

**PROTEIN FACTORS REGULATING THE HUMAN DEFENSE CELLS:
INVOLVEMENT OF GLYCOCONJUGATES****Lakhtin V. M.*, Lakhtin M. V. and Aleshkin V. A.**

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ABSTRACT

An analysis of the influence of glycoconjugates and carbohydrate moiety on protein factors modulating the activity of protective cells was carried out, including emphasis on the current state of problems in the relationship of protein macrophage-activating factors (MAFs). The prospects of using synthetic polymer glycoconjugates using the patterns of protein/glycoprotein-glycoconjugate interactions established by the authors and based on highly sensitive microanalysis methods developed with the participation of glycoconjugates of mono- and assembly molecular targets associated with membranes in the study of prophylactically and therapeutically significant (glyco)(lipo) proteins and (glyco)(lipo)peptides, including those affecting cellular population communications on the examples of MAFs and their peptide derivatives (beta-lactoglobulin, GcMAF, immunoglobulins, others) of colostrum, milk and probiotic cultures, are considered.

KEYWORDS: Cell activation factors, MAF, GcMAF, beta-lactoglobulin, macrophages, glycoconjugates, glycoproteins, recognition, assembly, protection, multiple forms, lectinbiotics, enzymebiotics.

ABBREVIATIONS

GC glycoconjugates

GcMAF Gc protein ((human group-specific component, a vitamin D-binding protein) derived MAF

GP glycoproteins

MAF macrophage-activating factor(s)

MIF	migration inhibitor factors
Mp	macrophages
PAA	polyacrylamide

1. INTRODUCTION

It is relevant to study the network biological activities of populations of protective cells and their subpopulations from the point of view of the positive effect of the protein factors of colostrum^[12,17,24-26,30,42,71, 60, 63, 65,69,78, 81], milk^[22,25,73,80,82], culture fluids and the cell surface of probiotics and other fermented milk bacteria-commensals^[4-7,9,35,48,54,55,58,62,83,87] on them with the result of strengthening the body's immunity.^[1,2,11,37,44,60] At the same time, there is an obvious need for systemic multifactorial approaches in the study of complex (combined and cascaded) biological activities of protective cells, including (sub)populations of macrophages (Mp) of the peritoneum and Mp-like blood cells^[1,3,7,8,10,11,29,37,41,52,55,58,62,81,83] Glycoconjugates (GC) play a special role in molecular and cellular communications of protection^[3-9,50,52,53] Their consideration can reveal a new potential of beneficial properties for humans at the supracellular, surface cell and intracellular (cytoplasmic and organelle) levels of the unified protection of the body from infections. However, this aspect remains the least studied.

The *goal* is to conduct a analysis of the potential of GC in the study, modulation and navigation of protein factors influencing the cells of the body's defense, including Mp, systemic GC as an important constituents of the overall adaptable defense of the body, based on our data.

2. THE MAIN CONTENT

2.1. *Aspects of the GC using in glycobiology*

GC include such carbohydrate-containing natural compounds as (lipo)polysaccharides, glycolipids, glyco(lipo)proteins (GP); their degradation products; as well as synthetic analogues of these natural categories. In this case, GC with a covalently attached carbohydrate part in the molecules are considered as containing a carbohydrate domain with additional functions such as modulating the communication activity of cells, the implementation of synergistic or antagonistic/ inhibitory combined reactions and resulting attacks, the manifestation of direct or indirect (mutual)actions against pathogens, pathogenic microbiocenoses, in support of non-antibody and antibody protection systems, innate and acquired immunity. GC can be submitted (and considered) systems of multiple forms, including combined ones, with varying biological activities, the manifestation of which

becomes even more diverse (the stability of the protective response, the response to stress is achieved) as part of complexes with other effectors and (co)factors - protein and non-protein, peptide and (glyco)(lipo)peptide nature. Such systems are characterized as co-functioning with a labile (navigationally configurable) system vector of potential resultant action/activities (combined activities).

The keys (according to our data) to the use of polymer synthetic GC as tools for the study of proteins, GP and receptors. ^[3-7, 9, 50-53]

Linear polymer based on the polyacrylamide (PAA) soluble in water and aqueous salt solutions were used-bark with lateral branches of carbohydrates (25% by weight) Gc (www.lectinity.com), imitating "mucins"-like polyvalent compounds with multiple clusters/sites of exposed homogeneous carbohydrate residues in the form of short (1-3 residues) lateral antennas. Such GC adapt their conformation ("random clumping" in solution) in accordance with the characteristics of carbohydrate-accepting protein sites, GP (and other types of GC) and their assembly complexes on the solid phase (membranes) simulating the cell surface. The GC used in the work imitated natural compounds of fundamentally different origin and could be used as natural models of reactions and processes in the body.

We have obtained the following results and conclusions about the patterns of interactions (glyco)(lipo)protein and synthetic polymer GC systems - soluble and solid-phase/surface-cell (in aqueous and solid phases):

- *synergism of the action of GC among themselves and with antibodies;
- *the ability of selected several (2-3) types of GC to visualize some/certain forms of protein factor as much as possible, to map the features of its structure;
- *the ability of synergistic sets of GC to recognize and bind protein factors /cytokines in the compositions of molecular complexes, nano- and microparticles;
- *preservation of factors in the composition of functionally active complexes and nanoparticles of the ability/activity for further sequential cascade binding of different types of GC;
- *sensitivity of GP targets to the topography of antenna clusters exposed in GC (the ability of the protein to recognize the availability and distribution in the space of clusters and antennas of glycans);
- *the presence of GC in the spatial field as systems (there is at least one "virtually binding")

parameter of the space coordination);

*asymmetry (generalized directionality and vectorality), mosaic systems of multiple forms of GP, including as complexes (their disparity in the severity of modulation of activities);

*expanding the potential of GC recognition by therapeutic protein/GP, the ability to form new recognition sites in protein-based complexes and supramolecular ensembles;

*correspondence between the systems "GC Recognition Proteins" and GC as a "One-Partner system/subsystem" (One GC is a set of proteins, One protein is a Set of GC), the presence of correspondence between protein and GC subsystems, the establishment of ranking orders (sub)Gc and protein systems, depending on the implementation of the types of activities;

*identification of the probiotic potential of the studied GC to support probiotic microbiocenoses, including those that increase the biocompatibility of lactobacilli, bifidobacteria, and their mixtures.

2.2. Basic properties of the protein factors influencing human protective cells^[10,39,40,56,68,74]

Protein MAFs exhibit all the properties of cytokine families, are, as a rule, GP and represent a set / system of multiple forms of various biological/ physiological purposes. Examples of some of the most studied protein factors (also as derivative effector peptides) are beta-lactoglobulin.^[16,20,21,27,28,32,36,45,46,66,70,77,86], GcMAF^[10,39,40,56,68,74], other types of milk proteins.^[72,73] As a rule, the properties and main forms of MAF are related (largely determined) by purification methods, for example, GcMAF using immobilized vitamin-D, since valuable antiviral properties are revealed in the line of anti(vitamin D/D3)-binding proteins.^[15,31,34,56,75,76,86] The microheterogenicity of the forms is determined not only by the action of environmental enzymes (including the effect of limited proteolysis), but is largely due to the presence of a highly variable carbohydrate part of the cytokine.^[24,25] The predominant forms corresponding to the specific responses of the Mp subpopulations are distinguished.^[10,11,15,24,25]

2.3. Biological activities of the MAFs

The main, most pronounced and studied protein factors affecting the activity of Mp include variants of MAFs, migration inhibitor factors (MIFs), chemotaxis reagents and other proteins, GP and their peptide derivatives - all of preventive and therapeutic significance, expanding the network of co-functioning between populations of Mp, Mp and other protective cells, tissues and organs, thereby increasing the reliability, stability and reactivity of the responses of the unified defense of the body. MAF activities are modulated by hydrolases present in the

environment (including participants in the probiotic compartment of biotope microbiocenoses), which strengthen or weaken the vector of the initial action of the protein factor at the peptide level.^[5-7,16,23,34,36,47,49,57,64,66,67,72,79,84]

2.4. Prospects for the discovery of new properties of MAFs, including taking into account interactions with GC, include (the basic aspects/ sides/ components of the issue are cited)

- genetic engineering aspects of studying and obtaining glycosylated MAFs^[24, 25];
- Lectin properties of MAFs, MAFs as lectinbiotics^[7,9,21,50,51,53,77], lectin sites in GC recognition during assembling milk protein factors^[20,21,45,77];
- The relationship of MAFs with enzymes and enzymbiotics^[53,73];
- MAF-regulation of biomarker enzymes in the serum of patients (increased activity of N-acetylgalactosaminidase - nagalase, which completely - not selectively deglycosylates vitamin D-binding protein and, thereby, blocks the formation of GcMAF, which leads to immune disorders)^[15,19];
- (glyco)(lipo)targeted peptides (including products of limited proteolysis, including those involving hydrolase systems of probiotics and/or strains and mixtures of lactic acid bacteria-comensals;
- postbiotic aspects (MAFs, including GcMAF, as sources and regulators of systemic postbiotic products, in synergistic combinations)^[5,6,83,87];
- directed purification of the "systemic MAFs" (through selected combined stages of affinity chromatography, with the achievement of a combined endogenous biocompatible/stable composition with the desired resultant action/actions);
- combined compositions of MAFs with additives, including probiotic and/or hydrolase nature;
- modulation of MAF communications through the formation of complexes with GP:
 - (inter)molecular, (inter)cellular, network;
- preventive and therapeutic aspects in relation to COVID-19 and concomitant diseases^[6-8,15,38,48,61,74-76,82], infectious diseases of microbial nature^[13,18,71,78,80]; in connection with tumors^[10,15,19], obesity and other metabolic disorders^[6,7,40,54,78];
- the use of GC/ oligosaccharides/ polysaccharides of milk and probiotics (their cultural hydrolysates) for disease-preventing purposes.^[1,13,18,19,29,33,43,59]

In connection with the fight against COVID-19, research and purification of angetensin-I-converting enzyme inhibitors are becoming important.^[14,15,35,85]

3. CONCLUSION

These data indicate a significant potential of GC in modulation, navigation and Mp training. The use of GC, the regularities of protein-GC interactions and GC-based technologies in the study of functionally active therapeutic proteins and factors on the examples of MAFs in aspects of immunology, biotechnology (microbiology), pharmacology, prevention and therapy of diseases is promising.

Disclosure of conflict of interest

The authors declare no conflict of interest.

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