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Spatially controlled glycocalyx engineering for growth factor patterning in embryoid bodies

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ABSTRACT: Growth factor (GF) patterning in stem cell spheroids, such as embryoid bodies (EBs), has been sought to guide their differentiation and organization into functional 3D tissue models and organoids. Current approaches relying on exposure of EBs to gradients of GFs suffer from poor molecular transport in the spheroid microenvironment and from high cost of production and low stability of recombinant GFs. We have developed an alternative method for establishing GF gradients in EBs utilizing stem cell surface engineering with membrane-targeting heparan sulfate-glycomimetic co-receptors for GFs. We have capitalized on the ability of amphiphilic lipid-functionalized glycopolymers with affinity for FGF2 to assemble into nanoscale vesicles with tunable dimensions and extracellular matrix penetrance. Upon size-dependent diffusion into EBs, the vesicles fused with the plasma membranes of stem cells, giving rise to concentric gradients of cells with enhanced FGF2-binding. The extracellular matrix-assisted cell surface remodeling process described is the first example of spatially-targeted glycocalyx engineering in multicellular systems to control GF localization. The glycopolymer structure, vesicle dimensions, and remodeling conditions determine the level of FGF2 adhesion and gradient slope. The increased chemical and thermal stability of the synthetic glycomimetics and the tunability of their GF-binding profile, which is defined by their glycosylation and may be extended to other recombinant or endogenous morphogens beyond FGF2, further increase the versatility of this method.

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Introduction.

Embryonic stem cells (ESCs) and induced pluripotent stem cells (iPSCs) have the capacity to differentiate into any cell type of the adult body.¹ They provide an important tool to study the process of embryogenesis and organismal development² and hold promise for a range of biomedical applications.³ A key challenge in realizing their full potential is in translating differentiation protocols established in 2D *in vitro* cultures to the highly complex 3D environments of tissues *in vivo*.^{4,5} 3D spheroids formed from ESCs and iPSCs, such as embryoid bodies (EB),⁶ offer a convenient laboratory system to bridge this gap. EBs have the capacity to undergo spontaneous self-patterning into ordered multicellular structures;⁷ however, harnessing this potential for reliable production of functional tissue and organ replacements will require achieving better spatial and temporal control over cell differentiation in these systems.⁸

Current methods for engineering the microenvironment and self-organization of stem cell spheroids focus primarily on directing the delivery, release and diffusion of differentiation cues, such as growth factors (GFs) and small molecule morphogens.^{9,10} For instance, spatial control over cell differentiation in EBs has been accomplished through their exposure to GF gradients in microfluidic devices^{11,12} or through encapsulation in hydrogels with controlled GF release (Fig 1A).^{13,14} The inherent challenge in these approaches is a restricted transport of morphogens into the spheroids through their outer shell composed of tightly associated cells and dense deposits of extracellular matrix (ECM).^{15,16} Imbedding of GFladen microparticles with stem cells during EB aggregation overcomes this challenge; however, it provides limited control over the localization of the GF source within the spheroid (Fig 1A).^{17,18}

Here, we present an alternative chemical method for generating gradients of GFs in EBs, which takes advantage of the restricted macromolecular transport across the spheroid boundary (Fig 1B). Our approach utilizes lipid-modified glycopolymers (**GPs**) with tunable size and affinity for GFs, which insert directly into the plasma membranes of ESCs and promote GF adhesion. Pre-assembly of the amphiphilic glycomimetics into nanoscale glycopolymer vesicles (**GPVs**) with defined dimensions enables tuning of their diffusion into the EBs, upon which they undergo fusion with the plasma membranes of ESCs to produce gradients of cells with enhanced GF-affinity (Fig 1B).

Results and discussion.

The glycomimetics are inspired by heparan sulfate (HS) glycosaminoglycans (GAGs), which are sulfated polysaccharide co-receptors for GFs on surfaces of cells and regulators of their activity,¹⁹ including during stem cell differentiation.²⁰ Synthetic GAG mimetics have emerged as readily accessible biomaterials with capacity to bind and regulate the activity of a range of HS-binding proteins. Comprising linear polymeric scaffolds decorated with variously sulfated synthetic disaccharides representing the elementary structural motif of HS, soluble HS-mimetic glycopolymers have been shown to attenuate B cell chemotaxis induced by the chemokine RANTES²¹ or inhibit breast cell metastasis in mouse models by blocking the activity of matrix heparinases.²² We have developed HS-mimetic polymers with affinity for various GFs comprising enzymatically derived HS disaccharides.^{23,24} Endowing the polymers with phospholipid tails allowed for their targeting to the plasma membranes of ECSs. There, the materials served as functional surrogates for native GAGs, activated signaling by FGF2 and BMP4, and promoted neural and mesodermal differentiation, respectively. We envisioned that the phospholipids, which allows for insertion of the HS mimetics into the plasma membranes of cells, also give the **GPs** amphiphilic character and should promote their assembly into **GPVs** (Fig 1B). Polymer length as well as the size and charge of their glycans should allow for tuning of **GPV** size and penetrance into the EBs (Fig 1B).

To test this hypothesis, we generated a panel of **GPs** ranging in size from short (**S**, degree of polymerization, $DP \sim 40$) to medium (**M**, $DP \sim 125$) and long (**L**, $DP \sim 300$) and glycosylated either with the 2,6-*O*-

disulfated HS-derived disaccharide, Δ UA2S-GlcNAc6S (D2A6), or the monosaccharides *N*-acetylglucosamine (GlcNAc) and its 6-*O*-sulfated derivative, GlcNAc-6S (Fig 2A). The **GPs** were derived from a Boc-protected poly(3-*N*-methylaminooxypropyl) acrylamide precursors (**P1**) generated by RAFT²⁵ polymerization of monomer **1** in the presence of a dipalmitoyl phospholipid-containing chain transfer agent **CTA** and the radical initiator, AIBN (Fig 2A).²³ The RAFT process furnished precursors **P1** with good control over molecular weight and dispersity (Đ), as determined by size exclusion chromatography (SEC, Figs 2B and S1).

The polymeric precursors **P1** were elaborated in four steps to produce glycopolymers **GP** labeled with AlexaFluor488 (AF488) for visualization (Fig 2A). First, the trithiocarbonate chain end groups in polymers **P1** were cleaved with *n*-butylamine to release a reactive thiol, which was capped immediately with AF488-maleimide. The protective Boc groups were then removed by treatment with trimethylsilyl chloride in phenol to yield polymers **P2** with exposed *N*-methylaminooxy side chains for subsequent ligation of reducing glycans.²⁶ The extent of polymer labeling (*f*_{AF488}) was determined by UV-VIS at this stage to be ~ 0.7 - 0.8 fluorophores per polymer chain (Table S3). Finally, heating of intermediates **P2** with the mono- and disaccharides at 50 °C under acidic conditions (acetate buffer, pH = 4.5), followed by size filtration yielded the desired **GPs** (Fig 2C). The fraction of glycosylated side chains was determined for each glycopolymer by ¹H NMR spectroscopy and ranged from ~ 40-50% for the charged glycans, D2A6 and GlcNAc-6S, to ~ 50-60% for the neutral monosaccharide, GlcNAc (Fig 2C and Table S4).

We evaluated the affinity of the negatively charged HS-mimetic glycopolymers for FGF2, which induces ECSs toward neural commitment.²⁰ Using ELISA with surface-immobilized FGF2 (Fig S3), we measured strong binding for polymers **GP-D2A6** carrying the 2,6-*O*-disulfated HS disaccharides ($K_{d,app} = 2-7$ nM). The observed affinity was comparable to that of heparin ($K_{d,app} = 6$ nM), which is a highly sulfated form HS. The monosaccharide-modified polymers **GP-GlcNAc-6S** showed no measurable FGF2 binding activity (Fig S3).

Dynamic light scattering (DLS) analysis of the **GP** solutions in PBS buffer (pH = 7.4) confirmed their assembly into **GPVs** with sizes of ~ 40-150 nm (Figs 3 and S4). Using the fluorescent lipophilic dye, DiI, we determined the critical vesicle-forming concertation (cvc) of the **GPs** in the range of 0.1 - 0.3 μ M regardless of polymer length (Figs S5). The molecular weight (Mw) of the polymers scaled linearly with their length (DP). The steepest rise in Mw was observed for the disaccharide containing polymers, **GP-D2A6**, followed by the monosaccharide polymers **GP-GlcNAc-6S** and **GP-GlcNAc** (Fig 3A). Accordingly, the short (**S**), medium (**M**), and long (**L**) glycopolymers **GP-D2A6** afforded the largest **GPVs** (Figs 3B and S4, Table S5) with diameters (*d*_{GPV}) of 44 ± 1, 106 ± 9, and 139 ± 12 nm, respectively (Fig 3B and 3C). The monosaccharide analogs **GP-GlcNAc-6S** and **GP-GlcNAc** gave **GPVs** of approximately equal sizes for each polymer length: 42 ± 8 and 36 ± 1 nm (**S**), 64 ± 5 and 65 ± 4 nm (**M**), and 74 ± 13 and 84 ± 24 nm (**L**). Polymers without the lipids remained dissolved and did not form **GPVs**. Our DLS data indicate that the **GPV** size is determined primarily by the DP and Mw of the glycopolymers but not their charge, which was reflected in the zeta potential of the vesicles (Fig 3D, Table S5).

The single short hydrophobic dipalmitoyl lipid tail attached to the much more extended hydrophilic glycosylated segment of the **GPs** is likely to drive the formation of vesicles rather than micelles,²⁷ with the latter expected to produce much smaller particle sizes than those observed by DLS. The **GPVs**, which form during the glycopolymer assembly, remain stable in solution over a range of concentrations (0.3 -10.0 μ M) according to DLS measurements (Fig S6). The **GPVs** can be disrupted by the addition a detergent, such as TritonX-100 (Fig S4E). Irreversible aggregation of the **GPVs** was observed after freezing, drying, or prolonged storage in solution. We envisioned that differential size-dependent diffusion of the **GPVs** through the ECM network of EBs, followed by their fusion with ESC membranes, would establish gradients of HS mimetic-remodeled stem cells with increasing FGF2 affinity (Fig 1B). To eliminate background FGF2 binding from endogenous HS glycans on ESCs, we used mutant cells missing the gene, exostosin-1 (*Ext1*), which encodes for a glycosyltransferase involved in HS chain polymerization.¹⁹ Unable to produce functional HS, the *Ext1*^{-/-} ESCs lack the ability to bind many HS-dependent GFs, including FGF2.²⁸

Since HS also contributes to cell adhesion and deposition of matrix proteins (e.g., fibronectin),²⁹ we first analyzed the effects of the *Ext1* gene deletion on EB formation and ECM organization. Wild type (*wt*) E14 and *Ext1*^{-/-} ESCs were aggregated into EBs under embryonic conditions in the presence of LIF using the hanging drop method (Fig 4A).³⁰ After 2 days, both cell lines formed EB spheroids with normal morphology and sizes of ~ 100-200 μ m (Figs 4B and S6). Scanning electron microscopy (SEM) imaging confirmed the formation of ECM deposits on the surface of EBs from both cell lines (Figs 4C and S8). Immunohistological analysis of the EBs confirmed the lack of HS expression by *Ext1*^{-/-} ESCs, which resulted in visible accumulation of fibronectin toward the EB periphery, while collagen distribution was affected to a much lesser degree (Fig 4D and Fig S9). The ability of *Ext1*^{-/-} ESCs to form EBs and deposit ECM, we reasoned, should influence **GPV** penetrance into the spheroids.

Next, we set to test whether the **GPVs** can remodel the glycocalyx of ESCs in a gradient pattern after diffusion into the EBs (Fig 5). In addition to their size, EB penetrance of the **GPVs** is likely to be determined by their concentration in the media and the duration of treatment. We examined these parameters by first incubating Day 2 *Ext1*^{-/-}ESC EBs with 0.1, 1.0 and 3.0 μ M fluorescently-labeled **GP-M/GlcNAc-6S** in serum-free media. After 2 hrs at 37 °C, the EBs were washed to remove excess **GPVs**, fixed, and imaged by confocal microscopy to assess glycopolymer distribution.

Fluorescence images of cross-sections at the EB midpoint showed robust and uniform labeling with 3.0 μ M **GP-M/GlcNAc-6S** (Fig S10). Dilution of the polymer resulted in overall lower levels of cell labeling and polymer accumulation toward the EB surface. By shortening the incubation time at the highest polymer concentration (3.0 μ M) to 1 hr, we retained robust cell labeling while also establishing a concentric polymer gradient across the spheroid (Fig S9). Treatment of the EBs with glycopolymers **GP-GlcNAc-6S** of all three lengths under the optimized conditions (3.0 μ M, 1hr, 37 °C) resulted in differential diffusion of their **GPVs** into EBs according to their size (Figs 5A and S16). We observed a transition from complete **GPV** penetrance for the shortest polymer **GP-S/GlcNAc-6S** (DP ~ 40, $d_{GPV} = 42$ nm) to increasing outer cell layer accumulation as the polymer size and **GPV** diameter increased (cf. **GP-L/GlcNAc-6S**, DP~300, $d_{GPV} = 74$ nm).

We sought to develop a set of quantitative measures to assess **GPVs** diffusion into the EBs and to characterize **GP** gradients after ESC remodeling. We collected fluorescence micrographs of EB cross-sections at their widest point and analyzed fluorescence intensity from the polymers (F_{488}) along a line passing through the EB center (Fig 5A). We then plotted the normalized intensities against distance from EB center (Fig 5A, *green*). The area under the curve provided a quantitative measure of **GPV** penetrance into the EB (Fig 5B, *bars*). To address signal variability arising from local microheterogeneities within the EB structures, we performed bin analysis of the normalized line fluorescence intensity and generated intensity histograms (Fig 5A, *bars*), where each bar represents mean pixel intensity over a 10 µm bin.

To evaluate how rapidly the gradient of **GP**-remodeled cells changes with **GPV** diffusion into EBs, we plotted the absolute values of line fluorescence intensity change (F'_{488}) against distance from EB center (Fig 5B, *line graph*). In agreement with our visual assessment of cross-section images, we observed the slowest change in fluorescence intensity for **GP-S/GlcNAc-6S**, consistent with its highest penetrance and

uniform distribution across the EB. The rate of gradient change increased with polymer length and **GPV** size. The neutral polymer **GP-L-GlcNAc** did not exhibit enhanced diffusion compared to the sulfated analogs GlcNAc-6S analog, indicating that **GPV** diameter rather than charge determines EB penetrance (Figs S14 and S15).

The polymer fluorescence after EB remodeling was localized to the ESC surface, indicating insertion of the **GPs** into the cell membranes via their lipid tails (Fig S13D). To confirm membrane tethering, EBs treated with **GP-S/GlcNAc-6S** and **GP-L/GlcNAc-6S** were dissociated into single cells by Accutase, washed to remove adsorbed **GPVs**, and analyzed by flow cytometry (Fig S16). Both polymers remained associated with the cell surface and the population of labeled cells increased with higher EB penetrance of the **GPs**, consistent with fluorescence image analysis (Fig 5B).

Focusing on our aim to exploit the size-limited diffusion of **GPVs** in the EB environment to engineer GF affinity gradients, we performed remodeling experiments with the FGF2-binding HS mimetics, **GP-D2A6**. We have previously shown that installation of glycopolymers carrying the D2A6 disaccharide at the surface of $Ext1^{-/-}$ ESCs restored FGF2 binding and signaling activity,²³ thus providing functional surrogates for cell-surface HS. While the *wt* E14 ESCs express endogenous HS structures that produce uniform FGF2 binding across the EB, the $Ext1^{-/-}$ ESCs HS mutants lack affinity for FGF2 and provide a clean slate for the pattering of the GF (Fig 5C).

Treatment of Day 2 *Ext1*^{-/-} ESC EBs with **GP-D2A6** polymers of all three lengths under the optimized conditions (3.0 μ M, 1hr, 37 °C) resulted in patterns of remodeled cells analogous to those observed for the model monosaccharide polymers **GP-GlcNAc-6S** (Figs 5D and 5A, *green*). The shortest polymers **GP-S/D2A6** (*d*_{GPV} = 44 nm) gave full coverage of the EB interior similar to its monosaccharide derivative **GP-S/GlcNAc-6S** which forms **GPVs** of comparable size (*d*_{GPV} = 42 nm). By contrast, the medium-sized polymer **GP-M/D2A6** (*d*_{GPV} = 106 nm) exhibited lower EB penetrance and generated a steeper fluorescence gradient compared to **GP-M/GlcNAc-6S** (with *d*_{GPV} = 64 nm). The same trend continued for the pair of longest polymers, **GP-L/D2A6** (*d*_{GPV} = 139 nm) and **GP-L/GlcNAc-6S** (*d*_{GPV} = 74 nm), which showed the lowest penetrance and remained localized primarily on the EB periphery. Our observations are consistent with **GP** diffusion into the EB spheroids being determined primarily by the size of their **GPVs** independent of the structure and charge state of their glycans, which provides an element of predictability and tunability.

The patterns of cell-surface anchored HS-mimetics, **GP-D2A6**, in turn, defined the distribution of FGF2 adhesion sites in the *Ext1*^{-/-} ESC EBs, which were visualized by incubation with recombinant FGF2 followed by immunostaining (Fig 5D, *red*, and Figs S14 and S15). No FGF2 signal was observed in EBs treated with polymers **GP-GlcNAc-6S** (Fig S19), confirming that the GF binding is glycan dependent and the display of the HS mimetics **GP-D2A6** on the surface of ESCs is required to constrain FGF2 localization.

Conclusions.

We have developed a novel, facile method for generating GF gradients in multicellular spheroids using cell membrane engineering with synthetic GF co-receptors. By exploiting the assembly of amphiphilic glycopolymers into nanoscale vesicles with tunable diameters, we were able to control their diffusion into EB spheroids. Upon entry, the **GPVs** fused with the membranes of nearby stem cells and introduced new adhesion sites for FGF2 at their surface. By using $Ext1^{-/-}$ ESCs lacking endogenous HS, we were able to limit FGF2 adhesion only to cells remodeled with appropriately glycosylated HS mimetics. The formation of vesicles and their fusion into the membranes of cells within EBs provides a unique delivery method for

glycan-based receptors to the surface of cells in complex systems. In departure from existing methods for gradient patterning in 3D spheroids, which rely on the controlled diffusion and release of recombinant GFs, the current approach achieves GF patterning by tailoring their interactions at the cell surface and, thus, provides control over the activity of endogenous GFs. The cell surface engineering method can be extended to other types of cells and HS-dependent signaling molecules, and we expect its utility to increase further with the continuing improvements in HS mimetic design^{21,22} and with the emergence of new tools for the genetic³¹ and chemical³² manipulation of HS expression in living cells. Further, we expect that the discovery of assembly of amphiphilic glycomaterials into tunable and stable vesicles capable of direct fusion with cell membranes will open new modalities for glycocalyx engineering in increasingly complex biological systems.

Conflict of interest

There are no conflicts to declare.

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Figures:



Figure 1. Growth factor (GF) gradient engineering in stem cell spheroids. A) Prior methods utilize exposure of spheroids to GF gradients in microfluidic devices and engineered hydrogels or through imbedding of GF-laden microparticles in the spheroid core. B) The present approach uses cell surface engineering with glycomimetics to tailor stem cell interactions with GFs. Amphiphilic GF-binding glycopolymers (**GPs**) assemble into glycopolymeric vesicles (**GPVs**) with tunable size and extracellular matrix penetrance. Upon entry into the spheroid, the **GPVs** fuse with cell membranes, forming gradients of stem cells with enhanced GF affinity.



Figure 2. Synthesis and characterization of lipid-terminated HS-mimetic glycopolymers, **GP**. A) Short (**S**), medium (**M**), and long (**L**) **GPs** were assembled from a RAFT-derived polymeric precursor **P1**. B) Size exclusion chromatography (SEC) traces of **P1-S**, **M** and **L** (DP = 40, 125 and 300, respectively). C) Structural characteristics of **GPs**. DP = degree of polymerization, D = dispersity, $f_{AF488} =$ fraction of AlexaFluor 488 (AF488)-labeled chains.



Figure 3. Assembly of amphiphilic HS-mimetic glycopolymers (**GPs**) into glycopolymeric vesicles (**GPVs**) with tunable diameter. A) Mw of glycopolymers **GP** increased proportionally with polymer **DP** and glycosylation. B) Short (**S**), medium (**M**), and long (**L**) **GPs** (3 μ M) assembled in water into **GPVs**. Polymer Mw, but not charge, determined **GPV** diameter. C) Size distributions for **GPVs** formed by polymers **GP-D2A6**. D) Zeta potential of **GPVs** (shown for glycopolymers **GP-L**), MQ = MilliQ water, ****p < 0.001.



Figure 4. Formation and characterization of embryoid bodies (EBs). A) E14 and $Ext1^{-/-}$ ESCs were aggregated in hanging drops in the presence of LIF for 2 days. B) Optical microscopy confirmed normal EB formation. C) SEM images show similar level of ECM deposition on the surfaces of EBs from both cell lines. D) Immunostaining with anti-HS antibody, 10E4, confirmed lack of HS expression by $Ext1^{-/-}$ ESCs. Loss of HS led to accumulation of fibronectin toward the outer regions of the EBs but had no effect on collagen deposition (*blue* = Hoechst 33342 nuclear stain).



Figure 5. Engineering FGF2 affinity gradients in EB spheroids. A) *Top:* Fluorescence confocal micrographs of EB cross-sections at midpoint after treatment with short (**S**), medium (**M**), and long (**L**) AF488labeled glycopolymers **GP-GlcNAc-6S**. *Bottom:* Normalized fluorescence intensity (*green*) and bin histogram analysis (*bars*, 1 bin = 10 µm) of AF488 signal (F_{488}) along a line passing through EB center. B) Total fluorescence intensity (F_{488}) and rate of intensity change with distance from EB center (F_{488}). C) FGF2 (*red*) binds to *wt* E14 but not $Ext1^{-/-}$ ESCs in EBs. D) Fluorescence micrographs and normalized line fluorescence for FGF2 (*red*) binding to EBs remodeled with glycopolymers **GP-D2A6** (*green*).