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## Comparative Immunological Studies of Tumor-Associated Lewis X, Lewis Y, and KH-1 Antigens

Jiatong Guo<sup>†</sup>, Wenjie Jiang<sup>†</sup>, Qingjiang Li, Mohit Jaiswal, Zhongwu Guo<sup>\*</sup>

Department of Chemistry, University of Florida, 214 Leigh Hall, Gainesville, Florida 32611, United States

### Abstract

Tumor-associated carbohydrate antigens Lewis X (Le<sup>x</sup>), Lewis Y (Le<sup>y</sup>), and KH-1 are useful targets for cancer immunotherapy. In this regard, an insight into the structure-immunogenicity relationships of these antigens is important but this has not been systematically investigated yet. In the current study, Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 antigens with a lactose unit at the reducing end as a spacer were synthesized and coupled with keyhole limpet hemocyanin (KLH) protein. Immunological evaluations of the resultant conjugates revealed that they all could elicit robust immune responses whilst the Le<sup>y</sup> conjugate could provoke the highest titers of total and IgG antibodies. The binding assays of their antisera to each antigen and to cancer cells showed that each antiserum had extensive cross-reaction with all three antigens as protein conjugates and strong but somewhat antigen-selective binding towards MCF-7 cancer cell. Moreover, none of these antisera had obvious binding to SKMEL-28 cancer cell that does not express Le<sup>x</sup>, Le<sup>y</sup> and KH-1. The results of assays of these antisera to mediate complement-dependent cytotoxicity (CDC) to MCF-7 and SKMEL-28 cancer cells were very similar to the results of binding assays. Thus, it was concluded that all three antigens could form effective conjugate vaccines whereas the Le<sup>y</sup> conjugate induced the most robust immune responses and the antiserum of Le<sup>x</sup> had the highest binding and cytotoxicity to target cancer cells. In addition, as the antibodies induced by each antigen had extensive cross-reaction with other two antigens, either Le<sup>x</sup> or Le<sup>y</sup> or the two combined can be

\*Corresponding author: phone: (+1) 352 392 9133; zguo@chem.ufl.edu.

<sup>†</sup>These authors contributed equally to the paper.

#### Author Contributions

JG, WJ and MJ contributed to biological and animal studies; QL contributed to the synthesis; ZG was responsible for the general design and management of the project; all of the authors were involved in the manuscript preparation.

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#### Approval of animal use

The animal use protocol for this research (#201609560) was approved by the Institutional Animal Care and Use Committee of University of Florida.

#### Consent for publication

All of the authors have read agreed to publish this article and declare that it is original, has never been published before, and has not been submitted to other journals.

#### Competing interests

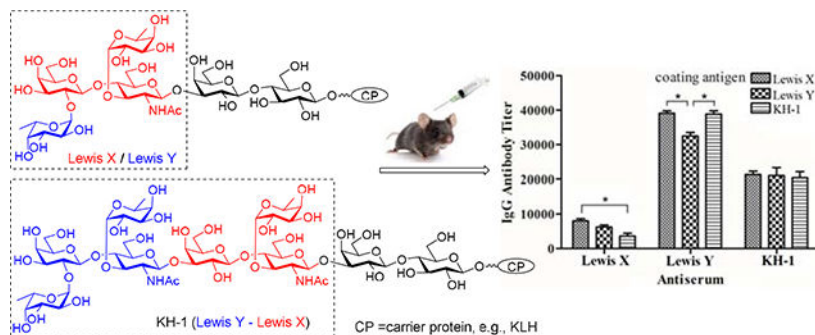
All of the authors disclose no competing interests.

#### Additional information

An additional file containing original ELISA, FACS and antibody-mediated CDC data and the MALDI-TOF mass spectra of glycoconjugates **2a**, **2b** and **2c** is available via the Internet at the journal's website.

utilized to formulate effective conjugate vaccines for cancer immunotherapy. Another paradigm-shifting discovery of this study is that the presentation of Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 antigens on cancer cell can be different from that in synthetic conjugates, which should be taken into consideration during the design and optimization of related cancer vaccines or immunotherapies.

## Graphical Abstract



## Keywords

carbohydrate; cancer antigen; Lewis X; Lewis Y; KH-1; glycoconjugate; cancer vaccine

## Introduction

Cancer cells display unusual glycosylation patterns in the oncogenic process; the abnormal glycans thus generated are called tumor-associated carbohydrate antigens (TACAs) [1–3]. TACAs are promising targets for the development of therapeutic cancer vaccines or cancer immunotherapies [4–7] that aim at activating the patients' immune system to cure cancer and are regarded as the ideal therapies due to their potentially high therapeutic efficacy and specificity [8,9]. Despite the great effort and progress in TACA-based cancer vaccine development [10–14], researches to gain a deeper understanding of the structure-immunogenicity relationships of TACAs and their immunorecognition are still difficult and largely deficient, whilst related information is especially important for the design of effective cancer immunotherapies. Systematic study and comparison of the immunological properties of TACAs and their analogs are arguably the most direct method to examine the structure-immunogenicity relationships of TACAs. For this purpose, it is necessary to have access to appropriate TACA derivatives and analogs to conduct comparative studies.

Recently, our group has developed some convergent strategies for efficient synthesis of Lewis type of TACAs, including Lewis X (Le<sup>x</sup>), Lewis Y (Le<sup>y</sup>) and KH-1 [15,16]. These structurally related antigens are expressed by several tumors [17,18] and thus have been extensively explored both as synthetic targets and as antigenic epitopes for cancer vaccine development [19–24]. Structurally, Le<sup>y</sup> is different from Le<sup>x</sup> only in that the former has an additional L-fucose (Fuc)  $\alpha$ -linked to the 3-*O*-position of the galactose (Gal) residue at the non-reducing end of the latter antigen, whereas KH-1 can be regarded as the heterodimer of Le<sup>x</sup> and Le<sup>y</sup> (Figure 1). Easy access to these antigens by the new synthetic strategies had provided us the opportunity to conduct unprecedented comparative study on their

immunological properties, including the relationships between the structure of Lewis antigens and their immunogenicity, the significance of a specific structural epitope in these oligosaccharide antigens for immune recognitions, etc. Consequently, in this work, the native form of Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 antigens that contained the common and natural lactose spacer, i.e., having these antigens linked to the 3'-*O*- position of lactose, were synthesized and conjugated with keyhole limpet hemocyanin (KLH) protein. The inclusion of the lactose spacer was intended to more closely mimic the natural form of these TACAs than in literature and, in the meantime, buffer any potential influence of direct conjugation of these antigens with carrier proteins on their structure and conformation. Furthermore, lactose is one of the most common disaccharide motifs in nature, thus we would not expect it to cause immune reactions, which was eventually proved by this study. The resultant conjugates **1a**, **1b**, and **1c** were immunologically evaluated in mouse, employing the corresponding human serum albumin (HSA) conjugates **2a**, **2b**, and **2c**, respectively, as coating antigens for the detection of TACA-specific antibodies in the mouse antisera by enzyme-linked immunosorbent assay (ELISA).

## Results

Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 derivatives **3**, **4**, and **5** (Scheme 1) were synthesized according to reported methods [15,16] and were coupled with KLH and HSA via the bifunctional glutaryl group. This linker was selected as it was proved efficient for carbohydrate-protein conjugations and exhibit no obvious immunogenicity or other adverse immunological properties by itself [25]. First, the linker was attached to glycans **3–5** by a reaction with the dually activated disuccinimidyl glutarate **6**. The resultant activated esters **7–9** were then reacted with KLH and HSA in phosphate-buffered saline (PBS) to give conjugates **1a–c** and **2a–c**, which were purified by gel filtration chromatography on a Biogel A 0.5 column. The carbohydrate loadings of these conjugates were analyzed [26] and were shown to be in the desired range (Table 1).

Immunological evaluations of the KLH conjugates **1a**, **1b**, and **1c** were performed in female C57BL/6J mice (6–8 weeks of age). The immunization protocol was the same as reported [15], i.e., to subcutaneously (s.c.) inject each conjugate as an emulsion with complete Freund's adjuvant (CFA) to a group of 5 mice on day 1 for initial immunization and then intraperitoneally (i.p.) inject the same conjugate as an emulsion with incomplete Freund's adjuvant (IFA) on day 8, 15, and 36 for boost immunizations. Blood samples were collected from mice on day 0 before initial immunization (as blank controls) and on day 21 and 43 after boost immunizations. The blood samples were clotted to obtain antisera by standard protocols, which were subjected to antibody titer analysis by ELISA. For ELISA, conjugates **2a–c** were utilized as capture antigens and 1:1,000 PBS-diluted alkaline phosphatase (AP)-linked goat anti-mouse kappa, IgG, and IgM antibodies were employed as secondary antibodies to detect total, IgG, and IgM antibodies, respectively. Antibody titers were defined as the dilution numbers of sera at which an optical density (OD) value of 0.2 was achieved at 405 nm wavelength, and each experiment was repeated three times.

ELISA results of the total antibodies (Figure 2A) indicated that after the second boost immunization (day 21 antisera) all three conjugates **1a**, **1b**, and **1c** had already elicited the

production of antigen-specific antibodies, whilst the total antibody titers of day 43 antisera obtained after the third boost immunization were significantly higher than that of day 21 sera. The results proved not only the efficiency of conjugates **1a**, **1b**, and **1c** to elicit robust immune responses but also the progress and strengthening of the induced immune responses as a result of boost immunizations. Furthermore, the total antibody titers of the antisera of conjugates **1a**, **1b**, and **1c** were significantly different from each other. The antiserum of conjugate **1b** exhibited the highest antibody titer; in turn, the antibody titer of antiserum **1a** was significantly higher than that of antiserum **1c**. Therefore, it seemed that the order of overall immunogenicity of the conjugates or related TACAs was **1b** ( $Le^y$ ) > **1a** ( $Le^x$ ) > **1c** (KH-1).

We also analyzed the isotypes of antibodies induced by conjugates **1a**, **1b**, and **1c**. The results (Figures 2B and 2C) revealed that the provoked antibody responses in both day 21 and 43 antisera were mainly of IgG type for conjugate **1b**, which was consistent with our previous observations that glycoproteins induce usually IgG antibodies against carbohydrate antigen [25,27,28]. On the other hand, the antibody responses were mainly of IgM type for conjugate **1a** and both IgG and IgM types for conjugate **1c**. It was interesting to find that although  $Le^x$ ,  $Le^y$  and KH-1 are structurally related, they had different immunogenicity and different immunological properties as IgG and IgM antibodies are produced through different pathways and mechanisms. For example, IgM antibodies are produced by plasma cells during the initial response to a specific antigen [29,30], whilst IgG antibodies are produced later to participate in primarily secondary immune responses. Usually, IgG antibody production indicates T cell-dependent immunity, antibody class switch and affinity maturation, and long-term immune memory [31], which are the properties desirable for cancer therapy. Therefore, compared to  $Le^x$  and KH-1 conjugates **1a** and **1c**,  $Le^y$  conjugate **1b** seemed to elicit more promising therapeutic immune responses.

Encouraged by the above discoveries, we investigated next the cross-reactivity of each antiserum with the other two antigens. For this purpose, we coated ELISA plates with HSA- $Le^x$ ,  $Le^y$  and KH-1 conjugates **2a**, **2b** and **2c**, respectively, and used them to test cross-reactive total and IgG antibodies in each pooled antiserum by ELISA as described above. ELISA results of total antibodies shown in Figure 3A suggested that the antiserum of  $Le^y$  (conjugate **1b**), which gave the highest total antibody titer as compared to other two antisera (Figure 2A), also showed the highest reactivity with  $Le^x$  (**2a**) and KH-1 (**2c**) antigens (Figure 3A). This was consistent with the conclusion that **1b** was more immunogenic than **1a** and **1c** to elicit the production of a higher level of antibodies. Most significantly, the antiserum of **1b** had similar reactivity with all three antigens, indicating that the antibodies elicited by **1b** could recognize  $Le^x$ ,  $Le^y$ , and KH-1 without significant discrimination. In addition, each of the antisera of conjugates **1a** and **1c** also exhibited essentially the same reactivity with all three antigens (Figure 3A).

Similarly, the antiserum of **1b** exhibited the highest titers of IgG antibodies for all three antigens (Figure 3B), which was also consistent with the above results that conjugate **1b** induced the strongest IgG antibody response (Figure 2B). Furthermore, each of the antisera of conjugates **1a**, **1b**, and **1c** showed comparable IgG antibody titers for all three different

antigens. The slightly higher reactivity of antiserum **1b** with **2a** and **2c** as compared to the reactivity with **2b** may be attributed to the higher carbohydrate loadings of **2a** and **2c** (10.3 and 7.5 vs 6.0 glycans/conjugate). The slightly decreased reactivity of the IgG antibodies in antiserum **1a** with conjugate **2c** (Figure 3B) may indicate that some of the IgG antibodies induced by Le<sup>x</sup> could not recognize KH-1. The antiserum of **1a** showed the similar trend of decreased reactivity, but less significantly, with conjugate **2b**.

The linker used in conjugates **1a**, **1b**, and **1c** was proved in other glycoconjugates to exhibit no obvious immunogenicity or other adverse immunological properties by itself [25]. Here, the reactivity of antisera **1a**, **1b**, and **1c** with the linker and the common lactose motif was assessed to exclude their contribution to the observed cross-reaction. In these experiments, mannose- and lactose-HSA conjugates containing the same linker as in conjugates **1a**, **1b**, and **1c** were used as the capturing antigen for ELISA. Both conjugates were proved to exhibit similarly low reactivity with all three antisera (Tables S5 and S6, Figures S1 and S2 of SI), suggesting that the observed different antibody titers shown in Figures 2 and 3 were indeed due to different carbohydrate antigen-antibody binding or cross-reaction.

Subsequently, we analyzed whether the carbohydrate hapten-specific antibodies induced by conjugates **1a**, **1b**, and **1c** could recognize and bind to Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 antigens present on the cancer cell surface by flow cytometry (FACS). In this study, MCF-7 cell, a human breast cancer cell line that was confirmed to express Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigens [17], was used as the positive control, whereas SKMEL-28 cell, a human melanoma cell line that was shown to not express Le<sup>x</sup>, Le<sup>y</sup> or KH-1 [20,32], was utilized as the negative control. Both cell lines were incubated with the normal mouse serum or a day 43 antiserum and then with FITC-linked goat anti-mouse kappa antibody. The cells were finally subjected to FACS study. As shown in Figure 4A, the fluorescent intensities of MCF-7 cells treated with the antisera of conjugates **1a**, **1b**, and **1c** were all much higher than that of cells treated with the normal mouse serum, and the increases in median fluorescence intensity (MFI) value were statistically significant ( $P < 0.05$ ). On the other hand, treating SKMEL-28 cells with antisera **1a**, **1b**, and **1c** caused little changes (statistically insignificant) in the fluorescent intensity when compared to cells treated with normal mouse serum (Figure 4B). The FACS results indicated that antibodies provoked by conjugates **1a**, **1b**, and **1c** could specifically bind to MCF-7 cell which expresses Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigens. Furthermore, the antisera of **1a** and **1b** exhibited similar binding activity to MCF-7 cell, and the activity was higher than that of the antiserum of **1c**.

Interestingly, the results of antibody binding to MCF-7 cells (Figure 4A) did not agree completely with the results of antibody titers (Figure 2A) and antibody-antigen cross-reactivity (Figure 3A) of the antisera. For example, the total antibody titer of antiserum **1b** was significantly higher than that of antisera **1a** and **1c**, but antiserum **1b** showed similar or even slightly lower binding to MCF-7 cell than antiserum **1a**. This result was also contradictory to the observation that antibodies in the antiserum of **1b** had extensive cross-reactivity with all synthetic antigens or conjugates (Figure 3A). Consequently, we propose that (1) at least a part of the antibodies in the antiserum of **1b** were specific to the Le<sup>y</sup> antigen on MCF-7 cells, i.e., they did not recognize and bind to Le<sup>x</sup> antigens on cancer cell, and (2) MCF-7 cell expresses a higher level of Le<sup>x</sup> antigen than Le<sup>y</sup> antigen and/or

antibodies in the antiserum of **1a** had more extensive cross-reaction with Le<sup>y</sup> and KH-1 antigens on the MCF-7 cell surface to result in elevated binding for the antiserum of **1a**. Similarly, the difference in binding of the antisera of **1b** and **1c** to MCF-7 cell was obviously less significant than the difference in their total antibody titers, which also suggested a higher expression level of KH-1 than Le<sup>y</sup> and/or more cross-reactivity of the antibodies in the antiserum of **1c** with Le<sup>x</sup> and Le<sup>y</sup> antigens on MCF-7 cell. Currently, there is no reported qualitative analysis of the expression levels of Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 antigens on MCF-7 cell, thus we cannot make conclusions about the cross-reactivity of antibodies raised by conjugates **1a** and **1c** with Le<sup>y</sup> on the MCF-7 cell surface. However, the results may suggest that at least some antibodies raised by conjugate **1b** could not recognize and bind to Le<sup>x</sup> and KH-1 antigens on the cancer cell surface although antibodies against **1b** showed extensive cross-reactivity with the protein conjugates of synthetic Le<sup>x</sup> and KH-1 in ELISA. Previously, Kitamura *et al.* also observed that the antibodies raised with synthetic Le<sup>y</sup> conjugate could not effectively recognize natural Le<sup>y</sup> antigens expressed on cancer cells [33].

To investigate the correlation between the cell-binding abilities of antibodies raised by conjugate **1a**, **1b** or **1c** and their potential anticancer activities, we evaluated further the activities of antisera **1a**, **1b**, and **1c** to mediate complement-dependent cytotoxicity (CDC) against MCF-7 and SKMEL-28 cells. Therefore, after the cells were treated with normal mouse serum (negative control) or the antiserum of **1a**, **1b** or **1c**, as described above, they were incubated with rabbit complements and then subjected to cell lysis analysis by the lactate dehydrogenase (LDH) test. The positive control of 100% cell lysis was obtained by treating cells with 1% triton X-100. As indicated in Figure 5, all three antisera could mediate significant lysis of MCF-7 cell that expresses Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 antigens, and the results correlated well with the antibody-cell binding results described above (Figure 4A). In contrast, under the same conditions none of the three antisera induced significant lysis of SKMEL-28 cell that does not express Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigens and showed no binding with the antisera (Figure 4B).

## Discussion

TACAs expressed on cancer cells are useful targets for the development of new cancer therapies, such as therapeutic cancer vaccines or cancer immunotherapies. In this regard, it is critical to appreciate the relationship between the structure of TACAs and their immunogenicity, the correlation between synthetic TACA conjugates and natural ones on cancer cells, the interaction between TACAs and their antibodies, etc. To gain a better understanding of Lewis type of TACAs, Le<sup>x</sup>, Le<sup>y</sup> and KH-1, we prepared their KLH conjugates and studied their immunological properties, the interaction of their antisera with synthetic and natural antigens and the efficacy of their antisera to mediate CDC to cancer cells.

Le<sup>x</sup> antigen is remarkably upregulated in colon cancer and several other tumors [34] and was shown to be associated with cancer progression. Le<sup>y</sup> antigen is an oncodevelopment-associated TACA, which has been detected in almost all colonic carcinoma tissues. Notably, Le<sup>y</sup> is a specific marker for malignancy [35]. KH-1 antigen is relatively cancer-specific, which has not been observed in normal tissues yet [36]. Structurally, these three antigens are

related. Le<sup>y</sup> and Le<sup>x</sup> are different in that the former has an additional fucosyl residue at the glycan non-reducing end, whereas KH-1 is a heterodimer of Le<sup>y</sup> and Le<sup>x</sup> (Figure 1). Consequently, a systematic and comparative study on these antigens may provide interesting insights into their immunology.

Our immunological evaluation of the KLH-Le<sup>x</sup>, Le<sup>y</sup> and KH-1 conjugates in mice revealed that they all provoked robust immune responses but had indeed different immunogenicity. The Le<sup>y</sup> conjugate induced significantly higher total antibody titers than the Le<sup>x</sup> and KH-1 conjugates, and the Le<sup>x</sup> conjugate elicited stronger immune responses than the KH-1 conjugate. Furthermore, it was shown that Le<sup>y</sup> induced mainly IgG antibody responses while Le<sup>x</sup> induced mainly IgM antibody responses. These results indicated that the structure of TACAs had a major impact on their immunogenicity and other immunological properties, although they are structurally similar. On the other hand, the Boons group found that the protein conjugate of a Le<sup>y</sup>/Le<sup>x</sup> heterodimer elicited a higher antibody titer than the Le<sup>y</sup> conjugate [37]. A major difference between Boons conjugate and our KH-1 conjugate was that the former had Le<sup>y</sup> and Le<sup>x</sup> antigens stitched together by an artificial linker whilst in our conjugate, Le<sup>y</sup> and Le<sup>x</sup> antigens were directly coupled together without any spacer. Therefore, it seems that in addition to the principal structure of haptens, other factors such as linkage form, conjugation method, immunization protocol, etc. may also have a significant impact on the immunogenicity of resultant glycoconjugate vaccines [38].

Theoretically, a complex carbohydrate antigen can induce a diversity of antibodies against not only its intact structure but also its partial sequences. Therefore, the induced antibodies may have cross-reactions with other closely related antigens [39]. Our results about the cross-reactivity between different antisera and antigens have revealed that antibodies in each antiserum could recognize all three antigens without significant difference, thus the cross-reactivity was strong and extensive. Furthermore, for each antiserum, the titers of cross-reactive total and IgG antibodies in Figure 3 corroborated the ELISA results of antigen-specific antibodies in Figure 2. The Boons group also reported that antibodies elicited by a Le<sup>y</sup>/Le<sup>x</sup> dimer could recognize both Le<sup>y</sup> and Le<sup>x</sup> although their titers were significantly lower than that of the binding to the Le<sup>y</sup>/Le<sup>x</sup> dimer [37]. Since all of the Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigens contain the fucosyl residue, we propose that the observed cross-reactivity may be at least partially related to this epitope, which is worthy further investigation in the future. On the other hand, because over-fucosylation is common among different tumors and is closely related to carcinogenesis [40,41], immunity centered on the fucose epitope may be of general significance for cancer immunotherapy.

Interestingly, the results of antibodies in the antisera to bind with Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigens expressed on cancer cells were significantly different from the ELISA results of antibody-antigen binding or cross-reactivity obtained with synthetic conjugates. In this study, first, we proved that the induced antibodies were specific to Lewis type of TACAs as Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigen negative cells did not have obvious binding with the antisera. Next, we disclosed that the antiserum of Le<sup>x</sup>, which contained a significantly lower concentration of antibodies than the antiserum of Le<sup>y</sup> (Figure 2A), showed similar or even slightly higher binding to Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigen-expressing MCF-7 cell (Figure 4A). These results suggested that at least some of the antibodies induced by the KLH-Le<sup>y</sup>

conjugate were specific to Le<sup>y</sup> antigen and did not recognize or bind to Le<sup>x</sup> and KH-1 antigens on the cancer cell surface; otherwise, we would have observed the highest binding of anti-Le<sup>y</sup> serum to MCF-7 cell. These results were further corroborated by antibody-mediated CDC study (Figure 5). The CDC study also confirmed the feasibility of Lewis antigen-based cancer immunotherapy.

There are two potential explanations for the increased binding of anti-Le<sup>x</sup> serum to cancer cells, which are: (1) a higher expression level of Le<sup>x</sup> antigen than Le<sup>y</sup> antigen and (2) stronger antibody-antigen binding and/or more cross-reaction of the antibodies in the anti-Le<sup>x</sup> serum with Le<sup>y</sup> and KH-1 antigens on cancer cells. If the differences between ELISA and antibody-cell binding assays (Figures 3A and 4A) were not simply because of the different expression level of antigens, which is likely to be the case, our results may indicate the potential discrepancies of these antisera to bind and cross-react with synthetic and natural Le<sup>y</sup> antigens. This should be perceivable because TACAs may have different interactions with molecules in artificial environment on ELISA plate and in the cancer cell matrix to affect the structure and organization of TACAs, namely that the antigens may have adopted different conformations on ELISA plates and cells. This conclusion was in agreement with the seminal discovery that antibodies raised by a specific TACA could distinguish the same antigen in different expression levels on cancer and normal cells [41–43], which has been the molecular foundation for the specificity and the therapeutic efficacy of TACA-based cancer vaccine or immunotherapy. In general, carbohydrates are flexible molecules that can adjust their structure to accommodate the environment [44,45]. Therefore, it should be easily perceivable that TACAs may adopt different forms of organization and presentation under different conditions, which can thereby affect the molecular recognition process [46].

## Conclusion

In conclusion, the KLH conjugates of Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 antigens were synthesized, immunologically evaluated, and compared. Whereas all of the conjugates could induce robust immune responses, they had indeed different immunogenicity and the KLH-Le<sup>y</sup> conjugate elicited the highest titers of IgG antibodies. Therefore, Le<sup>y</sup> seemed to be a more promising antigen for the development of cancer vaccines. Intriguingly, whereas the antiserum of each antigen had extensive cross-reactivity with the remaining two antigens in the form of synthetic HSA conjugates, these antisera exhibited some selectivity toward natural antigens on cancer cells. The antiserum-cell binding and the antibody-mediated CDC studies further indicated that MCF-7 cell may express different levels of Le<sup>x</sup>, Le<sup>y</sup> and KH-1 antigens, which is a research topic worthy further investigation. Overall, our findings combined with literature reports suggested that for the design of new TACA-based therapeutic cancer vaccine, in addition to the target antigen, we should be also careful about the selection of carrier proteins, conjugation methods, and so on, as these factors may affect TACA presentation and recognition and thereby the therapeutic efficacy of resultant vaccines.



## Experimental Section

### Synthesis of TACA-protein conjugates 1a-c and 2a-c (Scheme 1).

A mixture of oligosaccharide **3**, **4** or **5** (6.0 mg) and disuccinimidyl glutarate **6** (15 equiv) in DMF and PBS buffer (0.1x) (4:1, 0.5 mL) was stirred at rt for 4 h. The solution was condensed in vacuum, and then EtOAc (9 volumes) was added. The precipitates were isolated and washed with EtOAc 10 times and dried in vacuum to result in crude active ester **7**, **8** or **9**, respectively, which was directly used for the next step. Each product was mixed with HSA or KLH at a ratio of 1:1 (W/W) in 0.1 M PBS buffer (0.4 mL). The solution was stirred at rt for 3 days and then applied to a Superdex® 200 column, which was eluted with 0.1 M PBS buffer ( $I = 0.1$ , pH = 7.8). The glycoprotein-containing fractions, which were characterized by the bicinchoninic acid (BCA) assay for protein and charring with 15% (v/v) H<sub>2</sub>SO<sub>4</sub> in EtOH for carbohydrates, were combined, dialyzed against distilled water for 2 days, and finally lyophilized to afford the desirable glycoconjugate **1a**, **1b**, **1c**, **2a**, **2b** or **2c** as a white solid, respectively.

### Analysis of the carbohydrate loading of glycoconjugates 1a–c and 2a–c.

Chemical analysis to determine the carbohydrate loadings of glycoconjugates was performed according to a literature method [26]. First, a calibration curve was prepared for sugars utilizing a mixture of fucose, galactose, *N*-acetylglucosamine, and glucose in molar ratios equal to that in each antigen. Aliquots of the standards dissolved in distilled water (1 mg/mL) were transferred into 10 dry 10-ml tubes in 5  $\mu$ L increments ranging from 5 to 50  $\mu$ L. To the tubes were sequentially added 500  $\mu$ L of 4% phenol and 2.5 mL of 96% sulfuric acid. The glycosyl bonds were cleaved and a colored complex was developed in this step. Solutions were transferred from the test tubes to cuvettes and measured at the 490 nm wavelength. The calibration curve was obtained by plotting A<sub>490</sub> absorbance against the total weight ( $\mu$ g) of sugars in each standard sample. To analyze the carbohydrate loadings of **1a–c** and **2a–c**, three accurately weighed samples of each glycoconjugate (final concentration of the sugar should be in the range of the calibration curve) were dissolved in distilled water in 10-mL test tubes and then examined by the exactly same protocol, whereas free KLH and HSA proteins were used as the blank controls, respectively. The amount of sugars present in each sample was calculated based on its A<sub>490</sub> against the calibration curve, and the carbohydrate loading of each glycoconjugate was calculated according to the following equation:

$$\text{carbohydrate loading(\%)} = \frac{\text{carbohydrate content in the sample (mg)}}{\text{weight of the glycoconjugate sample (mg)}} \times 100\%$$

The carbohydrate loadings of HSA conjugates **2a–c** were confirmed by MALDI-TOF mass spectrometry (Supporting Information), which were calculated according to the following equation:

$$\text{carbohydrate loading(\%)} = \frac{\text{Molecule weight of HSA conjugate} - \text{Molecule weight of HSA}}{\text{Molecule weight of HSA conjugate}} \times 100\%$$

Apical points of the MS peaks were used for estimation of the average molecule weights of HSA and its conjugates **2a–c**.

### Immunization of animals.

Fifteen female C57BL/6J mice (6–8 weeks of age) were assigned into three groups randomly (five mice per group). On day 1, mice were subcutaneously injected with 0.1 mL of the CFA emulsion of a specific glycoconjugate containing 3 µg of the carbohydrate antigen. Thereafter, mice were immunologically boosted three times on day 8, 15 and 36, respectively. For boosting immunization, mice were intraperitoneally injected with 0.1 mL of the IFA emulsion of the same glycoconjugate at the same dose. Blood samples were collected from the mice via the saphenous vein on day 0 before the initial immunization and day 21 and 43 after immunization to prepare blank sera and antisera after clotting and centrifugation at 5000 rpm for 7 min at 4 °C. The animal use protocol for this research (#201609560) was approved by the Institutional Animal Care and Use Committee of University of Florida.

### ELISA.

ELISA was performed according to a previously report protocol with minor modifications [47]. ELISA plates were incubated with conjugate **2a**, **2b** or **2c** (2 µg/mL, 100 µL per well) in coating buffer (0.1 M bicarbonate, pH 9.6) at 4 °C overnight and then at 37 °C for 1 h. After washing with PBST (PBS containing 0.05% of Tween-20) 3 times, the plates were treated with blocking buffer (1% BSA in PBST) at rt for 1 h. Pooled antisera diluted from 1:300 to 1:72900 in PBS in serial half-log manner were added to the plates (100 µL/well), which was followed by incubation at 37 °C for 2 h. Then, AP-linked goat anti-mouse kappa, IgG and IgM antibodies (1:1,000 dilution in PBS) were added and the plates were incubated at rt for 1 h. After washing with PBST 3 times, 100 µL of a *p*-nitrophenylphosphate (PNPP) solution (1.67 mg/mL in PBS) were added and the plates were then incubated at rt for 30 min. The optical density (OD) values of the ELISA plates at 405 nm wavelength were read using a microplate reader (Cytation 1, Bio-Tek instruments Inc.). OD values were plotted against the serum dilution numbers to obtain a best-fit logarithm line and the equation of this line was used to calculate the antibody titer. The dilution number at which the OD value of 0.2 was achieved was taken as the antibody titer.

### FACS.

The binding of antibodies in the antisera to cancer cells was detected by FACS assay as reported previously [48,49]. MCF-7 and SKMEL-28 cells were cultured in DMEM containing 10% of fetal bovine serum (FBS) and 1% of antibiotics. Cells were harvested, washed twice with FACS buffer (PBS containing 5% of FBS) and counted ( $4.0 \times 10^5$  cells for each experiment). Thereafter, cells were incubated with 50 µL of 10 time-diluted (in FACS buffer) normal mouse serum or pooled day 43 antiserum at 4 °C for 30 min. Cells were washed twice and incubated with fluorescein isothiocyanate (FITC)-linked goat anti-mouse kappa antibody (50 µL, 1:25 dilution in FACS buffer) at 4 °C for 30 min. Then, cells were washed twice and re-suspended in 400 µL of FACS buffer for detection. The fluorescence intensity of cells were analyzed with Attune Nxt Acoustic Focusing Cytometer.

### Evaluation of antibody-mediated CDC.

Antibody-mediated CDC analysis was performed as reported previously [49,50]. MCF-7 and SKMEL-28 cells were cultured as described above and then plated in 96-well plates ( $1.0 \times 10^4$  cells/well for MCF-7 and  $1.5 \times 10^4$  cells/well for SKMEL-28) to allow for attachment overnight. After washing with DMEM without FBS, attached cells were treated with normal mouse serum or a day 43 pooled antiserum (100  $\mu$ L/well, 1:50 dilution in DMEM without FBS) at 37 °C for 2 h. Then, cells were washed and incubated with rabbit complement serum (100  $\mu$ L/well, 1:10 dilution in DMEM without FBS) at 37 °C for 1 h. The low control was the spontaneous LDH release from cells treated with rabbit complement serum only without mouse serum. For the high control, 1% triton X-100 (100  $\mu$ L/well, dilution in DMEM without FBS) was added instead of the rabbit complement serum. After incubation, 20  $\mu$ L of the supernatant from each well was relocated and mixed with 80  $\mu$ L of PBS in each well on another 96-well plate. Thereafter, the LDH activity in each well was detected with LDH Cytotoxicity Detection Kit (Pierce LDH Cytotoxicity Assay Kit, Thermo Scientific Co.) according to manufacturer's instructions. LDH cytotoxicity detection reagent (100  $\mu$ L/well) was added and incubated at rt in the dark for 1 h. The OD value of each well at 490 nm and 680 nm (instrument background signal) wavelengths was read with a microplate reader. The antibody-mediated CDC was indicated by the percentage of cell lysis, calculated according to the equation below:

$$\text{cell lysis}(\%) = \frac{\text{Experimental A} - \text{Low control A}}{\text{High control A} - \text{Low control A}} \times 100\%$$

where "experimental A" is the OD<sub>490</sub>-OD<sub>680</sub> value of cells treated with normal mouse serum or a pooled day 43 antiserum, "low control A" is the OD<sub>490</sub>-OD<sub>680</sub> value of cells treated without any serum, and "high control A" is the OD<sub>490</sub>-OD<sub>680</sub> value of cells completely lysed with 1% triton X-100.

### Statistical analysis.

All data were analyzed by one-way analysis of variance (more than 2 groups) or independent t test (between 2 groups) using SPSS software.  $P < 0.05$  was considered as statistically significant.

### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

### Funding

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### Abbreviations

AP	alkaline phosphatase
BCA	bicinchoninic acid

<b>CDC</b>	complement-dependent cytotoxicity
<b>CFA</b>	complete Freund's adjuvant
<b>DMEM</b>	Dulbecco's modified eagle medium
<b>DMF</b>	<i>N,N</i> -dimethyl formaldehyde
<b>ELISA</b>	enzyme-linked immunosorbent assay
<b>FACS</b>	flow cytometry
<b>FBS</b>	fetal bovine serum
<b>FITC</b>	fluorescein isothiocyanate
<b>Fuc</b>	fucose
<b>Gal</b>	galactose
<b>HSA</b>	human serum albumin
<b>IFA</b>	incomplete Freund's adjuvant
<b>i.p.</b>	intraperitoneal
<b>KLH</b>	keyhole limpet hemocyanin
<b>LDH</b>	lactate dehydrogenase
<b>Le<sup>x</sup></b>	Lewis X
<b>Le<sup>y</sup></b>	Lewis Y
<b>OD</b>	optical density
<b>PBS</b>	phosphate-buffered saline
<b>PBST</b>	PBS containing 0.05% of Tween-20
<b>PNPP</b>	<i>p</i> -nitrophenylphosphate
<b>s.c.</b>	subcutaneous
<b>SEM</b>	standard error of mean
<b>TACA</b>	Tumor-associated carbohydrate antigen

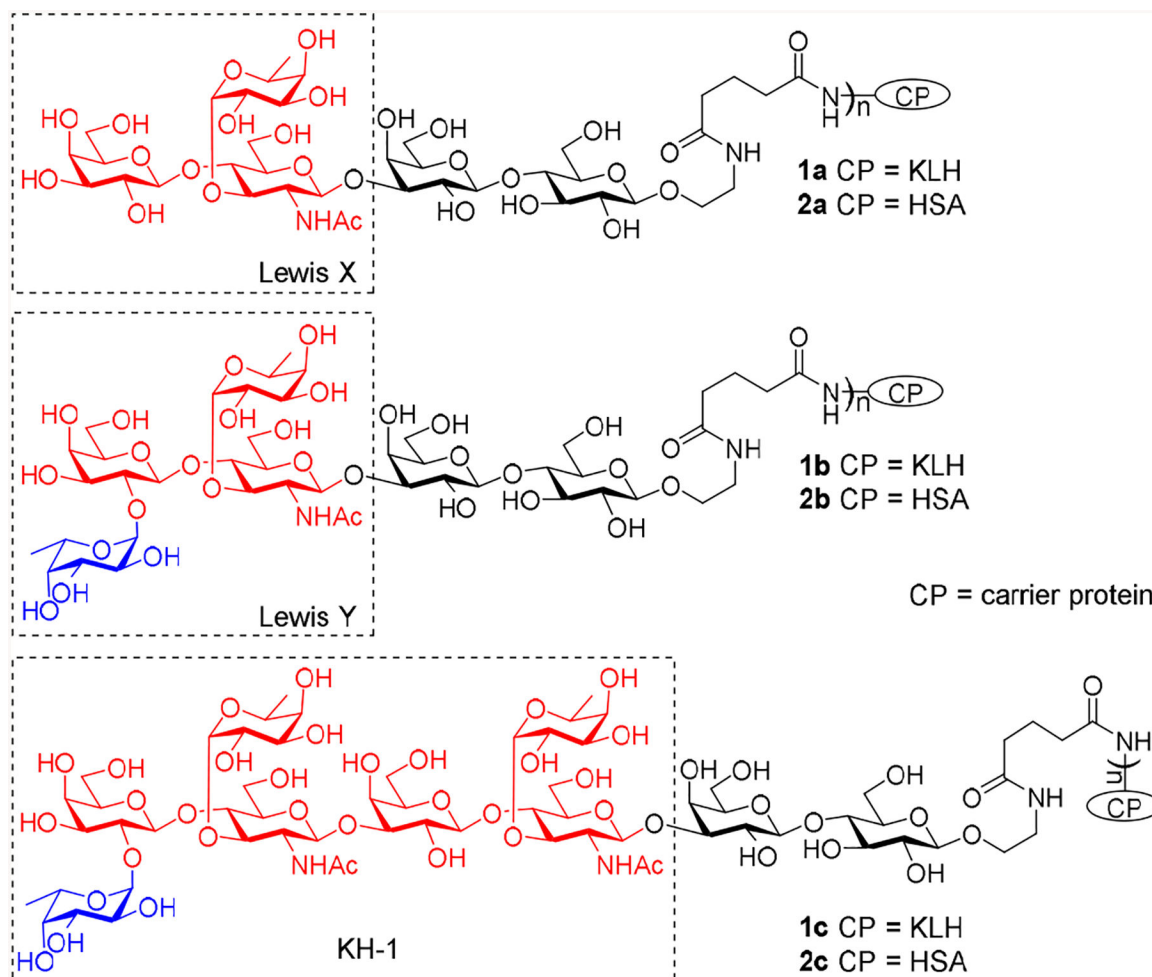
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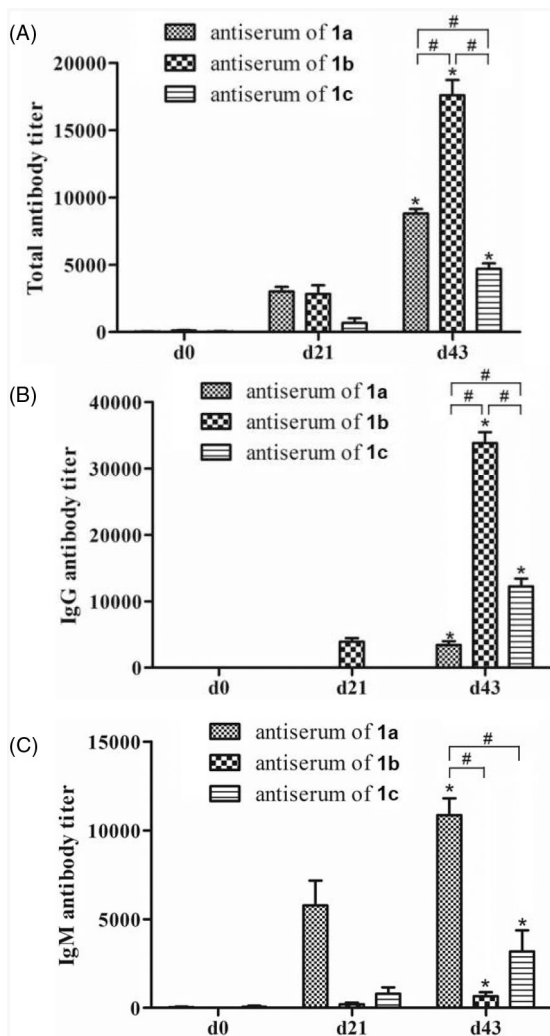
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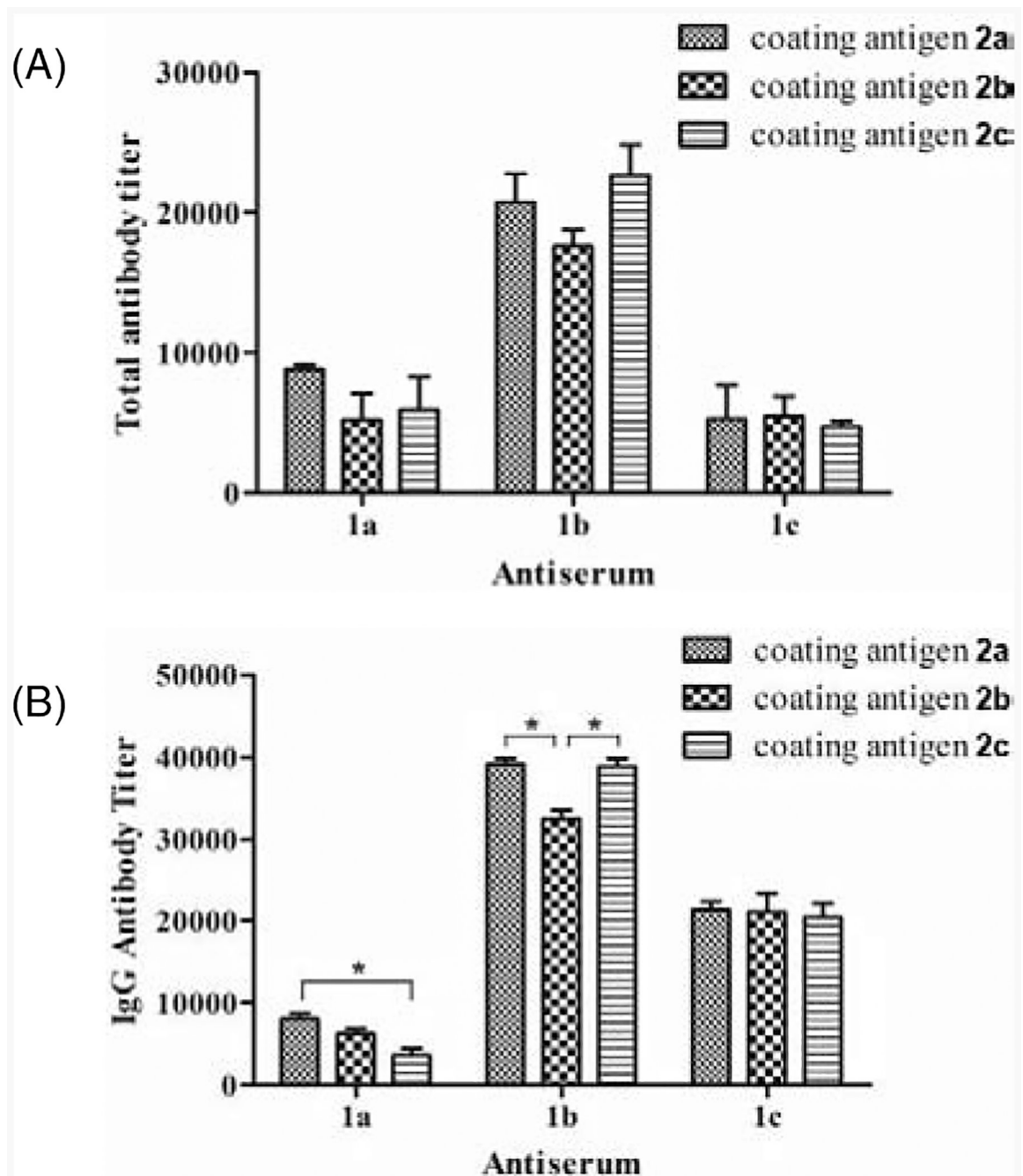
**Figure 1.** Structures of Le<sup>x</sup>, Le<sup>y</sup>, and KH-1 and their KLH and HSA conjugates **1a-c** and **2a-c**





**Figure 2.**

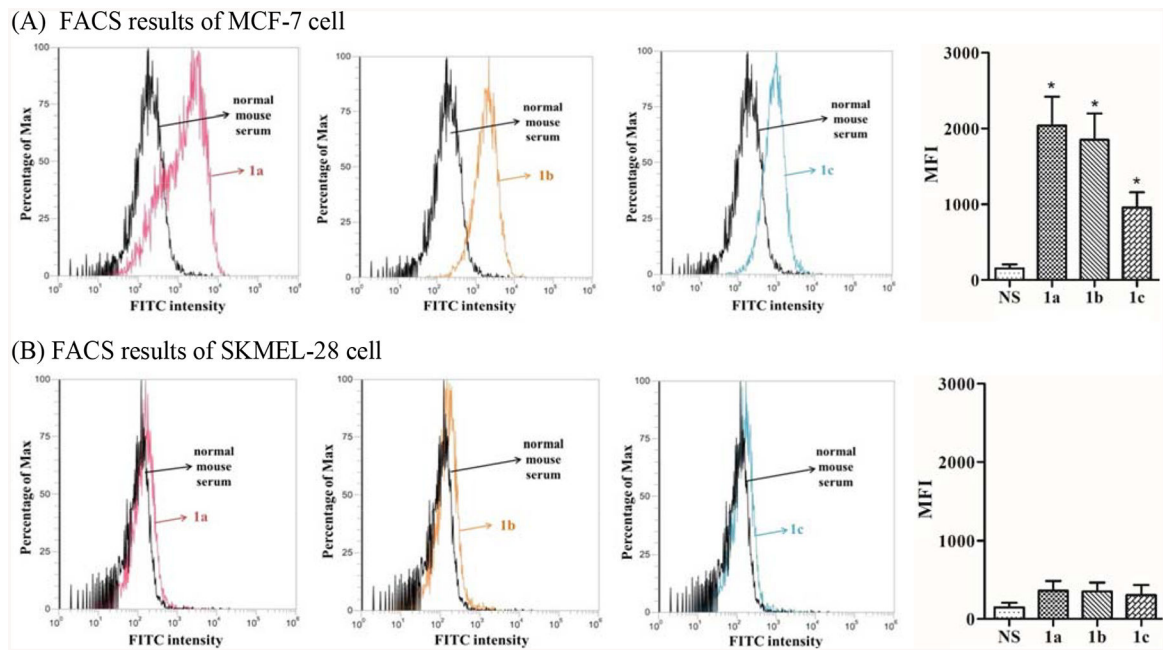
Total antibody (A), IgG antibody (B), and IgM antibody (C) titers of pooled day 0, 21, and 43 sera from mice before and after immunization with conjugates **1a**, **1b**, and **1c** as determined by ELISA with conjugates **2a–c** as coating antigens and 1:1,000 diluted AP-linked goat anti-mouse kappa (for total antibody detection), IgG and IgM antibodies as the secondary antibodies, respectively. Antibody titers were defined as the dilution numbers of sera at which the OD<sub>405</sub> value was 0.2. The mean antibody titer of three parallel experiments was presented for each sample, and the error bar showed the standard error of mean (SEM). \*Statistically different ( $p < 0.05$ ) as compared to the day 0 serum; #statistically different ( $p < 0.05$ ) between the two compared groups.



**Figure 3.**

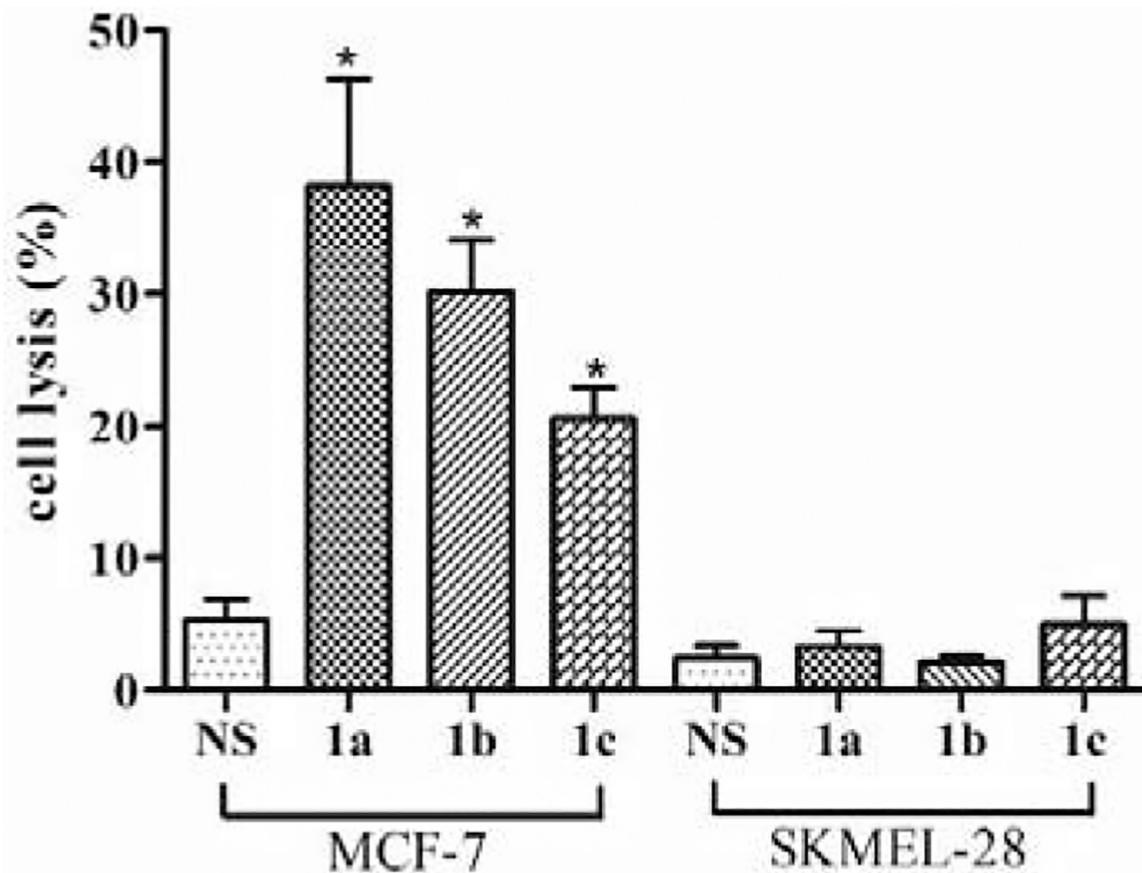
ELISA results showing the reactivity of each day 43 antiserum with all three different antigens. After ELISA plates were coated with HSA conjugates **2a**, **2b** and **2c**, the pooled day 43 serum from each group of mice immunized with conjugates **1a**, **1b** or **1c** was added to the plates for ELISA using 1:1,000 diluted AP-linked goat anti-mouse kappa and IgG secondary antibodies to detect total and IgG antibodies. Antibody titers were defined as the dilution numbers of sera at which an  $OD_{405}$  value of 0.2 was reached. The mean of antibody

titers from three parallel experiments was presented for each sample, and the error bar showed the SEM. \*Statistically different ( $p < 0.05$ ) between the two compared groups.

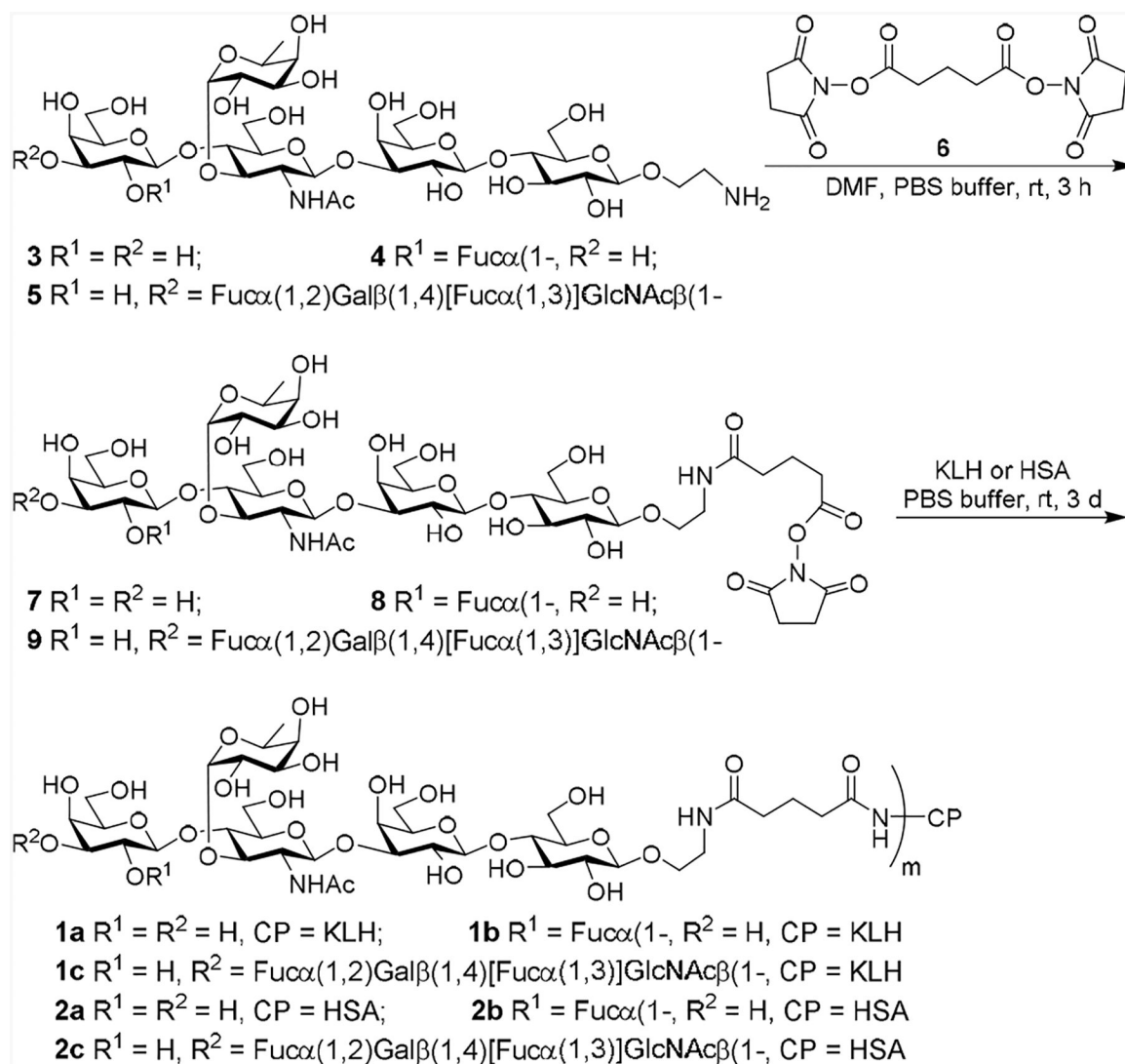


**Figure 4.**

FACS results about the binding of antibodies in normal mouse serum (black) or in the pooled antiserum of conjugate **1a** (red), conjugate **1b** (orange) and conjugate **1c** (blue) with MCF-7 cell (A) and SKMEL-28 cell (B), respectively, as well as the calculated MFI values. The error bar represents standard error of the mean (SEM) of three independent experiments. \*Statistically different ( $P < 0.05$ ) compared to that of cells treated with normal mouse serum.



**Figure 5.** The results of antibody-mediated CDC to MCF-7 and SKMEL-28 cells presented as cell lysis caused by treatment with normal mouse serum (NS) or a pooled antiserum of conjugate **1a**, **1b** or **1c** and then with rabbit complements. The error bar shows the SEM of 3 independent experiments. \*Significantly different ( $P < 0.05$ ) compared to the NS group.

**Scheme 1.**Synthesis of conjugates **1a–c** and **2a–c**

**Table 1.**Carbohydrate loadings of conjugates **1a–c** and **2a–c**

Glycoconjugate	1a	1b	1c	2a	2b	2c
Carbohydrate Loading (%)	18.3	13.9	11.4	11.1	7.9	14.1
Glycans/Glycoconjugate	105.0	64.6	33.8	10.3	6.0	7.5

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