus-Like Particles via isopeptide bonds for modular immunization. *Sci Rep* 2016;6:19234.
- [59] Thrane S, Janitzek CM, Matondo S, Resende M, Gustavsson T, de Jongh WA, et al. Bacterial superglue enables easy development of efficient virus-like particle based vaccines. *J Nanobiotechnol* 2016;14:30.
- [60] Peyret H, Gehin A, Thuenemann EC, Blond D, El Turabi A, Beales L, et al. Tandem fusion of hepatitis B core antigen allows assembly of virus-like particles in bacteria and plants with enhanced capacity to accommodate foreign proteins. *PLoS ONE* 2015;10:e0120751.
- [61] Ravindran R, Maji M, Ali N. Vaccination with liposomal leishmanial antigens adjuvanted with monophosphoryl lipid-trehalose dicorynomycolate (MPL-TDM) confers long-term protection against visceral leishmaniasis through a human administrable route. *Mol Pharm* 2012;9:59–70.
- [62] Kulkarni SP, Thanapati S, Arankalle VA, Tripathy AS. Specific memory B cell response and participation of CD4+ central and effector memory T cells in mice immunized with liposome encapsulated recombinant NE protein based Hepatitis E vaccine candidate. *Vaccine* 2016;34:5895–902.
- [63] Bobbala S, Hook S. Is there an optimal formulation and delivery strategy for subunit vaccines? *Pharm Res* 2016;33:2078–97.
- [64] Davis D, Gregoriadis G. Liposomes as adjuvants with immunopurified tetanus toxoid: Influence of liposomal characteristics. *Immunology* 1987;61:229–34.
- [65] Nagill R, Kaur S. Enhanced efficacy and immunogenicity of 78 kDa antigen formulated in various adjuvants against murine visceral leishmaniasis. *Vaccine* 2010;28:4002–12.
- [66] Mašek J, Bartheldyová E, Turánek-Knotigová P, Škrabalová M, Korvasová Z, Plocková J, et al. Metallochelating liposomes with associated lipophilised norAbuMDP as biocompatible platform for construction of vaccines with recombinant His-tagged antigens: Preparation, structural study and immune response towards rHsp90. *J Control Release* 2011;151:193–201.
- [67] Rao M, Wassef NM, Alving CR, Krzych U. Intracellular processing of liposome-encapsulated antigens by macrophages depends upon the antigen. *Infect Immun* 1995;63:2396–402.
- [68] Taki A, Smooker P. Small wonders—the use of nanoparticles for delivering antigen. *Vaccines (Basel)* 2015;3:638–61.
- [69] Teng X, Tian M, Li J, Tan S, Yuan X, Yu Q, et al. Immunogenicity and protective efficacy of DMT liposome-adjuvanted tuberculosis subunit CTT3H vaccine. *Hum Vaccin Immunother* 2015;11:1456–64.
- [70] Ma T, Liu Y, Cheng J, Liu Y, Fan W, Cheng Z, et al. Liposomes containing recombinant E protein vaccine against duck Tembusu virus in ducks. *Vaccine* 2016;34:2157–63.
- [71] Shariat SB, Badiie Ali, Jaafari Mahmoud Reza, Mortazavi Seyed Alireza. Optimization of a method to prepare liposomes containing HER2/Neu-derived peptide as a vaccine delivery system for breast cancer. *Iran J Pharm Res* 2014;13:15–25.
- [72] Wang CH, Huang YY. Encapsulating protein into preformed liposomes by ethanol-destabilized method. *Artif Cells Blood Sub, Biotechnol* 2003;31:303–12.
- [73] Hickman DT, Lopez-Deber MP, Ndao DM, Silva AB, Nand D, Pihlgren M, et al. Sequence-independent control of peptide conformation in liposomal vaccines for targeting protein misfolding diseases. *J Biol Chem* 2011;286:13966–76.
- [74] Muhs AH, Hickman David T, Pihlgren Maria, Chuard Nathalie, Giriens Valerie, Meerschman Carine, et al. Liposomal vaccines with conformation-specific amyloid peptide antigens define immune response and efficacy in APP transgenic mice. *Proc Natl Acad Sci U S A* 2007;104:9810.
- [75] Chen W, Huang L. Induction of cytotoxic T-lymphocytes and antitumor activity by a liposomal lipopeptide vaccine. *Mol Pharm* 2008;5:464–71.
- [76] Zaman M, Ozberk V, Langshaw EL, McPhun V, Powell JL, Phillips ZN, et al. Novel platform technology for modular mucosal vaccine that protects against streptococcus. *Sci Rep* 2016;6:39274.
- [77] Tyne AS, Chan JG, Shanahan ER, Atmosukarto I, Chan HK, Britton WJ, et al. TLR2-targeted secreted proteins from *Mycobacterium tuberculosis* are protective as powdered pulmonary vaccines. *Vaccine* 2013;31:4322–9.
- [78] Marques-Gallego P, de Kroon AI. Ligation strategies for targeting liposomal nanocarriers. *Biomed Res Int* 2014;2014:129458.
- [79] Hills T, Jakeman PG, Carlisle RC, Klenerman P, Seymour LW, Cawood R. A rapid-response humoral vaccine platform exploiting pre-existing non-cognate populations of anti-vaccine or anti-viral CD4+ T helper cells to confirm B cell activation. *PLoS ONE* 2016;11:e0166383.
- [80] Sullivan SJ, Jacobson RM, Dowdle WR, Poland GA. 2009 H1N1 influenza. *Mayo Clin Proc* 2010;85:64–76.
- [81] Singer M. The spread of Zika and the potential for global arbovirus syndemics. *Glob Public Health* 2017;12:1–18.
- [82] Buckland BC. The process development challenge for a new vaccine. *Nat Med* 2005.
- [83] Lopes AG. Single-use in the biopharmaceutical industry: a review of current technology impact, challenges and limitations. *Food Bioprod Process* 2015;93:98–114.
- [84] Tissot AC, Renhofs R, Schmitz N, Cielens I, Meijerink E, Ose V, et al. Versatile virus-like particle carrier for epitope based vaccines. *PLoS ONE* 2010;5:e9809.
- [85] Vitti A, Piazzolla G, Condelli V, Nuzzaci M, Lanorte MT, Boscia D, et al. Cucumber mosaic virus as the expression system for a potential vaccine against Alzheimer's disease. *J Virol Methods* 2010;169:332–40.
- [86] Zhao Y, Hammond RW. Development of a candidate vaccine for Newcastle disease virus by epitope display in the Cucumber mosaic virus capsid protein. *Biotech Lett* 2005;27:375–82.
- [87] Arora UT, Tyagi Poornima, Swaminathan Sathyamangalam, Khanna Navin. Chimeric Hepatitis B core antigen virus-like particles displaying the envelope domain III of dengue virus type 2 <Arora 20 12 .pdf>. *J Nanobiotechnol* 2010;10.
- [88] De Filette M, Martens W, Smet A, Schotsaert M, Birkett A, Londoño-Arcila P, et al. Universal influenza A M2e-HBc vaccine protects against disease even in the presence of pre-existing anti-HBc antibodies. *Vaccine* 2008;26:6503–7.
- [89] Dhanasooraj D, Kumar RA, Mundayoor S. Vaccine delivery system for tuberculosis based on nano-sized hepatitis B virus core protein particles. *Int J Nanomedicine* 2013;8:835–43.
- [90] Haynes JRC, Cunningham Janet, Von Seefried Adolph, Lennick Michael, Garvin Robert T, Shen Shi-Hsiang. Development of a genetically-engineered, candidate polio vaccine employing the self-assembling properties of the tobacco mosaic virus coat protein. *Bio/Technology* 1986;4:637.
- [91] Wu L. Expression of foot-and-mouth disease virus epitopes in tobacco by a tobacco mosaic virus-based vector\*1. *Vaccine* 2003;21:4390–8.

- [92] Jegerlehner A, Zabel F, Langer A, Dietmeier K, Jennings GT, Saudan P, et al. Bacterially produced recombinant influenza vaccines based on virus-like particles. *PLoS ONE* 2013;8:e78947.
- [93] Jegerlehner A, Schmitz N, Storni T, Bachmann MF. Influenza a vaccine based on the extracellular domain of M2: weak protection mediated via antibody-dependent NK cell activity. *J Immunol* 2004;172:5598–605.
- [94] Jemon K, Young V, Wilson M, McKee S, Ward V, Baird M, et al. An enhanced heterologous virus-like particle for human papillomavirus type 16 tumour immunotherapy. *PLoS ONE* 2013;8:e66866.
- [95] Watson DS, Platt VM, Cao L, Venditto VJ, Szoka Jr FC. Antibody response to polyhistidine-tagged peptide and protein antigens attached to liposomes via lipid-linked nitrilotriacetic acid in mice. *Clin Vaccine Immunol* 2011;18:289–97.
- [96] Takagi A, Kobayashi N, Taneichi M, Uchida T, Akatsuka T. Coupling to the surface of liposomes alters the immunogenicity of hepatitis C virus-derived peptides and confers sterile immunity. *Biochem Biophys Res Commun* 2013;430:183–9.
- [97] Hamborg M, Kramer R, Schante CE, Agger EM, Christensen D, Jorgensen L, et al. The physical stability of the recombinant tuberculosis fusion antigens h1 and h56. *J Pharm Sci* 2013;102:3567–78.
- [98] Barnier-Quer C, Elsharkawy A, Romeijn S, Kros A, Jiskoot W. Adjuvant effect of cationic liposomes for subunit influenza vaccine: influence of antigen loading method, cholesterol and immune modulators. *Pharmaceutics* 2013;5:392–410.