

During biological evolution two essential steps were necessary.

The first was the emergence of cells via molecular self assemblies, and the second subsequent one was the development of multicellular assemblies.

Multicellular life is the most sophisticated form of changing patterns of matter and energy known so far.

The emergence of multicellularity has been tightly coupled to the ability of an organism to retain its own anatomical integrity and to distinguish self from non-self.

This process required the simultaneous development of cell adhesion and recognition properties.

Cell adhesion defines the form, physical state and consequently the functioning of all multicellular living systems.

Self is commonly consider in biology as syngeneic specific (genetically identical).

Non-self as either xenogeneic specific or allogeneic specific.

Xenogeneic specific: derived from an organism of a different species – genetically not identical.

Allogeneic specific: derived from separate individuals of the same species - genetically not identical.

Intermolecular binding forces between cell adhesion molecules (CAMs) are intrinsic properties of such cohesive structures.

Differences in the degree of binding strengths between diverse types of CAMs at the given environmental thermodynamic conditions determine selectivity of their associations.

Therefore, distinguishing self from non-self must operate via alogeneic and/or xenogeneic differences in types and/or spatial and temporal expression of cell recognition molecules and CAMs. Information technologies are using principles of biological self/non-self discrimination in order to ensure the security of computer systems by fast detection of unauthorized access, assuring the integrity of data and preventing the spread of computer viruses.

These protection problems are instances of the more general problem of distinguishing self (legitimate users, corrupted data, etc.) from other (unauthorized users, viruses, etc.).

A change-detection algorithms based on the way that natural immune systems distinguish self from other were developed. The sensor molecules guiding such recognition and adhesion should be present at the outermost cell surface.

What is the nature of the molecules operating in self/non-self discrimination?

Such molecules are biological polymers: proteins, glycans, lipids and their naturally covalently linked combinations glycoproteins and glycolipids including few organic monomeric molecules. To understand the molecular basis of primordial self-recognition and non-self discrimination we focused our attention on the role of proteoglycanlike glyconectins in Porifera xenogeneic cell interactions, as the evolutionary most compatible model system for ancestors of Metazoans.

The molecular mechanism of self/non-self discrimination and adhesion in sponges should be most similar to the mechanism that operated during the emergence of multicellularity.

Living Marine Sponges

Clathria prolifera



Halichondria panicea



Cliona celata



Our earlier investigations provided direct evidence that a novel class of primordial proteoglycans, named by us GLYCONECTINS (GNs), can mediate cell adhesion via a new alternative molecular mechanism of polyvalent glycan-glycan binding.



Atomic force microscopy measurements demonstrated that the binding strength between a single pair of Porifera cell surface glyconectin 1 (GN1) glycoconjugates from *C. prolifera* can hold the weight of 1,600 cells, proving their adhesion functions.

Ca²⁺-dependent self-interactions between glyconectin 1 molecules provide the major driving force for cell adhesion.







Using a quantitative functional approach with <u>color-coded beads</u>, <u>cells</u>, and <u>blotting</u>, we have shown that highly specific cell surface glyconectin to glyconectin interactions indeed mediate cell recognition and adhesion in the three Porifera species.

Grazt dependent alyconectin to glycohestin interactions mediate species-specific cell-cell recognition and adhesion without Ca²⁺ 29 110 110 10 mM Ca²⁺

←H. panicea

.C. celata

-C. prolifera

400 µm

400µm

10 mM Ca²⁺

without Ca²⁺ Simultaneous aposics aposific glyce CLYCONECTIM ition in RECOGNITION ay

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Physicochemical and biochemical properties of glycans from three GNs, were established by extensive multidisciplinary approach:

- gel electrophoresis,
- ultracentrifugation,
- carbohydrate analyses,
- glycosaminoglycan-degrading enzyme treatment,
- NMR,
- mass spectrometry.

Physicochemical Properties of Sponge Glyconectins (GNs)

	GN1	GN2	GN3
M _r x 10 ⁶	19	10	8
S _{20,w} (S)	58	42	46
Carbohydrate/ protein ratio (w/w)	63/37	21/79	36/64
Fucose (mol %)		7	231
Py(4,6)Gal (mol %)	4		0
GICA (mol %)		3	6
SO ₄ ²⁻ (mol/mol)	<mark>820</mark>	620	700

Monosaccharide Composition of GN Glycans

Monosac-	GN1 glycans	GN2 glycans	GN3 glycans
charides	(mol %)	(mol %)	(mol %)
Ara		0	7
Fuc	213	5	
Man	6	13	13
Gal	30	37	16
GalNAc	2	0	8
GIcNAc	23	15	32
Py(4,6)Gal	8	26	
GIcA			6

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The GN1, GN2, and GN3, showed speciesspecific carbohydrate compositions.

GN1 was characterized by a fucose content of one-third of the total monosaccharide content and minute amounts of Py(4,6)Gal.

GN2 had lower proportion of fucose and a molar ratio of Py(4,6)Gal three times higher than GN1.

GN3 had high quantities of fucose but no Py(4,6)Gal at all and was the only GN to contain arabinose (7 mol %).

NMR Analyses of Glyconectin Glycans - one-dimensional spectra -



Methyl signals of



NMR Analyses of Glyconectin Glycans - two-dimensional COSY90 spectra -



ES/MS profiling of ethanol-soluble oligosaccharides from purified glyconectins

ES/MS-MS fragmentation patterns in negative mode

Mass (m/z)

Structural organization of Glyconectin 1 adhesion glycans

The sequencing work presented here provides the first detailed comparative structural information on the glycan moieties of the glyconectins isolated from the *C. prolifera*, *H. panicea*, and *C. celata*.

The sequential and selective chemical degradation of these three GN glycans, followed by mass spectrometric and NMR analyses of the fragments, resulted in chemical fingerprints revealing new types of species-specific structures:

- sulfated glycan sequences in GN1,
- branched pyruvilated sequences in GN2,
- linear sulfate structures in GN3.

These data show also that GNs define a new family of proteoglycan-like molecules exhibiting **species-specific structures** with complex and repetitive acidic carbohydrate motives different from the classical proteoglycans and mucins.

Clearly the evolution from primordial Metazoans to complex organisms required the development of additional cell recognition and adhesion mechanisms mediated via lectin, immunoglobulin, integrin, and cadherin families of molecules.

The specificity of carbohydratemediated homophilic GN interactions in Porifera approaches the binding selectivity of the evolutionarily advanced immunoglobulin (IG) superfamily. However, the structural differences between these 2 systems imply conceptually distinct molecular mechanisms.

First, GNs are 100 times larger and extend 10 times farther from the cell surface than IG molecules.

Second, the GN1 specificity and tight binding of 10⁹ M⁻¹ reside in polyvalent glycan-glycan interactions of 1,000 sites, with a low affinity for the single site (10³ M⁻¹), whereas IGs recognize antigens via higher affinity ranging from 10⁴ to 10⁹ M⁻¹, with low valency binding.

Despite this evolving complexity of the components participating in cellular interactions, glyconectin-like structures have been preserved in mammalian systems, suggesting their versatility in recognition and adhesion.

In conclusion, xenogeneic class of glyconectin glycans, as the most peripheral cell surface molecules of sponges (today's simplest living Metazoa), are proposed to be the primary cell adhesive molecules possessing self/non-self discrimination essential for the evolution of the multicellularity.

The GN-GN interactions, a veritable nano-velcro system, may thus provide a new model for molecular self-recognition.

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