Multivalent glycoconjugates as vaccines and potential drug candidates

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Pathogens adhere to the host cells during the first steps of infection through multivalent interactions which involve protein–glycan recognition. Multivalent interactions are also involved at different stages of immune response. Insights into these multivalent interactions generate a way to use suitable carbohydrate ligands that are attached to a basic scaffold consisting of e.g., dendrimer, polymer, nanoparticle, etc., with a suitable linker. Thus a multivalent architecture can be obtained with controllable spatial and topology parameters which can interfere with pathogen adhesion. Multivalent glycoconjugates bearing natural or unnatural carbohydrate antigen epitopes have also been used as carbohydrate based vaccines to stimulate an innate and adaptive immune response. Designing and synthesizing an efficient multivalent architecture with optimal ligand density and a suitable linker is a challenging task. This review presents a concise report on the endeavors to potentially use multi- and polyvalent glycoconjugates as vaccines as well as anti-infectious and anti-inflammatory drug candidates.

1. Introduction

Multi- and polyvalency play a crucial role in various biological processes like self-organization of matter, cell–cell recognition leading to signal transduction, detection, and immune response.1 Interactions between an m-valent receptor and an n-valent ligand (m, n > 1; also m ≠ n) are considered to be multivalent interactions. The term polyvalency is especially used in the context of ligands binding to receptors on interfaces such as cell membranes, which offer a large number (n = 10) of two-dimensionally distributed binding sites.1 The surface of a cell is covered with various carbohydrates which are usually attached to other biomolecules like proteins or lipids. Different pathogens like bacteria, fungi, viruses, toxins, etc., adhere to the host cells through multivalent interactions between carbohydrates and proteins as a first step in the infection process. The individual interactions between proteins and carbohydrates are weak but multiple simultaneous interactions lead to strong functionally useful avidities. The term avidity is defined in biochemistry as the accumulated strength of the multiple affinities of individual non-covalent binding interactions. Avidity was originally introduced to describe the binding behavior of immunoglobulins with different (multi) valencies (antibodies, IgG, IgEm, and IgG monomers: divalent; IgA homodimer: tetravalent; IgM homopentamer: decavalent).2 The recent advent of carbohydrate microarray technologies5,6 as well as advances in glycoimmunology and glycochemistry have paved the way for insights into these multivalent interactions. This new knowledge about protein–glycan interactions between pathogens and mammalian host cells has made multivalent glycoconjugates attractive scaffolds for targeting different pathogens. Therefore, several unnatural glycoconjugates with different valencies and the spatial arrangement of ligands based on scaffolds like proteins,5–7 polymers,8–10 fullerenes,11,12 calixarenes,13–15 dendrimers,16–20 nanoparticles,21,22 etc., have been reported. The anti-adhesive or cell targeting properties of these multivalent glycoconjugates can competitively inhibit the pathogens from adhering to the host cells which ideally would prevent or treat the disease caused by these pathogens. For example, a synthetic glycolipid Gb3 (α-D-Gal[1,4]β-D-Gal[1,4]β-D-Glc[1,0-ceramide]) analog that was covalently attached to insoluble silica particle has emerged as an anti-adhesive carbohydrate-based drug candidate for adsorbing shiga-like toxins of E. coli. This synthetic analog of the Shiga toxin (Stx) receptor (Synsorb Pk) is currently undergoing clinical trials.23 Furthermore, it has been found that surface sugar densities, as well as the length and nature of the linker used for conjugation to the basic scaffold, play a significant role in the optimum avidities and selectivities. Therefore, designing an efficient multivalent glycoconjugate remains a challenging task.24,25 Advances in the identification and synthesis of glycan epitopes on different pathogens have made carbohydrates attractive vaccine targets and immunomodulators. The structurally defined glycoconjugates displaying these unique glycan antigen structures can follow a lectin-mediated cellular uptake mechanism to generate an immune defense mechanism in the body against chronic infections. In spite of the existing challenges in the isolation and identification of the carbohydrate antigen itself and in the preparation of structurally similar glycan

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epitopes, a fruitful effort has been made in this area and a number of conjugate versions of polysaccharides vaccines are now commercially available or in clinical trials. An octavalent O-polysaccharide (OPS) with toxin A conjugate, for example, has been found to be immunogenic and clinically effective against *Pseudomonas aeruginosa* and is currently undergoing phase III clinical trials. This review is a concise overview on the different attempts to apply multivalent glycoconjugates as drugs in pathological conditions like infections or cancer. It highlights the efforts being made by various research groups to design multivalent glycoconjugates that can be used to inhibit the adhesion of bacteria, bacterial toxins, and viruses to the host cells or to stimulate the immune systems against pathogens like HIV. The endeavor to apply multivalent glycoconjugates as anti-inflammatory drugs by targeting leukocyte trafficking will also be discussed.

2. Vaccines

The unique glycoalyx and its dense surface distribution on different pathogens and malignant cells have made carbohydrates attractive vaccine targets. Glycan antigens are diverse and range from capsular polysaccharides to small monosaccharide antigens. Capsular polysaccharides have been in use for decades for providing immunity against different bacterial infections. The main bottleneck in the development of a carbohydrate vaccine, however, is to identify carbohydrate antigen itself. In recent years, the identification and synthesis of the unique glycan epitope present on the various pathogens and malignant cells has been more efficient. This has paved a way for the development of carbohydrate-based vaccines for eliciting the immune response to the different infections and cancer as well. Another obstacle associated with the development of carbohydrate-based vaccines is the poor antibody response to carbohydrates which is usually due to the T-cell independent immune responses triggered by repetitive carbohydrate antigens. Carbohydrate specific antibodies have a low affinity with dissociation constants in the micromolar range and glycans interact mainly rely on the avidity effects of multivalent interactions. Avery and Goebel reported that immunogenicity of carbohydrates can be increased by covalent attachment of glycans to a suitable immunogenic protein which has led to the discovery of conjugate vaccines. In the carbohydrate-based vaccines glycan antigens are conjugated with a source of helper T-cell epitope, usually with an immunogenic carrier protein. Taxoids, metalloproteins like keyhole limpet haemocyanin (KLH) or virus capsids are also used as carriers. Several strategies can be used for linking the carrier to the sugar but one needs to carefully investigate the immunogenicity of these linkers to the glycan antigens. Adjuvants are also included in the vaccine to increase the immunogenic response to antigens and/or modulates it towards the desired immune responses. Advantages of adjuvants include the reduction of the antigen amount needed for a successful immunization, the reduction of the frequency of booster immunizations needed, and an improved immune response in elderly and immunocompromised patients. Adjuvants include, for example, mineral salts, oil emulsions, surfactant based formulations, microbial derivatives, and particulate adjuvants (virosomes). With a few exceptions, alum (potassium aluminium sulfate) is still the only adjuvant that has been approved for human use in the majority of countries worldwide. Therefore, it is found in numerous vaccines, including diphtheria–tetanus–pertussis, human papillomavirus, and hepatitis vaccines. QS-21, a saponin derived from the tree bark of *Quillaja saponaria*, is also being clinically tested. Tripalmitoyl-S-glyceryl-cysteinylerine (Pam₃Cys) has also been discovered to have adjuvant properties. Fig. 1 depicts some of the different strategies for preparing glycoconjugate vaccines, i.e., multivalent synthetic conjugates, protein conjugates, and lipid conjugates.

2.1. Multivalent glycoconjugate vaccines against different pathogens

A diverse and dense array of protective capsular and lipopolysaccharides is present on the surface of bacterial pathogens. Heterogeneity and complexity of the capsular polysaccharides on bacterial pathogens have always been a great challenge in the development of multivalent carbohydrate-based vaccines. For example, more than 90 serotypes are known for *S. pneumonia*. The first polysaccharide based vaccine, Pneumo Vax from Merck and Co. launched in 1984 was composed of unconjugated capsular polysaccharide isolated from 14 pneumonia serotypes; the current development provides active immunization against 23 serotypes. In order to develop a multivalent conjugate vaccine against different bacterial pathogens, polysaccharides are generally isolated from most clinically relevant serotypes and then degraded into smaller products for activation and conjugation to immunogenic carrier proteins. For example, Prevnar from Wyeth/Pfizer is a heptavalent pneumococcal conjugate.

While developing the synthetic carbohydrate-based vaccines, the effect of chain length and saccharide density on the potent antibody response can be addressed. For example, conjugates with different saccharide densities of tetra-, octa-, dodeca-, and hexadecasaccharides were prepared for vaccine against *S. dysenteriae*. These conjugates were based on a tetrasaccharide repeat unit of *O*-specific oligosaccharides. In the process it was found that octa-, dodeca-, and hexadecasaccharides were immunogenic and that the chain length determined the optimal loading densities of saccharides. Furthermore, the nonreducing end of *Shigella dysenteriae* type 1 *O*-specific oligosaccharides had an effect on their immunogenicity as conjugates in mice. For many bacterial infections, multivalent glycoconjugate vaccines have been developed based on the fragments of their capsular polysaccharides. Examples include *Streptococcus pneumoniae*, *Neisseria meningitides*, and *Haemophilus influenzae*. Table 1 shows some examples of commercially available multivalent glycoconjugate vaccines.

Infection by parasitic protozoans and helminths is still a big threat to the human population. Over 15 million deaths occur worldwide per year due to parasitic infections and diseases, and no commercial vaccine has been developed as yet for any of
the parasitic infections like malaria, leishmaniasis, African trypanosomiasis, amoebiasis, schistosomiasis, and lymphatic filariasis. This is due to the complex immune evasion mechanisms used by the different parasites and to the lack of understanding of molecular interactions between host immune cells and parasites. The unique glycan antigens are abundant and accessible on the surface of the multiple developmental stages of different parasitic protozoans and helminthes which makes glycans attractive vaccine targets. Difficulties in obtaining enough parasitic material for study also limit the progress in this area. Several efforts are, however, being made to investigate potential vaccine antigens using advanced diagnostic techniques for glycan detection and chemical or chemo-enzymatic methods.

### Table 1: Some commercially available multivalent glycoconjugate vaccines

| Vaccine                                      | Manufacturer (trade name)                                      |
|----------------------------------------------|--------------------------------------------------------------|
| Pneumococcal vaccine, polyvalent            | Merck & Co, Inc. (Pneumovax 23)                              |
| Pneumococcal 7-valent conjugate (Diphtheria CRM197 protein) | Wyeth Pharmaceuticals Inc. (Prevnar)                          |
| Pneumococcal 13-valent conjugate (Diphtheria CRM197 protein) | Wyeth Pharmaceuticals Inc.                                   |
| Meningococcal (groups A, C, Y, and W-135) oligosaccharide diphtheria CRM197 conjugate | Novartis Vaccines and Diagnostics, Inc. (Menveo)             |
| Meningococcal (groups C and Y and *Haemophilus b*) tetanus toxoid conjugate | GlaxoSmithKline Biologicals (MenHibrix)                      |
| Meningococcal (groups A, C, Y and W-135) polysaccharide diphtheria toxoid conjugate | Sanofi Pasteur, Inc. (Menactra)                               |
| Meningococcal polysaccharide vaccine, groups A, C, Y and W-135 combined | Sanofi Pasteur, Inc. (Menomune-A/C/Y/W-135)                   |

*The information in the table has been taken from a complete list of vaccines licensed for immunization and distribution in the United States, which is available from the US Food and Drug Administration website [http://www.fda.gov/BiologicsBloodVaccines/Vaccines/ApprovedProducts/ucm093833.htm]. CRM197 is a nontoxic mutant of diphtheria toxin.*
with *Schistosoma* spp. and *Leishmania* spp. have been identified and are being evaluated as immunogens.\(^{71,42,44}\)

It is known that each type of parasitic protozoa produces its own unique and highly complex glycan structure. African trypanosomes, for example, keep genetically switching their highly immunogenic glycosylphosphatidylinositol (GPI)-anchored surface glycoprotein which is called variant surface glycoprotein (VSG).\(^{45}\) Virtually all protozoans, including *Plasmodium* parasites that cause malaria synthesize GPI. The GPI synthesized by *Plasmodium* increases, further infection could not be prevented.\(^{57}\)

Preclinical studies reported by Seeberger and coworkers showed that a synthetic version of GPI conjugated to keyhole limpet haemocyanin (KLH) (Fig. 2) elicited high titres of antibodies in mouse models increased, further infection could not be prevented.\(^{46}\)

The same group also presented a versatile approach based on a synthetic GPI glycan array to correlate anti-GPI antibody levels and protection from severe malaria but the question remains whether an antitoxic immune response to malarial GPI provides protection against malaria.\(^{46}\)

Some glycoconjugate biosynthetic studies done on *Leishmania* protozoans have revealed that they also generate a surface lipophosphoglycan (LPG) with unique phosphodiester linkages and mannose in the backbone and furanosyl galactoside modifications that might be contained within other based phospholipids, GPI-anchored glycoproteins, mucin-type membrane, and secreted glycoproteins.\(^{44,49-51}\) The recent automated syntheses of *Leishmania* related and other antigenic glycans reported by Seeberger and coworkers will give a tremendous boost to the generation of libraries of conjugate vaccine candidates for screening.\(^{52}\)

### 2.2. HIV vaccine development and challenges

The dense conserved cluster of oligomannose glycans on the envelope glycoprotein, gp120 subunit of Human Immunodeficiency Virus type 1 (HIV-1), masks conserved protein epitopes and facilitates virus entry via interaction with glycan binding proteins on susceptible host cells. The broadly neutralizing monoclonal antibody 2G12 neutralizes a wide range of HIV-1 strains in animal models by specifically binding to the cluster of oligomannose glycans on gp120.\(^{53-56}\) New broadly neutralizing antibodies that also recognize and penetrate HIV glycan shield have recently been isolated from patients.\(^{57}\) Highly dense clusters of oligomannose on gp120 are stabilized by a network of intermolecular hydrogen bonding and thus led to a close spacing between the carbohydrates which is unusual in mammalian glycoproteins. These observations provide the basis for immunological discrimination by 2G12.\(^{58}\) Using the information available by structural and modeling studies on how 2G12 interacts within the oligomannose clusters of gp120,\(^{59}\) efforts have been made to design an immunogen that elicits 2G12 like antibodies.\(^{54}\) This is a very challenging task due to the unique recognition mode of 2G12.

Several multivalent oligomannoside ligands containing Man\(_4\) (1,2)Man, including GlcNAcMan\(_9\) for 2G12, have been synthesized for mimicking the densely conserved oligomannose clusters in the glycan shield on gp120. The synthetic tetramannoside (Man\(_9\)) corresponding to the D1 arm of GlcNAcMan\(_9\) was found to be the minimum recognition motif and as effective as GlcNAcMan\(_9\) itself in binding to gp120.\(^{52}\) Synthetic strategies for the multivalent presentation of oligomannose have involved the use of oligodendrons, QB bacteriophage, and generation of cyclic glycopeptides\(^{60-63}\) (Fig. 3). A synthetic Man\(_4\) on bovine serum albumin (BSA) molecule by Burton and coworkers,\(^{44}\) and a cyclic glycopeptides conjugate of GlcNAcMan\(_9\), further conjugated with Outer Membrane Protein Complex (OMPC) by Danshisky and coworkers,\(^{64,65}\) could be used to induce a 2G12 like antibody response, but antibodies did not bind to the gp120. In another study by Geng and coworkers,\(^{66}\) a mutant strain of *Saccharomyces cerevisiae* containing a large number and high density of N-linked glycans elicited gp120 cross-reactive antibodies in rabbit, but these antibodies did not neutralize HIV. The current challenge lies in breaking the immune tolerance against HIV-1 strains by eliciting the antibodies with proper specificity. This is due to an inappropriate mimicry of the glycan shield (flexible glycans) and the unique architecture of the current model antibody, 2G12. Moreover, there is a need for more fundamental studies on B cell immunology to break the immune tolerance towards the HIV-1 carbohydrate antigens.

### 2.3. Cancer vaccine

Tumor associated carbohydrate antigens (TACAs) are over-expressed self glycans in cancer cells which occur because of the altered glycosylation.\(^{69}\) There is more glycosyltransferase expression in cancer cells than in normal tissues which leads to an increase in the size and branching of N-glycans, as well as to more sialylation,\(^{70,71}\) and linking terminal residues to glycans which results in an overexpression of certain terminal glycan epitopes on tumors, such as sialyl Lewis X, sialyl Lewis A, sialyl 2-6-α-N-acetylgalactosamine (sTn), sialyl Lewis Y, globo-hexaosylceramide, and PSA.\(^{72-74}\) Although TACAs may potentially

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**Fig. 2** Synthetic *Plasmodium*: glycosylphosphatidylinositol-keyhole limpet haemocyanin (GPI-KLH).\(^{45}\)
serve as vaccine antigens, they are poorly immunogenic and it is difficult to isolate the unique TACA glycan structure from natural sources. A common way to break tolerance towards TACAs is to use a monovalent carbohydrate antigen in multiple copies on an immunogenic carrier such as KLH in the presence of a strong adjuvant like QS21. This strategy describes the first generation monomeric cancer vaccine. Investigations on antigen conjugates have established that KLH and QS21 are the most potent carrier and adjuvant pair for breaking tolerance towards TACAs. There are several studies where natural (glyco-sphingolipid containing sialic acids like GD2, GD3, GM2, and fucosyl GM1) and unnatural or synthetic (N-propionylated PSA, GD2- and GD3-lactone, globohexaosylceramide and Lewis Y) carbohydrate antigens that have been conjugated with KLH in the presence of QS21 have been found to be immunogenic in cancer patients. A novel approach to break the immune-tolerance to TACAs is to combine cell glycoengineering with vaccines prepared using unnatural TACAs. The critical step in this new strategy is metabolic engineering of cancer, namely, to induce expression of an artificial form of a TACA by supplying tumors with an artificial monosaccharide precursor. For this purpose, tumor cells were incubated with different N-acyl-d-mannosamines, and modified forms of GM3 expressed on tumor cells were analyzed by flow cytometry using antigen-specific antisera. Recent reports by Guo and coworkers showed that tumor cells expressed unnatural GM3 analog, i.e., GM3NPhAc instead of natural GM3 when incubated in the presence of N-phenylacetyl-D-mannosamine. The GM3NPhAc-protein conjugates are particularly immunogenic and produce strong T-cell dependent antibodies. Therefore, the new glyco-engineered melanoma cells expressing unnatural GM3NPhAc were selectively killed by GM3NPhAc-specific antibodies produced by GM3NPhAc–KLH conjugates. These results suggest the feasibility of selectively eradicating cancer in vivo without affecting the normal cells. This immunotherapy may be applicable to other tumors as long as they express GM3.

The development of multivalent vaccines has been encouraged by the heterogeneity of TACA expression on malignant cells. Danishefsky and coworkers have prepared pentavalent and hexavalent carbohydrate antigenic constructs on a single peptide backbone, conjugated with KLH and Pam3Cys. The early immunogenicity studies in mice showed that pentavalent unimolecular conjugates (on KLH or Pam3Cys) were much more immunogenic than the corresponding pool of monomeric KLH conjugates (Fig. 4). Recent clinical studies, however, showed that lower IgM and IgG antibodies titres were produced against individual antigens in high risk patients who had been immunized with hexavalent vaccines including GM2, Globo H, Lewis Y, glycosylated MUC-1-32mer, Tn, and TF. These vaccines were prepared in a clustered formation, conjugated to KLH, and mixed with QS-21. Mucin type glycans such as α-N-acetyl-β-galactosamine (αGalNAc, the Tn antigen glyccan) are overexpressed on the surface of breast cancer cells. Cameron and coworkers have recently reported the preparation of Tn-glycan antigen-based glycopolymers which were conjugated to gold nanoparticles and yielded ‘multicopy multivalent’ glyco-nanoparticles with a size range of 5–25 nm. These glyco-conjugated gold nanoparticles produced a strong and long-lasting production of antibodies that were selective to the Tn-antigen glycian and cross-reactive towards mucin proteins displaying Tn.

3. Multivalent glycoconjugates as antiviral compounds

Multivalent binding interactions between glycoproteins on the surface of different pathogens and cellular receptors/oligosaccharides play a crucial role in the initial stages of cellular infection. For example, Fig. 5 shows how the multivalent interactions between hemagglutinin glycoproteins on influenza virus and sialic acid residues on cells lead to endocytosis and
then infection. In the recent past, multivalent glycoconjugates on dendritic/globular or linear scaffolds have emerged as potent inhibitors of infection by different viruses like HIV-1, Ebola, influenza, etc., which will be discussed in the following sections by focusing on the most relevant studies.

### 3.1. DC-SIGN mediated infection inhibition

Dendritic cells on mucosal surfaces carry a calcium dependent (C-type) lectin that is a dendritic cell-specific intercellular adhesion molecule (ICAM 3) grabbing non-integrin (DC-SIGN). This lectin can bind to a broad spectrum of pathogens: viruses (HIV-1, HIV-2, SIV, Ebola virus, hepatitis C virus, cytomegalovirus and dengue virus), bacteria (*Mycobacterium tuberculosis*, *Helicobacter pylori* and *Klebsiella pneumonia*), yeasts (*Candida albicans*), and parasites (*Leishmania* and *Schistosoma*).\(^\text{90}\) The carbohydrate recognition domain (CRD) of DC-SIGN recognizes the highly mannosylated and fucosylated glycoproteins which are found on the surface of certain pathogens. For example, highly mannosylated glycan shields on gp120 of HIV-1 are specifically recognized by DC-SIGN of dendritic cells on mucosal surfaces. The HIV-1 virus is taken by these dendritic cells to the lymph nodes where they effectively transfer the virus by a so-called trans infection of T-lymphocytes where the viral replication occurs. Multivalent glycoconjugate systems have been designed to mimic the surface of the virus to compete with natural ligands for binding with receptors for the development of carbohydrate-based antiviral drugs against HIV and Ebola virus infections. Recent investigations by Bernardi, Clerici, and coworkers have shown that a tetravalent dendron with four copies of linear trimannoside inhibits DC-SIGN mediated trans HIV infection process of CD4+ T lymphocytes in the presence of elevated viral loads in cellular and cervical explants models (Fig. 6).\(^\text{91,92}\) They found that multivalent presentation of linear di- and trimannosides on G3 Boltron-type dendrimers (32 copies) strongly inhibited the cell infection by Ebola pseudotyped viral particles by blocking DC-SIGN receptor showing IC\(_{50}\) values in the nanomolar range (Fig. 6).\(^\text{93}\) In another effort to mimic dense mannose clusters on HIV-1 envelope, Penades and coworkers prepared a series of gold nanoparticles with different spacers and different densities of oligomannoside. The mannoGNPs containing the disaccharide Manz1-2Manz were found to

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**Fig. 4** Structure of uimolecular pentavalent vaccine.\(^\text{87}\)

**Fig. 5** Multivalent interactions between trimeric haemagglutinin (HA\(_3\)) on influenza virus and sialic acid (SA) residues on cells leading to endocytosis and infection.
be the best inhibitors, because they increase inhibition 20 000-fold (100% inhibition at 115 nM) compared to the corresponding monomeric disaccharide (100% inhibition at 2.2 mM).\(^{44}\) Furthermore, these multivalent manno-glyco nanoparticles inhibited DC-SIGN-mediated HIV-1 trans-infection of human T cells at nanomolar concentrations in experimental setting that mimicked the natural route of virus transmission.\(^{35}\)

3.2. Influenza virus inhibition

Influenza viral infection is one of the biggest challenges for medicinal science and a big threat to the human population. It often causes epidemics and pandemics. Three glycoenzymes present on the virus surface, i.e., neuraminidase (NA), M2 ion channel protein, and hemagglutinin (HA) are important targets for fighting influenza. HA glycoprotein is trimeric and most abundant (~600 copies per virus particle) on the surface of influenza virus.\(^{99}\) There is no multivalent drug available till now which can target HA glycoprotein. The two glycoenzymes HA and NA determine the viral subtype. Commercially available unimolecular drugs target either NA (zanamivir/Relenza, Oseltamivir/Tamiflu) or M2 ion channel protein (amantadine and rimantadine) activity. However, the emergence of stable and transmissible drug-resistant influenza strains can render these drugs ineffective. The three-dimensional structure of HA complexed with SA reported by Wiley and coworkers\(^{46}\) led to the discovery that HA glycoproteins on the influenza virus play a significant role in adhering virus to the susceptible cells in a multivalent interaction between three sialic acid binding sites on its globular domain with sialyloligosaccharide moieties of cellular glycoconjugates. Therefore, macromolecular scaffolds bearing multiple copies of sialic acid residues can be of great use for a competitive inhibition of the virus attachment to the cells. Efforts are in progress for the design, synthesis, and investigation of such multivalent glycoconjugates which may lead to the discovery of anti-influenza drugs.

So far the synthetic HA inhibitors mainly consisted of low molecular weight scaffolds that form clusters of mono,\(^{77,96}\) bi,\(^{99}\) tetra,\(^{100}\) and octavalent sialosides.\(^{101}\) The octavalent sialosides were based on the polyzaido-calix[4]arene scaffold but they showed only a moderate glycoside cluster effect and minimum inhibitory concentration in the millimolar range. Both natural and synthetic multivalent carriers have been explored to prepare multivalent sialosides conjugate. Natural carriers include the usage of self-assembling sialo-glycopeptides,\(^{102}\) proteins (specifically equine α2-macroglobulin and guinea pig α2-macroglobulin),\(^{100}\) and polysaccharides.\(^{103}\) Synthetic carriers that have been typically used are spherical macromolecules like dendrimers\(^{104}\) and liposomes,\(^{105}\) linear polymers,\(^{106}\) and nanostructures.\(^{107}\) Sialylglycopolymers have been studied the most in detail. Whitesides and coworkers have extensively studied α-sialoside-polyacrylamide copolymers to inhibit the agglutination of erythrocytes by influenza A virus. The best copolymer in the series inhibited hemagglutination \(10^4\) to \(10^5\) times more strongly than did similar concentrations of α-methyl sialoside.\(^{108}\) They also prepared C-glycoside by coupling amino-sialosides with preactivated esters, i.e., poly[N-(acryloyloxy)succinimide] (pNAS).\(^{109}\) Using the C-glycoside of sialic acid made it possible to investigate hemagglutinin inhibition at different temperatures from 4–36 °C because C-glycosides cannot be hydrolyzed by the neuraminidase glycoproteins present on influenza virus. Extensive studies were performed to optimize sialylglycopolymers by varying the chemical structure and content of the ligand in the polymer and by altering the structure of the side chains in the carrier which has led to inhibitors with inhibitory concentrations in the nanomolar range and were thousand-fold more effective than the most potent natural inhibitor, equine α2-macroglobulin. These polymeric sialosides were quite effective but they failed to become drugs due to the severe toxicity of their polyacrylamide backbone. Whiteside and coworkers only tested these polymeric inhibitors for a single influenza virus strain X31. Other groups, however, observed a significant variation in the sensitivity of these inhibitors when tested against different virus strains.\(^{109}\) Haag and coworkers chemically functionalized gold nanoparticles sized between 2 nm and 14 nm with sialic acid-terminated glyceral dendron. The 14 nm sialic acid conjugated AuNPs were nontoxic to cells and had a high affinity for HA on the influenza virus surface as observed by electron microscopy techniques and biochemical analysis (Fig. 7).\(^{110}\) The infection inhibition amounted to 30% which led to the hypothesis that virus-sized particles could further improve the inhibition of the infection.\(^{111}\) Recently, Haag and coworkers investigated a new class of biocompatible dendritic multivalent sialo conjugates based on appropriately functionalized polyglycerol nanogels (nPG) which were varied in size between 40–100 nm, respectively (Fig. 8). The size and ligand density on the nanoparticles were optimized, whereby the 50 nm sized sialic acid conjugated
nPG with only 12% ligand density was found to be the best conjugate showing 80% inhibition of the influenza virus A virus (strain A/X31).

Chen and coworkers recently reported that poly-L-glutamine conjugated with multiple copies of drug zanamivir (PGN-ZA) via a flexible linker inhibited influenza viral fusion and release at subnM concentrations of zanamivir. These PGN-ZA polymers were still effective against isolated ZA- and oseltamivir-resistant influenza virus.

4. Antibacterials

Bacteria and bacterial toxins infect host cells via multivalent interactions involving protein–glycan recognition in the first steps of infection. Bacteria constitute a large domain of prokaryotic microorganisms which can be typically a few micrometers in length, whereas bacterial toxins are small poisonous substances secreted by living bacteria during exponential growth. Millions of deaths per year are caused by shiga and cholera toxin infections. Both these toxins belong to the clinically significant subset of bacterial AB₅ toxins, where A subunit is enzymatically cleavable and B₅ homopentamer has receptors for oligosaccharide recognition on the cellular surface. Fig. 9 shows, for example, an inhibited cellular uptake of a pentameric bacterial toxin by a sugar-conjugated pentacyclen core. Bundle and co-workers designed and improved some oligovalent, water soluble carbohydrate ligands called “Starfish” and “Daisy” to inhibit Shiga toxin (Stx) 1 and Stx2 which were generated by *E. coli* strains. The inhibition potential was confirmed with *in vivo* experiments. Multivalent glycoconjugates based on different scaffolds like pentacyclen, dendrimers, polymers, fullerenes, cyclodextrins, cyclopeptides, and gold nanoparticles have been developed which can interfere with the pathogen (bacteria or bacterial toxin) adhesion process. However, the resulting sugar density at the glycoconjugate surface as well as the chemical structure of the linker groups are important factors for an effective inhibitor. To determine the corresponding partner for the glycan-binding protein (GBP) of the pathogenic microorganisms, glycan microarrays were also developed to scan a great number of glycan candidates in a short time. The combination of active glycans covalently linked to various carrier platforms resulted in powerful compounds which should be able to treat infectious diseases. These multivalent glycoconjugates may play a role in fighting antibiotic resistant bacteria strains as well.

4.1. Cholera toxin inhibition

Cholera is an intestinal infection caused by the cholera toxin (CT) produced from *Vibrio cholera* results in fast
dehydratation and shock caused by watery diarrhea. The seven cholera pandemics over the past 185 years affected nearly the whole world. Cholera deaths are still common in remote areas where effective treatment is unavailable.\textsuperscript{121} Conjugative-plasmid mediated multiply antibiotic-resistant (including to tetracycline) \textit{V. cholerae} O1 (MARV) has also emerged as a major problem, first in Tanzania then in Bangladesh. Antibiotic strains that are resistant to tetracycline, ampicillin, kanamycin, streptomycin, sulphonamides, trimethoprim, and gentamicin have appeared in several cholera endemic countries during the past two decades.\textsuperscript{121}

CT belongs to a family of bacterial AB\textsubscript{5} toxins with an enzymatically active A subunit which is uptaken by the mammalian cells \textit{via} an oligosaccharide recognition of the B\textsubscript{5} subunit. CT’s first mode of action is to strongly bind to the receptor cells on the cell surface. The receptor for the CT is a specific glycolipid, the ganglioside GM1, a regular component in the outer leaflet of many cell membranes. The A subunit is connected to the B-pentamer \textit{via} a disulfide bridge. After cellular uptake, the A subunit is activated and released in the cytosol by reduction of the disulfide bond.\textsuperscript{24,122,123} Several multivalent glycoconjugates have been explored to target the B\textsubscript{5} subunit of cholera toxin to prevent its cellular uptake.

Different multivalent glycoconjugates, bearing pseudo-trisaccharides and four to eight oligo-GM1 units, have been prepared based on calixarenes, poly(propylene imine), and PAMAM dendrimers. The B\textsubscript{5} pentamer unit of CT had a much higher affinity for these conjugates than native GM1.\textsuperscript{124-126} Pieters and coworkers synthesized multivalent versions of the GM1os using glycodendrimers using the click chemistry approach. These multivalent versions strongly inhibited the CTB\textsubscript{5}-subunit. For example, the most complex multivalent GM1os conjugate, which was an octavalent glycodendrimer, was 380 000-fold more potent with IC\textsubscript{50} = 50 pM than the monovalent GM1-mimic\textsuperscript{127} (Fig. 10). This GM1-mimic was compared with natural GM1 gangliosides and showed a similar binding to the CTB\textsubscript{5}-subunit in an ELISA-type assay.\textsuperscript{128} Multivalent galactose inhibitors were prepared as simplified versions of multivalent GM1 derivatives. Multivalent binding interactions compensated the low binding affinity of galactose for CTB\textsubscript{5}-subunit. Tetravalent and octavalent galactose inhibitors with long spacer arms, for example, competed well with natural GM1os.\textsuperscript{129}

Fan and coworkers explored a series of multivalent ligands based on different glycoside moieties with linkers of various lengths for CT inhibition. In the first approach, a semi-rigid pentacyclen core was used to bind 5 copies of \textit{m}-nitrophenyl-\textit{d}-galactoside (MNPG) \textit{via} a semiflexible linker. The MNPG unit represents the active ligand which can be bound to the CT receptor. The monovalent MNPG finger showed 100-fold greater inhibition in comparison to the weaker binding monovalent galactose ligand. In terms of multivalency, the pentavalent MNPG ligand exhibited a 260-fold better affinity enhancement (IC\textsubscript{50} = 0.9 \textmu M) in comparison to the monovalent MNPG finger.\textsuperscript{130} The second approach involved the design of branched multivalent ligands with 5-fold symmetry. The branched bivalent galactose ligands were attached to the pentacyclen core \textit{via} flexible arms which led to branched decavalent conjugates. A million-fold improvement in inhibition power (IC\textsubscript{50} = 0.04 \textmu M) was observed with the best branched decavalent ligand with a repeating unit of the linker \(n = 4\) in comparison to the monovalent galactose. The compound with a lower number of linker units showed less inhibition affinity. DLS studies proved that the bivalent ligand can complex the CT in a 1 : 2 manner. The results of Fan and coworkers showed that the rational design of these multivalent ligand and their linker size is very important for CT inhibition\textsuperscript{131} (Fig. 11).

Richards and coworkers examined several glycopolymers based on a poly(pentafluorophenyl methacrylate) precursor with varying saccharide density, linker length, and chain length due to a 1,3 dipolar cycloaddition. Compounds with a longer

\begin{figure}
\centering
\includegraphics[width=\textwidth]{fig9.png}
\caption{Inhibition of the cellular uptake of a pentameric bacterial toxin by a sugar-conjugated pentacyclen core. (a) Cellular uptake of a pentameric bacterial toxin. The whole toxin binds to cellular receptor, followed by endocytosis. (b) Inhibition of a toxin binding to cellular receptors using pentavalent pentacyclen based glycoconjugate.}
\end{figure}
linker length showed higher CT inhibition. Polymers with the highest and lowest sugar density (100% and 10%) showed the highest activity, which demonstrates the complexity of the glycoconjugate design for CT inhibition.  

4.2. Escherichia coli (E. coli) inhibition  

E. coli is a Gram-negative bacterium that easily adheres to the target cells in the urinary tract through a mannose-binding lectin FimH. The FimH lectin is located at the tip of type 1 fimbriae, also called pili, which are the most abundant structures on the surface of Gram-negative bacteria. The uropathogenic E. coli strains can bind to the carbohydrate receptors in the urinary tract through FimH, colonize, and then cause bladder infections. This bacterial adhesion can be prevented by competitive mannosidic inhibitors. Effective inhibitions have been shown by compounds with terminal α-D-mannosyl residues.  

Lindhorst and coworkers developed various saccharide modified nanodiamond (ND) conjugates to detect and remove E. coli bacteria from aqueous solutions. The surface of the thermally annealed ND was functionalized via a Diels–Alder cycloaddition of the π-bonds on the diamond surface (dieneophil) and an in situ generated o-quinonimethane (dienen). The covalently aryalted ND can easily be sulfonated and the sulfonic acid groups can be subsequently partially reduced to target thiol groups for more flexible functionalization of the ND. These modified NDs were used as platforms and were conjugated via a “thiol–ene reaction” with different monosaccharides. These NDs are also applied to inhibit biofilm formation on biotic surfaces or medical devices due to E. coli attachment. These biofilms result in an increasing source of infections in clinical settings. Barras and coworkers developed ND that were modified with mannose moieties via an azide–alkyne Huisgen cycloaddition. These multivalent glycol–nanodiamonds can inhibit the E. coli type 1 fimbriae-mediated adhesion to eukaryotic cells (T24 bladder cells) and reduce the E. coli biofilm formation.  

A second carbon-based platform inhibiting pathogens with a lower valency is provided by multivalent fullerenes. Vincent and co-workers developed “fullerene sugar balls” with twelve mannose moieties per fullerene. Different sugar units were grafted on the fullerene core via a copper-catalyzed 1,3-dipolar
cycloaddition to achieve a C_{60} core which was completely surrounded by sugar residues. Durka and coworkers used this method to couple twelve mannose moieties with different linker lengths to the fullerene core to inhibit the FimH of E. coli. The affinity of the mannoside fullerences to the FimH was determined by ITC and SPR. Those experiments showed that the increased distance between the active mannose moieties and the C_{60} core can considerably improve the binding affinity to FimH. However, the nanomolar affinity values were measured for both methods. The mannoside fullerene had a benzylic aromatic ring at the anomeric position of the sugar as well as an amide group in the linker structure, which seems to be beneficial for lectin binding and can accommodate up to seven FimH molecules.

A third class of glycoconjugates which can inhibit E. coli bacteria are multivalent sugar functionalized pillar[5]arenes. Nierengarten and coworkers developed a pillar[5]arene glycocluster with ten peripheral acetylated mannosyl groups via a 1,3 dipolar cycloaddition. The final product, which is a promising compound for FimH inhibition, was achieved by deacetylation. The inhibition capacity was determined in a hemagglutination inhibition assay. The assay determines the concentration of the glycoconjugate, which is necessary to prevent adhesion of uropathogenic E. coli with guinea pig’s red blood cells. The inhibition titre was in a low μM range (5.91 μM).

Huang and coworkers used a similar approach and “clicked” galactose groups (hydrophilic compartment) and alkyl chains (hydrophobic compartment) via an azide–alkyne Huisgen cycloaddition to a pillar[5]arene core. Self-assembled nanotubes were formed in water due to van der Waals interactions of the hydrophobic and hydrophilic parts of the glycoconjugate and intermolecular hydrogen bonds after one week. TEM and SEM measurements as well as fluorescence microscopy confirmed the structure. Furthermore, TEM and fluorescence microscope measurements exhibited a high agglutination rate of the multivalent nanotubes with E. coli, which confirms the high affinity of these glycoconjugates to the pathogenic bacteria. The length of the self-assemblies here also played a key role in the formation of bacterial clusters and was a crucial factor in controlling agglutination. Because the self-assembled nanotubes were several micrometers long, they could interconnect more bacteria, which resulted in an enhanced agglutination ability.

Other scaffolds of multivalent glycoconjugates which can inhibit E. coli were also established. Gouin and co-workers designed heptavalent glycoconjugates where N-heptyl α-D-mannoside was covalently linked to β-cyclodextrin core via a “click” procedure. 2 μg of the multivalent FimH antagonist instilled in the mouse bladder significantly reduced the urinary tract infection. Furthermore, recent advances in functional supramolecular/polymeric structures like bigger nanogels, graphene sheets, micrometer sized fibres, etc. could be promising scaffolds for effective inhibition of large bacteria particles.

4.3. Inhibition of the lipopolysaccharide synthesis

Lipopolysaccharides (LPS) are thermostable compounds that protect and stabilize the bacterial membrane can be divided into three parts. The core region of LPS consists of oligosaccharides with a 3-deoxy-D-manno-oct-2-ulosonic acid (Kdo) and two L-glycero-D-manno-heptose (heptose) units in the inner core and hexoses in the outer core. E. coli with only a few heptose units are significantly more permeable to certain hydrophobic antibiotics. The inhibition of the heptose biosynthetic pathway is just one approach which can be used to circumvent the conventional methods of targeting the bacterial receptors.

Vincent and coworkers developed some multivalent glycosylated fullerenes that can inhibit the LPS heptosyltransferase.

![Fig. 12 Preparation of three multivalent glycocluster via copper-catalyzed 1,3 dipolar cycloaddition with a series of fullerene hexa-adducts to inhibit the LPS heptosyl-transferase WaaC.](image-url)
WaaC. This glycotransferase catalyzes the incorporation of the first L-heptose into LPS. A series of fullerene hexa-adducts with 12 copies of peripheral sugar units displaying the mannosyl-glycine core structure of bacterial L,L-heptoses was synthesized via a copper-catalyzed 1,3 dipolar cycloaddition (Fig. 12). The inhibition of WaaC was determined by an assay which measures the IC50 values of ADP-2-heptose. The multivalent fullerene glycoconjugate of 1, 2, and 3 showed an inhibition potential in a low micromolar range (IC50 = 11; 47 and 6.7 μM). The monomeric glycosides, on the other hand, displayed IC50 values above 400 μM. Another effect of the “fullerene sugar balls” is that the inhibition data evaluated on a “per-sugar” basis showed an increase of the multivalent single glycosides in comparison to the monovalent ones.148

5. Anti-inflammatory drugs

The recruitment of circulating leukocytes is crucial at the sites of inflammatory diseases including hypersensitivity reaction as well as autoimmune diseases. The leukocyte recruitment from the bloodstream occurs through a multistep process in which the leukocytes captured on the endothelium, roll along, and interact with the endothelium in postcapillary venules. The transmigration of the leukocytes through the vessel wall and further migration into the extracellular tissue takes place subsequently. Leukocyte migration is initiated by interactions between E-, L-, and P-selectins (cell adhesion molecules) and their associated carbohydrate ligands. L-selectin is expressed by the most leukocytes. E-selectin and P-selectin are expressed by inflamed endothelia cells. However, P-selectin is also expressed by activated platelets. One approach for an anti-inflammatory therapy is to target the leukocyte trafficking. Therefore multivalent glycoconjugates systems have been developed that mimic the naturally occurring SLex6 ligands and inhibit the selectin-moieties.149–151

Patton and coworkers developed a series of multivalent polysyne conjugates to inhibit the interactions of E-selectin. The polysyne was conjugated with tetracarboxylic acid Lewis X and showed a highly active E-selectin inhibition and reduced the neutrophil rolling on activated endothelium cells with IC50 values in the nanomolar range. Furthermore, the increased size of the multivalent molecule as well as the ligand loading (35% and 50%) at the surface yielded the best inhibition data and a 700-fold increase in inhibition compared to the monovalent compound.152 Haag and coworkers synthesized several multivalent glyco-conjugates to inhibit L- and P-selectin binding.153 These functionalized galactose moieties were then coupled to a dendritic polyglycerol (dPG) scaffold via a 1,3 dipolar cycloaddition. The inhibition rate of the multivalent compound was compared to a tetravalent one based on a pentaerythritol core. The multivalent glycoconjugates showed a significant increase in the inhibition of L- and P-selectin binding compared to the tetravalent ones. The multivalent compound with the sulfated sugar moieties exhibited the highest inhibition rate.153 These studies were expanded and other carbohydrates were coupled to the PE and dPG cores. The multivalent dPG-conjugates with sulfated sugar moieties showed the best inhibition results with IC50 values in the high picomolar to the low nanomolar range and confirmed the previous results.154 Other scaffolds of multivalent glycoconjugates which can inhibit L-selectin binding were also established. A series of multivalent polyacrylamide-based conjugates155 as well as some neoglycopolymers156 confirms the multivalent effect. Furthermore, Haag and coworkers demonstrated that single dendritic polyglycerol sulfates (lacking the sugar unit) are also excellent candidates to inhibit L-, P-selectin binding and damp inflammation in vivo.151

6. Conclusion

In this article we have reviewed the current applications of multivalent glycoconjugates for a diverse set of diseases and discussed the challenges involved and endeavors made to address them. The development of multivalent glycoconjugate based vaccines has been fruitful, and a number of carbohydrate-based vaccines for various bacterial infections are already commercially available or under clinical trial. Vaccines against Shigella dysenteriae, Streptococcus pneumoniae, Neisseria meningitides, and Haemophilus influenzae are commercial examples. However, significant issues remain to be addressed. Development of vaccines against parasitic infections is still a big challenge due to the lack of understanding of complex immune evasion mechanisms used by different parasites. The key difficulty in developing HIV vaccines is mimicking the complex, dense cluster of glycans that are capable of eliciting neutralizing antibodies, specifically against HIV. On the other hand, researchers have been successful in identifying tumor associated carbohydrate antigens (TACAs) and then using them in multivalent forms, along with immunogenic carrier proteins, like keyhole limpet haemocyanin (KLH) to elicit strong T-cell dependent antibodies. Several synthetic antigen candidates have reached clinical trials, although more precise criteria for their efficacy and clinical impact remain to be defined. Besides using carbohydrates as vaccines, several multivalent glycoconjugates with controlled topology and valency have been explored to better understand multivalency, and their potential as anti-infectious agents for several viral and bacterial infections has been demonstrated in multiple examples. However, very few approaches consider biocompatible scaffolds in the appropriate dimension of the biological pathogen for the most effective shielding. Developing compatible polyvalent glycoconjugates with strong binding and infection inhibition efficiencies to reach the market as drugs still remains a challenging goal.

Abbreviation

| Abbreviation | Definition |
|--------------|------------|
| ADP-2-heptose | Adenosine diphosphate-2-heptose |
| Alum | Potassium aluminium sulfate |
| AuNPs | Gold nanoparticles |
| BSA | Bovine serum albumin |
| CRD | Carbohydrate recognition domain |
| CD4+ | Cluster of differentiation 4+ |
| CT | Cholera toxin |
| DLS | Dynamic light scattering |
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