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To cite this article: Henderson Zhu, Christine S. Rollier & Andrew J. Pollard (2021) Recent advances in lipopolysaccharide-based glycoconjugate vaccines, Expert Review of Vaccines, 20:12, 1515-1538, DOI: [10.1080/14760584.2021.1984889](https://doi.org/10.1080/14760584.2021.1984889)

To link to this article: <https://doi.org/10.1080/14760584.2021.1984889>



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Published online: 25 Oct 2021.



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REVIEW



Recent advances in lipopolysaccharide-based glycoconjugate vaccines

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ABSTRACT

Introduction: The public health burden caused by pathogenic Gram-negative bacteria is increasingly prominent due to antimicrobial resistance. The surface carbohydrates are potential antigens for vaccines against Gram-negative bacteria. The enhanced immunogenicity of the O-specific polysaccharide (O-SP) moiety of LPS when coupled to a carrier protein may protect against bacterial pathogens. However, because of the toxic lipid A moiety and relatively high costs of O-SP isolation, LPS has not been a popular vaccine antigen until recently.

Areas covered: In this review, we discuss the rationales for developing LPS-based glycoconjugate vaccines, principles of glycoconjugate-induced immunity, and highlight the recent developments and challenges faced by LPS-based glycoconjugate vaccines.

Expert opinion: Advances in LPS harvesting, LPS chemical synthesis, and newer carrier proteins in the past decade have propelled LPS-based glycoconjugate vaccines toward further development, through to clinical evaluation. The development of LPS-based glycoconjugates offers a new horizon for vaccine prevention of Gram-negative bacterial infection.

ARTICLE HISTORY

Received 09 May 2021
Accepted 21 September 2021

KEYWORDS

Bioconjugation; carrier protein; chemical conjugation; glycoconjugate; lipopolysaccharide; vaccine

1. Introduction to bacterial carbohydrate conjugate vaccines

Outer surfaces of blood-invading bacteria are formed from many components, including glycoproteins and glycolipids, which serve as virulence factors. For Gram-negative bacteria, these include lipopolysaccharide (LPS), capsular polysaccharide (CPS), and exopolysaccharide (EPS) [1] (Figure 1).

The development of vaccines derived from bacterial surface carbohydrates were initially based upon evidence accumulated in the early 20th century. In the 1920s, carbohydrates were first described as immunogens when capsular polysaccharides extracted from *Streptococcus pneumoniae* were shown to precipitate in immunized horse serum [2]. In the 1930s, observations that the loss of responses to skin tests using purified *S. pneumoniae* CPS in participants who recovered from pneumococcal infection pointed to the potential application of bacterial surface carbohydrates as vaccine antigens, which led to the introduction of *S. pneumoniae* CPS-based polysaccharide vaccines [3].

Following on, CPS-based polysaccharide vaccines against *Neisseria meningitidis*, and *Haemophilus influenzae* type b (Hib) were introduced in the late 20th century [4]. The expansion of CPS-based polysaccharide vaccines revealed several limitations through time: (1) carbohydrate vaccines are less immunogenic in infants and the elderly compared with adolescents and adults [5–7]; (2) antibody responses produced by the CPS vaccines waned quickly [8,9]. To address these problems, bacterial surface carbohydrates were chemically coupled to carrier

proteins to enhance immunogenicity and to induce immunological memory [10]. These glycoconjugate vaccines induce a T cell-dependent immune response which involves the formation of memory B cells against the antigen [11]. Excellent safety and immunogenicity of CPS-based glycoconjugate vaccines in infants were shown in several earlier clinical studies [12–14].

Vaccines against bacterial pathogens had been undergoing a continuing shift from carbohydrate-based vaccines to glycoconjugate vaccines over the past decades [7]. Glycoconjugate vaccines against Hib received licensure in industrialized countries including the UK and the USA in the late 1980s and the early 1990s, with the inclusion into routine immunization for infants [15]. To date, many CPS-based conjugate vaccines were licensed for childhood vaccinations in the EU and the US (Table 1). Broad immunization in low-and-middle-income countries was made available later with support from the Gavi Alliance, which impacted significantly on the reduction of Hib- and pneumococcus-related mortality across the African meningitis belt [16,17].

While CPS-based glycoconjugate vaccines have a positive impact on global health, many other Gram-negative bacteria do not have a capsule and therefore a non-CPS-based glycoconjugate vaccine is needed. Also, antimicrobial resistance (AMR) driven by the use of broad-spectrum antibiotics has highlighted the need for vaccines against Gram-negative bacteria [20]. LPS stands out as an alternative vaccine antigen to address these problems. LPS is an outer membrane glycolipid of Gram-negative

Article highlights

- Lipopolysaccharide (LPS) is a weak vaccine antigen but can be made immunogenic by chemically conjugating with a carrier protein. Glycoconjugate vaccines have enhanced immunogenicity compared to plain polysaccharides in young and elderly age groups due to T-cell mediated immunity.
- New technologies for obtaining LPS, as well as novel carrier proteins, have generated LPS-based glycoconjugate vaccine candidates with promising yields and immunogenicity.
- Conjugation methods are increasingly well-understood, while some conjugation methods will result in glycoconjugates of random conformations, conformation-specific conjugation methods are gaining more attention.
- Several immunological caveats are present with glycoconjugate vaccines, such as pre-existing immunity to the carrier protein and the reduced immunogenicity when concomitantly administering glycoconjugate vaccines.

bacteria that facilitates immune evasion, inflammation, and anti-microbial resistance [21]. While LPS alone elicits weak immune responses and shares the same immunological inadequacies of other plain polysaccharides, LPS-based glycoconjugate vaccine candidates have demonstrated promising levels of immunogenicity against unencapsulated bacteria that have been confirmed to develop AMR, including *Shigella* spp, enteropathogenic *Escherichia coli*, and *Pseudomonas aeruginosa* in clinical and pre-clinical investigations [22–25]. Further details on the mechanisms of glycoconjugate-induced immunity and the rationale of using LPS as a vaccine antigen are discussed in the following sections.

2. The role of conjugation in immune response

Two working models are currently present to explain glycoconjugate-induced immunity, both involving the interactions between the glycoconjugate, B cells, and CD4 + T cells (Figure 2). Both working models initiate with the carbohydrate moiety of the glycoconjugate binding to the B cell receptor (BCR). Then the glycoconjugate is endocytosed to be destined for the lysosomal compartments. In the classic working model, glycoconjugates disassemble within the lysosomal compartments, in which the liberated carrier proteins break down into smaller peptides. These peptides bind to class II major histocompatibility (MHCII) complexes in the B cell cytosol and traffic anterogradely to present these peptides for fusion with T cell receptors (TCR) on CD4 + T cells [26]. The binding to TCR facilitates the binding of CD80/86 and CD84, thus activating the production of cytokines such as IL-4 and IL-2, activating the proliferation of carbohydrate-specific B cells and the maturation into memory B cells, resulting in the production of anti-polysaccharide IgG antibodies by B cells [27,28].

The classic working model is based on the knowledge that MHCII complexes cannot bind carbohydrates [29], however, in recent years, a newer working model had gained increased attention. The model suggested that carbohydrates can bind indirectly to MHCII complexes. To achieve that, it was proposed that the carrier protein within the glycoconjugate is enzymatically degraded in the lysosome, generating fragmented glycopeptides held together by the covalent linkage formed in chemical conjugation. The peptide moiety of the

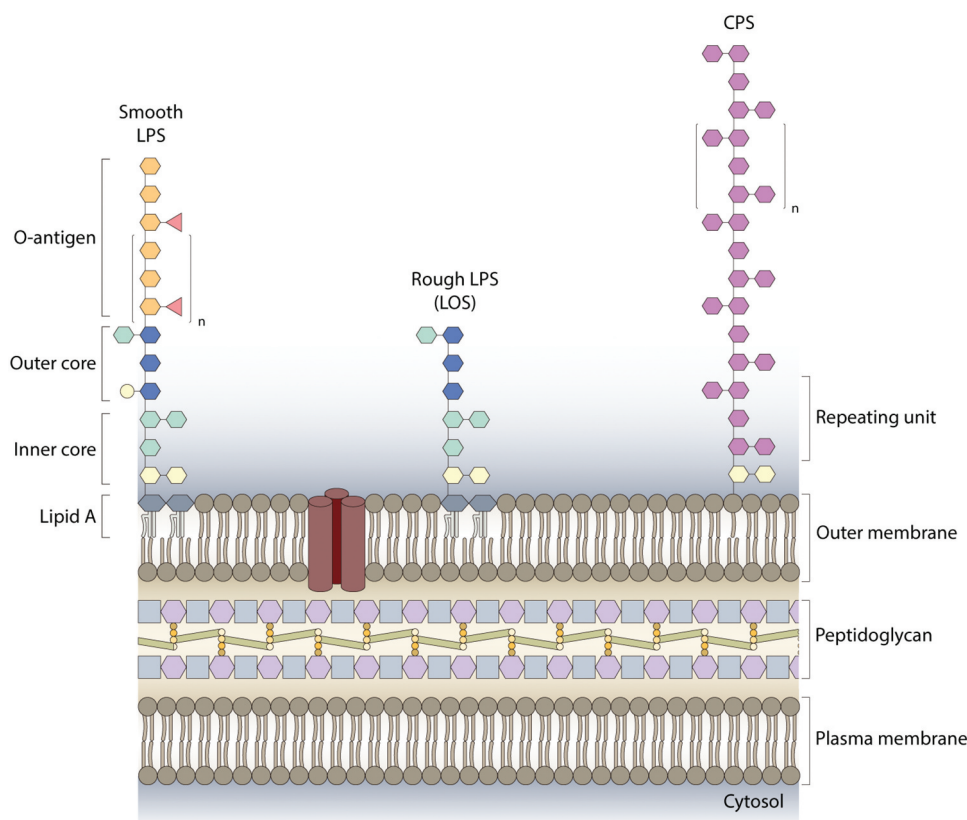


Figure 1. Outer membrane structure of gram-negative bacteria.

Table 1. Licensed CPS-based glycoconjugate vaccines in the EU and the US [18,19].

Pathogen(s) protected with CPS-conjugates	Disease(s) protected not with CPS-conjugates	Carrier protein	Tradename	Manufacturer	Approved in EU and/or US
<i>Neisseria meningitidis</i> serogroups A, C, Y, and W-135	None	CRM197	Menveo	GlaxoSmithKline	US/EU
<i>Neisseria meningitidis</i> serogroups A, C, Y, and W-135	None	Diphtheria toxoid	Menactra	Sanofi Pasteur	US
<i>Neisseria meningitidis</i> serogroups A, C, Y, and W-135	None	Tetanus toxoid	Nimenrix	Pfizer	EU
<i>Neisseria meningitidis</i> serogroups A, C, Y, and W	None	Tetanus toxoid	MenQuadfi	Sanofi Pasteur	US/EU
<i>Streptococcus pneumoniae</i> serotypes 1, 3, 4, 5, 6A, 6B, 7 F, 9 V, 14, 18 C, 19A, 19 F and 23 F	None	CRM197	Prevnar 13	Pfizer	US/EU
<i>Streptococcus pneumoniae</i> serotypes 1, 4, 5, 6B, 7 F, 9 V, 14, 18 C, 19 F and 23 F	None	<i>Haemophilus influenzae</i> proteind D, tetanus toxoid, and diphtheria toxoid	Synflorix	GlaxoSmithKline	EU
<i>Haemophilus influenzae</i> type b	Diphtheria, tetanus, pertussis, poliomyelitis, and hepatitis B	Outer membrane protein complex (OMPC)	Vaxelis	Merck/Sanofi Pasteur	US/EU
<i>Haemophilus influenzae</i> type b	Diphtheria, tetanus, pertussis, poliomyelitis, and hepatitis B	Tetanus toxoid	Hexacima	Sanofi Pasteur	EU
<i>Haemophilus influenzae</i> type b	Diphtheria, tetanus, pertussis, poliomyelitis, and hepatitis B	Tetanus toxoid	Hexyon	Sanofi Pasteur	EU
<i>Haemophilus influenzae</i> type b	Diphtheria, tetanus, pertussis, poliomyelitis, and hepatitis B	Tetanus toxoid	Infanrix Hexa	GlaxoSmithKline	EU
<i>Haemophilus influenzae</i> type b	Diphtheria, tetanus, pertussis, and poliomyelitis	Tetanus toxoid	Pentacel	Sanofi Pasteur	US
<i>Haemophilus influenzae</i> type b	None	Outer membrane protein complex (OMPC)	PedvaxHIB	Merck	US
<i>Haemophilus influenzae</i> type b	None	Tetanus toxoid	ActHIB	Sanofi Pasteur	US
<i>Haemophilus influenzae</i> type b	None	Tetanus toxoid	Hiberix	GlaxoSmithKline	US

glycopeptides acts like an anchor that binds to MHCII complexes, allowing the glycopeptide-bound MHCII to present its carbohydrate moiety to T cell receptors on a unique subset of CD4⁺ T cells, designated as carbohydrate-specific T cells (Tcarbs). Thereafter, the T cell receptor-binding can induce B cell proliferation, maturation, and the production of anti-polysaccharide IgG in an analogous way to the classical working model [27,30]. This newer working model suggested that glycoconjugate vaccines may benefit from a 'sun'-type conformation as supposed to a 'lattice'-type conformation, as the prior would allow the formation of more glycopeptides due to the abundant covalent linkage. Reports have been emerging to suggest that the two mechanisms may be co-existent [27,31]. Thus, further investigations on the biochemistry of Tcarbs can be helpful in elucidating glycoconjugate-induced immunity and optimizing vaccine design [28].

3. LPS as a vaccine antigen

The outer membrane of most Gram-negative bacteria is predominantly made up of LPS [32], which can be further subcategorized into smooth LPS and rough LPS. Smooth LPS is the most common form of LPS, composed of three moieties, a hydrophobic lipid A, a non-repeating rough core oligosaccharide, and a hydrophilic O-specific polysaccharide (O-SP). Rough LPS differentiates from smooth LPS by the absence of O-SP [33] (Figure 1). LPS functions as a virulence factor in several ways. Firstly, the continuum of LPS along the bacterial surface forms a non-fluid permeability barrier that blocks the entrance of certain antimicrobial molecules [34,35]. Also, the long and

outward-projecting O-SP moiety prevents the deposition of complement on the surface of smooth strain bacteria, contributing to immune evasion [36].

In many instances, antibodies targeting LPS confer protection against bacterial infections. *In vitro* studies showed antibodies to the O-Ag (O-SP and the core oligosaccharide) induced protection against *Klebsiella pneumoniae*, *Vibrio cholerae*, *N. meningitidis*, and *Salmonella enterica* by mediating opsonophagocytosis, complement-mediated lysis, and agglutination [37–40]. It has been recently shown that anti-LPS IgG within the mucus layer of the gastrointestinal mucosa can crosslink with mucin to capture *S. Typhimurium* on the mucin mesh, thus inhibiting the flagellum-based bacterial motility [41]. This capturing of bacteria highlighted a new, but potentially conserved mechanism of protection, likely involved in the protection against other enteric pathogens. The neutralizing effect of the O-Ag-specific antibodies is weaker in encapsulated bacteria, such as encapsulated serotypes of *Klebsiella pneumoniae*, because the shielding created by CPS can lower the accessibility of anti-LPS antibodies to the LPS [42].

In higher organisms, the toxicity of LPS is conferred by the lipid A moiety, which can bind to a protein complex formed by Toll-like receptor 4 (TLR4) and myeloid differentiation factor 2 (MD2) [43]. Upon binding with LPS, TLR4 oligomerizes to allow the recruitment of adaptor proteins including myeloid differentiation primary response 88 (Myd88) and TIR domain-containing adaptor inducing IFN- β (TRIF) through the Toll-interleukin-1 receptor (TIR) domains of TLR, which subsequently lead to the activation of the transcription factor

nuclear factor- κ B (NF- κ B), inducing the secretion of proinflammatory cytokines including IL-1, IL-6, and IFN- γ . Release of these cytokines activate signaling cascades which typically cause fever, inflammation, and tissue damage, however in severe cases, might cause hypotension and septic shock [43–45]. Thus, detoxification techniques were developed to convert LPS into vaccine candidates, including alkaline and acid detoxification to degrade or remove lipid A, respectively [46,47]. Between these detoxification methods, alkaline detoxification may result in the incomplete removal of lipid A, which

may also cause incomplete detoxification [48]. Due to this drawback, detoxification of LPS is now more commonly performed by acid hydrolysis, which removes the whole lipid A moiety from the LPS, producing O-Ag. Earlier attempts to detoxify LPS isolated from *E. coli* and *S. Typhimurium* showed low immunogenicity in mice challenge models, highlighting the need for conjugation to suffice potency [49,50].

The development of LPS-based vaccines was also stagnated by the low harvesting efficiency of LPS from bacterial culture. In the 20th century and early 21st century, the extraction of LPS

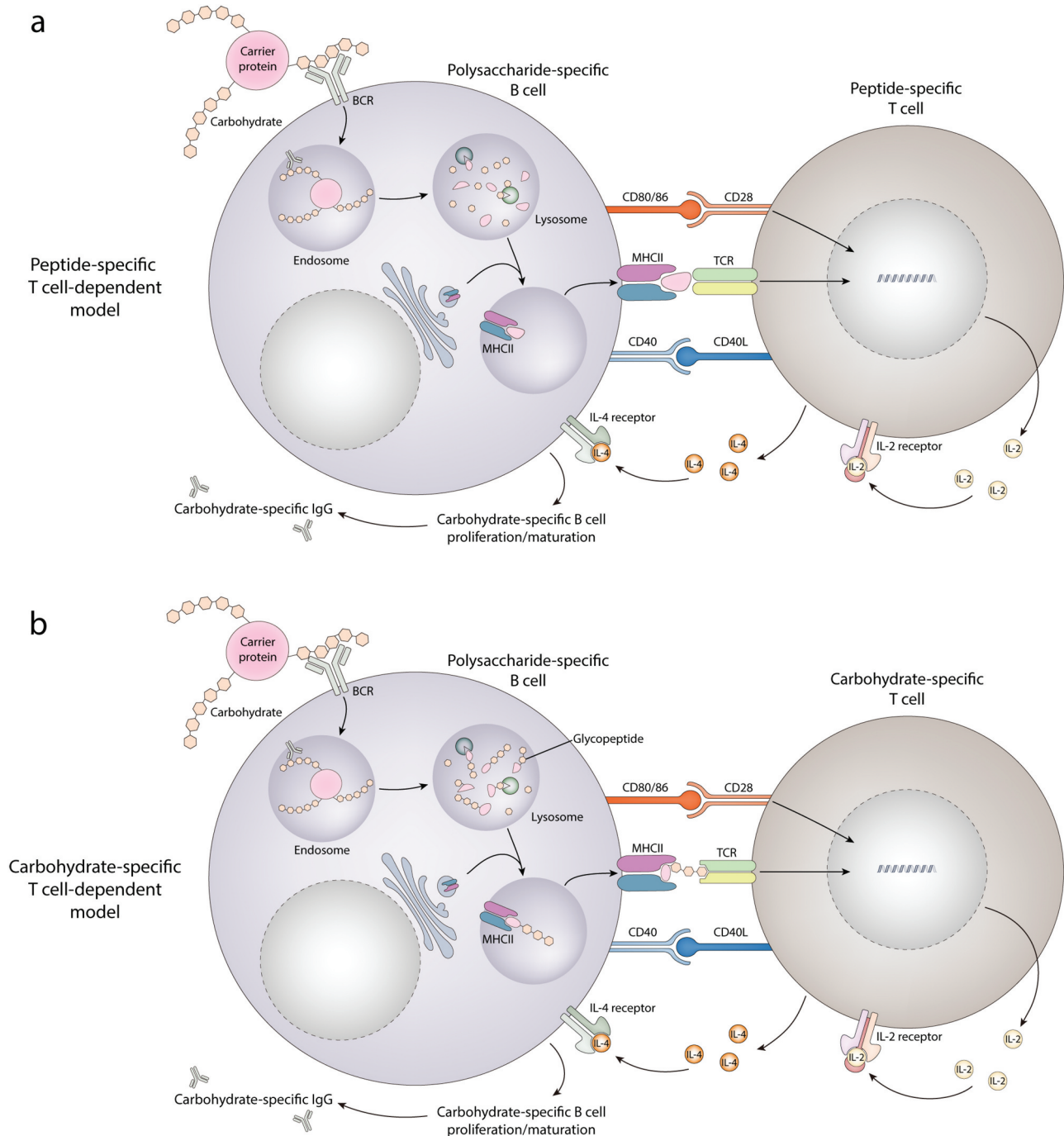


Figure 2. Working models for T-cell dependent immunity induced by glycoconjugate vaccines. Both mechanisms involve BCR-mediated intake of glycoconjugates resulting in the activation of T cells, resulting in the secretion of cytokines inducing B cell proliferation and maturation. (a) The peptide-specific T cell-dependent model involves the presentation of peptide to a peptide-specific T cell. (b) The carbohydrate-specific T cell-dependent model involves the presentation of glycopeptides to carbohydrate-specific T cells. This figure is adapted from Avci et al. 2011.

is a laborious process that made the industrial-scale production of LPS-based derivatives costly. It was only more recently that LPS extraction had been adapted to industrial-scale operations, which will be discussed in the next section [51].

4. Obtaining LPS for generating glycoconjugates

4.1. Harvesting LPS from bacterial culture

The most common method of isolating smooth LPS from whole bacteria in both industrial and laboratory settings is the hot phenol-water method developed by Westphal & Jann [52]. This process involves an initial thermal lysis of suspended ground freeze-dried bacteria, followed by centrifugation and resuspension into a phenol-water mixture (typically 1:1) [53]. Due to the varied hydrophobicity of carbohydrates and proteins, the bacterial material dissolved in the phenol-water mixture partition to either phenol or water depending on their hydrophobicity. The amphipathic LPS, comprising the hydrophilic O-SP domain, is enriched in the aqueous phase, whereas proteins partition into phenol [52]. Most preclinical and clinical LPS-based vaccine candidates using LPS harvested bacterial culture use hot phenol-water method in their production, including O-SP-conjugate vaccines for *V. cholerae*, *S. dysenteriae* type I, *S. flexneri* 2a, and *S. sonnei* [54,55].

Refinements to the hot phenol-water method were made over time to address several aims: to adapt to different scales of operation, to improve yield, and to improve purity [56,57]. Notably, the initial drawback of the hot phenol-water method is the enrichment of contaminants in the preparation, including RNA, DNA, and lipoproteins. These contaminants are difficult to isolate and can confound downstream applications relying on the biological properties of the LPS, such as the production and validation of glycoconjugates. To minimize these contaminants, enzymatic pre-treatments with RNase A, DNase I, and proteinase K before the phenol-water extraction are now standard [58–60]. To achieve an even higher purity of LPS with the aim of removal of remaining lipoproteins, repurification methods has been developed using phenol extraction [61].

Despite achieving a high yield and purity using hot phenol-water extraction, the method is demonstrated to be time-consuming (requires three to four days), costly (involving multiple steps including enzymatic pre-treatment, multiple centrifugations, and detoxification to remove lipid A), and can lead to toxic remnants of phenol. Preliminary work done in the late 1980s and early 1990s proposed another method suitable for the industrial-scale isolation of LPS [62]. The method worked on the principle that the ketosidic bond between the core oligosaccharide and the lipid A moieties is acid-labile; thus, the O-Ag can be liberated after persistent heating under an acidic condition. Due to the increasing demand for LPS in glycoconjugate vaccine production, recent works had adapted the technique to industrial operations by making it compatible with industrial sterilize-in-place fermenters, allowing microbial fermentation and acetic acid hydrolysis to happen sequentially *in situ*, followed by harvesting via centrifugation [51]. The subsequent supernatant can then be purified using a series of tangential flow filtration, cation-exchange chromatography, and acid/base precipitation to remove residual peptides and nucleic acids. The

method was successful in isolating the O-Ag from *S. Paratyphi* A (O:2), *S. Typhimurium* (O:4,5), and *S. Enteritidis* (O:9) in higher quantities [51,63]. This method was proven viable to be used in LPS-based glycoconjugate vaccine production as demonstrated by a vaccine candidate against *S. Typhimurium* (O:4,5), which contained O-Ag conjugated to CRM197. Administering the vaccine candidate in mice induced >10-fold greater increase in anti-O-Ag IgG titers compared with pre-immune serum, while also showing a two-log reduction in bacterial load in the spleen and liver of immunized mice after bacterial challenge [63].

Alternatively, efficient extraction methods for isolating rough LPS were established based on phenol/chloroform/light petroleum ether extraction developed by Galanos, Luderitz & Westphal [64]. This method was used in the pre-clinical development of glycoconjugate vaccines against rough mutant strains of *Escherichia coli*, including strains Rc, R1, R2, R3, R4, and K-12 [65]. However, the outcome of the study suggested that the rough LPS conjugates against *E. coli* strain with the R1 core were not immunogenic in cows. Other methods were also developed to isolate both smooth and rough LPS depending on the bacterial strain used, including the Darveau & Hancock method, which relied on the precipitation of LPS with magnesium chloride and ethanol [66].

4.2. Cell-free harvesting of O-SP through chemical synthesis

The chemical synthesis of LPS mimics is a cell-free, standardized, and scalable alternative to the isolation of LPS from bacterial cell cultures. This approach brings several critical advantages for vaccine production and development, including high reproducibility, avoidance of batch-batch variation seen in bacterial cultures, and free of bacterial remnants [67]. Beyond acting as a vaccine candidate, synthetic O-SP can also act as a tool to determine the immunogenicity of different lengths of O-SP in glycoconjugates and assess the level of immunogenicity elicited by different non-reducing end sugars [68]. Recently, Colombo, Pitirollo & Lay constructed an informative review on the advances in synthetic glycoconjugate vaccines over the past decades [69].

The first breakthrough in glycoconjugates with LPS mimics was demonstrated by an O-SP-mimic-tetanus toxoid conjugate vaccine candidate SF2a-TT15 developed by the Institut Pasteur. The vaccine candidate had three repeating units of a pentasaccharide mimicking the O-SP of *S. flexneri* 2a. In a recent phase 1 study, the vaccine candidate induced a strong antibody response, demonstrated by anti-LPS IgG seroconversion (\geq four-fold rise in antibody titer 28 days after vaccination) in all vaccine recipients, while sustaining a high level of anti-LPS IgG with no significant decline 84 days after vaccination in the alum-adsorbed cohort. The vaccine candidate was also well-tolerated, with no serious or severe adverse events [23]. Currently, synthesized LPS mimics are implemented in pre-clinical studies to propose vaccine candidates against other enteric pathogens. A glycoconjugate vaccine candidate incorporating synthetic core oligosaccharide of *Neisseria meningitidis* demonstrated IgG agglutination *in vitro* and showed strong recall responses after reimmunisation 260 days from the initial priming [70]. Another glycoconjugate vaccine candidate,

utilizing the hexasaccharide fragment of *Vibrio cholerae* O:1 (Ogawa) O-SP conjugated to BSA induced a robust anti-LPS IgG response and showed a vaccine efficacy of 86% in a mice challenge model with one 50% lethal dose (LD50) [71].

To date, there are four preclinical and one phase 1 LPS based glycoconjugates incorporating chemically synthesized glycans, including vaccine candidates against *S. dysenteriae*, *Vibrio cholerae*, *Neisseria meningitidis*, *Bordetella pertussis*, and *S. flexneri* [23,70–74]. The emergence of these synthetic carbohydrate-based glycoconjugate vaccine candidates results from evolving automated glycan assembly (AGA) technologies [75]. The introduction of the first commercial AGA platforms (Glyconeer 2.1 synthesizer) [76] in 2017 showed the synthesis of hexasaccharides with purities >99%. To synthesize the oligosaccharide, the AGA platform functions by executing reaction cycles that assemble appropriate fluorenylmethyloxycarbonyl (Fmoc) protecting group-conjugated building block sugars to polystyrene-based resins. The oligosaccharide is then cleaved by ultraviolet (UV)-irradiation, while quality control is carried out by HPLC, MS, and NMR. Recent advances in AGA quality control allowed real-time monitoring of reaction cycles and may improve the platform's efficiency further, preparing it for commercial-scale production of high-quality oligosaccharides [77].

More recent reports had also described the chemoenzymatic synthesis of oligosaccharides, which involves the enzymatic polymerization of chemically synthesized monosaccharides. This method had been applied to generate meningococcal CPS for glycoconjugate vaccine production [26,78]. However, while results have been published describing the chemoenzymatic synthesis of core oligosaccharide of *E. coli* O157 for enzyme activity assays [79], the adaptation of this method to generate the O-SP of LPS is still underway.

5. Recent developments in carrier molecules

Conjugation technology allowed previously T-cell independent antigens to become potent T-cell dependent immunogens. Carrier proteins seen extensively in LPS-based conjugate vaccines to date include tetanus toxoid (TT), diphtheria toxoid (DT), cross-reactive material 197 (CRM197, a modified nontoxic mutant of DT), and bovine serum albumin (BSA; in proof-of-concept phase only) (Table 2). These carrier proteins represent most of the carrier proteins present in clinical and preclinical LPS-based glycoconjugate vaccines. These carriers are time-proven and have been manufactured on large scales. Other conventional carrier proteins, which are well-documented in past reviews [80,81], include diphtheria toxoid (DT), meningococcal outer membrane protein complex (OMPC), keyhole limpet hemocyanin (KLH), ovalbumin (OVA), recombinant exoprotein A of *Pseudomonas aeruginosa* (rEPA), and *H. influenzae* protein D (HiD).

Aside from the well-documented carriers, novel carriers recently experimented in preclinical LPS-based glycoconjugate vaccine candidates include nanoparticle-based carrier proteins and peptide-based dual-role carriers, both being noteworthy considering their potentially higher immunogenicity compared with conventional carriers.

5.1. Nanoparticle-based carriers

5.1.1. Gold nanoparticles

Most currently licensed glycoconjugate vaccines with CPS-based carbohydrates have a 'lattice'-type conformation [91]. In contrast, 'sun'-type glycoconjugates were shown to induce higher IgG titer against *S. sonnei* and *S. dysenteriae* type 1 compared with the conventional 'lattice'-type configuration [72,92]. The advantage to AuNPs is that protein and carbohydrate linkers can conjugate onto AuNPs in a 'sun'-type conformation, where thiol ends of thiolated peptides/glycans attach to the gold particle [93]. This property allowed AuNP-peptide-glycan conjugates to elicit robust immune responses toward both the carbohydrate and peptide moieties [94].

The immunogenicity of AuNP-based glycoconjugates is based on the ability to enter antigen-presenting cells (APCs) such as the dendritic cells and B cells. The rate of internalization of AuNPs into DCs varies depending on the shape and size. Smaller, rod-shaped AuNPs are more readily internalized into DCs and perinuclearly localized [95]. In contrast to DCs where AuNPs are internalized more readily, AuNPs are 10-fold less internalized in B cells while remaining nontoxic to B cells when incubated in culture [96]. Improving uptake rates of AuNPs is of interest to develop more potent nanoparticle-based vaccine candidates.

Recent work on preclinical *Burkholderia mallei* LPS-based glycoconjugate candidates have incorporated AuNPs. The vaccine candidates were constructed in two steps: an initial conjugation was performed between the AuNP and a synthetic peptide antigen, followed by a second conjugation of LPS onto the peptides pre-decorated on the AuNPs [94]. By doing this, the AuNP conjugate carries both peptide and carbohydrate antigens. The vaccine candidates demonstrated high levels of protection shown by 100% survival in mice challenged with 50 times LD50.

Besides the application in LPS-based glycoconjugate vaccines, AuNP-based carriers has also been conjugated to CPS. CPS-AuNP vaccine candidates for *Streptococcus pneumoniae* type 14 have also induced anti-CPS IgG responses in murine models. The vaccine candidates were composed of AuNP conjugated to varied ratios of OVA 323–339 peptide, D-glucose, and synthetic branched tetrasaccharide representing a single repeat of the *Streptococcus pneumoniae* type 14 CPS. While the best candidate yielded an IgG response representative of one log10 less than the CPS-CRM197 positive control, mouse serum from CPS-AuNP-administered mice had a 75% phagocytic capacity compared with CPS-CRM197-administered mice. The results show that immunity can be induced from vaccines constructed with this platform, but further optimizations can be necessary [93].

5.1.2. Poly lactic-co-glycolic acid nanoparticles

PLGA-NPs are biodegradable polymeric esters formed from a 1:1 ratio of lactic acids and glycolic acids [97]. Due to the intake of PLGA-NPs into APCs such as DCs, these constructs were used to explore the functionality both as a vaccine adjuvant carrier to deliver peptide and carbohydrate-based adjuvants, and as a vaccine candidate itself, either by the conjugation of antigens on its surface or by encapsulating

Table 2. Clinical and preclinical LPS-based glycoconjugate vaccine candidates.

Pathogen	Source of carbohydrate	Carbohydrate used	Carrier protein	Linker	Phase	Efficacy	If terminated	Reference
<i>Bordetella pertussis</i>	Synthesis	pentasaccharide of O-SP	bacteriophage Qb, KLH		Preclinical	anti-LPS IgG produced for all vaccine candidates		[74]
<i>Bordetella pertussis</i> / <i>bronchiseptica</i>	Hot phenol-water extraction	O-Ag	BSA		Preclinical	Both produced 750 reciprocal bactericidal titer		[146]
<i>Brucella abortus</i>	Hot phenol-water extraction	O-Ag/O-SP	Poly lactic-co-glycolic acid (PLGA) nanoparticles		Preclinical	68% (O-Ag) and 64% (O-SP) inhibition of OPK activity; 8X IgG response compared to O-Ag/O-SP (unconjugated) alone		[98]
<i>Burkholderia mallei</i>	Hot phenol-water extraction	Whole LPS	Gold nanoparticles – OmpW, OpcP, HA, Combo2 (OmpW+OpcP+HA)		Preclinical	Challenge with LD50 – OmpW (80%), OpcP (80%), HA (50%), Combo2 (100%)		[94]
<i>Burkholderia pseudomallei</i>	Bioconjugation	O-Ag	<i>E. coli</i> multidrug efflux pump subunit AcrA		Preclinical	Murine infection model with 12 times LD50 showed delayed death than group vaccinated with killed whole-cell		[184]
Enteropathogenic <i>E. coli</i> serotype O1A, O2, O6A, O25B	Bioconjugation	O-Ag	Exotoxin A from <i>Pseudomonas aeruginosa</i> (EPA)		Phase 2	75% – 100% demonstrated an ELISA titer increase of ≥ 2 -fold (depending on dose groups); 57.5% – 83.6% ≥ 4 -fold.		[24]
<i>Escherichia coli</i> O111	Commercial	O-Ag	cytochrome C, EtxB (a recombinant B subunit of heat-labile toxin of ETEC)		Preclinical	IgG detected in both blood and stool		[109]
<i>Escherichia coli</i> O111:B4 (J5 mutant)	Phenol/chloroform/light petroleum ether	Core oligosaccharides	group B meningococcal outer membrane protein (OMP)		Phase 1	1 of 8 vaccine-only recipient had four-fold increase in IgG titer		[82]
<i>Escherichia coli</i> rough mutants (K12/R1-R4/J5)	Phenol/chloroform/light petroleum ether	Core oligosaccharide (retaining deacetylated lipid A)	Hemocyanine		Preclinical	Failed to protect cows from <i>E. coli</i> O157 challenge		[65]
<i>Escherichia coli</i> serotype O1, O2, O4, O6, O7, O8, O12, O15, O16, O18, O25, O75	Hot phenol-water extraction	O-Ag	<i>Pseudomonas aeruginosa</i> toxin A		Preclinical	IgG detected in immunized rabbit serum; Significant protection in mice challenge model for O4, O15, O25		[126]
<i>Francisella tularensis</i>	Hot phenol-water extraction	O-Ag	TT		Preclinical	Challenge with 100x LD50 – VHMW (75%), HMW (0%), LMW (20%)		[134]
<i>Francisella tularensis</i>	Bioconjugation	O-Ag	Exotoxin A from <i>Pseudomonas aeruginosa</i> (EPA)		Preclinical	50% survival in 100 CFU mouse infection models		[183]
<i>Helicobacter pylori</i>	Hot phenol-water extraction	O-Ag/O-Ag (retaining deacylated lipid A)	BSA/TT		Preclinical	O-SP-TT conjugate showed 5-fold decrease in stomach bacterial load compared to saline control;		[199]
<i>Helicobacter pylori</i>	Hot phenol-water extraction	O-Ag	BSA		Preclinical	Seroconversion in mice; limited cross-reactivity with serotypes O:1 and O:5 LPS and the core oligosaccharide		[106]

(Continued)

Table 2. (Continued).

Pathogen	Source of carbohydrate	Carbohydrate used	Carrier protein	Linker	Phase	Efficacy	If terminated	Reference
<i>Mannheimia haemolytica</i>	Hot phenol-water extraction	O-Ag (retaining deacylated lipid A)	CRM197		Preclinical	Bacterial killing with immunized rabbit serum		[83]
<i>Moraxella catarrhalis</i>	Hot phenol-water extraction	O-Ag (retaining deacylated lipid A)	CRM197		Preclinical	Bacterial killing with immunized rabbit serum		[84]
<i>Moraxella catarrhalis</i> serotype A, B, C	Hot phenol-water extraction	Core oligosaccharides	TT/CRM197		Preclinical	Bacterial killing with immunized mice serum		[197]
<i>Neisseria meningitidis</i>	Synthesis	Core oligosaccharides	CRM197		Preclinical	Successful re-immunization observed at d = 267; Confocal microscopy verifies bacteria bound to serum IgG		[70]
<i>Neisseria meningitidis</i>	Hot phenol-water extraction	Core oligosaccharides	CRM197		Preclinical	Post immunization rabbit sera able to perform > = 50% bacteria killing in bactericidal assay after >2048-fold dilution		[115]
<i>Pseudomonas aeruginosa</i> serotype 6,11	Phenol/chloroform/light petroleum ether	O-Ag	BSA		Preclinical	Survival with LD50 – up to 100%		[85]
<i>Pseudomonas aeruginosa</i> serotype 1,2,3,4,5,6,10,11	Hot phenol-water extraction	O-Ag	Exotoxin A from <i>Pseudomonas aeruginosa</i> (EPA)		Phase 1	Halved cases of <i>P. aeruginosa</i> chronic infection; seroconversion in all mice		[129]
<i>Salmonella enterica</i> serovar Enteritidis	Darveau and Hancock	Core oligosaccharides	FliC (flagellin)/CRM197		Preclinical	Challenge with 5X LD50 – vaccine efficacy 91.7% in mice		[144]
<i>Salmonella enterica</i> serovar Paratyphi	Hot phenol-water extraction	O-Ag	diphtheria toxoid	Adipic acid dihydrazide (ADH)	Preclinical	Conjugate IgG titer p < 0.05 compared to LPS administered alone	Terminated	[145]
<i>Salmonella enterica</i> serovar Paratyphi O2	Hot phenol-water extraction	O-Ag	CRM197		Preclinical	Seroconversion in all mice; significantly more immunogenic than O-SP alone		[120]
<i>Salmonella enterica</i> serovar Typhimurium	Darveau and Hancock	O-Ag	BSA, OVA, FliC		Preclinical	Survival with LD50 – BSA (70%), OVA (77.8%), FliC (80%); Vaccine efficacy – BSA (56.5%), OVA (60.5%), FliC (74.1%)		[141]
<i>Salmonella enterica</i> serovar Typhimurium	Direct acid hydrolysis	O-Ag (retaining deacylated lipid A)	CRM197		Preclinical	2-log reduction in bacterial burden in immunized mouse spleen and liver		[63]
<i>Salmonella enterica</i> serovar Typhimurium	Direct acid hydrolysis	O-Ag	CRM197		Preclinical	Seroconversion in most mice; all glycoconjugates induced bactericidal antibodies; 3 to 100-fold increase in anti-O-SP IgG titers depending on O-SP per CRM197		[148]
<i>Shigella dysenteriae</i> type 1	Synthesis/Hot phenol-water extraction	tetra-,octa-, dodeca-, and hexadecasaccharide of O-SP and isolated O-SP	HSA (human serum albumin)		Preclinical	anti-LPS IgG produced for all experimental vaccine candidates, lowest for isolated O-SP-HSA conjugate		[68]
<i>Shigella dysenteriae</i> type 1	Bioconjugation	O-Ag	Exotoxin A from <i>Pseudomonas aeruginosa</i> (EPA)		Phase 2	100% ≥4-fold increase in LPS-specific serum IgG in Phase 1 (Phase 2 unpublished)		[86]

(Continued)

Table 2. (Continued).

Pathogen	Source of carbohydrate	Carbohydrate used	Carrier protein	Linker	Phase	Efficacy	If terminated	Reference
<i>Shigella flexneri</i> 2a	Hot phenol-water extraction	O-Ag	CfaEB (recombinant ETEC adhesin)		Preclinical	4.5-fold anti-O-SP IgG compared to control		[106]
<i>Shigella flexneri</i> 2a	Bioconjugation	O-Ag	Exotoxin A from <i>Pseudomonas aeruginosa</i> (EPA)		Phase 2	92 to 100% of the volunteers showing a \geq 4-fold increase in LPS-specific serum IgG		[187]
<i>Shigella flexneri</i> 2a	Synthesis	15-mer oligosaccharide mimic of O-SP	TT	Maleimide spacers	Phase 1	Seroconversion in all human recipients		[23]
<i>Shigella sonnei</i>	Hot phenol-water extraction	O-Ag (low molecular mass fraction, 3.5 repeat units/1.3 repeat units)	BSA, recombinant DT		Preclinical	Higher anti-LPS IgG level than full-length O-SP conjugates		[92]
<i>Shigella sonnei</i>	Hot phenol-water extraction	O-Ag	recombinant exoprotein A from <i>Pseudomonas aeruginosa</i> (rEPA)		Phase 3	35.5% in children >2-3 years; 71.1% in children >3-4 years	Terminated	[22,54]
<i>Shigella flexneri</i> 2a	Hot phenol-water extraction					22.5% in children >2-3 years; -3.6% in children >3-4 years		
<i>Shigella flexneri</i> 2a <i>Shigella flexneri</i> 6 <i>Shigella dysenteriae</i> type 1	Hot phenol-water extraction	O-Ag	BSA		Preclinical	anti-LPS IgG produced for all three experimental vaccine candidates		[87]
<i>Vibrio cholerae</i> O1 (Inaba)	Hot phenol-water extraction	O-Ag	BSA/chitin-binding-protein A (CBP-A)	ADH	Preclinical	mouse lethality cholera challenge model – 100% survival		[132]
<i>Vibrio cholerae</i> O1 (Ogawa)	Filtering of released LPS	O-Ag (retaining deacylated lipid A)	TT	17-atom with linear hydrazide and alkyl bonds	Preclinical	anti-LPS IgG produced against both Inaba and Ogawa		[133]
<i>Vibrio cholerae</i> O1 (Ogawa)	Hot phenol-water extraction	O-Ag/O-Ag (retaining deacylated lipid A)	BSA	ADH	Preclinical	Seroconversion in mouse	Terminated	[88]
<i>Vibrio cholerae</i> O1 (Inaba) <i>Vibrio cholerae</i> O1 (Ogawa)	Hot phenol-water extraction	O-Ag	BSA		Preclinical	Recognized by acute and convalescent patient serum		[138]
<i>Vibrio cholerae</i> O139	Hot phenol-water extraction	O-Ag	BSA		Preclinical	ELISA response seen in O139 convalescent serum		[139]
<i>Vibrio cholerae</i> O139	Hot phenol-water extraction	O-Ag (retaining deacylated lipid A)	BSA		Preclinical	about 60% bacterial killing in vibriocidal assay by undiluted sera d = 77		[89]
<i>Vibrio cholerae</i> O1 (Inaba)	Hot phenol-water extraction	O-Ag	TThc (tetanus toxoid heavy chain fragment)		Preclinical	mouse lethality cholera challenge model – 60% survival		[90]
<i>Vibrio cholerae</i> O1 (Ogawa)	Synthesis	hexasaccharide fragment of O-SP	BSA		Preclinical	Survival with LD50 – 92%; Vaccine efficacy – 86%		[71]
<i>Vibrio cholerae</i> O1 (Ogawa)	Hot phenol-water extraction	O-Ag	TThc (tetanus toxoid heavy chain fragment)		Preclinical	mouse lethality cholera challenge model – 95% survival		[137]

the antigens [98,99]. Antigens encapsulated by PLGA-NPs were shown to be released in the cytosol of DCs followed by binding MHCII complexes, allowing the presentation of the bound antigens to CD4 + T cells [100]. Several vaccine candidates were developed using PLGAs-NPs entrapping peptide-based antigens, however the immunogenicity varied significantly between one vaccine candidate and another [101–103].

Recent PLGA-NP-LPS/O-PS vaccine candidates for *Brucella abortus* showed that PLGA-based platform could produce glycoconjugate vaccine candidates in a 'sun'-like conformation [98]. The vaccine candidate comprised LPS/O-SP of *Brucella abortus* conjugated onto hollow PLGA-NPs. In mice, the vaccine candidates elicited an eight-fold greater increase in anti-LPS IgG response when administered in mice than LPS or O-SP

alone, demonstrating the proof-of-concept for producing glycoconjugate vaccines with this platform [98].

5.2. Dual-role peptide-based carriers

Modern bioinformatics allows the identification of pathogen-specific peptide antigens in which some can act both as antigen and as a carrier protein, thus termed dual-role peptide-based carriers. Dual-role peptide-based carriers allow antibody responses to be directed toward both the carrier and the carbohydrate, potentially enabling an additional layer of protection by anti-carrier antibodies [104]. In theory, this approach can also induce protection against more than one pathogen when the carrier and the carbohydrate are derived from different pathogens [105]. Meanwhile, using dual-role peptide-based carriers is also beneficial in alleviating carrier-induced epitopic suppression (CIES), which will be described later in this review.

Recent LPS-based glycoconjugate vaccine candidates have demonstrated the potential of dual-role peptide-based carriers as both a protective antigen against additional pathogens, or as an extra layer of protection against the same pathogen as the O-SP. In the case of protection against additional pathogens, a recent glycoconjugate vaccine candidate against *S. flexneri* 2a with O-Ag conjugated to CfaEB, a recombinant carrier protein composed of the major pilin subunit (CfaB) and a minor adhesive subunit (CfaE) of enterotoxigenic *Escherichia coli* (ETEC), showed seroconversion with both anti-O-SP and anti-CfaEB IgG titers in mice [106]. The presence of anti-O-SP is a correlate of protection against *S. flexneri* 2a [107], while CfaB and CfaE are also protective antigens against ETEC [108]. Additionally, another enterohemorrhagic *Escherichia coli* O111 glycoconjugate comprising the O-SP of *E. coli* O111 conjugated to the nontoxic B subunit of heat-labile toxin of ETEC (EtxB) induced anti-IgG and anti-EtxB antibodies, which are both correlates to protection against ETEC [109,110].

Besides LPS-based glycoconjugates, dual-role peptide-based carriers have been used in many glycoconjugate vaccine candidates. Perhaps the most notable example is the non-typeable *Haemophilus influenzae* (NTHi) protein D incorporated as the carrier for eight out of ten serotypes covered by the licensed pneumococcal polysaccharide and NTHi protein D conjugate vaccine (Synflorix, GlaxoSmithKline Biologicals). In a phase 3 study, the pneumococcal polysaccharide and NTHi protein D conjugate vaccine demonstrated opsonophagocytic activity (OPA) titers of ≥ 8 -fold against the pneumococcal pathogens, while also eliciting a 3.5-fold increase of anti-protein D IgG in children that have not received pneumococcal vaccines previously, thus inducing an immune response against both the CPS moiety and the carrier protein [111]. Other examples of dual-role carriers include a group A *Streptococcus* vaccine candidate with group A *Streptococcus* carbohydrate (GAC) conjugated to *Streptococcus* streptolysin O (SLO), which elicited higher anti-GAC IgG titers compared with GAC-CRM197 conjugates [105]. Another example is a pneumococcal CPS-hepatitis B surface antigen (HBsAg) conjugate vaccine candidate, which showed potent T cell-dependent immune responses against both the CPS and the carrier protein [112].

The approach of combining carbohydrate antigen and carrier proteins from different origins is reminiscent of a previous study that showed TT and DT conjugates confer protection against lethal challenges of both toxins in mice [113]. However, there have not yet been investigations to determine if novel dual-role peptide-based carriers, besides protein D in pneumococcal polysaccharide and NTHi protein D conjugate vaccine, will induce antibody responses against both the carbohydrate antigen and the carrier protein in humans. The use of dual-role carriers may benefit from site-specific conjugation methods to retain important epitopes with the aim of eliciting a robust immune response to the carrier protein, which will be discussed in the following section.

6. Chemical conjugation methods

To date, multiple ways to conjugate carrier proteins and glycans have been used in the preclinical development of LPS-based glycoconjugate vaccines to achieve different conformations (Table 3; Figure 3). The high-value nature of carbohydrates and carrier proteins has led to a requirement for high conjugation efficiency. Thus, the ideal conjugation method would aim to maximize the preservation of carbohydrates and carrier proteins to ensure sufficient immunogenicity. To achieve this, conjugation techniques need to be carried out under temperature, pH, and media that best resemble physiological conditions [114]. Glycoconjugates can be present in two conformations, 'sun'-type conformation is resulted from single-point attachments between glycans and the central carrier protein, whereas the 'lattice'-type conformation is resulted from multiple-point attachment between a single glycan and several carrier proteins (Figure 3). Also, recent preclinical glycoconjugate vaccine candidates had reflected an increased awareness that the steric hindrance arising from directly conjugating carbohydrate to a carrier protein would require a linker or spacer to evade. At the same time, different linkers may also result in varied immunogenicity [115]. In this section, chemical conjugation methods used for past clinical and preclinical LPS-based vaccine candidates are discussed.

6.1. Reductive amination

One of the most implemented conjugation methods in glycoconjugate vaccine production is via reductive amination [118,119]. Licensed vaccines conjugating via this method include CPS-based conjugate vaccines against 13 serotypes of *S. pneumoniae* (Pneumovax13; Pfizer, USA).

Reductive amination involves the selective mild oxidation of reducing or non-reducing sugar residues to form aldehyde groups. The aldehydes form Schiff base intermediates upon interacting with lysine residues present on the carrier protein. The reduction of these Schiff intermediates via the addition of borate buffer generates stable glycoconjugates. The addition of linkers, such as adipic acid dihydrazide (ADH), can be incorporated in reductive amination to evade steric hindrance and facilitate coupling of protein to the sugar component [115,120].

The merit of this approach is two-fold. Firstly, the simplicity of the workflow allows minimization of costs and loss of materials during the process. At the same time, synthetic sugars can also

adapt to this workflow by treating with aqueous ammonium bicarbonate to activate the oligosaccharide by transforming other functional groups into aldehyde [121]. However, reductive amination is yet to be the most efficient: the accumulation of alcohol can form irreversibly from the aldehydes on the glycans, which are not stabilized with reducing agents, causing a reduction in efficiency during conjugation [122]. Stabilization of Schiff intermediates would require toxic chemicals, such as borohydride [123]. Also, the level of oxidation when activating the carbohydrates can directly impact carbohydrate:protein ratio of the resultant glycoconjugate [124]. It was documented that while a higher carbohydrate:protein ratio does not positively correlate with immunogenicity, an optimal carbohydrate:protein ratio is necessary to evoke the most robust immune response possible [72].

LPS-based conjugate vaccine candidates for *Pseudomonas aeruginosa*, *Escherichia coli*, and *Helicobacter pylori* have used

reductive amination for conjugation [125–127]. Notably, the vaccine candidate against *P. aeruginosa* entered phase 1 trials but was terminated because of a low level of immunogenicity in humans [128,129].

6.2. Active ester chemistry

Active ester chemistry is seen extensively in glycoconjugate vaccine candidates which contained synthetic carbohydrates [130]. The mechanism of this approach requires the synthetic oligosaccharides to carry a terminal amino group which allows attachment of di-*N*-hydroxysuccinimidyl adipate (DSAP) to form a half-ester in dimethyl sulfoxide (DMSO). The half-ester, or active ester can readily attach to lysine residues on carrier proteins [130].

Table 3. Glycoconjugate conformations resulting from different chemical conjugation methods applied in LPS-based glycoconjugate vaccine production.

Conjugation method	Glycoconjugate conformations	Advantages	Drawbacks	Reference
Reductive amination	Sun; Lattice	Well-documented workflow Have been used in licensed CPS-based glycoconjugate vaccines	Requires a LPS oxidation step which may damage sugars Carrier derivatization under low pH may alter epitopes	116,124
Active ester chemistry	Sun	Simple workflow when conjugating with synthetic glycans	Only convenient when working with synthetic glycans carrying a terminal amino group	130
CNBr/ADH activation	Lattice	Time-proven technique with well-documented workflow	CNBr is toxic	116,131
Squaric acid chemistry	Sun	Quick workflow	Conjugation under high pH may affect epitopes May introduce steric hindrance between glycan and carrier, causing reduction in immunogenicity	136,139
CDAP chemistry	Lattice	Easier transition from CNBr/ADH activation due to similar workflow CDAP is a nontoxic alternative to CNBr	Over-crosslinking can happen, causing reduction in immunogenicity	143
Aminoxyoxime chemistry	Sun; Lattice	Simple and well-documented workflow Does not require extra derivatization of LPS Mild conjugation conditions may help preserve epitopes	Have not been used in licensed vaccines	118
Click chemistry	Sun	Can be performed at room temperature	May have lower conjugation efficiency than random conjugation	117,147

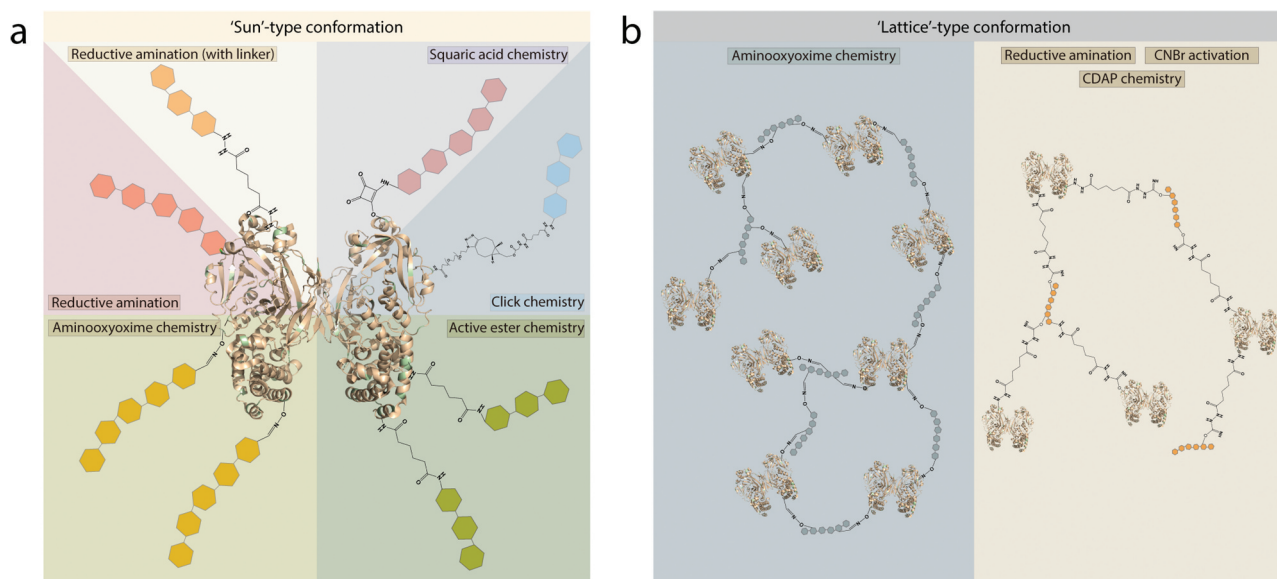


Figure 3. Conformations of chemically conjugated glycoconjugate vaccines. (a) 'Sun'-type glycoconjugates. (b) 'Lattice'-type glycoconjugates. Carrier protein is represented by crystal structure of CRM197 (PDB accession: 4AE1), lysine residues are highlighted in green, linkers are represented by their respective chemical structures attached on lysine residues. Each colored grid represents sugars conjugated to the carrier protein with their respective linkers.

Active ester chemistry was incorporated in two LPS-based vaccine candidates. More recently, a vaccine candidate comprising synthetic pentasaccharide of *B. pertussis* O-SP to bacteriophage Q β showed a three-fold higher anti-carbohydrate IgG response compared with control groups receiving Q β alone [74]. The method can also be adapted to conjugating O-Ag isolated from bacteria by additional linkers such as ADH, as seen in a *S. Paratyphi* A vaccine candidate demonstrated higher anti-O-Ag IgG titers in mice than administering O-Ag alone [120].

6.3. Cyanogen bromide (CNBr) activation

Activation of glycan with cyanogen bromide is a time-proven technique for sugar-protein conjugation, involving a simple and quick workflow to produce glycoconjugates. The technique has been implemented for several purposes, including conjugate vaccine production, enzyme purification, and biological assay development [131]. In brief, the O-Ag is activated with cyanogen bromide (CNBr) to generate cyanate esters that further reacted with vicinal hydroxyl groups to form cyclic imidocarbonates, which can couple to the carboxyl groups of carrier proteins following 1-ethyl-3-(3-dimethylaminopropyl) carbodiimide (EDC)-mediated condensation.

In vaccine production, CNBr activation is commonly accompanied by the use of ADH linkers. LPS-based vaccine candidates conjugated via CNBr activation include ones against *V. cholerae* O:1 Serotype Inaba [132], *V. cholerae* O:1 Serotype Ogawa [133], and *Francisella tularensis* [134]. All the CNBr-conjugated candidates were immunogenic in mice, while several also provided protection to lethal bacterial challenges [132,134]. The most notable drawback to this method is the toxicity of CNBr, which may contribute to difficulty in handling. Conjugation conditions using CNBr are also a potential problem as it requires a high pH (pH = 9–10), which may interfere with the stability of carbohydrates and proteins [135].

6.4. Squaric acid chemistry

Squaric acid chemistry presents a relatively quick and convenient way of conjugation. The use of squaric acid chemistry was first demonstrated in the synthesis of glycoconjugates from synthetic carbohydrates [136]. To carry out conjugation via squaric acid chemistry, the carbohydrate component is linked with a monoamide to the squaric diester. The squarate intermediate can then be reacted with lysine residues in borate buffer (pH = 9) through the formation of 1,2-bisamide of the squaric acid [137].

Reports have suggested that glycoconjugates produced through squaric acid chemistry may be less immunogenic due to the relatively small size of squaric acid, compared with other common linkers such as ADH, thus may result in steric hindrance [138]. Examples of LPS-based vaccine candidates using squaric acid chemistry for conjugation include ones for *V. cholerae* O1 serotypes Inaba and Ogawa, and for *V. cholerae* O139 [137–139]. Within these, the vaccine candidates for *V. cholera* O1 shown similar IgG titers to administering LPS alone [138]. Similarly, significantly lower anti-CPS IgG was elicited by pneumococcal squaric acid-linked

glycoconjugates compared with glycoconjugates using longer linkers, such as ADH [140].

6.5. 1-cyano-4-dimethylaminopyridinium tetrafluoroborate (CDAP) chemistry

While its mechanism is analogous to CNBr activation, CDAP chemistry is a newer and potentially better alternative to CNBr chemistry [141]. Unlike CNBr, CDAP is nontoxic and does not require a high pH to carry out carbohydrate activation. The mechanism of CDAP activation primarily involves the formation of isourea bonds between cyanoesters on CDAP-activated carbohydrates and ϵ -amino group of lysine residues on carrier proteins [142,143]. As in CNBr chemistry, ADH is commonly used as a linker in CDAP chemistry.

The primary concern of using CDAP is over-crosslinking. Commonly, over-crosslinked glycoconjugates can cause gelling of carbohydrate and peptides, thus reducing immunogenicity. To overcome this, CDAP chemistry can be performed under an acidic pH using nucleophilic linkers, such as adipic dihydrazine, which can in turn reduce gelling and restore immunogenicity of the glycoconjugate [51,143].

CDAP chemistry was used in the preclinical development of several LPS-based glycoconjugate vaccine candidates, which induced significantly greater IgG responses in mice against *S. enterica* serovar Enteritidis [144] and *S. enterica* serovar Paratyphi A than administering O-Ag alone [120,145].

6.6. Aminoxyoxime chemistry

Aminoxyoxime chemistry is a precise means of conjugation used in several LPS-based glycoconjugate vaccine developments, which included ones against *S. Enteritidis*, *B. pertussis*, *S. sonnei*, and *S. dysenteriae* [68,92,144,146]. Aminoxyoxime chemistry allows conjugation to take place at specific moieties of the carbohydrate and proteins, making the result predictable while also being capable of manufacturing glycoconjugates in both 'lattice'-like or 'sun'-like conformations [114].

Both proteins and carbohydrates can be functionalized with an aldehyde or with an aminoxy group. Under a mild acidic condition (pH = 5), the aldehyde-functionalized protein or carbohydrate can readily form oximes with the strong nucleophilic aminoxy group attached on the protein or carbohydrate counterpart [118]. Several ways have been developed to functionalize carbohydrates and polysaccharides for conjugation via aminoxyoxime chemistry, detailly described by Lee, Sen, and LopezAcosta [118].

6.7. Click chemistry

Click chemistry allows the attachment of glycans onto specific activated residues on the carrier protein in a 'sun'-like conformation [147]. Click chemistry gained popularity more recently through its increasingly extensive use in conjugating glycoconjugates in biochemical research [148,149]. The method works by the ligation between propargylamine-activated glycans and an alkyne-activated carrier protein.

The advantages to click chemistry include that the method can efficiently generate structurally defined ('sun'-type

conformation) glycoconjugates at room temperature while also having a relatively simple workflow. However, the conjugation efficiency of this method is also lower than those of random conjugation [147]. For LPS-based glycoconjugate vaccines, click chemistry was used to generate vaccine candidates against *S. Typhimurium*, showing an increase of more than 100-fold in anti-O-Ag IgG titers in murine models [147,150].

7. Immunological challenges for LPS-based glycoconjugates

7.1. Carrier-induced epitopic suppression (CIES)

The most considerable challenge faced by all glycoconjugate vaccines is likely CIES, which is the reduced immunogenicity of a glycoconjugate when preexisting immunity is present against the same carrier protein it comprises. CIES was well-documented in past clinical studies as a significant perturbing factor for vaccine efficacies [151,152].

Several mechanisms contributing to CIES have been proposed. One theory suggests the binding of preexisting antibodies against the carrier protein can interfere with the binding of glycoconjugates to B cells, thus disrupting the internalization and the subsequent initiation of antibody responses [153]. Another theory is that previous carrier protein priming would elicit clonal expansion of carrier-specific B cells. This would in turn increase the probability of the glycoconjugate being internalized and degraded by carrier-specific B cells, while preventing the necessary binding to glycan-specific B cells to induce immunity against the glycan moiety of the glycoconjugate [154].

Recent work by Pecetta et al. suggested the possibility that CIES is carrier-dependent. A cohort of mice primed with two different carrier proteins, CRM197 or DT, induced a low anti-CRM197 IgG titer and a high anti-DT IgG titer, respectively. The same cohort was then vaccinated with MenA-CRM197 and MenA-DT. As a result, CRM197-primed mice receiving the MenA-CRM197 conjugate showed both anti-carrier and anti-carbohydrate IgG responses, while the mice primed with DT before administering the MenA-DT conjugate showed a high anti-carrier and a low anti-carbohydrate IgG response [155]. Thus, the results likely reflected that priming with a less immunogenic carrier does not induce expansion of carrier-specific B cells. In contrast, immunogenic carriers elicit more robust carrier-specific B cell responses that compete with, and prevent, carbohydrate-specific B cell responses [156]. While the mechanisms of CIES are not fully understood, increasing vaccine doses and introducing alternative carrier proteins may help alleviate CIES [157]. In contrast, several studies suggest that priming with a small amount of carrier protein may enhance vaccine efficacy in a dose-dependent manner, highlighting a need for further investigations into the mechanisms of CIES [158–160].

7.2. Bystander interference

Like CIES, bystander interference (BI) presents an additional layer of complication in developing and implementing glycoconjugate vaccines. BI happens when multiple glycoconjugate

vaccines are introduced concomitantly, causing a decreased immune response to the carbohydrate antigen administered. The current mechanism of bystander interference is that the abundance of conjugates may introduce competition for cytokines and cytokine inhibitors present within the lymph node while resources remain limited [153,161,162].

A well-documented example of BI is the reduced vaccine efficacy of Hib conjugate vaccines in the UK. The Hib-TT conjugate vaccine, combined with diphtheria-tetanus-acellular pertussis vaccines (DTPa-Hib), was introduced as a routine in the UK in the 1992 [163]. The vaccine's initial results ranged from 0.83 incidents of Hib infection per 100,000 children of <5 years of age in 1996 to 0.84 in 1997, and 0.63 in 1998. However, the figures rose to 1.1 in 1999, followed by 1.8 in 2000, indicating an increase in Hib vaccine failures [163,164]. In November 1999, a MenC-CRM197 conjugate vaccine (MCC-CRM197) was introduced as a routine 3-dose infant immunization course in the UK [165]. Despite the success in reducing MenC cases in the UK from 1.81 per 100,000 population in 1998/1999 to 0.08 per 100,000 in 2015/16 [162], the concomitant administering of both the MenC conjugate and the Hib conjugate may have contributed to the reduced protection of the Hib conjugate compared with previous years. This was validated by later clinical studies showing the coadministration of MenC-CRM197 and DTPa-inactivated polio (IPV)-Hib in infants produced lower anti-polyribitylribitol phosphate (PRP; a constituent of Hib CPS) titers than receiving DTPa-IPV-Hib alone, while co-administering with MenC-TT resulted in an additional reduction in MenC-TT efficacy [166]. In addition, the coadministration of MenC-CRM197 and DTPa-IPV-Hib showed decreased immunogenicity for all three vaccines after a 3-dose immunization in infants [167]. Statistically significant interference was also reported in concomitant administration of Infanrix-hexa (DTPa-IPV-hepatitis B (HBV)/Hib; GlaxoSmithKline) and Prevenar7 (7-valent pneumococcal vaccine) [168]. While Hib glycoconjugate vaccines are particularly more prone to BI, other glycoconjugate vaccines also showed small reductions in immunogenicity when co-administered with other vaccines [162,167].

7.3. Administration and vaccine molecule trafficking

Several available polysaccharide-based and live attenuated vaccines can be administered subcutaneously, while all licensed glycoconjugate vaccines are administered intramuscularly in the US and the UK [169,170]. Despite similar levels of immune responses elicited by these two administration routes, it has been shown that vaccines traffic to physiologically different lymph nodes depending on the proximity to the injection sites, with intramuscular administered vaccine molecules trafficking to lymph nodes further away from the skin in rhesus macaques [171,172]. Additionally, it has been shown that antigen uptake in booster doses can benefit from intramuscular administration compared with subcutaneous administration [172].

While most vaccines are injected, alternative routes of vaccine administration to injection may lead to improved vaccine responses [173]. Oral and nasal administration of vaccines has

been successful of inducing mucosal immunity. Intramuscular injection is effective in inducing antibodies in the blood, including IgM and IgG, which are thought to be protective against the spread of bacteria within circulation, but, might offer weaker protection at the sites of primary entry, such as the respiratory tract and the gut mucosa [174]. These sites of primary entry would require the production of IgA or transfer of IgG from the blood to provide a robust mucosal immunity [173]. Intranasal administration of LPS-AuNP glycoconjugate against *Burkholderia mallei* showed complete protection against a mouse lethal challenge model, which has also highlighted that novel nanoparticle-based carriers can cross mucosal barriers and inducing a strong immune response [94]. Similarly, PLGA-NPs can induce immune responses intranasally; however, the immune responses correlated with the size of the molecule [175]. A potential caveat to the intranasal administration of LPS-based glycoconjugates is the role of LPS in asthma and pulmonary inflammation. In mouse models, inhaled LPS was shown to recruit eosinophils and neutrophils to the lungs and induce T helper 2 (Th2) lung inflammation in a dose-dependent manner [176,177].

As for the oral route, LPS-based glycoconjugates have been shown to act as potential boosters to oral live-attenuated vaccines when injected. A synthetic LPS-based glycoconjugate vaccine candidate against *V. cholera* O1 induced a three-fold higher level of stool anti-LPS IgA post-boost compared with oral priming with live attenuated *V. cholera* O1 alone [71]. Limited attempts were made to use glycoconjugates as an orally administered vaccine. However, a past study has shown that *E. coli* O-SP conjugated to horse serum albumin did not produce an immune response in mice [178].

8. Bioconjugation is a newer alternative for glycoconjugate production

In vivo protein glycan coupling technology (PGCT) involves the bioengineering of bacteria to enable the *in vivo* coupling of bacterial carbohydrates (like the O-SP of LPS) onto excreted carrier proteins by introducing exogenous genes, while disabling any genes that may perturb its efficiency. These genes include glycan synthesis loci from the carbohydrate of interest, genes encoding oligosaccharyl transferases (OTases), and genes encoding for carrier proteins [179]. Mechanisms, trends, and challenges of the PGCT technology is comprehensively documented in recent reviews [180–182].

The advantage of PGCT-derived bioconjugates compared with chemically conjugated glycoconjugates is that *in vivo* synthesis would eliminate intermediate steps, thus reduce costs associated with glycan synthesis/harvesting, detoxification, chemical conjugation, and quality control. To date, there are no licensed bioconjugate vaccines. Several LPS-based bioconjugate vaccine candidates are currently recruiting or are undergoing phase 2 clinical trials, this includes bioconjugates against *S. dysenteriae* type 1, *S. flexneri* 2a, and enteropathogenic *E. coli* (ClinicalTrials.gov Identifier: NCT04056117; NCT02646371; NCT03500679).

Several technical challenges of PGCT are yet to be addressed. The current technical challenges for PGCT can be summarized as (1) Current OTases are not optimally compatible with all bacterial glycans, producing ununiform bioconjugates; (2) Glycan biosynthesis loci can be challenging to identify and clone, thus not all PGCT can take place in nonpathogenic microbial cell factories as explained below.

8.1. Coverage and uniformity of chemically conjugated glycoconjugate vaccines compared with bioconjugate vaccines

LPS is commonly selected as a target for PGCT-derived bioconjugates [181]. Clinical and preclinical LPS-based bioconjugate vaccine candidate includes ones against *E. coli*, *F. tularensis*, *Burkholderia pseudomallei*, *S. dysenteriae*, and *S. flexneri* [24,183–187]. However, the spectrum of pathogens PGCT can cover is dependent on the compatibility between the glycan-based antigens and the OTases. The most widely used OTase, *Campylobacter lari* PglB, preferentially transfers glycans with a reducing end monosaccharide comprising an acetamido group [188]. Directed protein evolution applied to PglB enhanced its glycan compatibility, resulting in the successful transfer of glycans without a reducing end acetamido group; however, viability and efficiency of this variant may require further assessments [189]. Alternative OTases include *Neisseria meningitidis* PglL, which also transfers glycans in a preferential manner but has a more relaxed specificity than PglB [190]. Recent work had demonstrated the implementation of *Acinetobacter*-derived OTase, PglS, which demonstrated broader compatibility of glycans. PglS permits glycans containing glucose on the reducing end to be transferred [191].

However, loosening the glycan specificity can also introduce interference from other carbohydrates, thus contaminating the vaccine by making undesired glycoconjugates. Besides the nonspecific transfer of glycans, the uniformity of glycoconjugates is not guaranteed when synthesized *in vivo*. PGCT typically yields a pool of glycoconjugates with varied molecular masses depending on the length of the glycan, as seen in recent preclinical studies [191–193]. While the effects of this heterogeneity are not yet determined, classic chemical conjugation with synthesized carbohydrates bypasses this potential issue by producing uniform glycans for making glycoconjugates.

8.2. Safety and yield can be improved for PGCT systems

In most instances, the identification, cloning, and transfer of all genes relevant to glycan synthesis are necessary for PGCT-based bioconjugate production in well-studied microbial cell factories like *E. coli* [179]. As *E. coli* may encode genes for the same purpose as those present in the glycan biosynthesis loci, it is important to deduce the minimal genes required to synthesize the desired bioconjugate [194].

Typically, genes required for glycan biosynthesis are concentrated in a cluster and ubiquitously expressed. However, glycan synthesis genes can be scattered at several loci within

the genome for certain pathogens, making the identification and cloning of the genes necessary for bioconjugate production difficult. In these cases, in which the biosynthetic genes are not identified, or where it is difficult to transfer them to *E. coli*, studies have circumvented this complication by directly glycoengineering the pathogenic bacterial strains, which have shown to successfully yield bioconjugates. Generation of bioconjugate vaccine candidates against *S. enterica* serovar Paratyphi A and *S. flexneri* 2a was demonstrated directly from glycoengineered strains of these bacteria [192,195]. However, these bacteria retain pathogenicity and would require cautious handling of these in processes including fermentation, transformation, and harvesting.

As for yield, ~100-250 mg of LPS per liter of bacteria culture is typically isolated from the hot phenol-water method in small-scale preclinical production of chemically coupled glycoconjugates [51,63,196–198]. Conjugation efficiency of LPS to a carrier protein can vary depending on the conjugation chemistry and the desired conformation of the glycoconjugates but is typically >25% (based on the carbohydrate content) [196,198,199]. Thus, an overall estimate of glycoconjugates per liter of culture is >25 mg. In comparison, PGCT-derived bioconjugates typically provide >5 mg of bioconjugates per liter of culture, perhaps due to the need for media, culturing conditions, and bacterial strain optimization [195].

Additionally, in the case of introducing glycan biosynthesis genes via plasmids, the natural loss of plasmids may reduce the efficiency of PGCT. A previous study has indicated that PGCT efficiencies can be maintained by minimizing the time between cell culture inoculation and induction of PGCT [200].

9. Conclusion: toward promising LPS-based vaccines

Despite the increasing number of preclinical LPS-based glycoconjugate vaccine candidates that have arisen in recent years (Table 2), there is still a plethora of Gram-negative bacterial pathogens for which there is still no potent vaccine. These bacteria are viable targets for future LPS-based glycoconjugates. While antibodies to rough-LPS generally show weak protection, developing glycoconjugates which induce antibody responses to the O-SP moiety remains a feasible means of protection against many Gram-negative bacterial pathogens (Table 4).

In conclusion, it is likely that LPS-based conjugate vaccine candidates could begin to play an important role in protection against Gram-negative pathogens in the near future. The efficacy of these glycoconjugates could be improved through the use of conformation-specific conjugation methods, or the use of dual-role carriers. Also, it is important to be alert to the potential for CIES as more glycoconjugate vaccines are used and to monitor for evidence of BI. If the technical hurdles can be overcome, LPS-based conjugates are set to bring a new era of anti-Gram-negative vaccines that will improve human health.

10. Expert opinions

The need for broader immunization against Gram-negative bacterial pathogens has driven the rapid development of

LPS-based glycoconjugate vaccines over recent years. As an increasing number of LPS-based glycoconjugate vaccine candidates move toward clinical trials and potential licensure, the emerging demand for large-scale harvesting of LPS is on the horizon. While means to purify smooth LPS remain limited, the recently maturing *in situ* fermentation, purification, and detoxification techniques can deliver better economic prospects than classic hot phenol-water extraction followed by acid or alkaline detoxification [51,63]. In particular, alkaline detoxification has been dismissed in the production of LPS-derivatives due to their unknown toxicity [48]. As an alternative way to obtain LPS, synthetic and semi-synthetic glycan production shows promising yield and efficacy [69]. These platforms are likely to deliver O-SP for LPS-based conjugates with high purity and safety, which may in turn reduce the difficulty of meeting regulatory standards. However, efficacy studies are likely to be needed to elucidate the correlates of protection for LPS-based glycoconjugate vaccines. These studies may require costly investments and thus a potential regulatory hurdle to overcome for LPS-based glycoconjugates.

While novel carriers have shown promising preliminary results, using these carriers in the clinical development of vaccine candidates requires assessments of the efficacy, manufacturability, and scalability of the carriers, thus introducing additional regulatory hurdles in the process [80]. In particular, no licensed vaccines are incorporating nanoparticle-based carriers, which would require careful development and clinical validation. As for dual-role peptide-based carriers, the potential disruption in peptide epitopes when conjugating carbohydrates to carrier proteins can hinder the immunogenicity of the dual-role carrier proteins, requiring a careful selection of conjugation chemistries to avoid [81,104].

Several conjugation chemistries are available for LPS-based glycoconjugate vaccines, but some resulted in the loss of immunogenicity or manufacturability. In most instances, conjugation chemistries will aim to introduce suitable linkers to reduce steric hindrance between sugar and peptide moieties, while also preserving the relevant epitopes in the resultant glycoconjugates [81,119]. To satisfy these aims, methods including reductive amination, CDAP chemistry, and aminooxoxime chemistry have been more extensively incorporated in LPS-based glycoconjugate vaccine candidates than other techniques [114,118,120,144]. Compared to PGCT-derived glycoconjugates, chemically conjugated glycoconjugates bear higher yields while having fewer regulatory hurdles as they are based upon techniques developed over the past decades. However, the PGCT is a fast-evolving technology with rapidly increasing efficiencies, and thus further exploration of its potential is warranted.

Immunological factors are also essential to consider in glycoconjugate vaccine design, such as the interfacing between newly administered glycoconjugates with the pre-existing immunity to carrier proteins via CIES, or the potentially reduced immunogenicity with concomitant administration of glycoconjugate vaccines via BI. While CIES has been a persistent problem, novel carriers likely represent the potential solution. Nanoparticle-based carriers and dual-role peptide-based carriers are both likely to be viable avenues toward potential evasion of CIES [151,152]. As for BI, it is likely that

Table 4. Anti-LPS antibody as correlates to protection in selected pathogenic Gram-negative bacteria.

Pathogen	LPS moiety providing protectivity	Mechanism	Other antigens providing protectivity	Cross-reaction of anti-LPS antibodies	Level of protection	Reference
<i>Bordetella parapertussis</i>	O-SP	Affected the adherence of <i>B. parapertussis</i> to epithelial cells; may partially contribute to protection.	Unknown	Unknown	Partial protection	[201,202]
<i>Bordetella pertussis</i>	No	Rough LPS is a poor immunogen and fail to contribute to producing protective antibodies.	pertussis toxin (PT); filamentous hemagglutinin (FHA); pertactin (PRN); fimbriae (FIM 2/3)	with <i>Bordetella parapertussis</i>	Strong protection	[202–205]
<i>Brucella abortus</i>	O-SP	Complement-dependent bactericidal activities; Phagocyte-mediated immunity via opsonization.	Omp31	Unknown	Strong protection	[206–208]
<i>Burkholderia mallei</i>	O-SP	Phagocyte-mediated immunity via opsonization; may partially contribute to protection.	Unknown	with <i>Burkholderia pseudomallei</i> LPS	Partial protection	[94,209–211]
<i>Burkholderia pseudomallei</i>	O-SP	Phagocyte-mediated immunity via opsonization; IgG2 to LPS correlative to protection against lethal infection; may partially contribute to protection.	Unknown	Unknown	Partial protection	[211–214]
<i>Coxiella burnetii</i>	O-SP	Likely through phagocyte-mediated immunity via opsonization.	Unknown	Unknown	Strong protection	[215–217]
<i>Escherichia coli</i> rough strain J5	No	Rough LPS is a poor immunogen and fail to contribute to producing protective antibodies.	Unknown	Unknown	Strong protection	[218,219]
<i>Escherichia coli</i> O157	O-SP; R3 core oligosaccharide	Complement-dependent bactericidal activities; Phagocyte-mediated immunity via opsonization	Flagellin	with <i>Escherichia coli</i> O7, O116	Strong protection	[220–222]
<i>Escherichia coli</i> O111	O-SP		Unknown	Unknown		[223]
<i>Escherichia coli</i> O18	O-SP		Unknown	Unknown		[224]
<i>Francisella tularensis</i>	O-SP	Likely through phagocyte-mediated immunity via opsonization	outer membrane protein A (FopA)	Unknown	Strong protection	[225–228]
<i>Helicobacter pylori</i>	O-SP	Promote complement-dependent inflammation; may partially contribute to protection.	Adhesins	Unknown	Partial protection	[229–231]
<i>Legionella pneumophila</i>	O-SP	Phagocyte-mediated immunity via opsonization.	peptidoglycan-associated lipoprotein (PAL), macrophage infectivity potentiator (Mip), type IV pilin (PilE), and flagellin (FlaA)	Unknown	Strong protection	[232,233]
<i>Mannheimia haemolytica</i>	No	Rough LPS is a poor immunogen and fail to contribute to producing protective antibodies.	OmpA, SSA-1, Gs60, PlpE, TBPs, PlpF, OmpD15, and OmpP2	Unknown	Strong protection	[234–236]
<i>Moraxella catarrhalis</i>	No	Rough LPS is a poor immunogen and fail to contribute to producing protective antibodies; rough LPS can promote inflammation.	UspA	Unknown	Strong protection	[237–239]
<i>Neisseria meningitidis</i> group B	No	May weakly contribute to complement-dependent bactericidal activities; likely not contributing to protection.	Neisseria adhesin A (NadA); H-binding protein (fHbp); Neisseria heparin-binding antigen (NHBA); porin (PorA)	Unknown	Partial or no protection	[240–242]
<i>Pseudomonas aeruginosa</i>	O-SP; core oligosaccharide	Complement-dependent bactericidal activities; Phagocyte-mediated immunity via opsonization.	IROMPs, OprC, esterase, OprP, OprD1, D2, OprE, OprF, OprG, OprH1, H2, and OprI.	Unknown	Strong protection	[243–246]
<i>Salmonella enterica</i> serovar Typhimurium	O-SP	Complement-dependent bactericidal activities; Phagocyte-mediated immunity via opsonization; may partially contribute to protection.	Outer membrane proteins	Unknown	Strong protection	[39,247–250]
<i>Salmonella enterica</i> serovar Paratyphi	O-SP		FliC	with <i>S. Typhi</i> LPS		[251–253]
<i>Salmonella enterica</i> serovar Typhi	O-SP		FliC	with <i>S. Paratyphi B</i> LPS	Partial protection	[253,254]

(Continued)

Table 4. (Continued).

Pathogen	LPS moiety providing protectivity	Mechanism	Other antigens providing protectivity	Cross-reaction of anti-LPS antibodies	Level of protection	Reference
<i>Shigella flexneri</i> 2a	O-SP	Complement-dependent bactericidal activities	type III secretion system proteins (lpaB, lpaC); pan Shigella surface protein-1 (PSSP-1)	with <i>Shigella dysenteriae</i> LPS	Strong protection	[54,107,255,256]
<i>Shigella sonnei</i>	O-SP		type III secretion system proteins (lpaB, lpaC); pan Shigella surface protein-1 (PSSP-1)	Unknown		[54,255–257]
<i>Vibrio cholerae</i> O1 (Inaba)	O-SP	Complement-dependent bactericidal activities; antibody-mediated inhibition of adherence and colonization (in intestine)	Unknown	Unknown	Strong protection	[40,258–260]
<i>Vibrio cholerae</i> O1 (Ogawa)	O-SP		Unknown	Unknown		
<i>Vibrio cholerae</i> O139	O-SP		Unknown	Unknown		

Bl may emerge as LPS-based glycoconjugate vaccines start to reach licensure. It will also be of interest to identify any vaccine antigens prone to Bl, like that seen in CPS-based glycoconjugates [153,161,162]. Investigating the potential mechanisms for Bl would help avoid this problem, hence increasing the efficiency of vaccinations. However, in the cases of Bl, a reduction in immunogenicity can be acceptable if sufficient levels of efficacy are retained, while additional booster doses may help compensate for the reduction. Exploring different administration routes for LPS-based glycoconjugate vaccines is of interest. However, the pro-inflammatory properties of LPS might bring additional difficulties in developing LPS-based glycoconjugates for nasal administration.

Overall, we think that with the development of novel techniques in obtaining LPS, novel carrier proteins, and a better understanding of conjugation chemistries, LPS-based glycoconjugate vaccines are about to emerge as a new category of vaccines that will improve human health.

Funding

This manuscript is supported by the Innovate UK Vaccines for global development grant (project reference 971619). A J Pollard and C S Rollier are Jenner Investigators.

Acknowledgments

We acknowledge the support from the National Institute for Health Research (NIHR) Oxford Biochemical Research Centre and Innovate UK.

Declaration of interest

A J Pollard is Chair of UK Dept. Health and Social Care's (DHSC) Joint Committee on Vaccination & Immunisation (JCVI), and is a member of the WHO's SAGE. The views expressed in this article do not necessarily represent the views of DHSC, JCVI, or WHO. The University of Oxford has entered into a partnership with AstraZeneca on coronavirus vaccine development. The authors have no other relevant affiliations or financial involvement with any organization or entity with a financial interest in or financial conflict with the subject matter or materials discussed in the manuscript. This includes employment, consultancies, honoraria, stock ownership or options, expert testimony, grants or patents received or pending, or royalties.

Author contribution

We confirm all authors have (1) substantially contributed to the conception and design of the review article and interpreting the relevant literature, and (2) been involved in writing the review article or revised it for intellectual content.

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