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Imaging the Glycome in Living Systems

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Abstract

The glycome, the full complement of glycans that cells produce, is an attractive target for molecular imaging. Imaging of the glycome in living systems has recently been enabled via bioorthogonal chemical reporter-based approaches. In this chapter, we describe two approaches to introduce bioorthogonal chemical reporters (tags) onto cell surface fucosylated glycans and glycans bearing LacNAc disaccharides, respectively. The tagged glycans can then be conjugated to imaging probes via bioorthogonal click chemistry. Similar approaches can be extended to image other sectors of the glycome in living systems.

1. Introduction

The surfaces of eukaryotic cells are covered with complex glycans that participate in a variety of physiological processes, including angiogenesis, fertilization, embryogenesis, cell adhesion, and neuronal development (Gupta *et al.*, 2009; Ma *et al.*, 2006; Murrey and Hsieh-Wilson, 2008; Varki *et al.*, 2008). Inside the cell, glycans regulate transcription, translation, and protein trafficking (Gouyer *et al.*, 2001; Hart *et al.*, 2011). Glycoconjugates and glycoproteins are also found in certain prokaryotes, including pathogenic bacterium *Campylobacter jejuni* and human gut symbionts within the order Bacteroidales (Coyne *et al.*, 2005). Studies have shown that these glycoproteins are either associated with virulence factors of clinically significant pathogens (Schmidt *et al.*, 2003) or play critical roles in host-symbiont interactions (Comstock, 2009).

The full complement of glycans that a cell produces is collectively termed the cell's glycome. Many cellular factors, including the cell's genome, transcriptome, and proteome, and environmental cues and nutrients may have influence on the glycome (Freeze, 2006). Thus, the glycome responds to and reports on the physiological state of the cell. In humans, changes in cells' glycome are associated with developmental disorders and defects (Varki *et al.*, 2008), and can mark the onset of cancer and inflammation (Brooks *et al.*, 2008; Dube and Bertozzi, 2005). The ability to visualize and characterize these changes in living systems would advance our understanding of the detailed roles of glycans in these processes and provide new clinical tools for disease diagnosis. However, glycans are assembled in a step-wise fashion by multiple enzymes, that is, monosaccharide kinase, nucleotidyltransferase, glycosyltransferase, and thus by multiple genes. Therefore, they are not amenable to imaging techniques that rely on genetic reporters (i.e., green fluorescent protein, GFP).

Conventional methods to detect cell surface glycans rely on lectins and antibodies (Comer *et al.*, 2001; David *et al.*, 1992; Duijvestijn *et al.*, 1988; Pilobello and Mahal, 2007). Lectins

have been used extensively for the detection of both monosaccharides and oligosaccharides (Hirabayashi, 2008), and many of them have been commercialized. However, lectins typically have low affinity for their glycan epitopes and are often toxic (Ohba and Bakalova, 2003). Likewise, most antibodies generated against glycans are of the low-affinity IgM subtype and are tissue-impermeant. Moreover, lectin- and antibody-based imaging approaches only provide a snapshot of the labeled glycans at a particular time-point, and cannot be used for dynamic studies in a cellular environment. Glycans can also be metabolically labeled with radioisotope-bearing monosaccharides or the corresponding nucleotide sugars (Becker and Lowe, 2003; Jork *et al.*, 1984), but the subsequent visualization of the radioisotopically labeled glycans in intact cells is not well resolved or amenable to current fluorescence microscopy techniques.

It is now possible to image the glycome in live cells or in living organisms using new tools from the emerging field of bioorthogonal click chemistry (Baskin and Bertozzi, 2007; Laughlin and Bertozzi, 2009a,b). By hijacking a cell's glycan biosynthetic machinery, a monosaccharide building block functionalized with a bioorthogonal chemical tag is incorporated into target glycoconjugates. Subsequently, a tailor-made click reaction is employed to conjugate a complementary biophysical probe, enabling visualization (Laughlin and Bertozzi, 2009a,b), or enrichment of the target glycoproteins for molecular identification (Fig. 21.1) (Hanson *et al.*, 2007; Wang *et al.*, 2010).

In this chapter, we first briefly describe the most frequently used bioorthogonal click chemistry that has been applied to imaging the glycome *in vivo*. Next, we focus on two bioorthogonal click chemistry-based techniques developed in our laboratory for imaging fucosylated glycans and glycans bearing *N*-acetyl-D-lactosamine (LacNAc, Gal β 1,4GlcNAc) in living systems.

2. Bioorthogonal Chemistry in Glycan Labeling: Merits and Limitations

2.1. The condensation of ketones/aldehyde with aminoxy and hydrazide reagents

The first bioorthogonal click reaction that has been exploited to label surface glycans for imaging studies is the condensation of ketones and aldehydes with aminoxy and hydrazide-bearing reagents to form stable hydrazones and oximes adducts, respectively (Fig. 21.2A). The optimal pH of these reactions is 5–6, which works for some cell lines, but is not compatible with many living systems.

In 1997, Bertozzi and coworkers reported that an *N*-acetyl mannosamine analog derivatized with a levulinoyl side chain can be metabolized and incorporated into cell surface sialylated glycans in Jurkat cells. The ketone group allows subsequent reaction with a biotin hydrazide probe, and the labeled cells can then be detected using flow cytometry (Mahal *et al.*, 1997).

To improve the biocompatibility of this reaction, Dawson and coworkers introduced aniline as a catalyst to accelerate the reaction so that the condensation can be performed at physiological conditions (Dirksen *et al.*, 2006). The Paulson group, in a joint effort with Dawson and coworkers, used this optimized reaction to image sialylated glycans on the surface of mammalian cells (Zeng *et al.*, 2009). In their experiment, cells were first

subjected to mild periodate oxidation to selectively introduce an aldehyde at the C-7 position of sialic acid, which then underwent a condensation reaction with aminooxybiotin catalyzed by aniline (Fig. 21.2A). Subsequently, the treated cells were stained with a streptavidin conjugated with a green fluorophore. As analyzed by confocal microscope, robust labeling was achieved on the cell surface without jeopardizing membrane integrity.

2.2. Azide-associated bioorthogonal reactions

To date, azide is the most utilized bioorthogonal chemical tag for labeling glycans due to its small size and inertness to most components in a biological environment (Sletten and Bertozzi, 2009). Three bioorthogonal click reactions have been reported for labeling azide-tagged biomolecules. The Staudinger ligation covalently links the azide and an ester-functionalized triphenylphosphine via an amide bond (Fig. 21.2B) (Saxon and Bertozzi, 2000). Recently, this reaction has been successfully applied for imaging sialylated tumor cell glycans *in vivo* (Neves *et al.*, 2011). As demonstrated by Brindle and coworkers, peracetylated azidoacetyl mannosamine was injected intraperitoneally to label sialic acids with an azide tag in tumor-implanted nude mice. The sialic acid-associated azides were then reacted, by Staudinger ligation, with a biotinylated phosphine probe and the biotin was detected by subsequent intravenous injection of a far-red fluorophore or a DOTA-¹¹¹In-conjugated neutravidin. At 24 h after administration of neutravidin derivatives, the mice were imaged using optical imaging or single-photon-emission computed tomography (SPECT), respectively. Positive signal resulting from the azido-dependent labeling was primarily detected in tumors. As upregulated sialylation is strongly correlated with the transformed phenotype of many cancers, this technique has the potential to be translated into a clinical setting to monitor the progression of cancer.

Though highly specific, the Staudinger ligation suffers from slow reaction kinetics and competing oxidation of the phosphine reagents (Baskin *et al.*, 2007). By contrast, the Cu(I)-catalyzed azide-alkyne cycloaddition (CuAAC) (Fig. 21.2C) (Rostovtsev *et al.*, 2002; Tornøe *et al.*, 2002), promoted by the Cu(I)-stabilizing ligand TBTA (Fig. 21.2E) (Chan *et al.*, 2004), enjoys readily available coupling reagents and improved kinetics (Hein and Fokin, 2010; Wu and Fokin, 2007). It has been used extensively by chemical biologist for labeling glycans (Laughlin and Bertozzi, 2009a,b) and detecting glycoproteins in cell lysates (Chang *et al.*, 2009; Hsu *et al.*, 2007). However, the current Cu(I) catalyst formulation has two major problems: toxicity, hindering its use in living systems (Prescher and Bertozzi, 2005; Sletten and Bertozzi, 2009), and slow kinetics in aqueous solutions at micromolar substrate concentrations (Soriano del Amo *et al.*, 2010), resulting in incomplete cycloaddition reaction (Kaltgrad *et al.*, 2007), which hampers its use in modifying biomolecules of limited quantities.

The third reaction, the strain-promoted cycloaddition of azides and cyclooctynes (Agard *et al.*, 2006; Jewett *et al.*, 2010), inherits the bio-benign characteristics of the Staudinger ligation but is further endowed with improved kinetics (Fig. 21.2D) (Baskin *et al.*, 2007; Jewett and Bertozzi, 2010; Ning *et al.*, 2008). Among the cyclooctynes developed by the Bertozzi and Boons group, a difluorinated cyclooctyne (DIFO) (Baskin *et al.*, 2007), and a biarylazacyclooctynone (BARAC) (Jewett *et al.*, 2010) showed rapid kinetics in

biomolecular labeling experiments. Currently, DIFO-fluorophore conjugates are the first choice for imaging azide-tagged glycans in living systems (i.e., zebrafish embryos and *Caenorhabditis elegans*) (Laughlin and Bertozzi, 2009a,b; Laughlin *et al.*, 2008). However, labeling requires a 1-h incubation time (Baskin *et al.*, 2010; Laughlin *et al.*, 2008). Moreover, *in vivo* studies revealed that cyclooctyne-based probes bind to mouse serum albumin nonspecifically, presumably via covalent reactions with cysteine residues (Chang *et al.*, 2010). In addition, the construction of these cyclooctynes involves a multistep linear synthesis that can be technically challenging (Baskin *et al.*, 2007; Poloukhine *et al.*, 2009).

Most recently, our lab discovered a new ligand for CuAAC—BTTEs (Fig. 21.2E) (Soriano del Amo *et al.*, 2010). Not only does BTTEs dramatically boost the reactivity of CuAAC, it also confers the canonical CuAAC with biocompatibility. The BTTEs-Cu(I) catalyst allowed, for the first time, noninvasive imaging of fucosylated glycans during zebrafish early embryogenesis. Robust labeling of the enveloping layer, the embryos' outermost layer of cells, was achieved within 2–3 min. Using this bioorthogonal reaction, we also developed a chemoenzymatic approach for labeling cell surface glycans bearing the LacNAc disaccharide for imaging and glycomic analysis (Zheng *et al.*, 2011).

3. Metabolic Labeling of Cell Surface Fucosylated Glycans in Zebrafish Embryos for Fluorescence Imaging

In the past 14 years, the metabolic approach combined with bioorthogonal chemistry has been successfully used for the detection and imaging of several sectors of the glycome, including mucin-type O-linked glycans (Hang *et al.*, 2003), sialylated (Baskin *et al.*, 2007; Chang *et al.*, 2009; Hong *et al.*, 2010), fucosylated glycans (Hsu *et al.*, 2007; Rabuka *et al.*, 2006; Soriano del Amo *et al.*, 2010), and cytosolic O-GlcNAcylated proteins (Vocadlo *et al.*, 2003).

In this section, we describe the detailed procedures for imaging cell surface fucosylated glycans in zebrafish embryos by microinjection of GDP-6-ethynylfucose (GDP-FucAl) followed by conjugation of azide-bearing fluorophores via the biocompatible CuAAC and imaging via confocal fluorescence microscopy.

L-fucose, the signature monosaccharide possessed by all fucosylated glycans, is a determinant of many functional glycans that play key roles in numerous physiological and pathological processes (Becker and Lowe, 2003; Lu and Stanley, 2006; Ma *et al.*, 2006). Specific terminal glycan fucosylation confers unique properties to cell surface glycoconjugates and is often regulated in cellular differentiation and embryogenesis (Becker and Lowe, 2003). Fucosylated glycans are also critical mediators of cell–cell recognition, neurite outgrowth, and neuronal migration during central nervous system development (Brito *et al.*, 2007). Therefore, fucosylated glycans represent an attractive target for molecular imaging. We use the zebrafish embryo as a vertebrate model system for imaging fucosylated glycans due to its rapid embryonic development, amenability to genetic and embryological manipulations, and its optical clarity.

In nature, fucosylated glycans are synthesized by fucosyltransferases, enzymes that transfer the activated fucose from GDP-fucose (GDP-Fuc) to the acceptor substrates. GDP-Fuc is biosynthesized through two independent pathways: a *de novo* biosynthetic pathway and a salvage pathway (Ma *et al.*, 2006). In the salvage pathway, fucose is first activated by fucokinase to form Fuc-1-Phosphate, which is then converted to GDP-Fuc by GDP-fucose pyrophosphorylase. In vertebrates, the *de novo* biosynthetic pathway usually produces 90% of the total GDP-Fuc and the remaining 10% of GDP-Fuc is contributed by the salvage pathway (Becker and Lowe, 2003). Discovered by Wong and Bertozzi, the salvage pathway of cultured mammalian cells can be hijacked by unnatural fucose analogs, such as 6-azidofucose (FucAz) and 6-ethynylfucose (FucAl) (Rabuka *et al.*, 2006); (Hsu *et al.*, 2007).

To bypass the low yielding salvage pathway, we designed a strategy to inject GDP-FucAl, directly as the metabolic precursor into the yolk sack of one-cell stage embryos (Fig. 21.3A). When GDP-FucAl diffuses into daughter cells, the alkyne-bearing fucose will be incorporated into cell surface fucosylated glycans, allowing a click reaction with a fluorophore-bound azide via the BTES-mediated CuAAC (Soriano del Amo *et al.*, 2010).

3.1. Incorporation of alkyne groups to the cell surface fucosylated glycans in zebrafish embryos

3.1.1. Required materials

Microinjection apparatus: World Precision Instruments (Sarasota, FL) provides the main device (PV 820 Pneumatic PicoPump). The microinjection is done under Nikon SMZ1500 with lens Plan Apo 1 × WD70 (Nikon, Tokyo, Japan).

Microinjection needle preparation materials: Sutter Instrument Co. provides both needle glass and puller. Needle puller is flaming/brown micropipette puller, Model P-9. Needle glass is borosilicate glass with filament, OD: 1.0 mm, ID: 0.5 mm, 10 cm length, fire-polished.

Zebrafish embryos: Any wild-type zebrafish can be used in this study, for example, WIK, AB, and Tüebigen (for detailed information, see <http://zebrafish.org/zirc/fish/lineAll.php>). We recommend using Casper mutant (ZIRC catalog ID: ZL1714), a homozygous double mutant that lacks melanocytes and iridophores due to the mutations in both the genes of *mifta* and *roy*, because of its physical transparency. Fish lines are maintained under the recommended instruction of Zebrafish International Resource Center (ZIRC).

GDP-FucAl injection solution: 20 mM GDP-FucAl in 0.2 M potassium chloride with either Alexa Fluor 594-dextran (5%, w/v) or phenol red loading dye (0.1%, w/v) as tracer. FucAl is chemically synthesized based on the reported procedure (Sawa *et al.*, 2006) and is converted into GDP-FucAl using l-fucokinase/GDP-fucose pyrophosphorylase (FKP), a bifunctional enzyme isolated from *Bacteroides fragilis* 9343 (Wang *et al.*, 2009).

Other reagents

Fish water: 60 mg “Instant Ocean” per liter distilled water.

E3 embryo medium (5 mM NaCl, 0.17 mM KCl, 0.33 mM CaCl₂, 0.33 mM MgSO₄, pH 7.4).

Pronease E (Fisher Scientific) solution: 1 mg/mL prepared with E3 embryo medium. Store at 4° C.

Agarose (Invitrogen).

Others: Fish plastic mating cage set.

35 and 100 mm tissue culture petri dish.

Fire-polished glass Pasteur pipette.

3.1.2. Single-cell zebrafish eggs preparation for microinjection

3.1.2.1. One day prior to microinjection: In the evening, prepare the mating set by inserting the mating cage with mesh bottom inside the fish tank and filling with fish water. Add the divider. Put single pair per cage.

Petri dishes for egg transfer and embryo handling have to be all coated with agarose. To make those agar-coated petri dishes, enough warm agarose (1.2%, v/w, in E3 embryo medium) is poured onto the petri dish to cover the entire surface and immediately poured out, leaving a very thin layer of agarose. Air-dry until agarose is solidified.

Microinjection dishes need to be prepared following the reported protocol (Kemp *et al.*, 2009). Basically, a glass slide is inserted at a 45° angle across the widest part of 100 mm petri dish with three-quarters full of warm agarose (1.2%, v/w, in E3 embryo medium). Let the agarose solidify and remove the slide after complete solidification, leaving a beveled trough.

3.1.2.2. On the day of microinjection: In the morning, after the light is turned on, change the fish water and take out the divider. In this case, when fish spawn, eggs will fall through the mesh of the cage to the floor of fish tank, thus preventing the egg cannibalism by adult fish. Collect the eggs with a pipette and transfer them to a 35 mm agar-coated petri dish. Chorions have to be removed from the eggs before the microinjection by protease digestion. To achieve the complete release of the eggs, remove the water as much as possible and add 1 mg/mL pronease E in E3 embryo medium. Usually this enzymatic digestion lasts for 3–5 min, during which bubbling around chorion can be observed followed by the eggs withdrawn from the chorion. Immediately transfer those dechorionated eggs to a beaker filled with fish water by merging the petri dish into the water and gently sliding the petri dish down. Rinse the eggs with fish water for three times, then transfer the dechorionated eggs using fire-polished pipette to the microinjection dishes filled with E3 embryo medium.

3.1.3. Microinjection—Prepare the sterilized injection needle and load 2 µL GDP-FucAl injection solution. Break the tip of the needle carefully. Adjust the injection pressure and duration and inject each egg with 1 nL of injection solution. Then transfer the eggs back into the 100 mm agar-coated petri dishes filled with E3 embryo medium.

3.1.4. Zebrafish Embryo handling after microinjection—Incubate the eggs at 28 ° C in the incubator under the flow of air. Remove the unfertilized eggs within 5 hours postfertilization (hpf). Unfertilized eggs adopt abnormal cell division, if any; while fertilized eggs undergo asymmetrical divisions during this time period. Transfer the fertilized eggs to a new agar-coated petri dish filled with E3 embryo medium. The fish density should be no more than 60 per one 100 mm petri dish.

3.2. Conjugation of Azide-bound fluorophores by click chemistry

3.2.1. Required materials

Reagents for click reaction

Alexa Fluor-488 azide (Invitrogen): 2.5 mM stock solution in water. Store at 4 ° C in dark.

The CuSO₄-BTES catalyst: [BTES]:[CuSO₄] = 6:1, [CuSO₄] = 2.5 mM, in water. BTES is synthesized and purified based on our published work (Soriano del Amo *et al.*, 2010) and adjusted to pH 7.4 by 1 M NaOH before mixing with CuSO₄ solution. BTES stock should be stored at 4 ° C. Copper sulfate solution can be prepared from dissolving commercially available copper sulfate pentahydrate. Copper sulfate stock solution can be stored at room temperature and last for months. Seal the stock solution bottle with parafilm to avoid water evaporation and check the copper concentration by UV-vis spectrometry if necessary.

Tip: We recommend using fresh prepared catalyst if possible, but premixed CuSO₄-BTES catalyst can last for 1–3 weeks when stored at 4 ° C.

Sodium ascorbate: 100 mM stock in water. Sodium ascorbate is a strong reductant used to generate Cu(I) from Cu(II) *in situ*. In aqueous solutions, sodium ascorbate is easily oxidized so the stock solution has to be freshly prepared every time right before the experiment.

Bathocuproine sulphonate (BCS): 50 mM stock in water. BCS is a biocompatible copper chelator used to quench the CuAAC. It is sensitive to light and the stock solution should be stored at 4 ° C in dark.

Additional reagents: E3 embryo medium

Agarose

Disposable: 96-well plate

Fire-polished glass Pasteur pipette

Glass petri dish

3.2.2. Click chemistry performance—Coat the base of 96-well plate with agarose before transferring the embryos as described in Section 3.1.2. Add 92 µL E3 embryo medium to each well followed by addition of 4 µL Alexa Fluor-488 azide stock solution and 2 µL CuSO₄-BTES catalyst 1:6 complex. Gently shake the plate to mix. Then carefully

transfer the embryos at desired developmental stages (e.g., late gastrula, tissue segmentation, and early larva, etc.) into the well using a fire-polished glass Pasteur pipette. Each well should contain less than five embryos. To initiate the click reaction, add 2.5 μL freshly prepared sodium ascorbate stock and shake gently.

Tip: The final concentration for each reagent in click reaction is: Alexa Fluor-488 azide: 100 μM ; CuSO_4 : 50 μM ; BTES: 300 μM ; sodium ascorbate: 2.5 mM .

After 3 min, add 2 μL BCS (final concentration: 1 mM) to quench the reaction then dilute the reaction system immediately by adding 100 μL E3 embryo medium. Transfer the embryos to a glass petri dish and wash the treated embryos two times with 15 mL E3 embryo medium.

3.3. Imaging

3.3.1. Required apparatus—Confocal fluorescence microscope: Leica SP5 Confocal fluorescent microscope is used in our studies to obtain the images following the manual instruction.

Additional reagents: Ultralow melting point agarose (Invitrogen)

E3 embryo medium

Disposable: MatTek glass bottom microwell dish

3.3.2. Procedure—Prepare ultralow melting point agarose in E3 embryo medium at the concentration of 1.2% (w/v). Place a drop of the agarose solution on a MatTek glass bottom microwell dish. Use a fire-polished glass Pasteur pipette to transfer an embryo into the agarose drop. Position the embryos based on your experiment design (e.g., dorsally or laterally). Set the microwell dish on ice for 5 min to solidify the agarose drop, then add E3 embryo medium gently to the dish until it covers the agarose drop. Place the microwell dish onto the microscope workstation and start to acquire images. Fluorescence and bright field images are acquired sequentially using a 5 μm step interval. Composite figures are prepared using ImageJ (Collins, 2007) (Fig. 21.3B).

4. Labeling Cell Surface Glycans Bearing the LacNAc Disaccharide in Chinese Hamster Ovary(CHO) Cells for Fluorescence Imaging

Although the metabolic glycan labeling strategy we described in Section 3 gives us promising results *in vivo*, there are two downsides to this approach: (1) certain unnatural sugars, such as 6-azidofucose (Rabuka *et al.*, 2006) and Ac_4ManNAz impart different levels of toxicity to live cells and organisms; (2) the metabolic incorporation of a monosaccharide derivative cannot specifically label a oligosaccharide of defined composition due to the prevalence of the monosaccharide in various oligosaccharides on the cell surface.

In this section, we describe an alternative method to label cell surface oligosaccharide glycans in live mammalian cells using a chemoenzymatic strategy instead of hijacking the

glycan biosynthetic pathways with unnatural substrates. The sector of the glycome we target here is glycans bearing *N*-acetylglucosamine (LacNAc, Gal β 1,4GlcNAc). This protocol utilizes a recombinant *Helicobacter pylori* α -(1,3)-fucosyltransferase C168S mutant (α -1,3-FucTM) to transfer a C-6 azide- or alkyne-tagged fucose residue to the 3-OH of *N*-acetylglucosamine of the LacNAc disaccharide. The bioorthogonal tag can then be selectively derivatized with probes via CuAAC or the copper-free click chemistry as described in Section 3. for imaging (Fig. 21.4A).

Since LacNAc-bearing glycans are developmentally regulated and the upregulation of LacNAc is correlated with malignant phenotypes, such as colorectal cancer (Ichikawa *et al.*, 1999), this method for imaging LacNAc may serve as a powerful tool for tracing the changes in cell surface LacNAc-cylation level during the development and has the potential to be transferred to a clinical setting for disease diagnosis.

4.1. Incorporation of alkyne groups into the CHO cell surface glycans bearing the LacNAc disaccharide via a chemoenzymatic approach

We chose a mutant CHO cell line—Lec2 CHO cells as a model system for labeling glycans bearing the LacNAc disaccharide. The major glycan epitope on the surface of Lec2 CHO cells are polyLacNAc on N-linked glycans (North *et al.*, 2010). *H. pylori* α -(1,3)-fucosyltransferase C168S mutant (α -1,3-FucTM) is used to incorporate unnatural fucose analogs to LacNAc residues for the second step click modification. C168S mutation minimizes the dimerization of the parent enzymes while maintaining its activity.

4.1.1. Required materials

Cell line and media: *Lec2 CHO mutant* is established by Patnaik and Stanley (2006). This cell line can grow both in suspension and in monolayer. Incubation condition is 5.0% carbon dioxide and water-saturated at 37 °C.

Growing medium: α -Minimum essential medium (α -MEM) (Invitrogen) supplemented with 10% fetal bovine serum (FBS).

α -1,3-Fucosyltransferase-M (α -1,3-FucMT): α -1,3-FucTM can be generated using our established protocol (Zheng *et al.*, 2011). The enzyme can be store at high concentration (>5 mg/mL) with 10% glycerol at 4 °C and is stable for 3 months. The specific enzymatic activity of α -1,3-FucTM is ~6.0 U/mg protein (one unit is defined as the amount of enzyme that is required to convert 1 μ M of GDP-fucose per min at 37 °C).

Other reagents

GDP-FucAl stock solution: 5 mM, in water. Store at 4 °C (see Section 3.1.1).

GDP-FucAz stock solution: 5 mM, in water. Store at 4 °C. 6-azidofucose (FucAz) is chemically synthesized based on the reported procedure (Zheng *et al.*, 2011) and is converted into GDP-FucAz using FKP by the same method as described for the synthesis of GDP-FucAl in Section 3.1.1.

Fucosylation buffer: HBSS buffer supplemented with 20 mM MgSO₄, 3 mM HEPES and 1% FBS. Store at 4 ° C.

PBS + 1% FBS: Ca²⁺ and Mg²⁺ free supplemented with 1% FBS (v/v).

Store at 4 ° C.

Disposables: 96-well round-bottom plates

Regular and multiple-channel pipette and pipette tips (Rainin Instrument)

Tissue Culture tubes: TC-treated, 16 × 125 mm, screw cap (Corning Corp.)

4.1.2. Enzymatic incorporation of alkyne fucose derivatives on to LacNAc residues of the glycans on CHO cell surface

For flow cytometry: We grow Lec2 CHO cells in 15 mL screw-capped cell culture tube and harvest them when the cell density reaches $\sim 1.0 \times 10^6$ cells/mL. Cultured cells are washed three times with PBS + 1% FBS (200 μ L each time, for three times, centrifuge at $300 \times g$ for 3 min) and resuspended in fucosylation buffer at the concentration of 5.0×10^6 cells/mL. Place 90 μ L cell suspension in a 96-well plate (final density: $\sim 0.45 \times 10^6$ cells per well). Treat the cells with 100 μ M of GDP-FucAl and 30 mU of α -(1,3)FucT-M at 37 ° C for 10 min. After the reaction, wash the treated cells three times with PBS + 1% FBS then resuspend the cells in 100 μ L of PBS + 1% FBS.

For imaging: Seed the Lec2 CHO cells at a density of 3000 cells per well in an eight-well LabTek II chambered cover glass with 100 μ L α -MEM media (10% FBS). Allow the cells to grow for 72 h followed by washing for three times with PBS + 1% FBS (200 μ L each time, for three times, centrifuge at $300 \times g$ for 3 min). Then treat the cells with 200 μ M GDP-FucAz in the presence of 30 mU α -(1,3)FucT-M in fucosylation buffer for 10 min at 37 ° C. After the incubation wash the cells three times with α -MEM medium.

4.2. Conjugation of azide-bound probes by click chemistry

4.2.1. Required materials

Biotin-azide (Click Chemistry Tools): 2.5 mM stock solution in water. Store at -20 ° C.

Streptavidin-Alexa Fluor 488 (Invitrogen): 100 μ g/mL in water. Store at 4 ° C in dark.

Alexa Fluor-488 alkyne (Invitrogen): 10 mM in water. Store at 4 ° C in dark. *CuSO₄-BTES catalyst, Sodium ascorbate and BCS* are all prepared as described in Section 3.2.1.

Hoechst 33342 dye: 1 mg/mL in DMSO. Store at -20 ° C.

α -MEM medium: Store at 4 ° C.

4.2.2. Click chemistry performance

For flow cytometry: After *in situ* fucosylation, add 2.5 μL biotin-azide and 2 μL CuSO_4 -BTTEs catalyst to the Lec2 CHO cells suspension (final concentration: [biotin-azide] = 100 μM , $[\text{CuSO}_4] = 50 \mu\text{M}$, [BTTEs]: $[\text{CuSO}_4] = 6:1$) and ascorbate (final concentration = 2.5 mM). Mix well immediately after sodium ascorbate is added. Wait for 3 min and Quench the click reaction by the addition of 2 μL BCS. Wash the cells three times and resuspend the cells in 100 μL PBS + 1% FBS with streptavidin-Alexa Fluor 488 (final concentration = 1 $\mu\text{g}/\text{mL}$). Incubate in dark at 4 ° C for 30 min. Wash the cells three times with PBS + 1% FBS.

For imaging: The labeling of Lec2 CHO cells for fluorescence imaging is similar to what described above with the replacement of GDP-FucAl with GDP-FucAz, and biotin-azide with an alkyne-bearing fluorophore—Alexa Fluor-488 alkyne (this step can also be accomplished using GDP-FucAl and an azide-bearing fluorophore). In the cell suspension, add Alexa Fluor-488 alkyne (final concentration 1 μM), CuSO_4 -BTTEs catalyst ($[\text{CuSO}_4] = 50 \mu\text{M}$, [BTTEs]: $[\text{CuSO}_4] = 6:1$) and sodium ascorbate (final concentration = 2.5 mM). Mix well. After 3 min, quench the reaction with BCS (final concentration = 1 mM). Wash the cells with the α -MEM medium followed by treatment with Hoechst 33342 dye to stain the nucleus (1:1000 dilution in medium of the stock). Incubate the cells at room temperature in dark for 3 min and wash for three times with α -MEM medium (10% FBS), and image in the presence of 100 μL of the medium.

4.3. Labeling, characterization and imaging

4.3.1. Required materials

Device: Flow cytometry experiments are performed on a Becton Dickinson FACScan analog bench top analyzer using a 488 nm argon laser.

Reagents: *FACS buffer:* HBSS buffer, pH 7.4, 1% bovine serum albumin, 2 $\mu\text{g}/\text{mL}$ 7-AAD, 0.2% NaN_3 . Store at 4 ° C in dark.

4.3.2. Labeling efficiency characterization by flow cytometry—After Lec2 CHO cells are treated as described in Section 4.2.2, they can be resuspended in 400 μL of FACS buffer for flow cytometric analysis. At least 18,000 cells are recorded for each sample. Flow cytometry data were analyzed using Flowjo (TriStar Inc.). Mean fluorescence intensity (MFI) is calculated for live cells. Cell viability is ascertained by gating the sample on the basis of forward scatter (to sort by size) and FL3 (to sort by 7-AAD negative).

4.3.3. Imaging—Image the cells prepared in Section 4.2.2 using a Zeiss Axio Observer. The channels imaged are DAPI (for nucleus staining) and FITC (for membrane staining). Images acquired were processed using Axiovision (Carl Zeiss MicroImaging). Composite figures are prepared using ImageJ (download at: <http://rsbweb.nih.gov/ij/>) and Photoshop CS2 (Adobe) (Fig. 21.4B).

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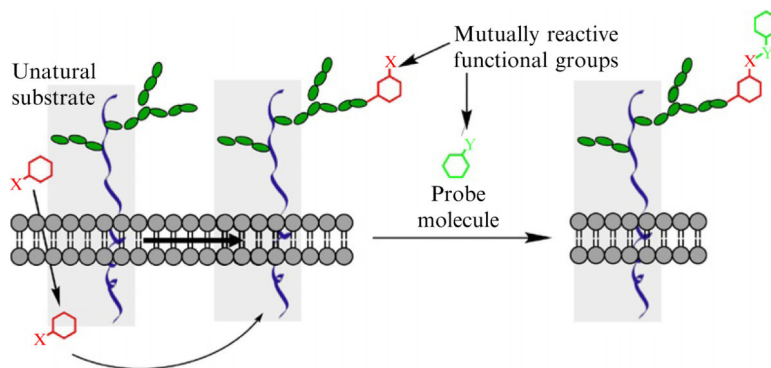


Figure 21.1.

A schematic description of metabolic labeling of cell surface glycans for fluorescence imaging. Red: an unnatural monosaccharide building block functionalized with a bioorthogonal chemical tag X; Green: a biophysical probe functionalized with a complementary chemical group Y. The covalent reaction between X and Y enables visualization or enrichment of the target glycoproteins for molecular identification.

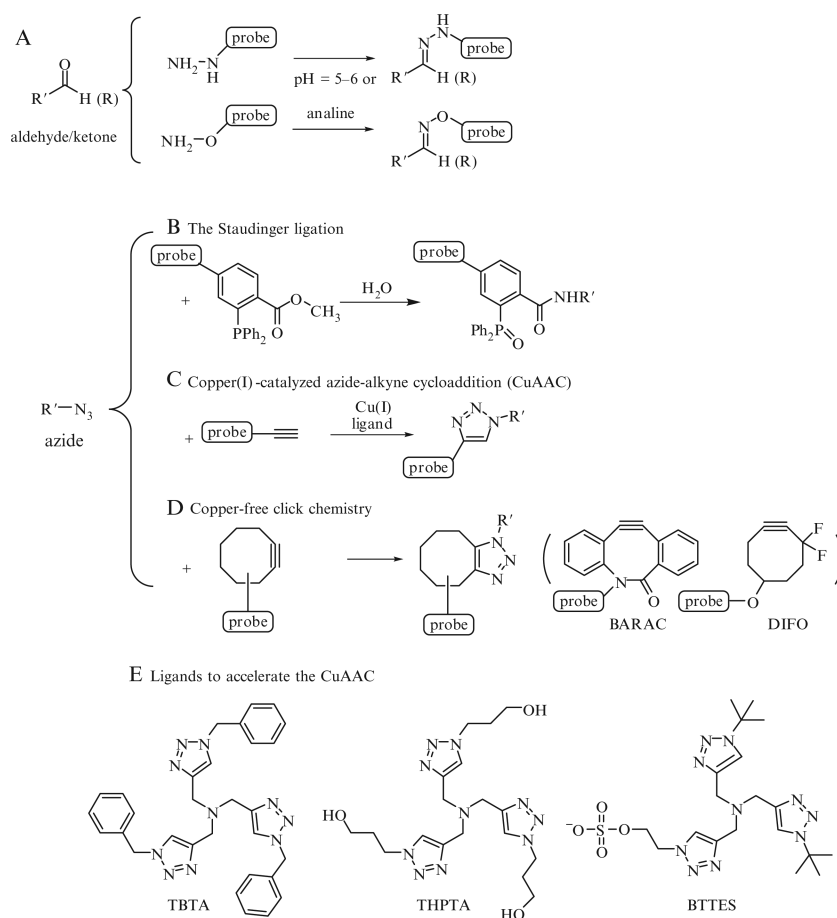


Figure 21.2. Most frequently used bioorthogonal click chemistry for imaging studies. (A) The condensation reaction of ketones and aldehyde with aminoxy and hydrazide-bearing reagents to form stable hydrazones and oximes adducts, respectively; (B) The Staudinger ligation; (C) Cu(I)-catalyzed azide-alkyne cycloaddition (CuAAC); (D) the strain-promoted cycloaddition of azides and cyclooctynes; (E) representative examples of Cu(I)-stabilizing ligands to accelerate the CuAAC. TBTA=tris[(1-benzyl-1H-1,2,3-triazol-4-yl)methyl]amine, THPTA=Tris-(hydroxypropyltriazolylmethyl)amine, BTES=2-(4-((bis((1-tert-butyl-1H-1,2,3-triazol-4-yl)methyl)amino)methyl)-1H-1,2,3-triazol-1-yl) ethyl hydrogen sulfate.

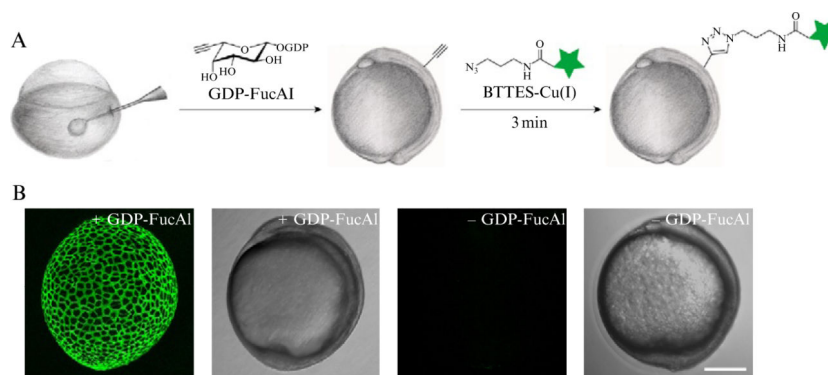


Figure 21.3.

A schematic description of metabolic labeling of cell surface fucosylated glycans in zebrafish embryos for fluorescence imaging. (A) workflow of metabolic labeling of fucosylated glycans followed by conjugation with fluorophores via CuAAC for imaging; (B) representative images of zebrafish embryos treated by this process: *from left to right*, fluorescence image of 10 hpf embryos treated with GDP-FucAl followed by a click reaction with Alexa Fluor 488-azide; the corresponding bright field image of 10 hpf zebrafish embryos treated with GDP-FucAl followed by a click reaction with Alexa Fluor 488-azide; fluorescence image of 10 hpf embryos treated without GDP-Fuc; the corresponding bright field image of 10 hpf embryos treated without GDP-Fuc followed by a click reaction with Alexa Fluor 488-azide.

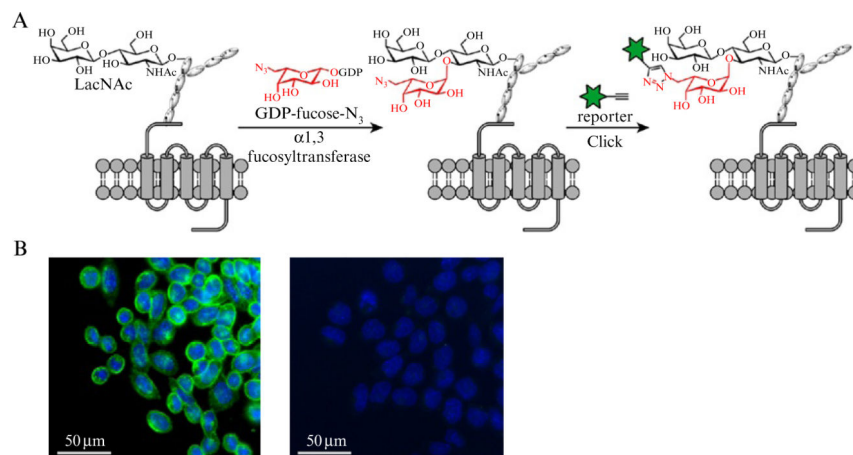


Figure 21.4.

A schematic description of chemoenzymatic labeling cell surface glycans bearing LacNAc in Lec2 CHO cells for fluorescence imaging. (A) workflow of the chemoenzymatic labeling LacNAc by GDP-FucAz followed by conjugation of fluorophores for imaging; (B) Fluorescence images Lec2 CHO cells labeled by this method: *left*: fluorescence image of Lec2 CHO cells treated with GDP-FucAz followed by a click reaction with Alexa Fluor 488-alkyne; *right*: fluorescence image of Lec2 CHO cells treated with GDP-Fuc followed by a click reaction with Alexa Fluor 488-alkyne. Green: cell surface labeled by Alexa Fluor 488; Blue: nucleus stained by Hoechst 33342 dye.