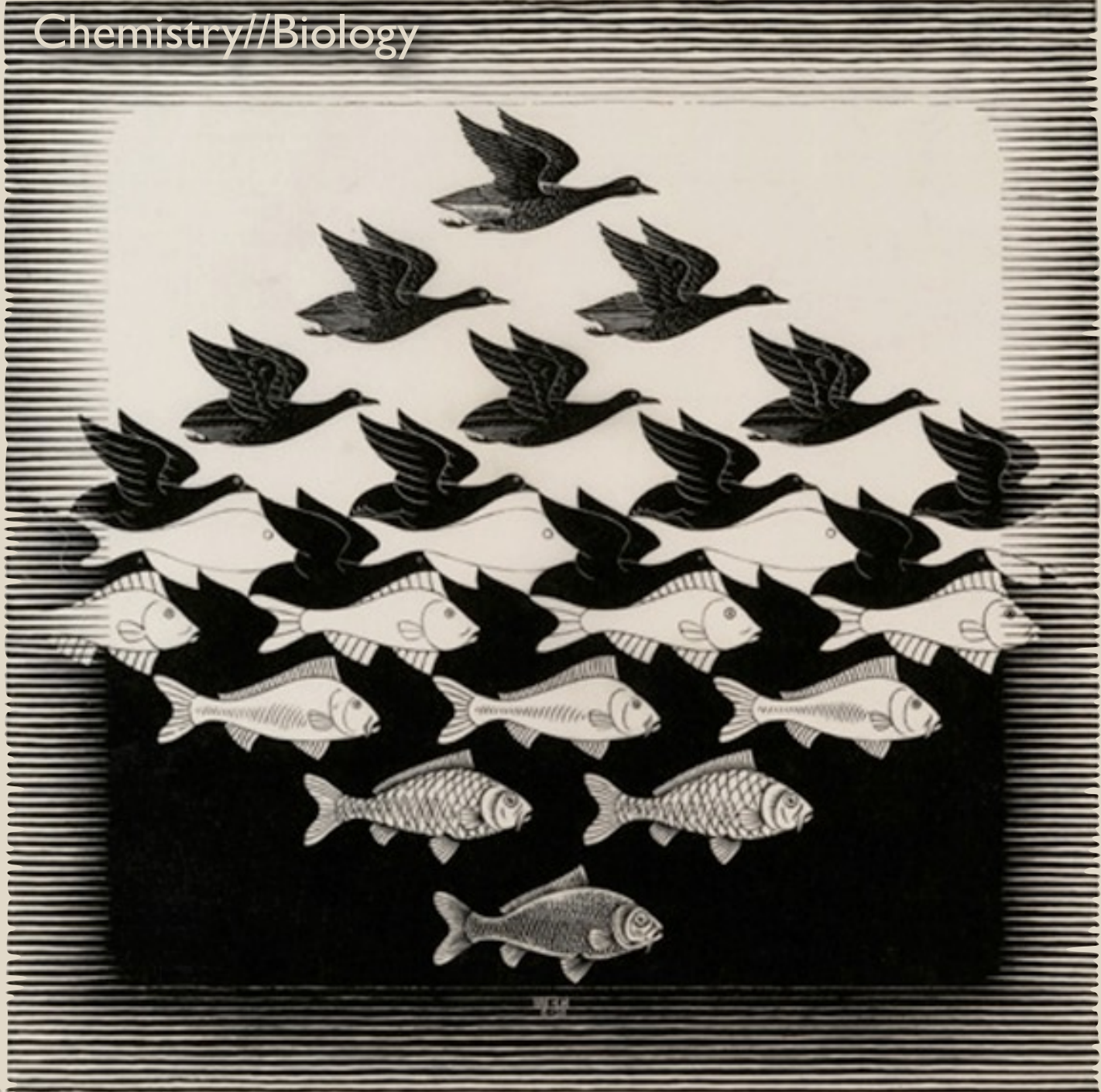


Activation  
et étiquetage  
des protéines  
et  
biomolécules

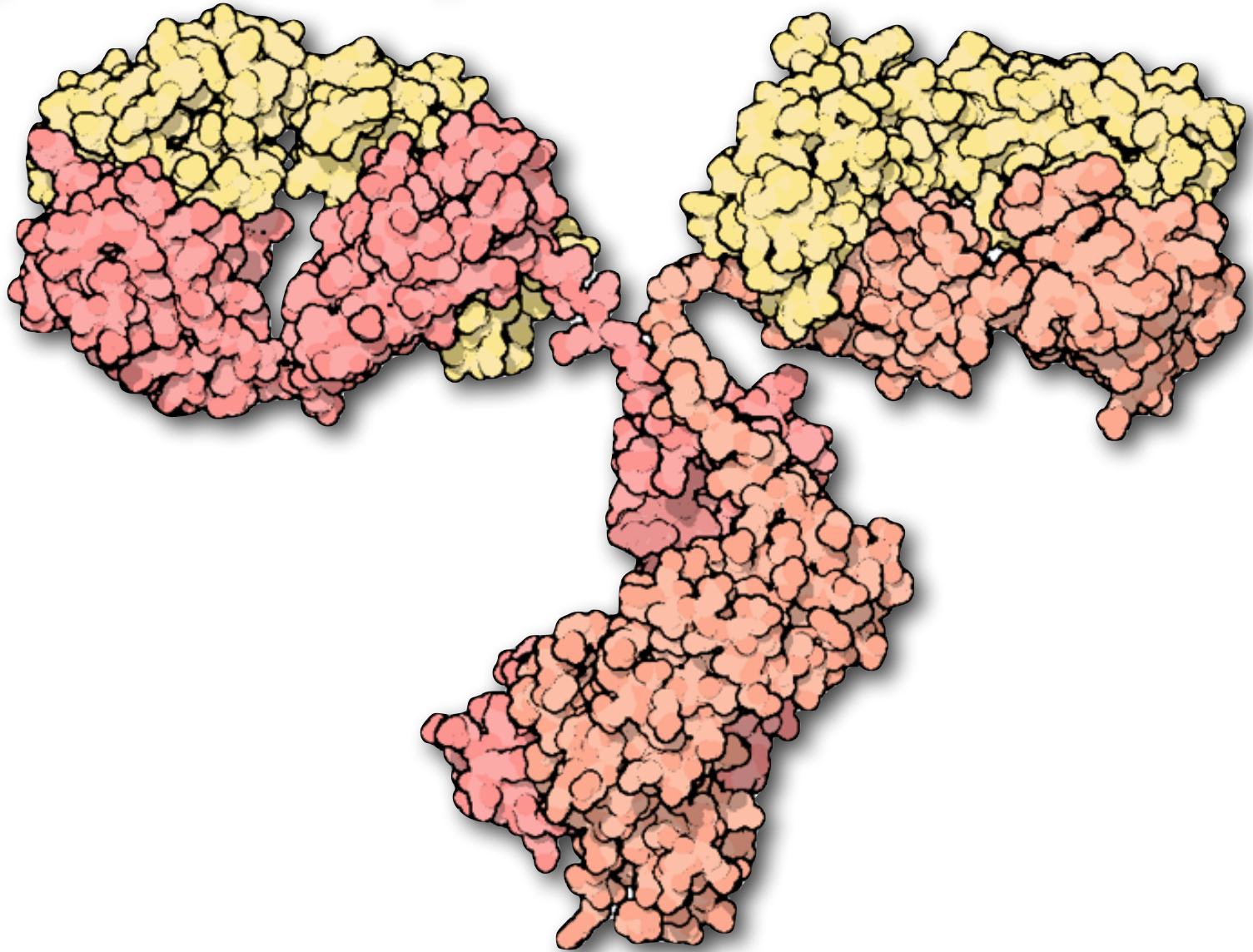


M. C. Escher  
Sky and Water I  
1938 2

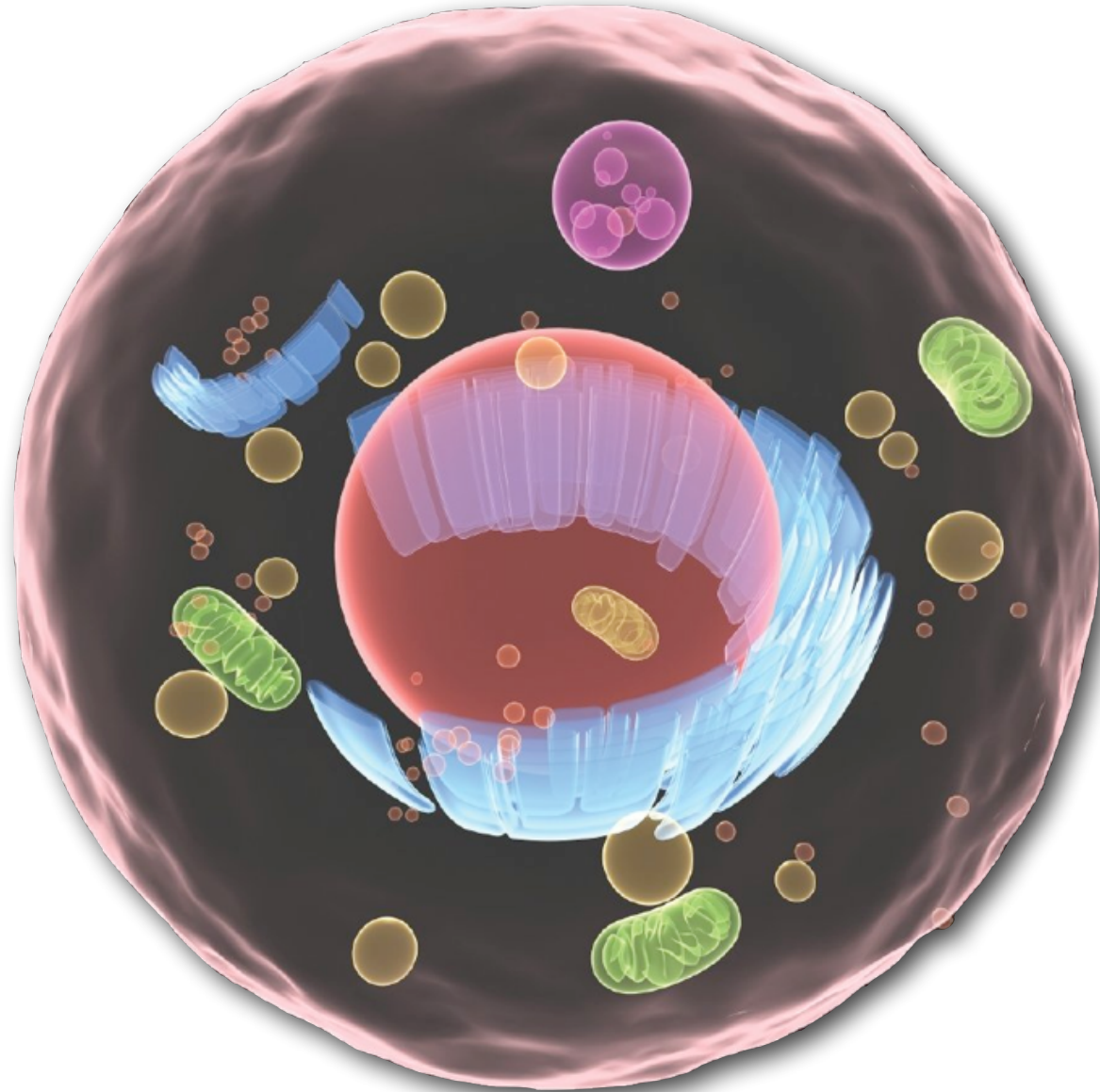
# Synthetic chemistry in a flask



# Synthetic chemistry on biomolecules



# Synthetic chemistry in a cell



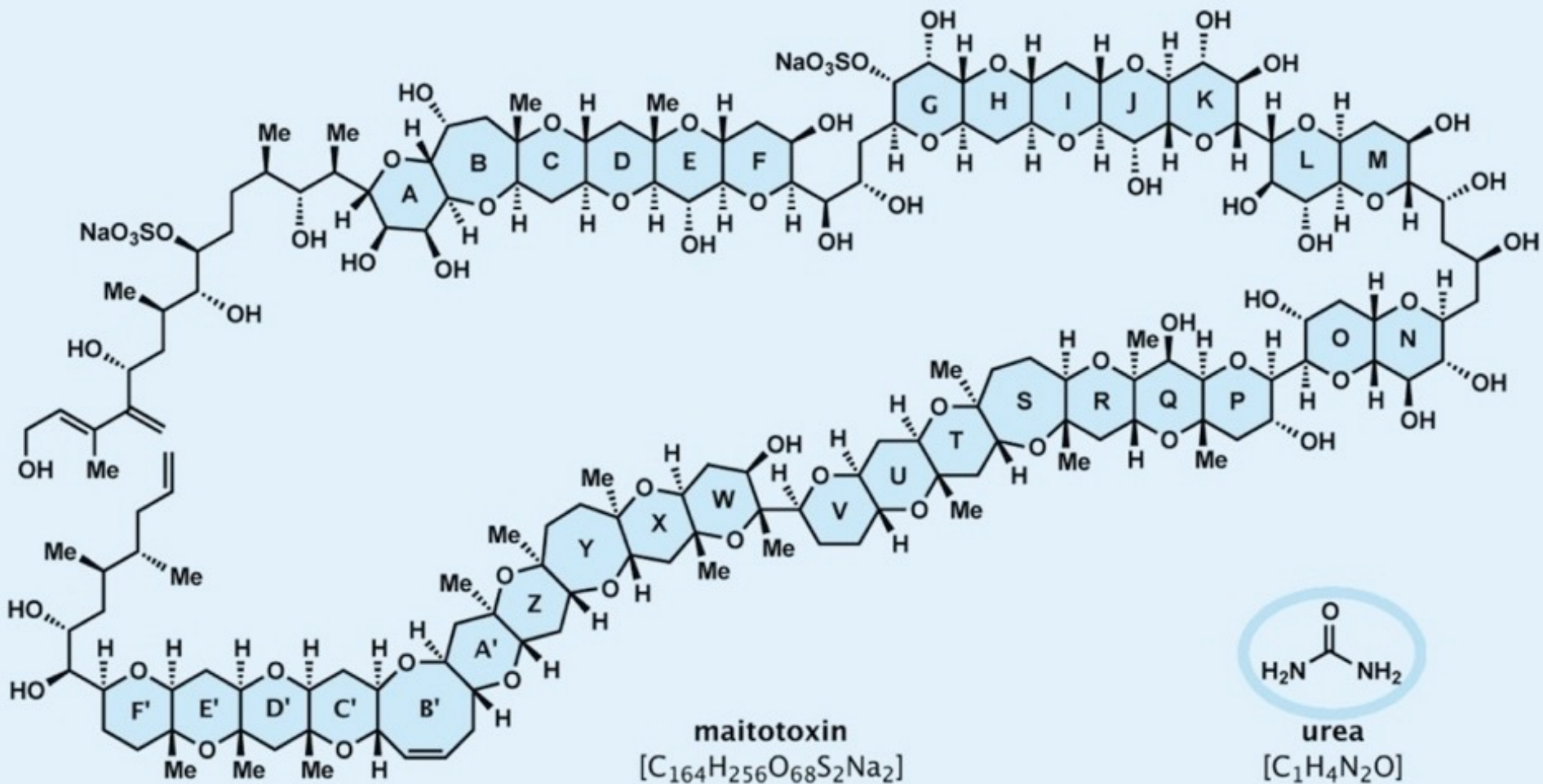
# Synthetic chemistry in living organisms



Bio

conjugation

# Natural products synthesis

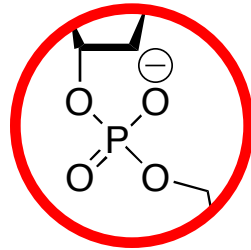




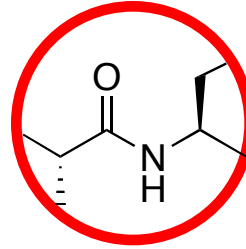
# Macromolécules biologiques



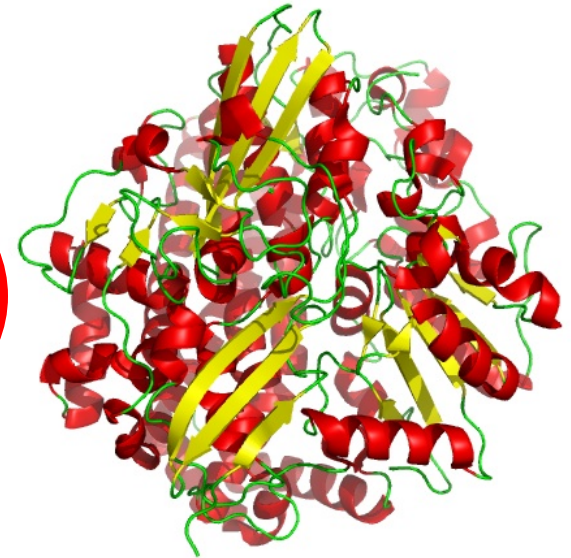
Acide nucléique



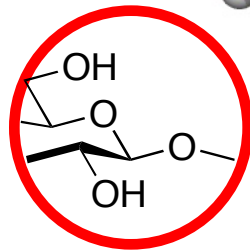
Phosphodiester



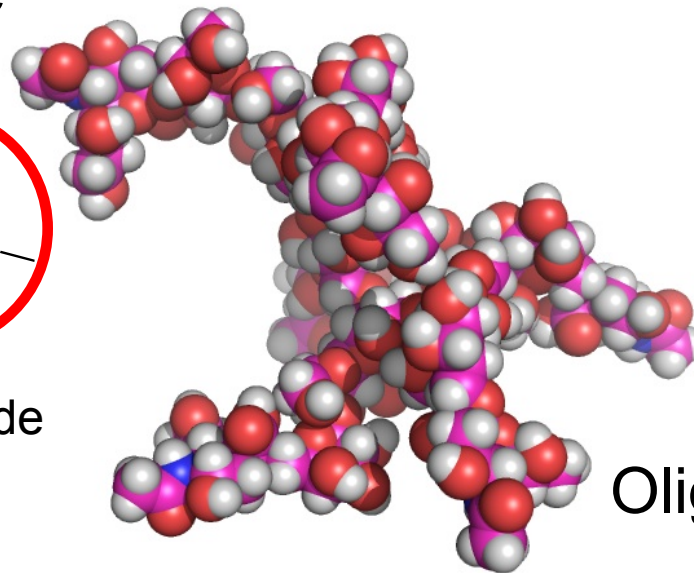
Amide



Protéine

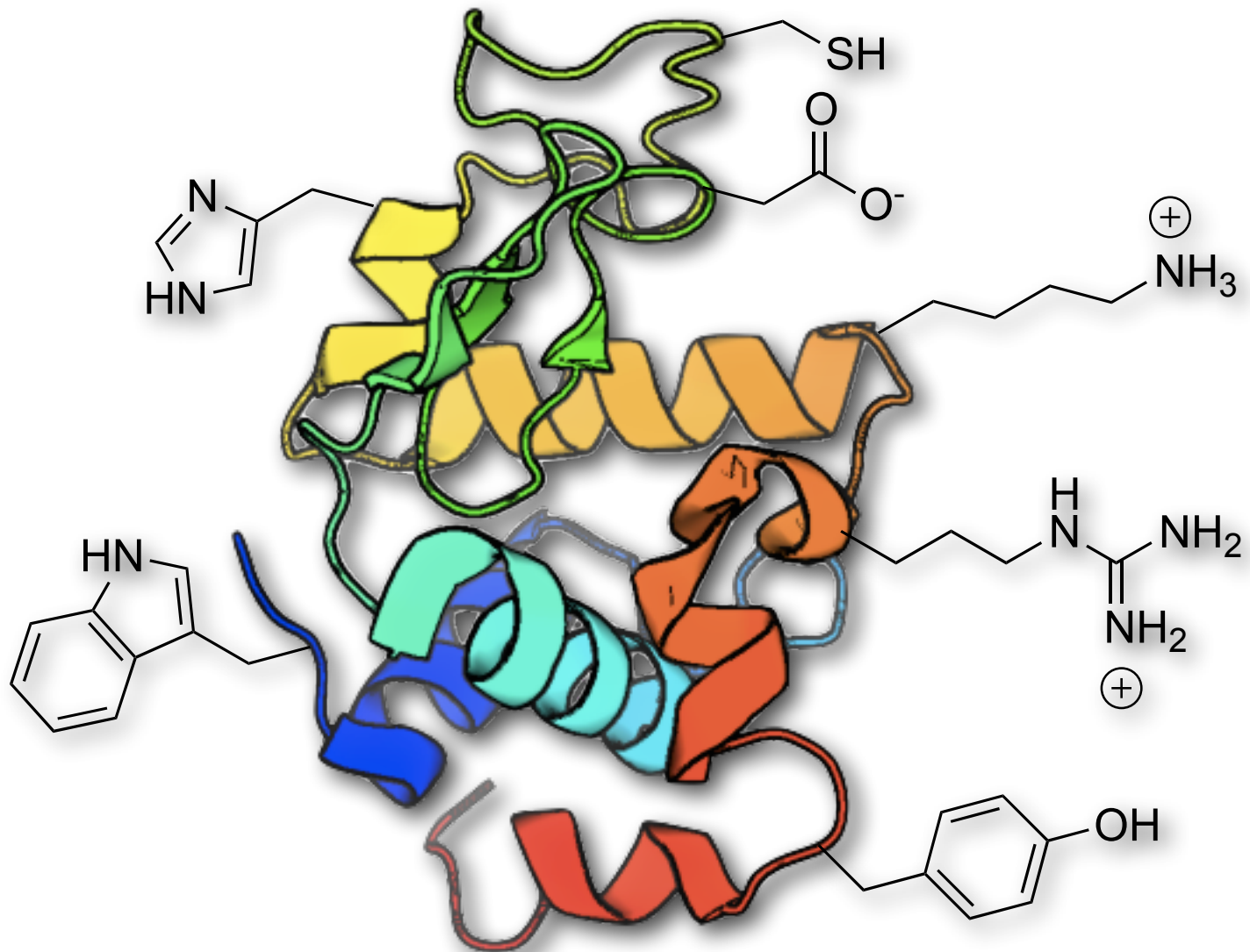


Glycoside



Oligosaccharide

# Bioconjugation

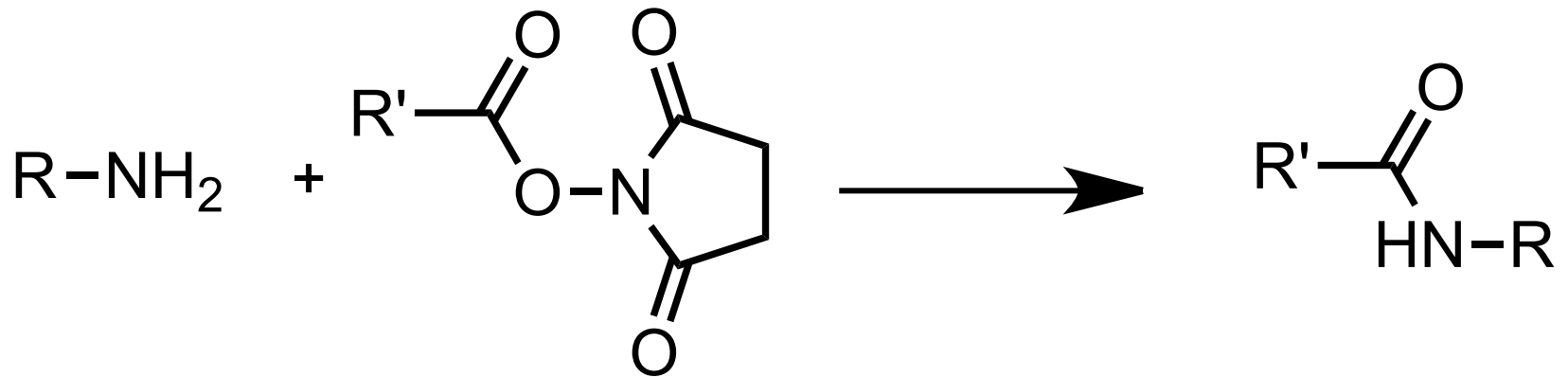


# Relevant chemical transformations ?

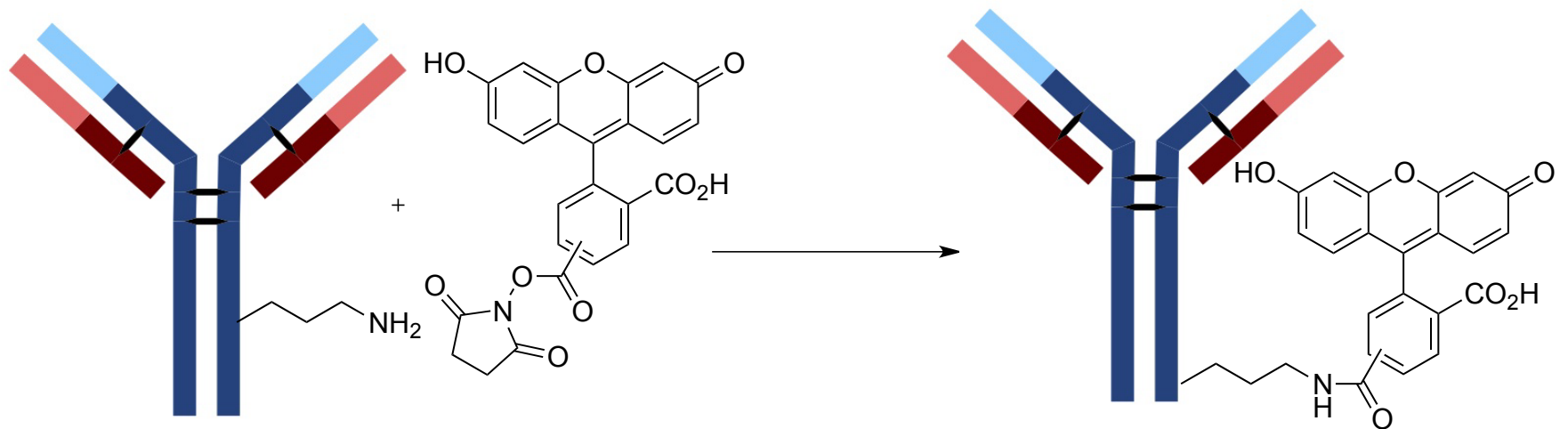
## Chemical Transformations Leading to Protein Covalent Modifications

- Water Is the Sole Solvent
- A Neutral pH Is Required
- Ambient Temperature (Up to 40°C)
- Kinetics, which Adapted to the Observed Phenomenon (on the Hour Scale)
- Low Reactant Concentrations
- Nontoxic Reagents

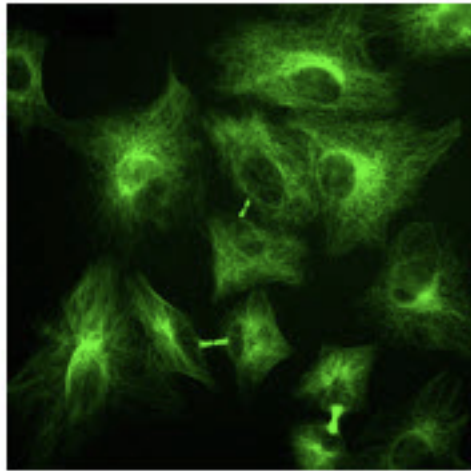
# Bioconjugation - Aminolysis



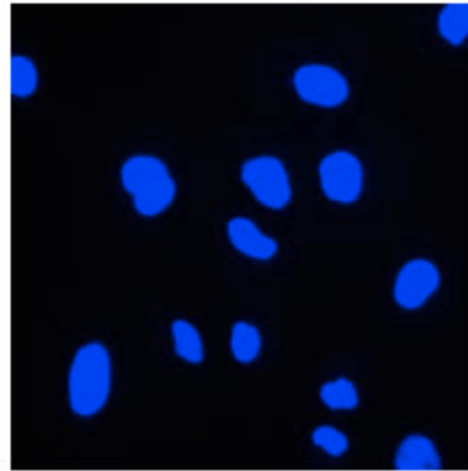
# Antibody labeling



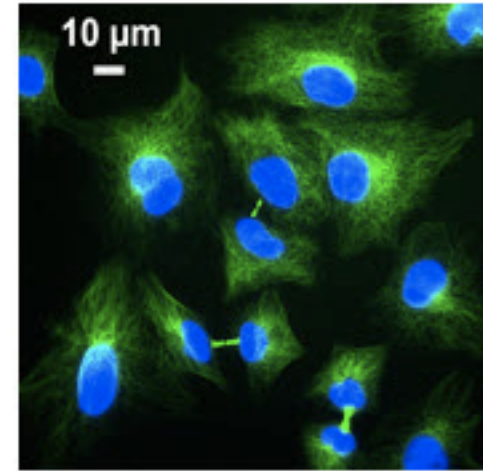
# Tubulin imaging



A. Fluorescein



B. Hoechst

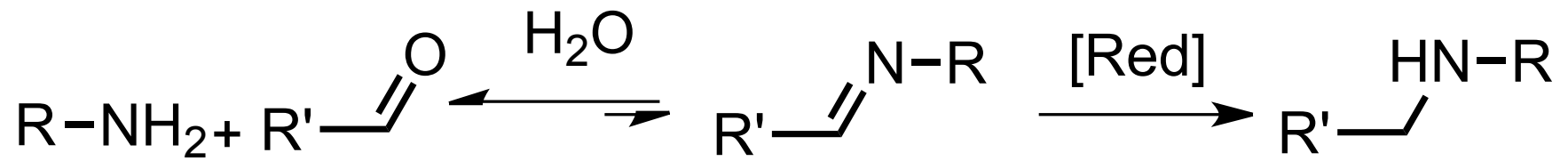


C. Merged

## Detection of $\alpha$ -tubulin in A549 cells demonstrates use of fluorescein-labeled secondary antibody

Cells were grown in 96-well microplates for 18-20 hrs, fixed with 4% paraformaldehyde (Part No. 28906) and permeabilized with 0.1% Surfact-Amps X-100 (Part No. 28314). Cells were then probed with a mouse anti- $\alpha$ -tubulin primary antibody (0.4 $\mu$ g/mL) and Fluorescein-goat anti-mouse secondary antibody (2 $\mu$ g/mL). Nuclei were labeled with Hoechst Dye. Images were acquired by fluorescence microscopy. **A.** Fluorescence image shows a delicate network of  $\alpha$ -tubulin (pseudo-colored green) located exclusively in the cytoplasm. **B.** Nuclear counterstain with Hoechst Dye (pseudo-colored blue) **C.** Merged image.

# Bioconjugation - reductive alkylation



# Vascular Lumen

Endothelial Glycocalyx

Endothelial Cell

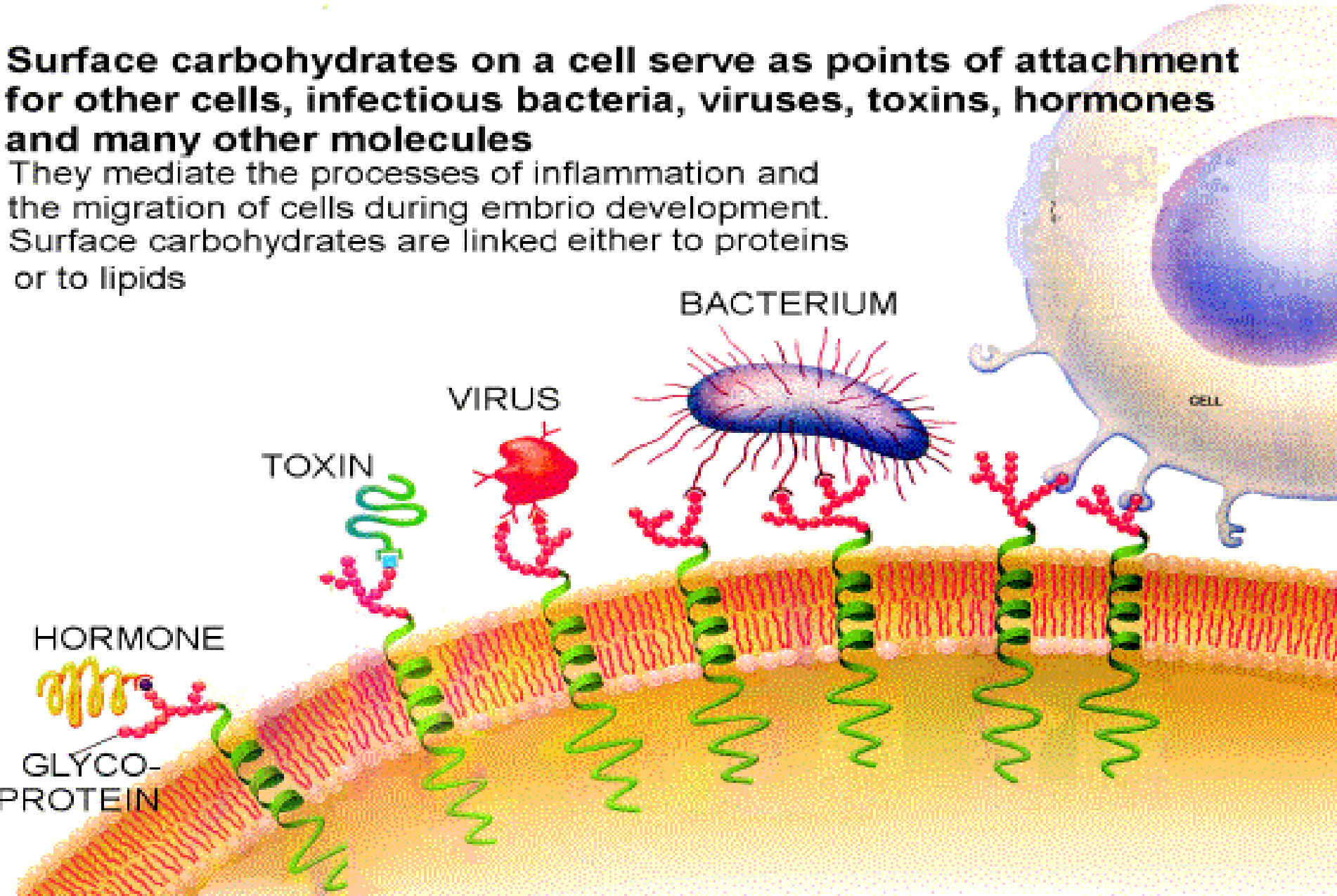
0.1  $\mu\text{m}$



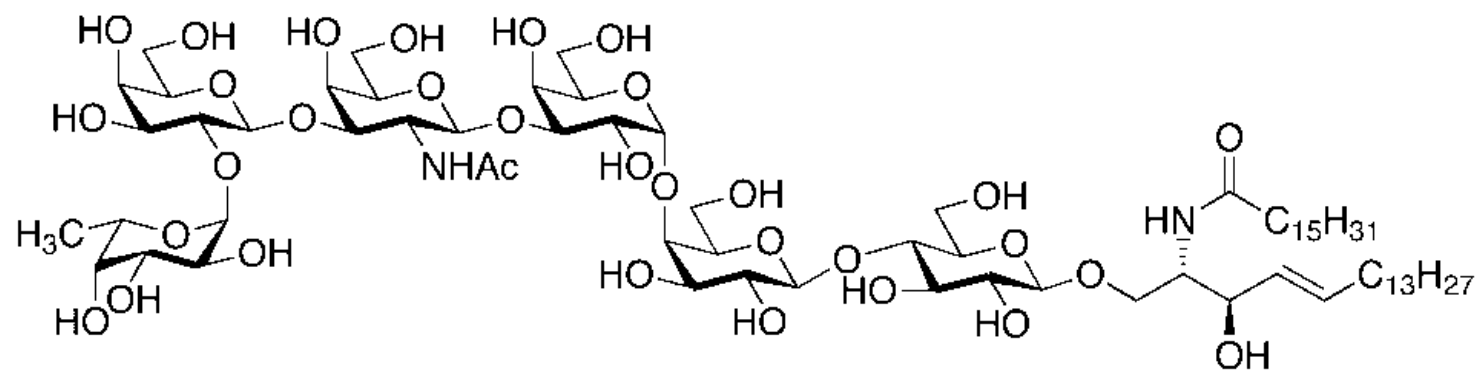


**Surface carbohydrates on a cell serve as points of attachment for other cells, infectious bacteria, viruses, toxins, hormones and many other molecules**

They mediate the processes of inflammation and the migration of cells during embryo development. Surface carbohydrates are linked either to proteins or to lipids

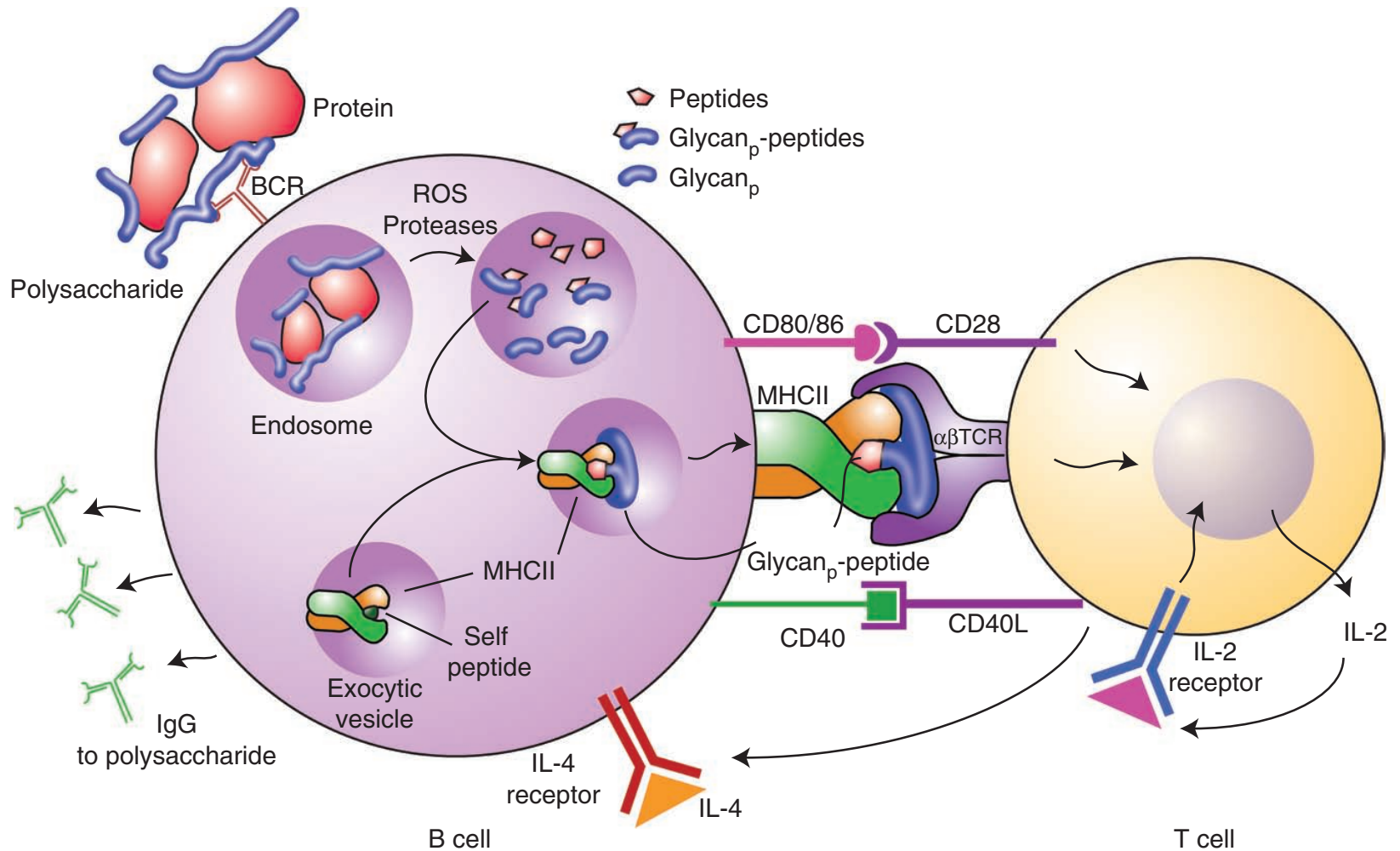


# Tumor associated antigens

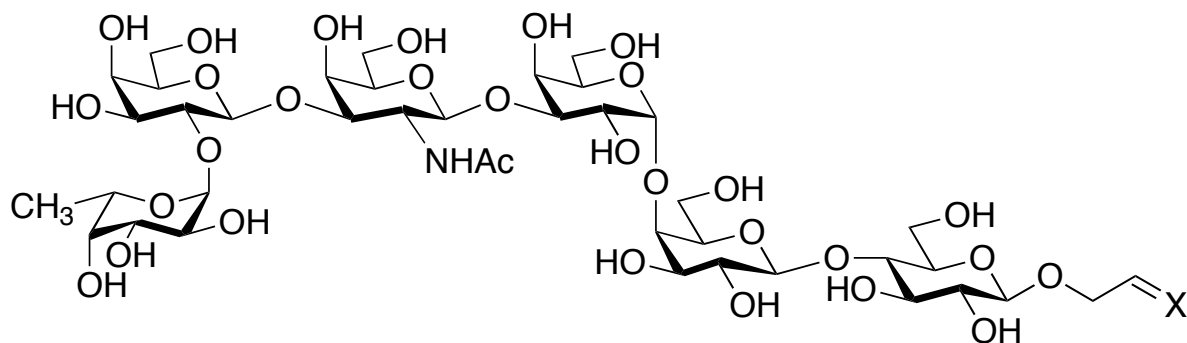


**1:** MBr1 antigen / Globo-H

# Glycoconjugate vaccines

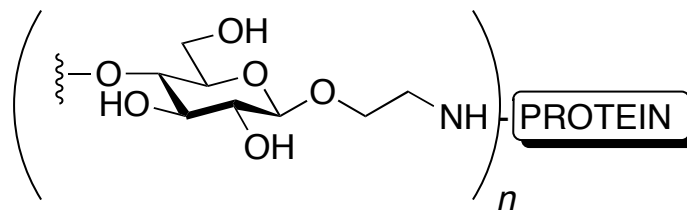


# Anticancer vaccines ?



O<sub>3</sub>; Me<sub>2</sub>S ( **40**: X = CH<sub>2</sub>  
**41**: X = O

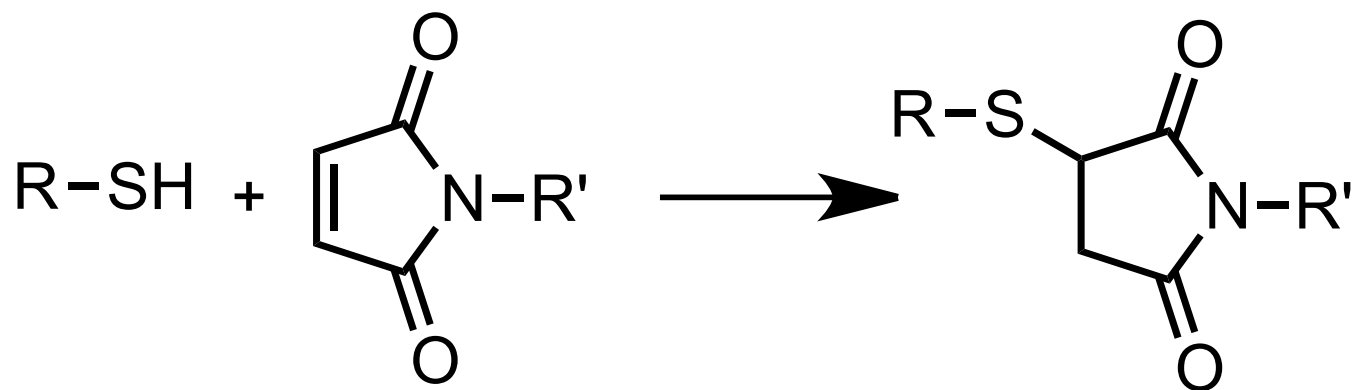
conjugate to protein  
via reductive amination



**42**: KLH conjugate  
**43**: BSA conjugate

Scheme 8. Synthesis of the Globo-H–KLH vaccine construct.

# Bioconjugation - Michael addition





Paul Ehrlich

César Milstein and George Köhler

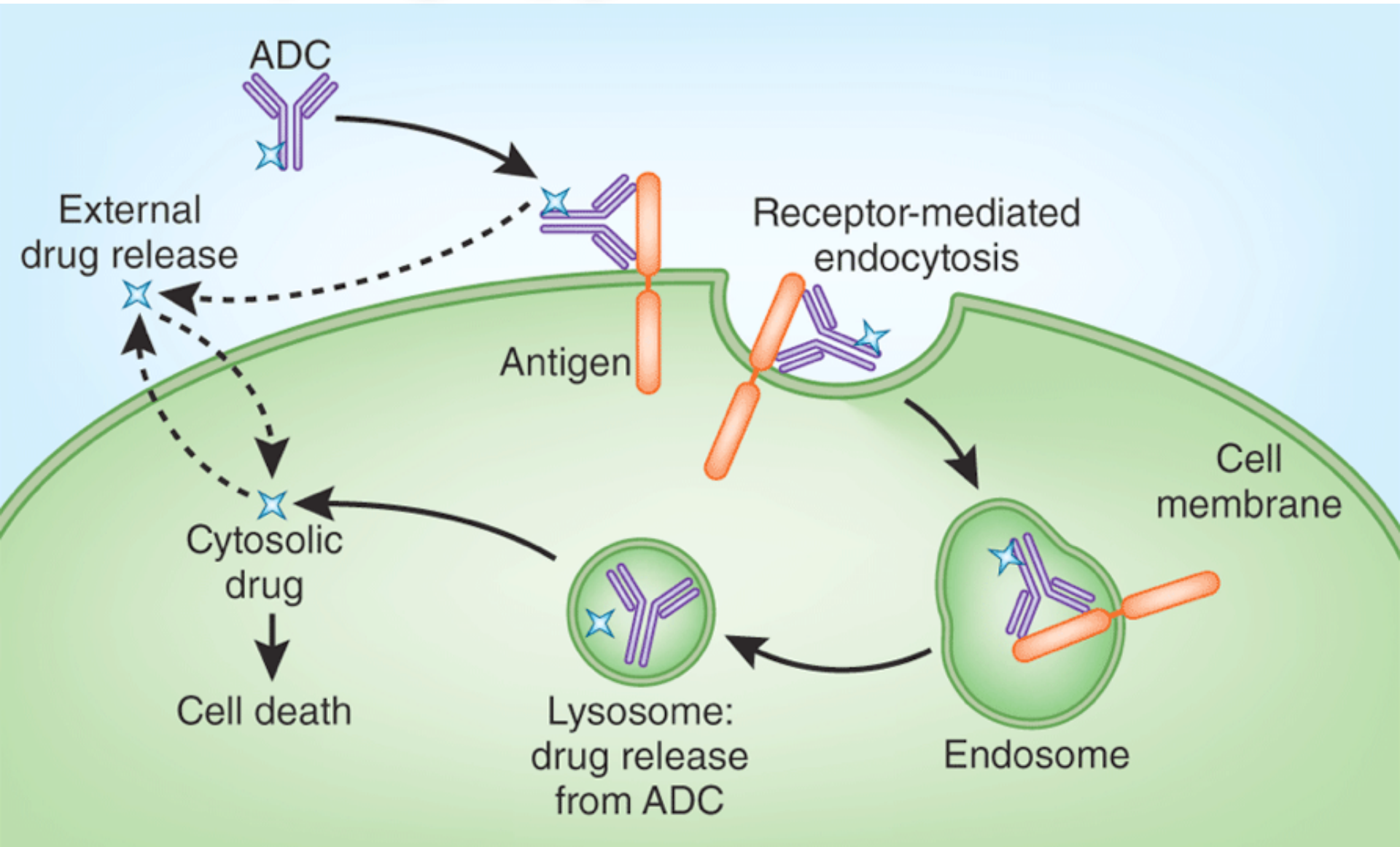


*Dolabela auricularia*

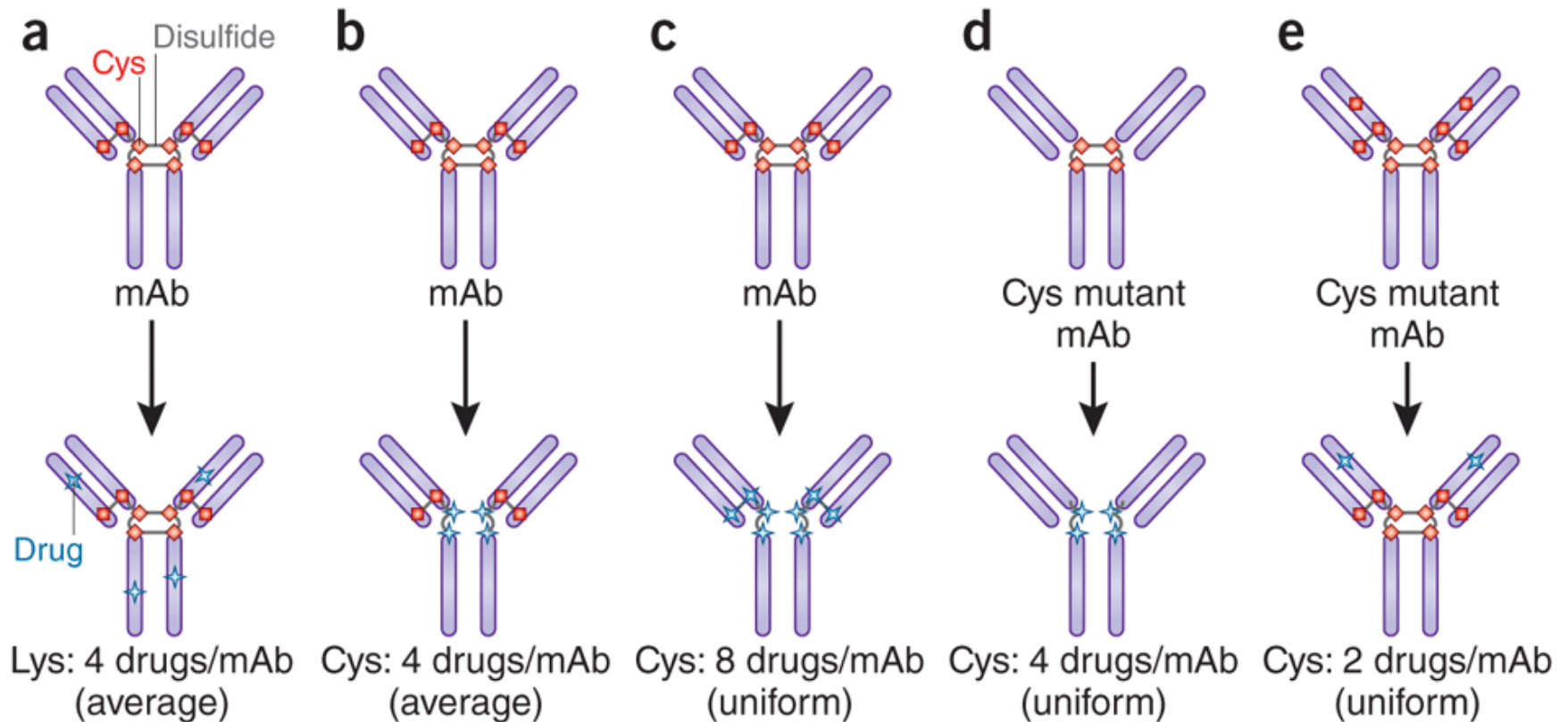


George Pettit

# Antibody-Drug conjugates

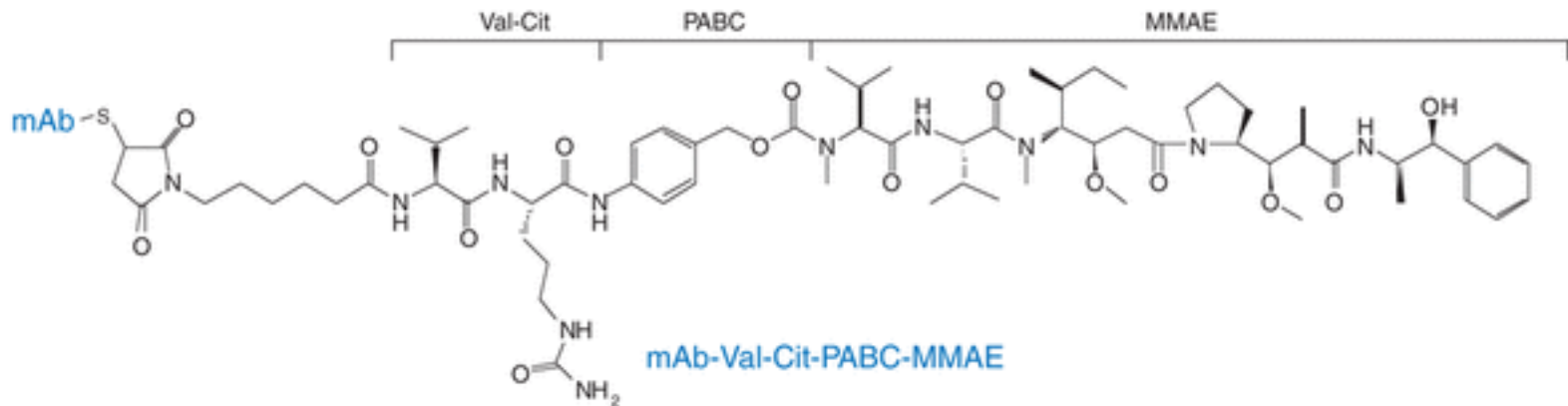
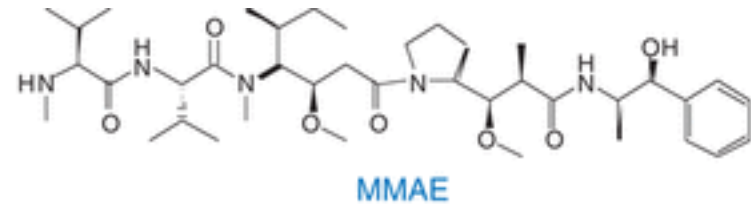
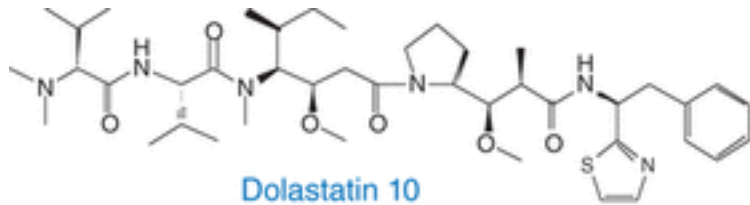


# Conjugation technology

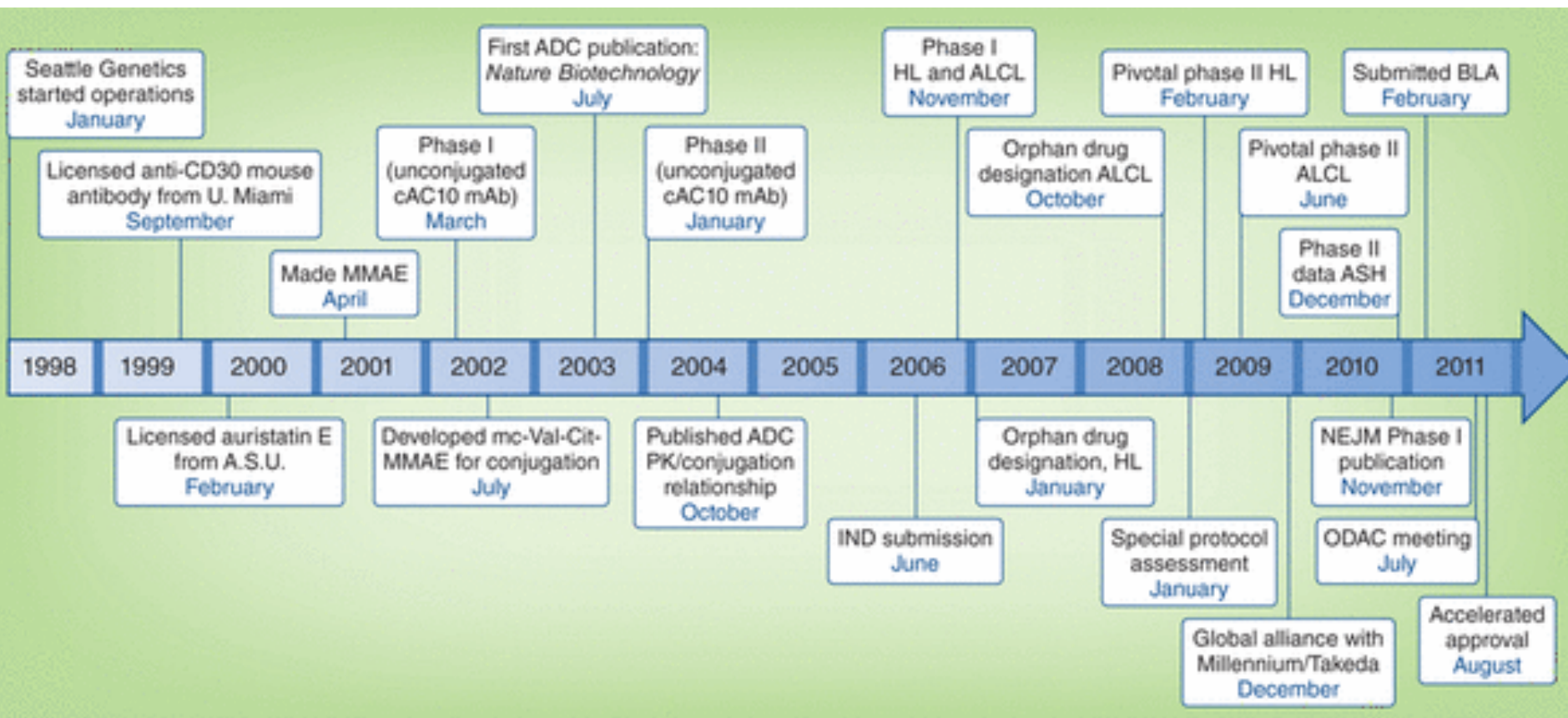




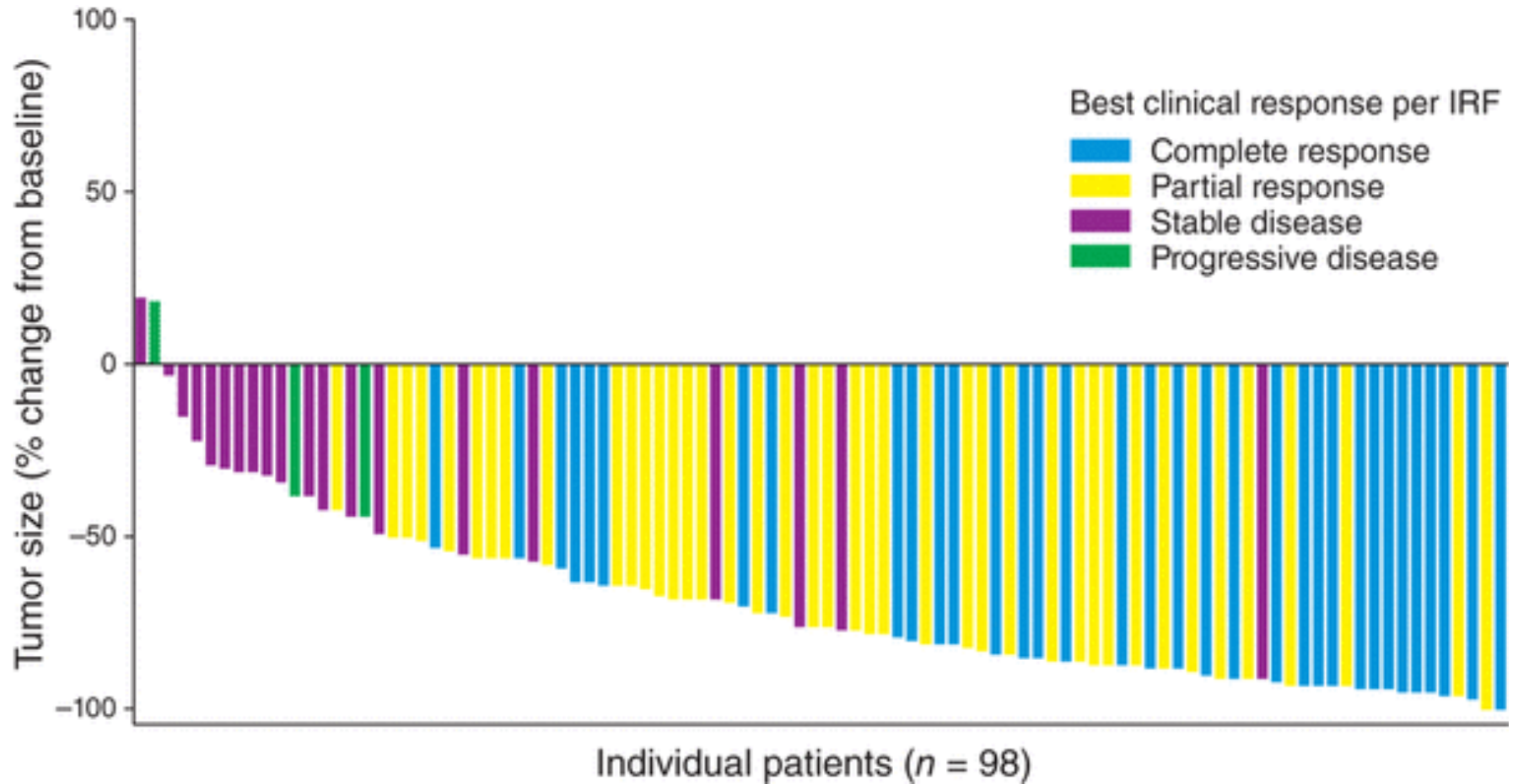
# Choosing the linker



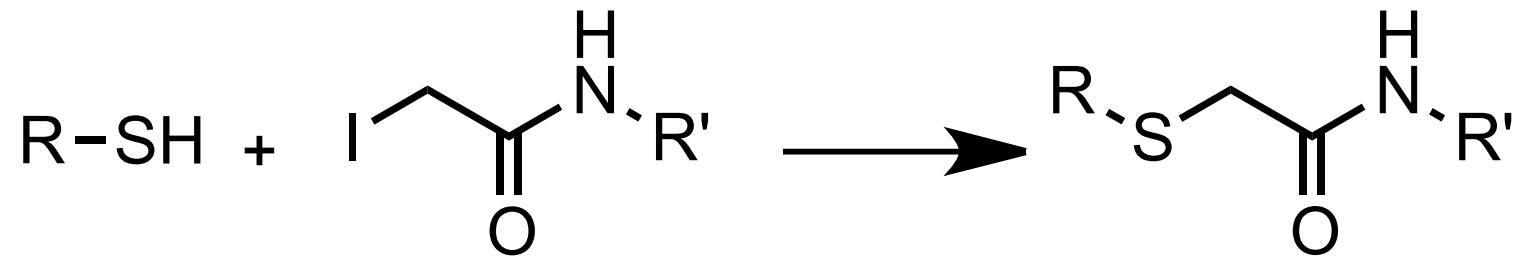
# Approval



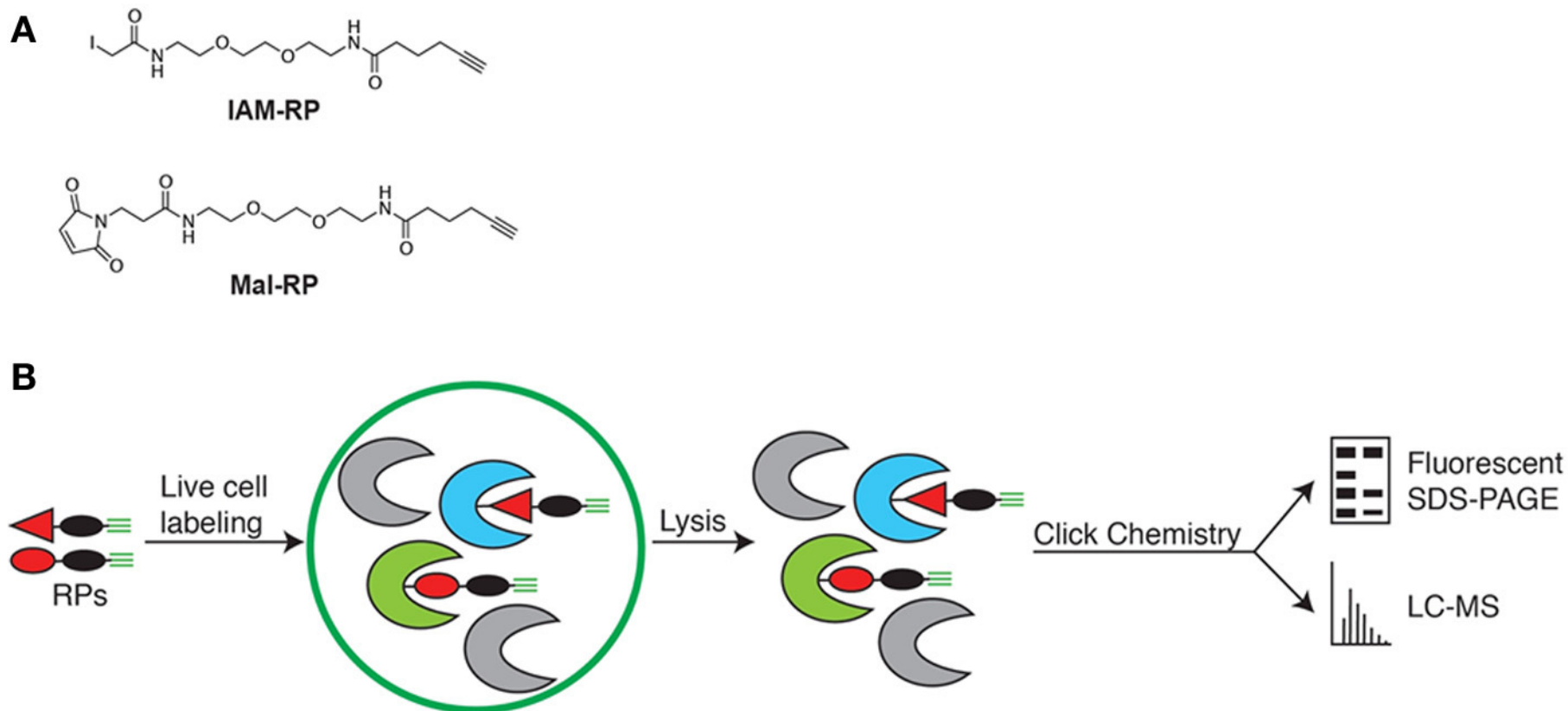
# Phase II clinical trials...



# Bioconjugation - Alkylation



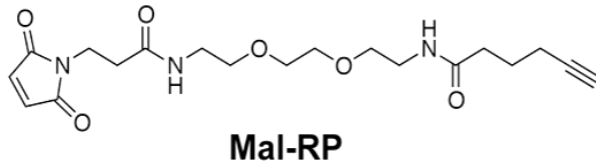
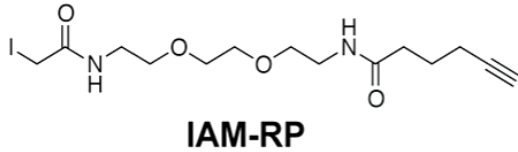
# Probing cysteine oxydation



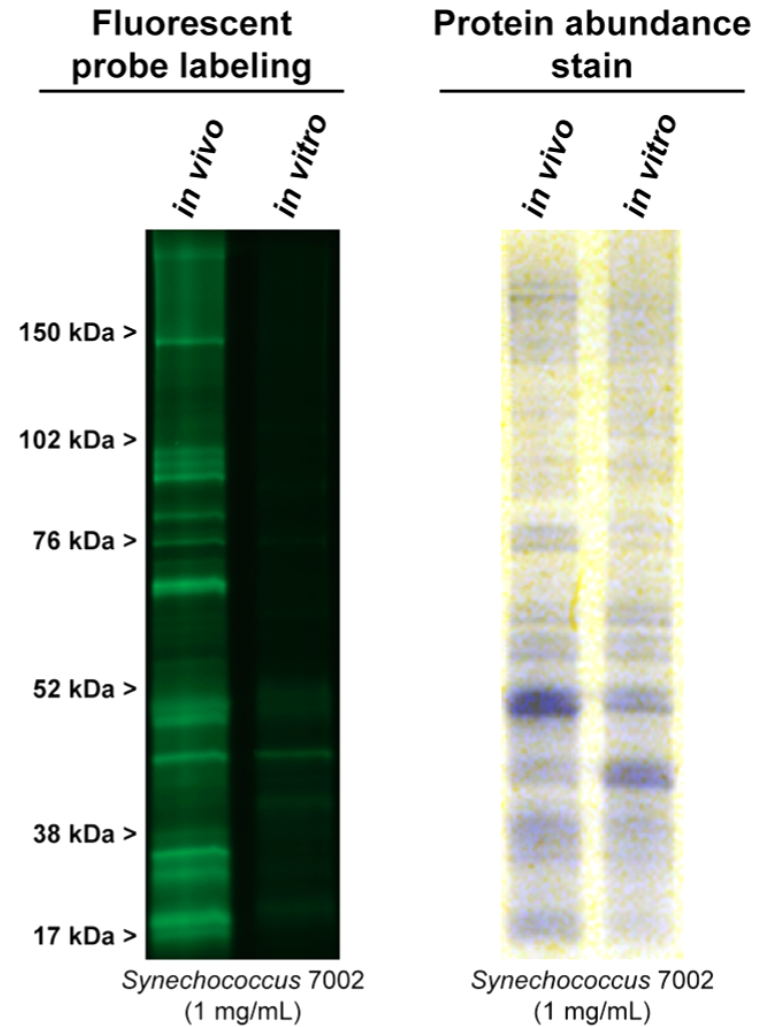
**FIGURE 1 | (A)** Chemical probes for *in vivo* labeling of reduced cysteine thiols. IAM-RP and Mal-RP. **(B)** Schematic of *in vivo* chemical proteomics approach for analysis of redox-sensitive *Synechococcus* 7002 proteins.

# Thiol reactive probes

**a**

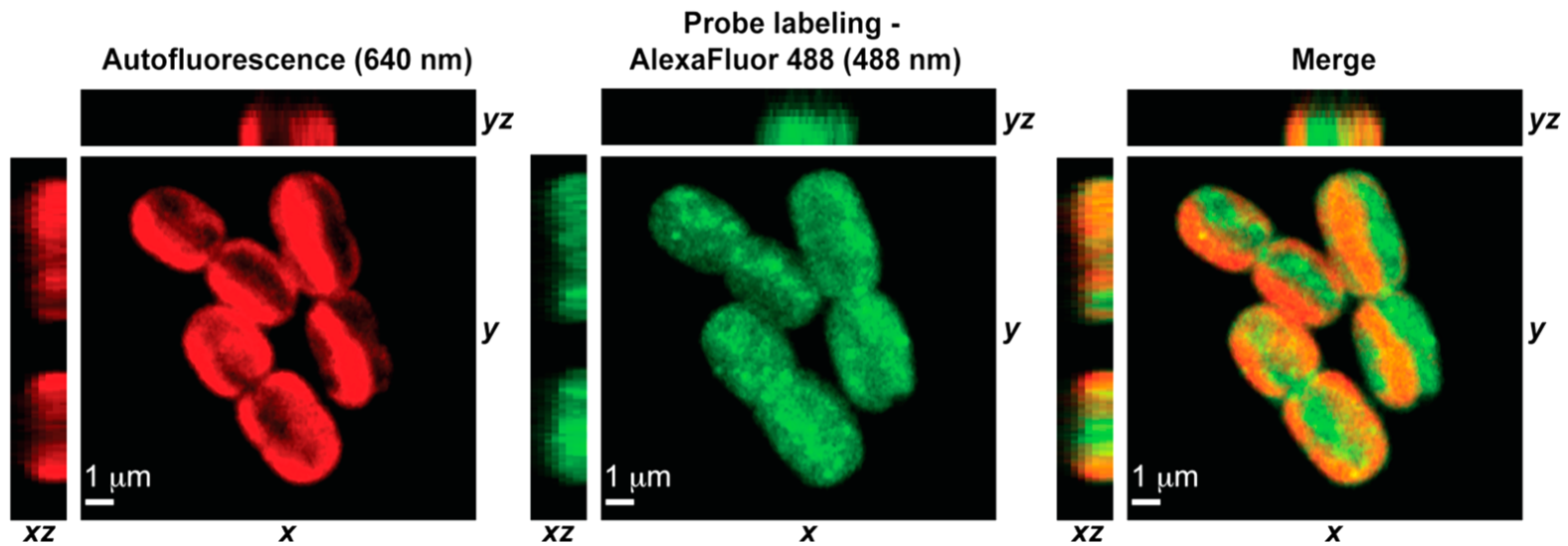


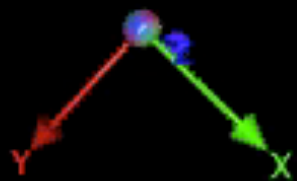
**b**



# Labeling thiols *in vivo*

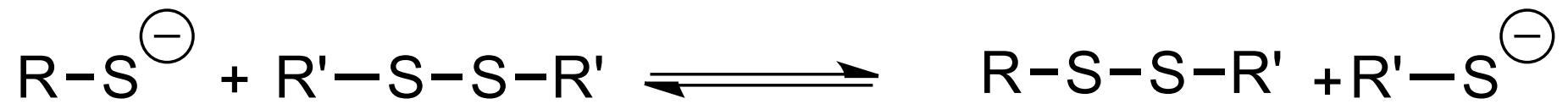
**C**



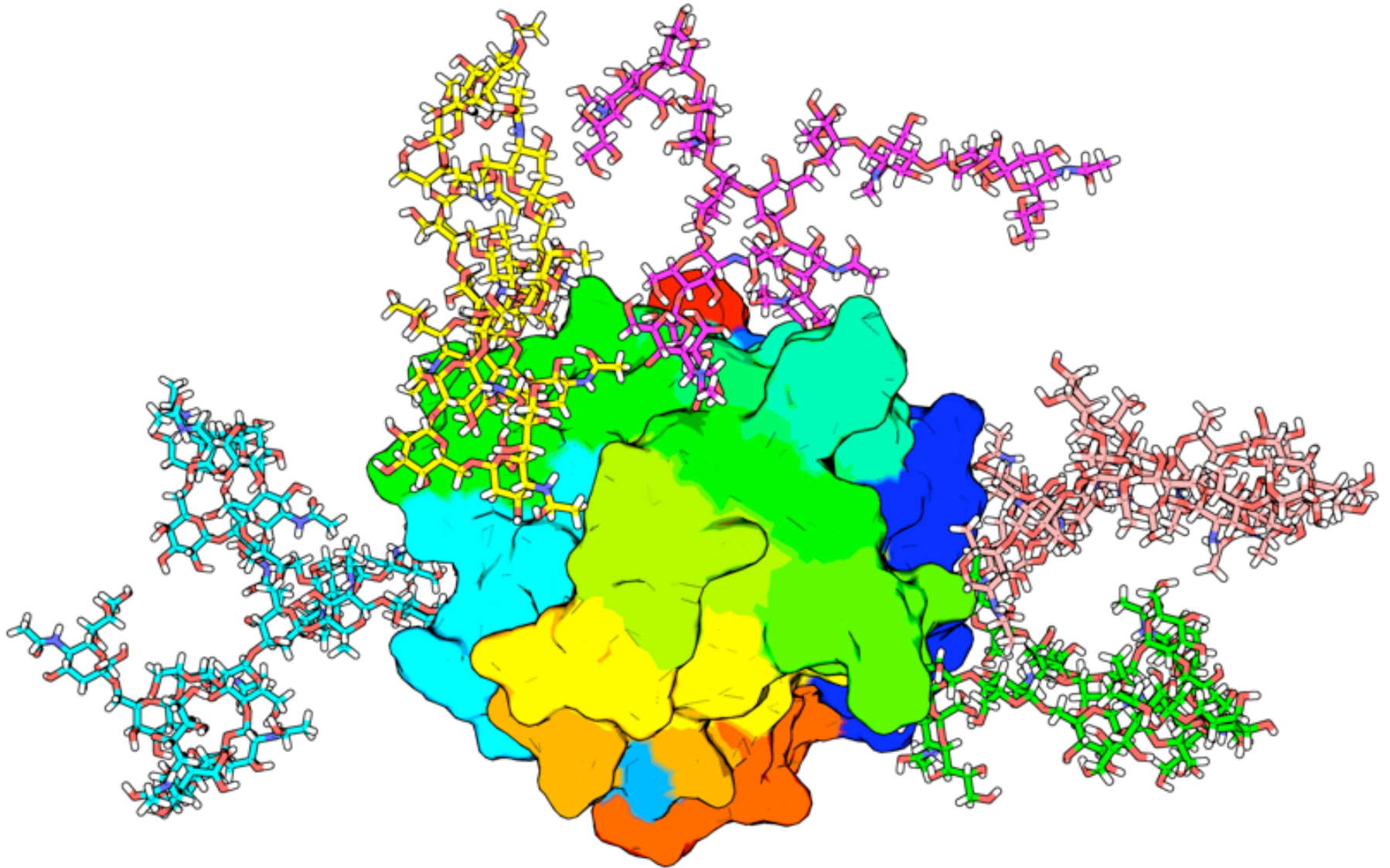




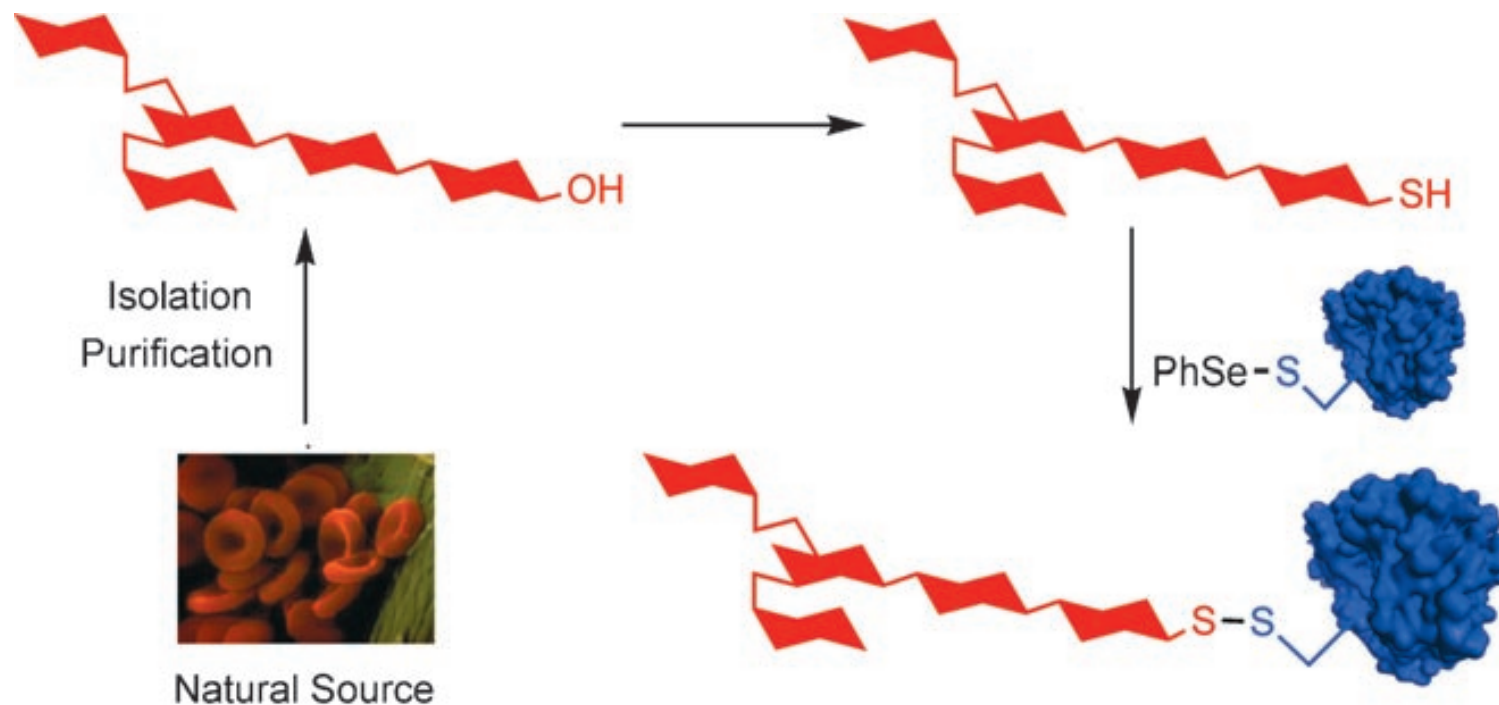
# Bioconjugation - Thiol exchange



# Glycoproteins



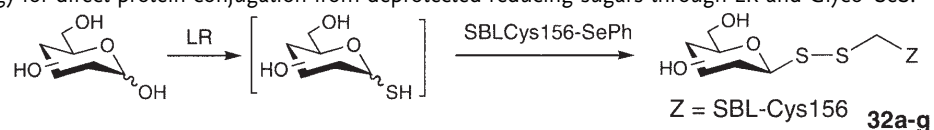
# Neoglycoproteins



**Scheme 2.** One-pot protein glycosylation with reducing sugars isolated from natural sources.

# Proof of concept

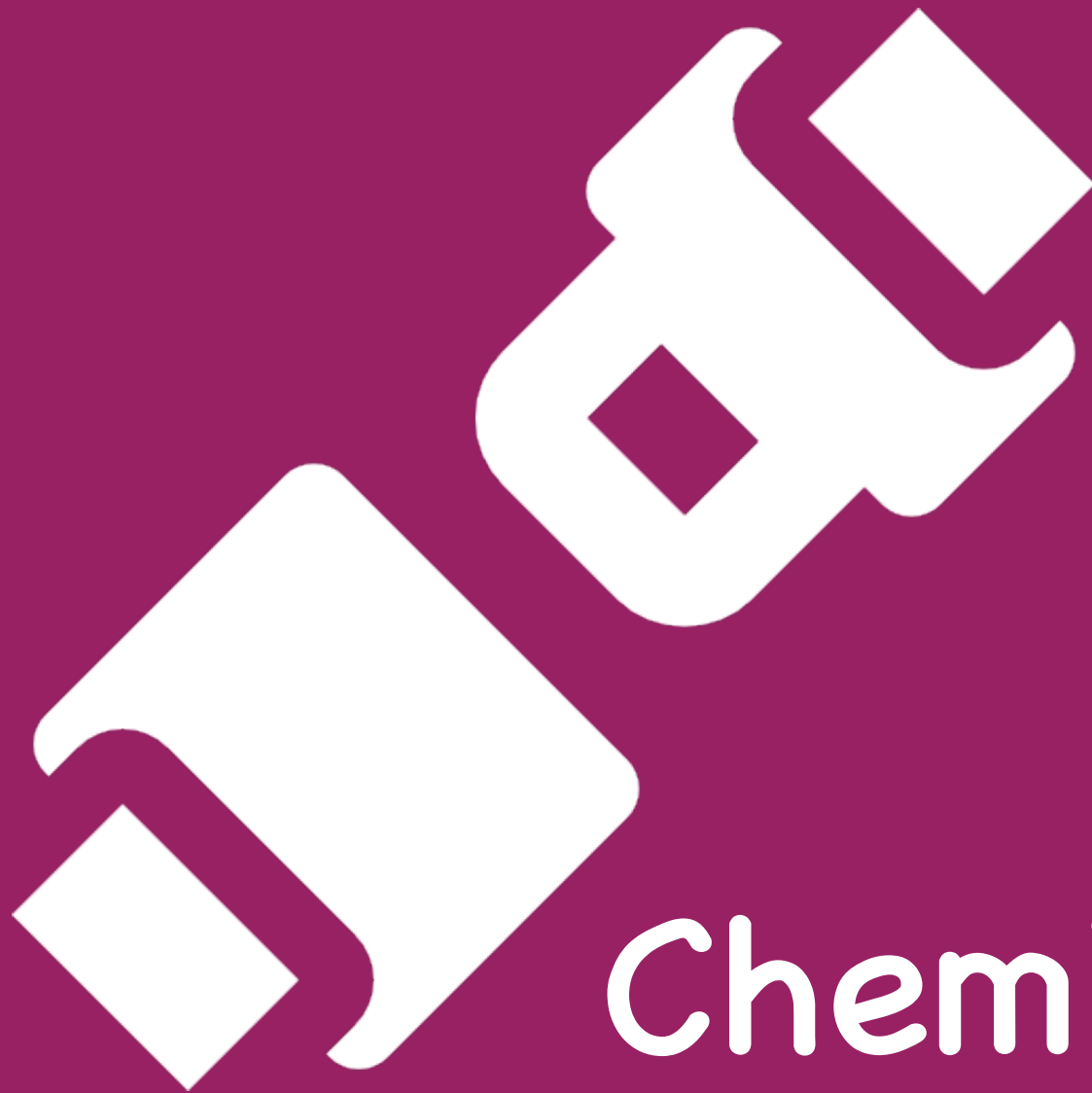
**Table 4:** Two-step strategy for direct protein conjugation from deprotected reducing sugars through LR and Glyco-SeS.<sup>[8]</sup>



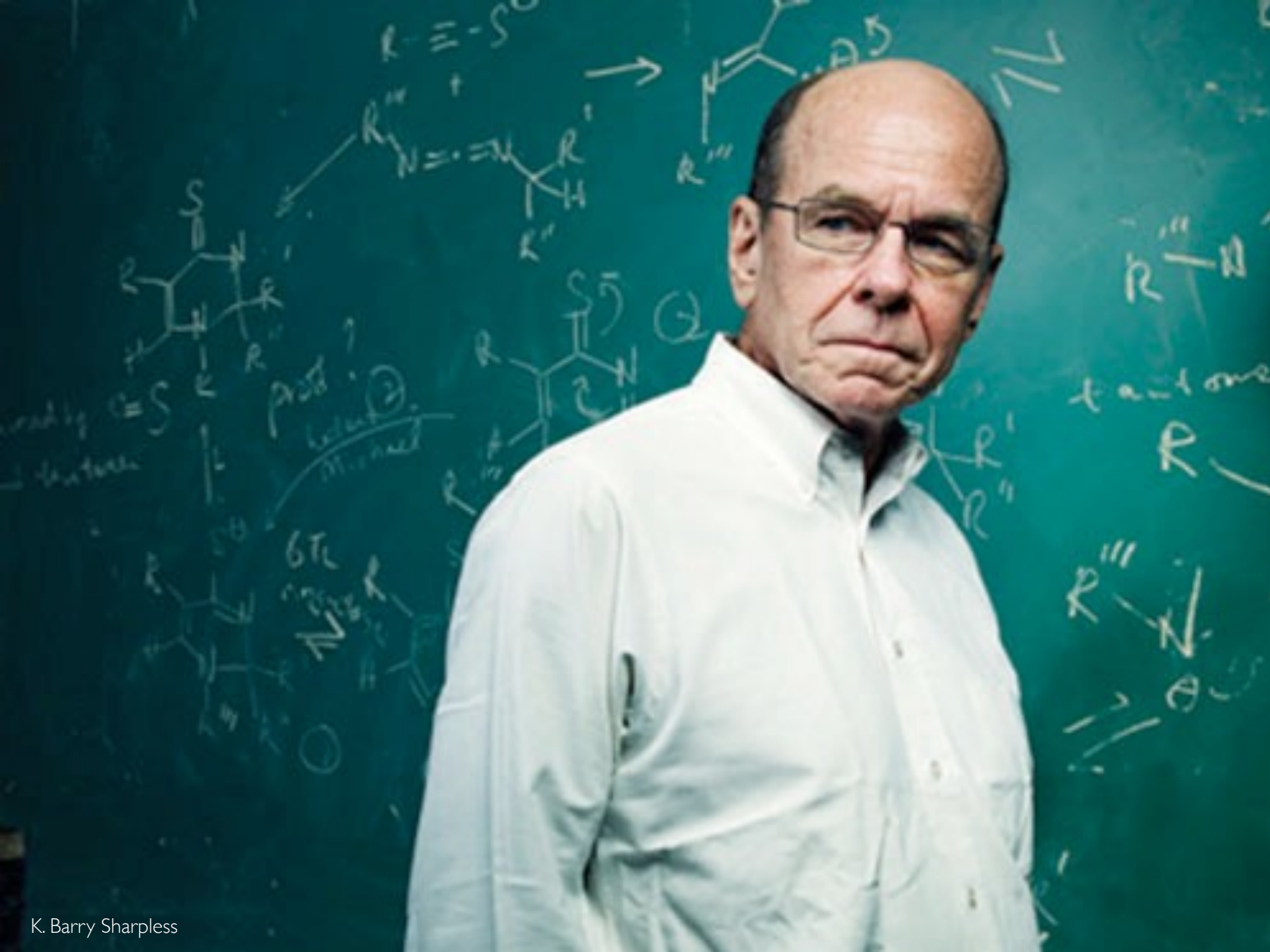
Sugar	Thiol product <sup>[a]</sup>	Protein product <sup>[b]</sup>	Conversion [%] <sup>[c]</sup>	ESI-MS Found (calcd)
Glc			<b>32a</b> > 95	27066 (27064)
Man			<b>32b</b> > 95	27064 (27064)
Gal			<b>32c</b> > 95	27064 (27064)
Lactose (Galβ1,4Glc)			<b>32d</b> > 95	27234 (27225)
Maltopentaose (Glcα1,4) <sub>4</sub> Glc			<b>32e</b> > 95	27722 (27711)
Galabiose (Gacα1,4Gal)			<b>32f</b> > 95	27224 (27225)
(Xylα1,3Glc)			<b>32g</b> > 95	27194 (27194)

[a] Typically, LR (1.2 equiv) was added to a solution of the deprotected sugar in anhydrous dioxane and left to stir at 110 °C for 48 h; see the Supporting Information for more details. [b] Typically, crude thiol (20–50 equiv), in water, was added to preactivated SBL156C-SePh in CHES (70 mM), MES (5 mM), CaCl<sub>2</sub> (2 mM); pH 9.5. After 30 min at RT, the reaction was analyzed by LC-MS; [c] Conversion determined by ESI-MS.





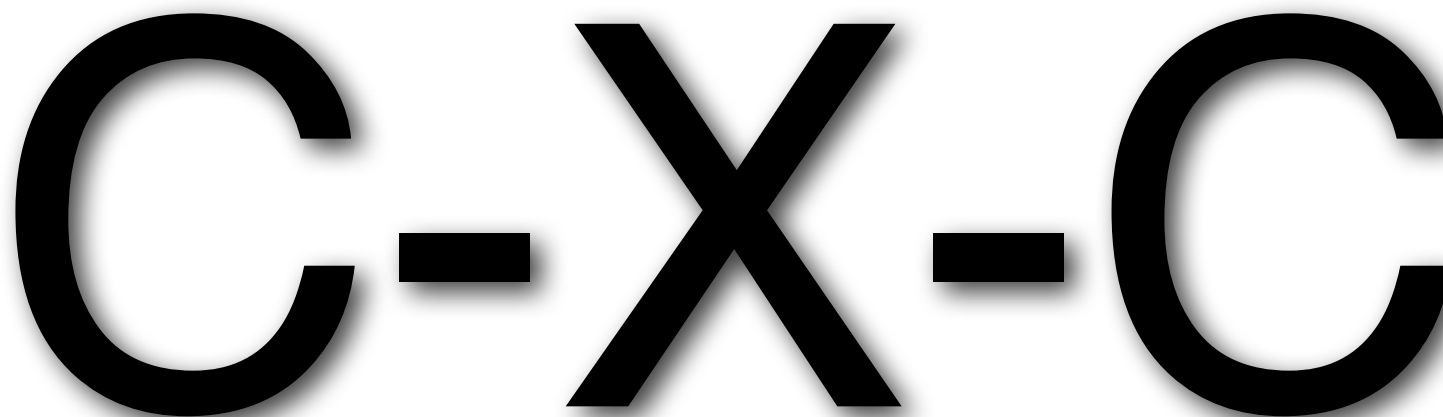
# Click Chemistry



K. Barry Sharpless

# Click Chemistry

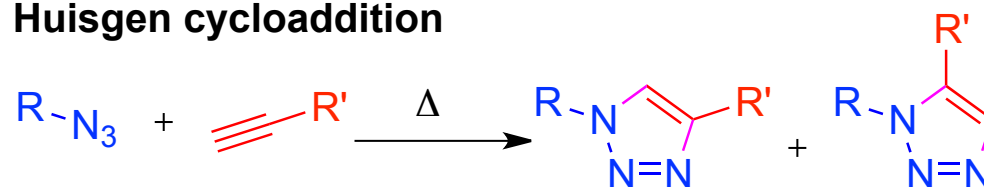
Diverse Chemical Function from a Few Good Reactions



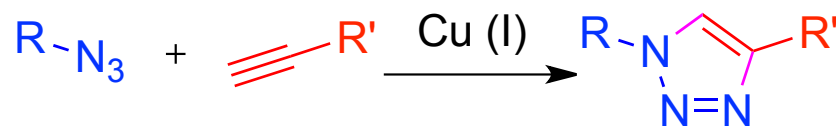


# Click chemistry - 2+3 Cycloadditions

## Huisgen cycloaddition

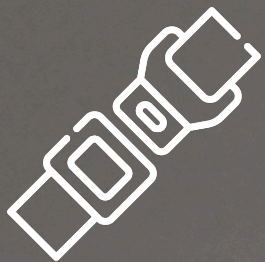


## CuAAC

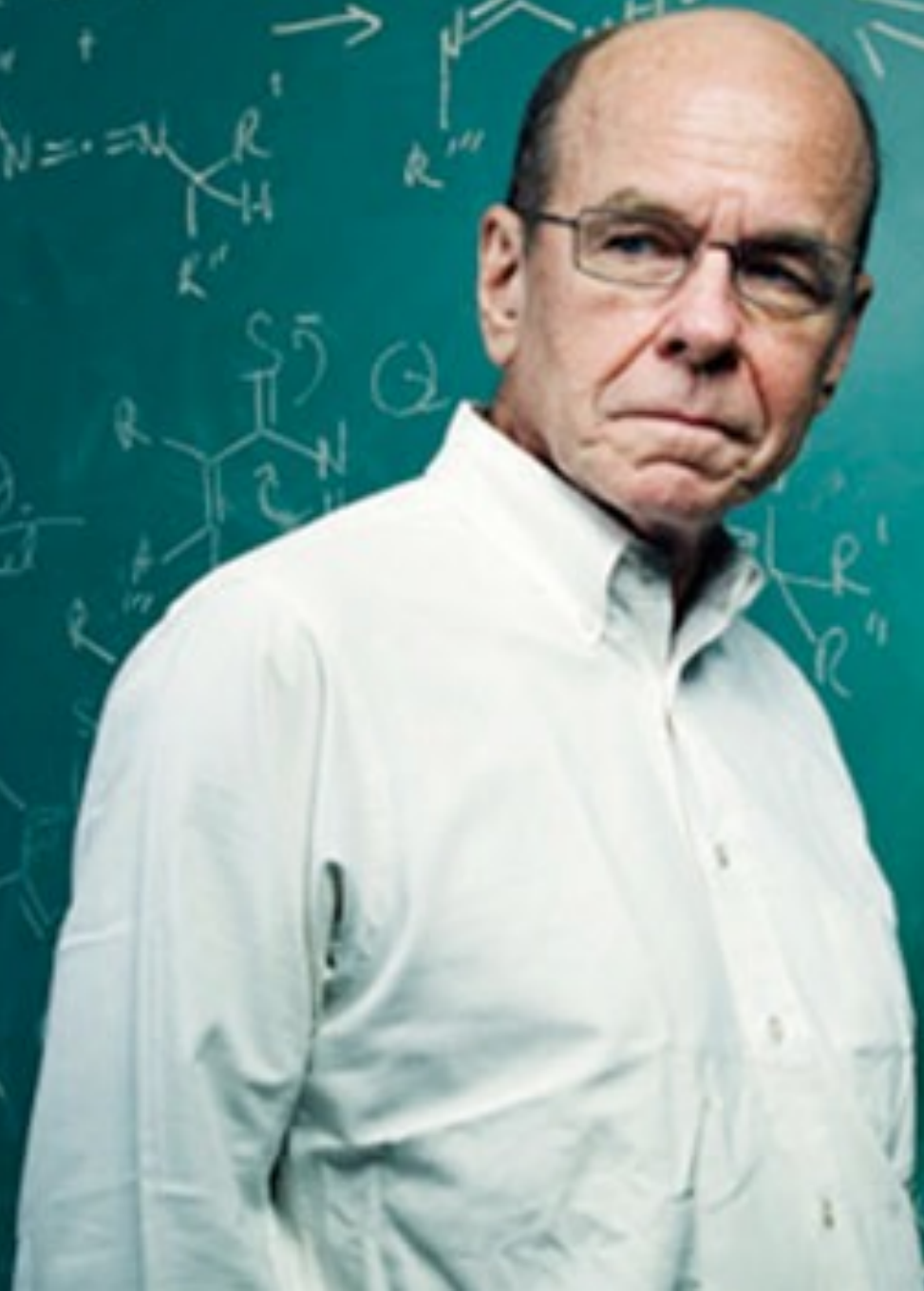


Meldal et coll., *J. Org. Chem.*, **2002**, 67, 3057-64

Sharpless et coll., *Angew. Chem. Int. Ed.*, **2002**, 41, 2596-99



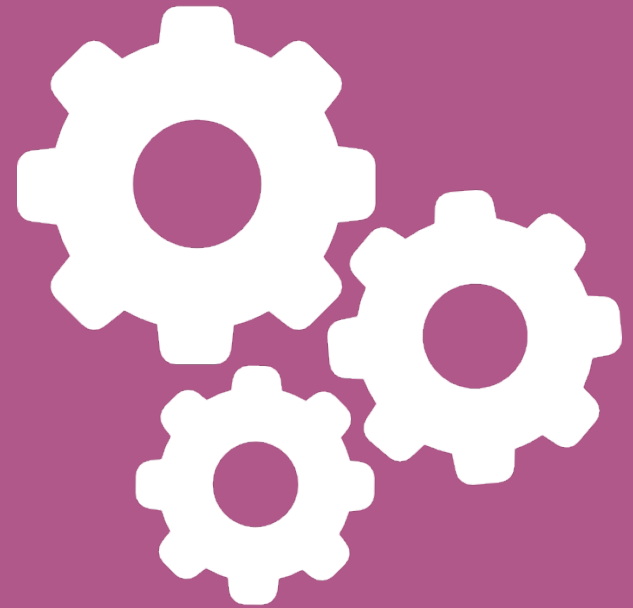
Morten P. Meldal



K. Barry Sharpless



Structure



Function



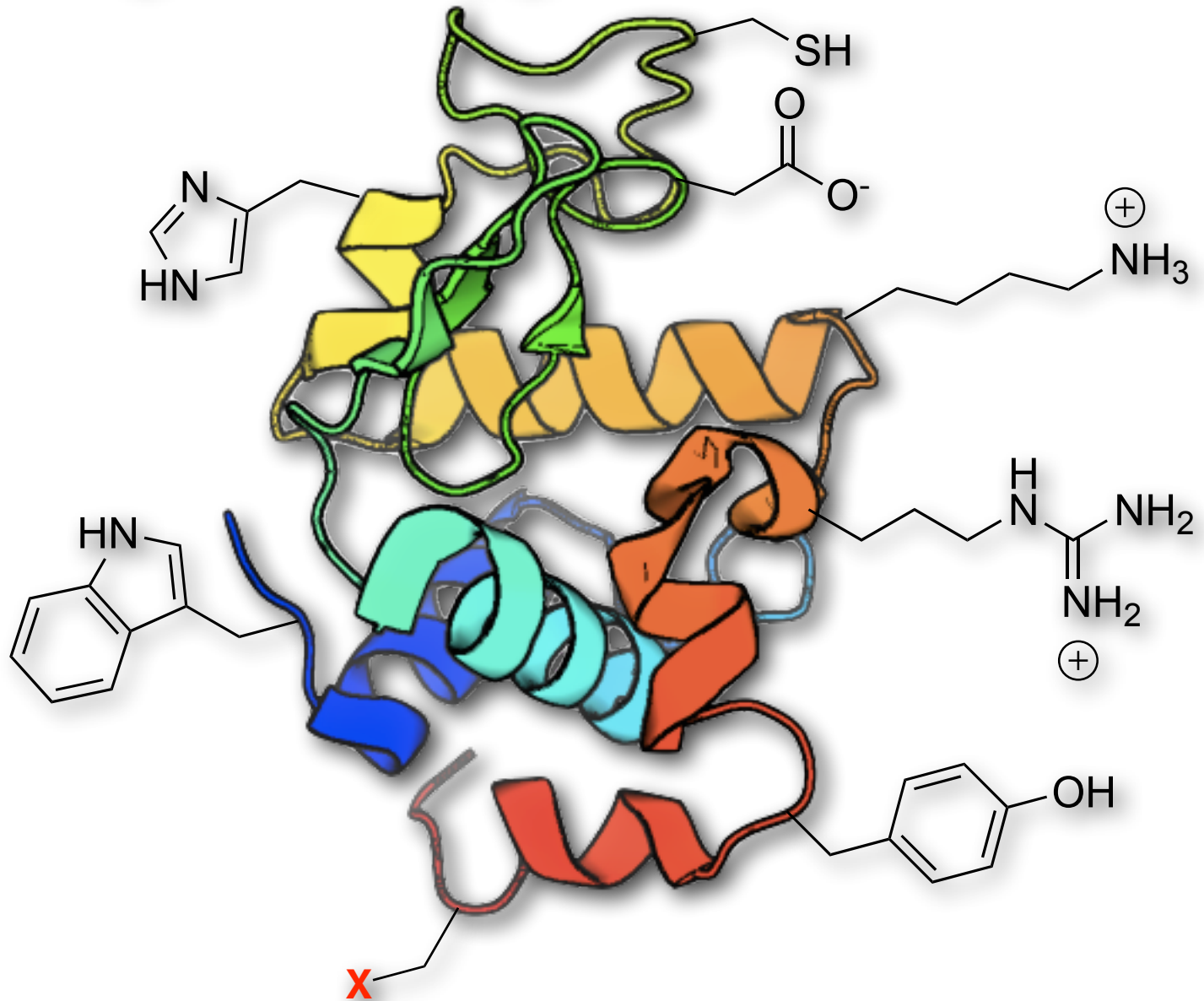
# Bioorthogonal Chemistry





Carolyn Bertozzi

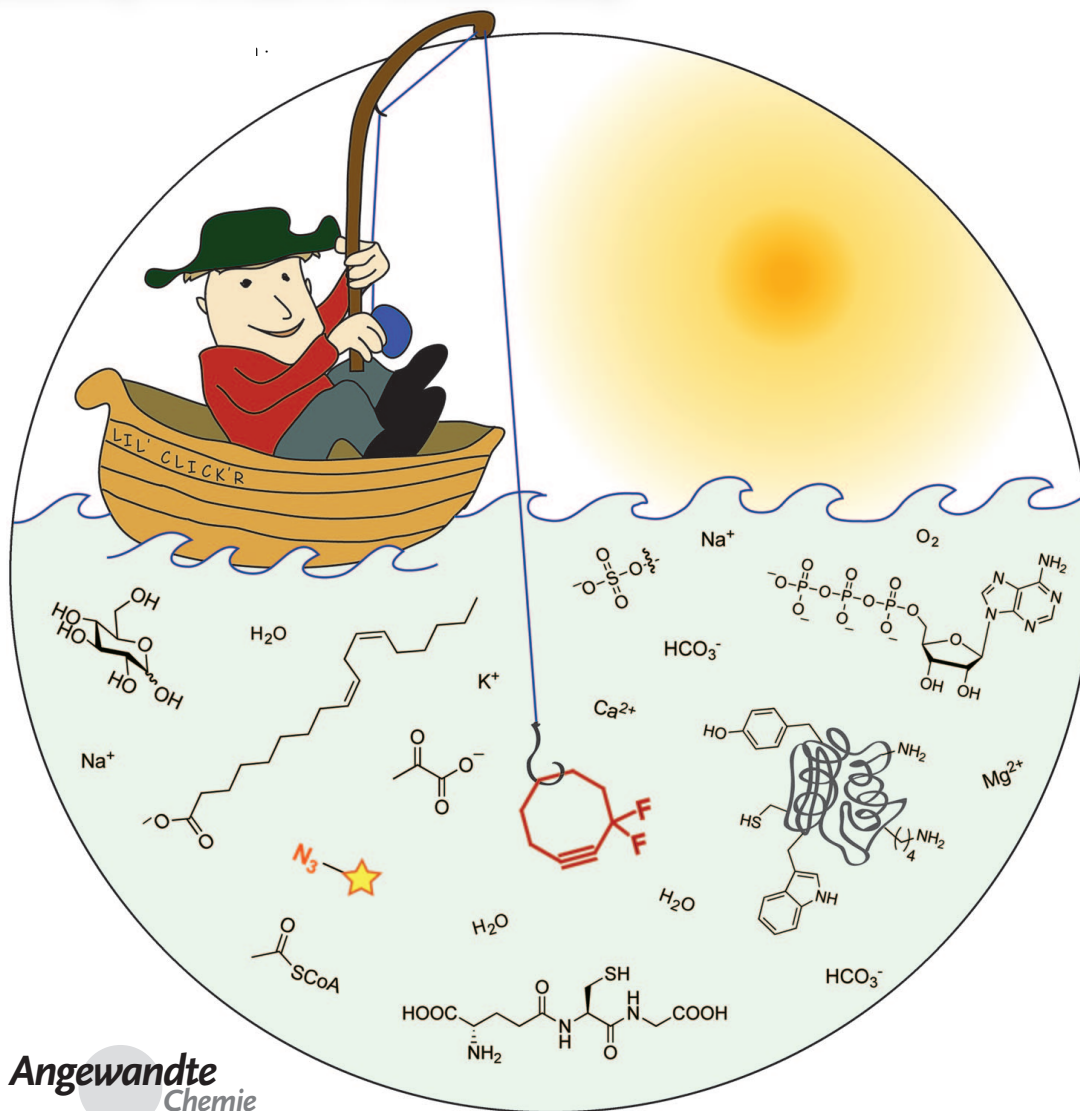
# Bioorthogonal bioconjugation





# Bioorthogonal chemistry

Fishing for Selectivity in a Sea of Functionality



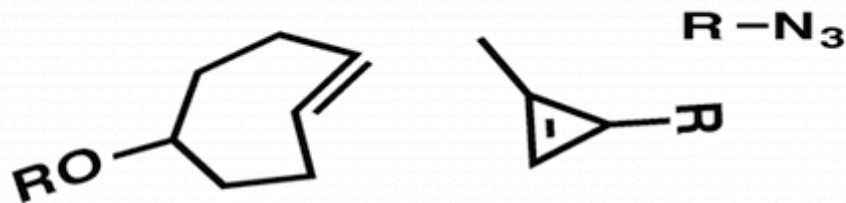
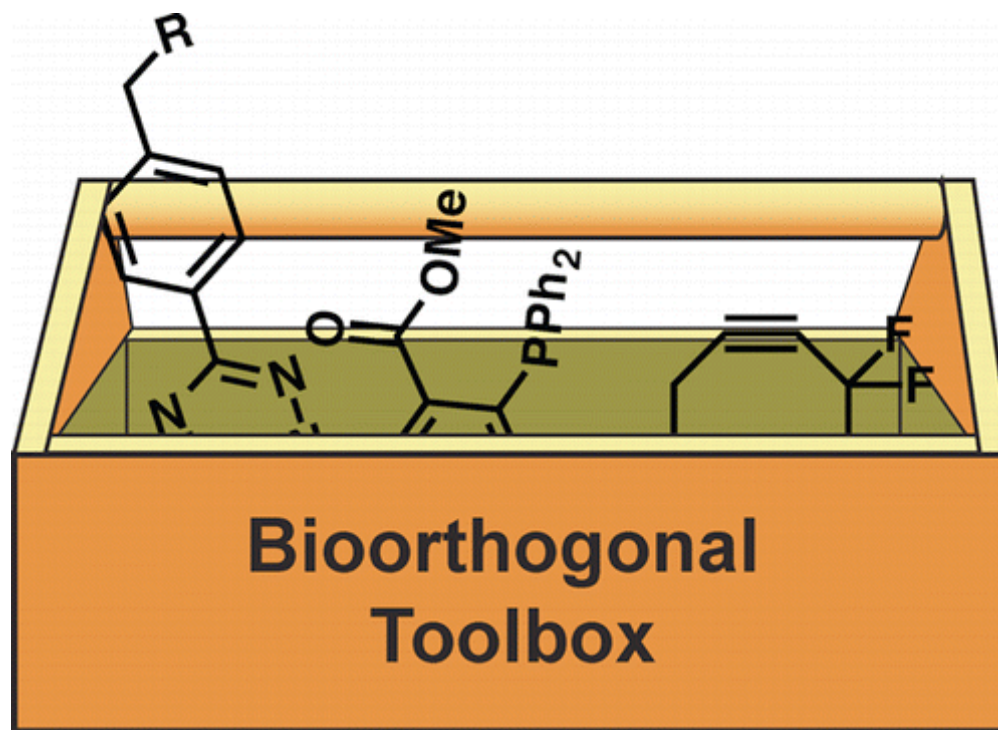
Angewandte  
Chemie

# Relevant chemical transformations ?

## Chemical Transformations Leading to Protein Covalent Modifications

- Water Is the Sole Solvent
- A Neutral pH Is Required
- Ambient Temperature (Up to 40°C)
- Kinetics, which Adapted to the Observed Phenomenon (on the Hour Scale)
- Low Reactant Concentrations
- Nontoxic Reagents

# Finding the right (bioorthogonal) chemistry



# Finding the right bioorthogonal reaction

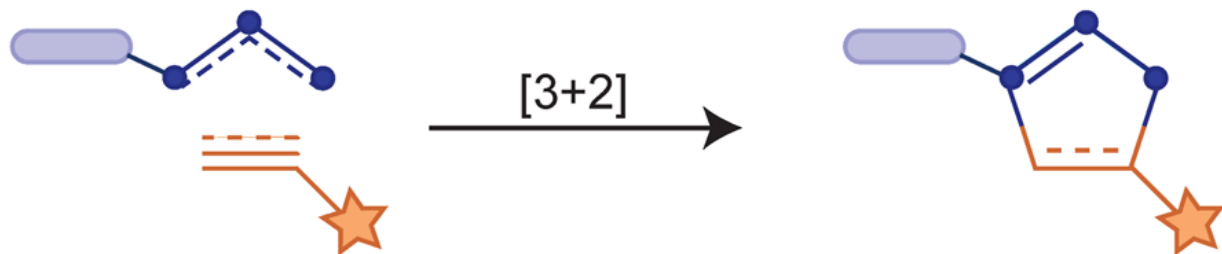
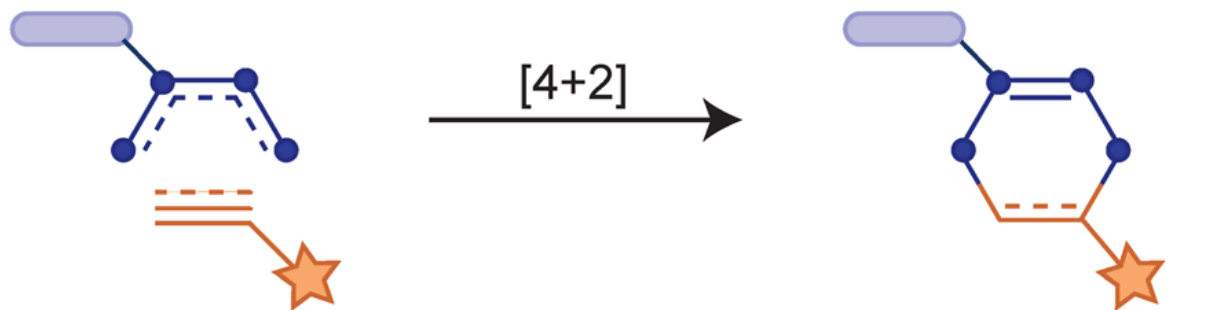


# Finding the right bioorthogonal reaction

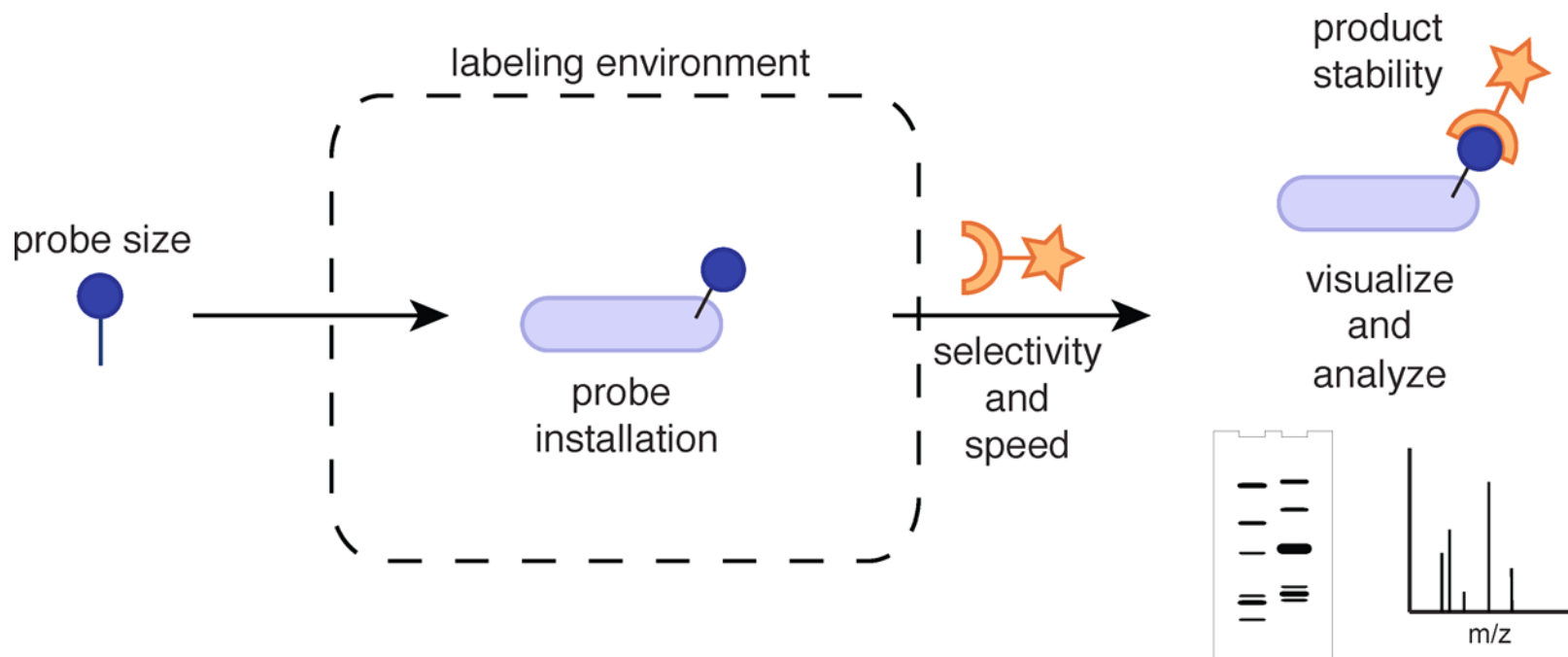
B polar reactions



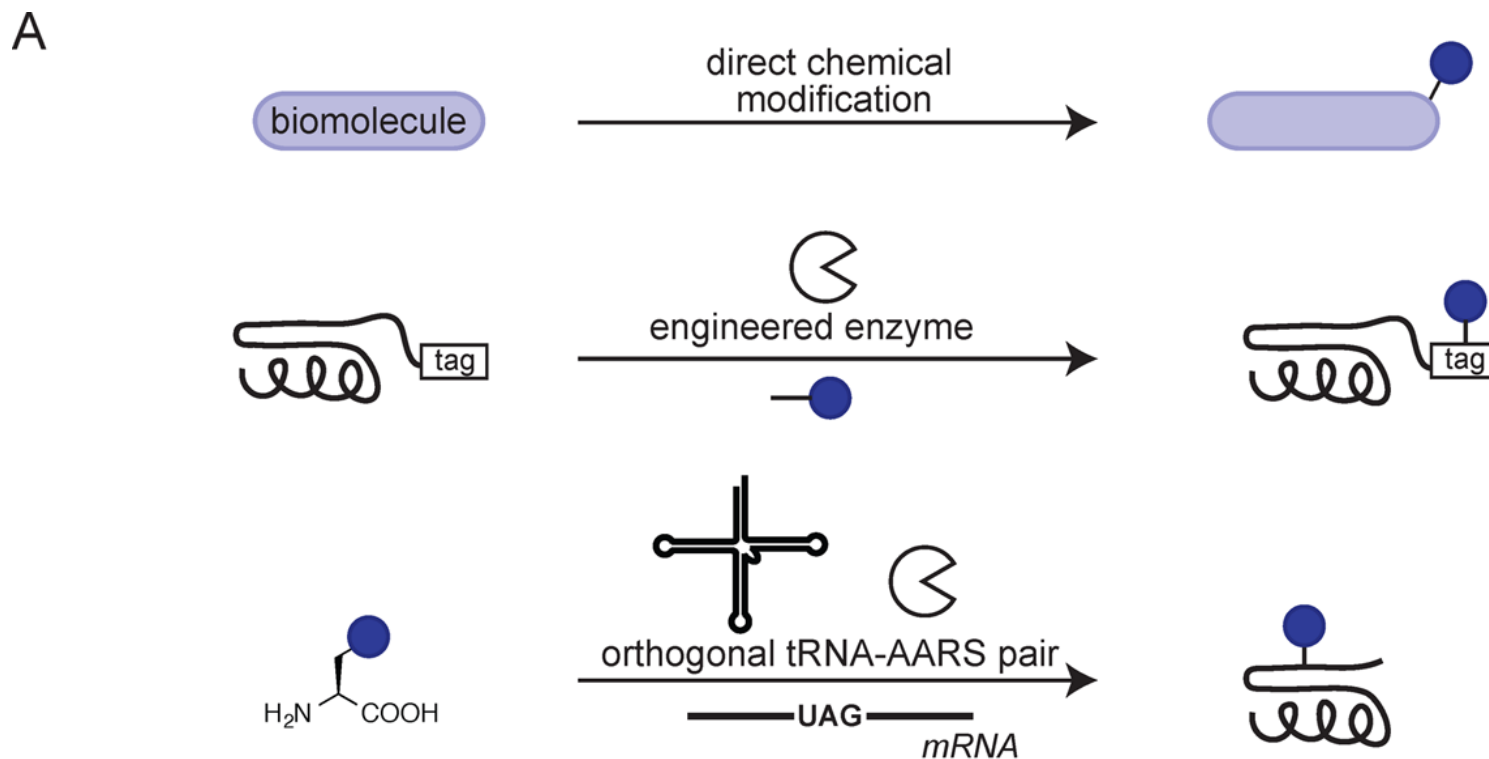
cycloadditions



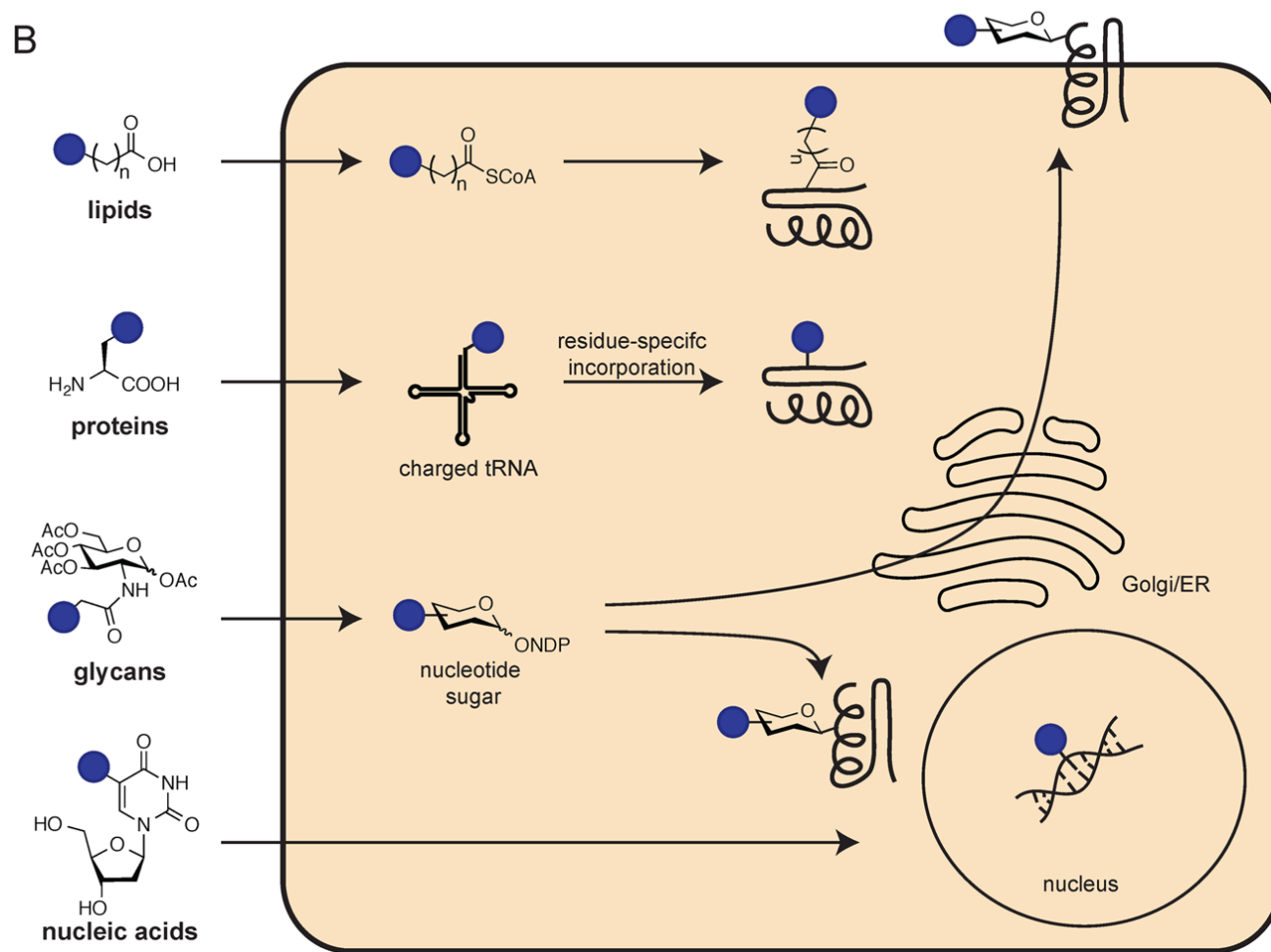
# Labeling biomolecules using bioorthogonal chemistry



# Modification of protein targets



# Metabolic modification of biomolecules





Metabolic  
glycan  
labeling

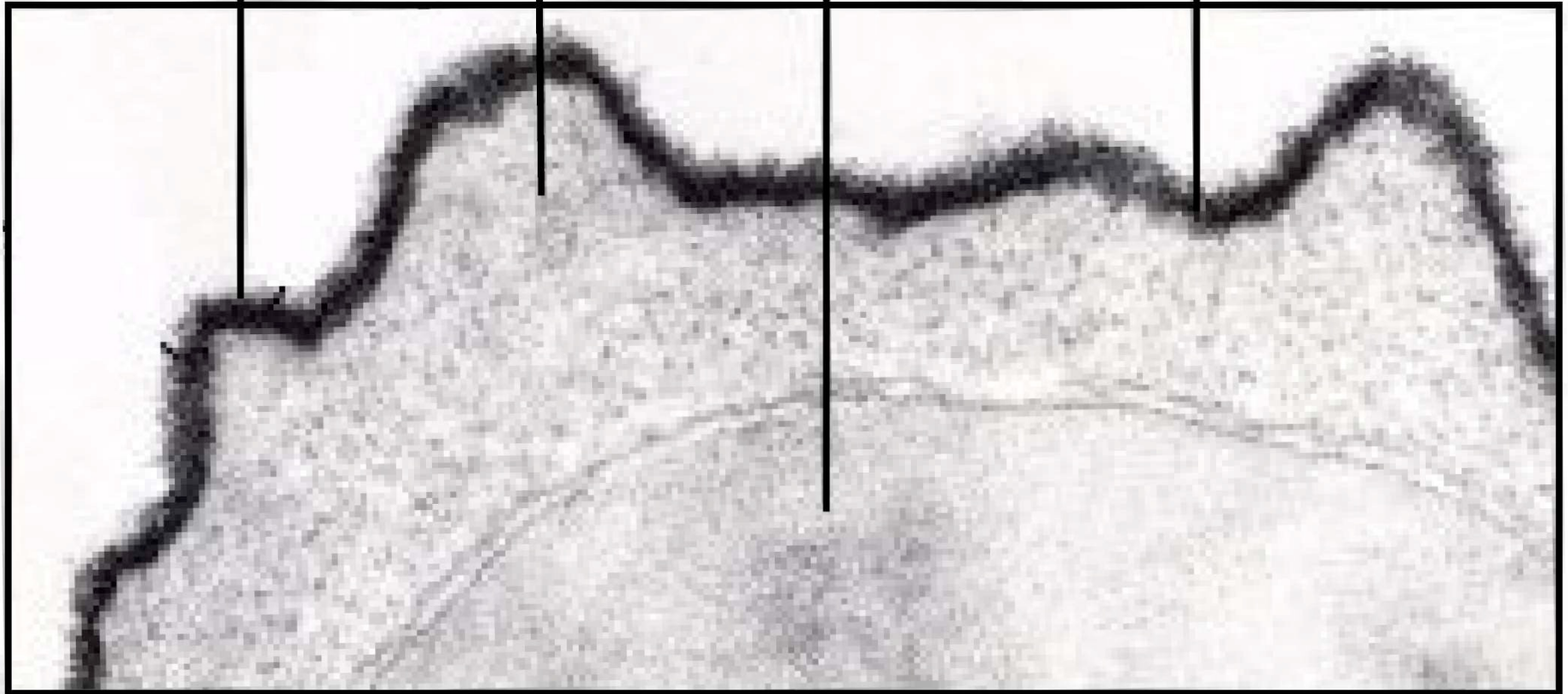
# Cell surface

Glycocalyx

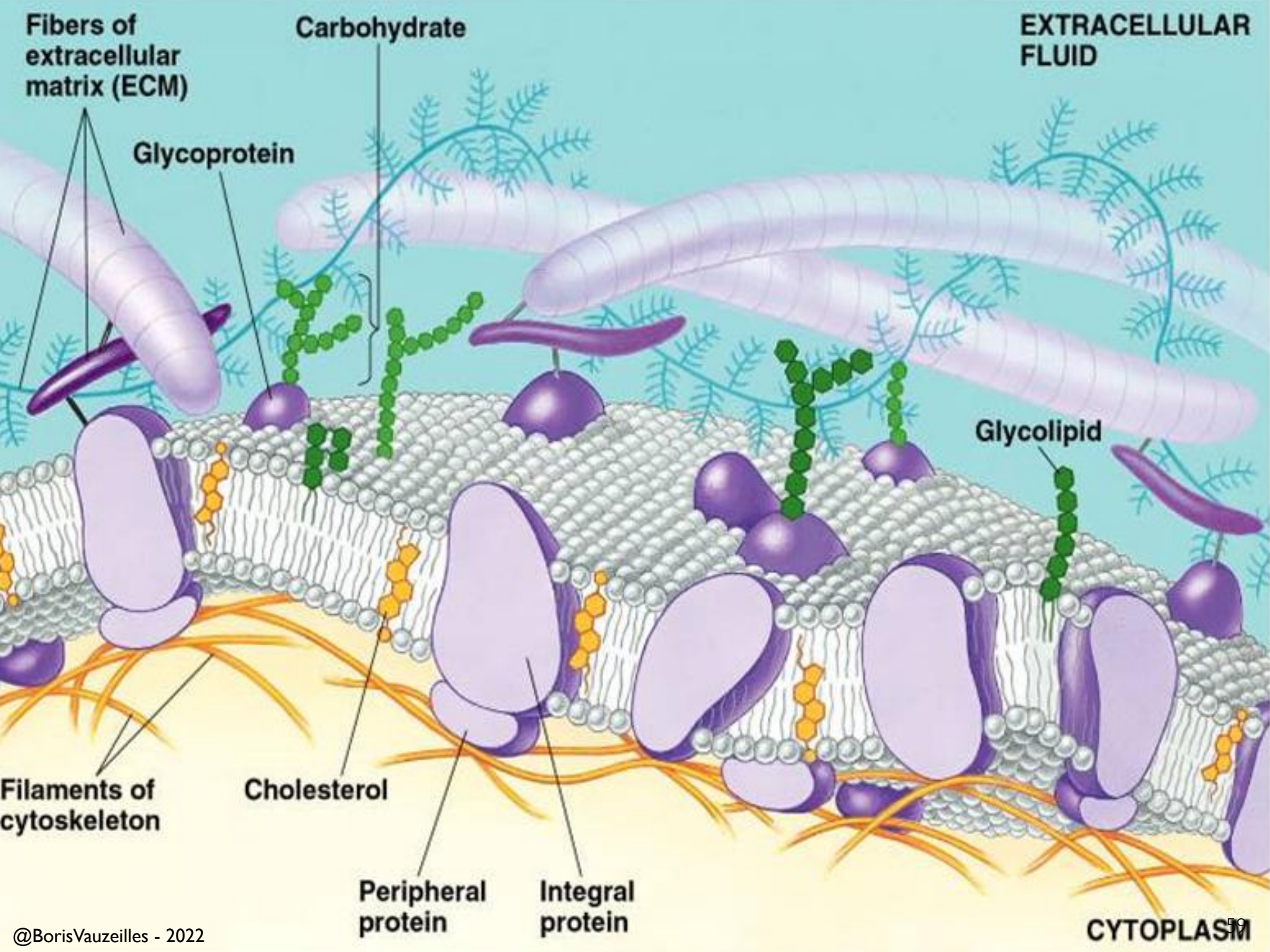
Cytoplasm

Nucleus

Plasma membrane

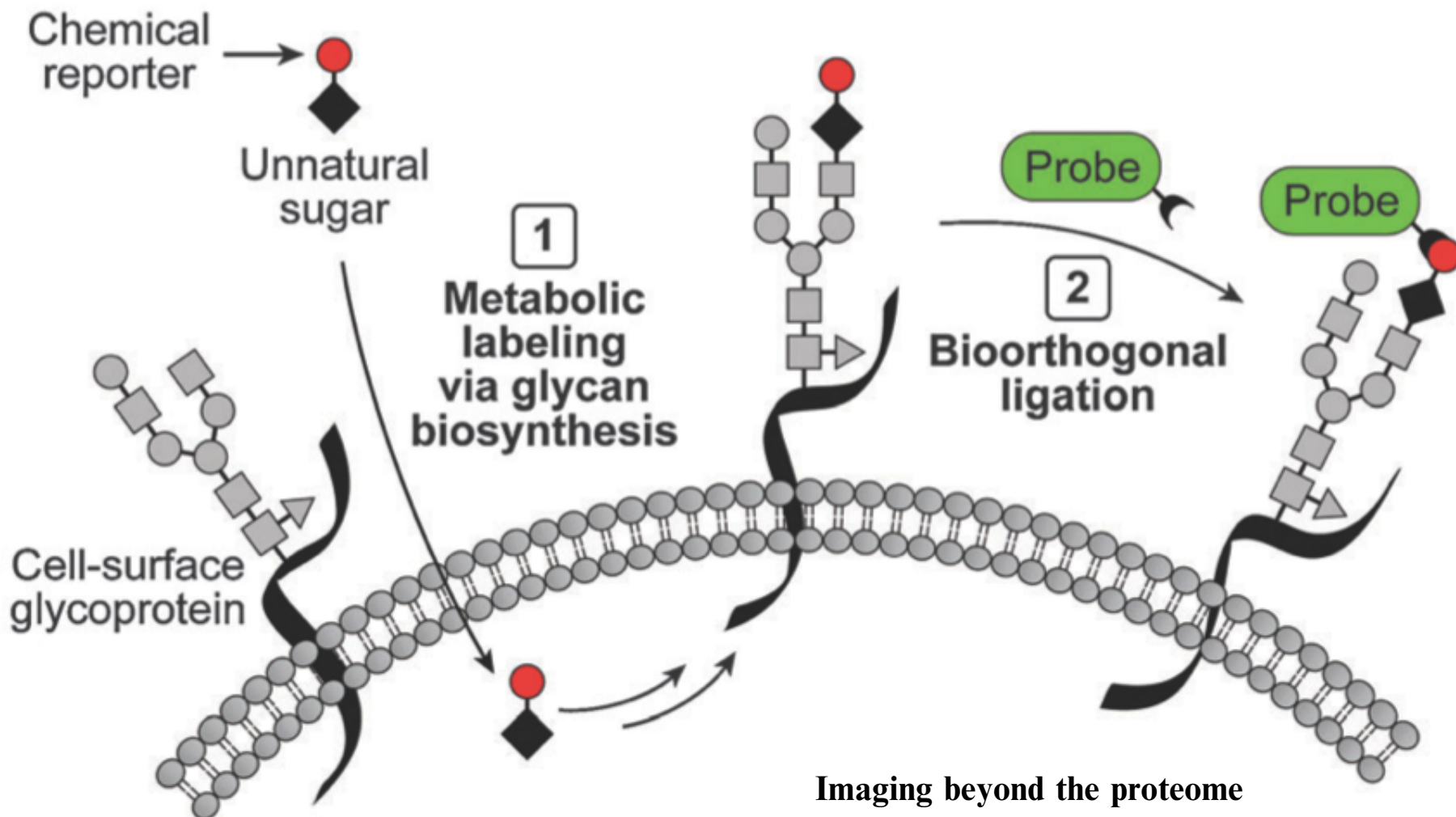


200 nm



# Metabolic Glycan Labeling

**A**

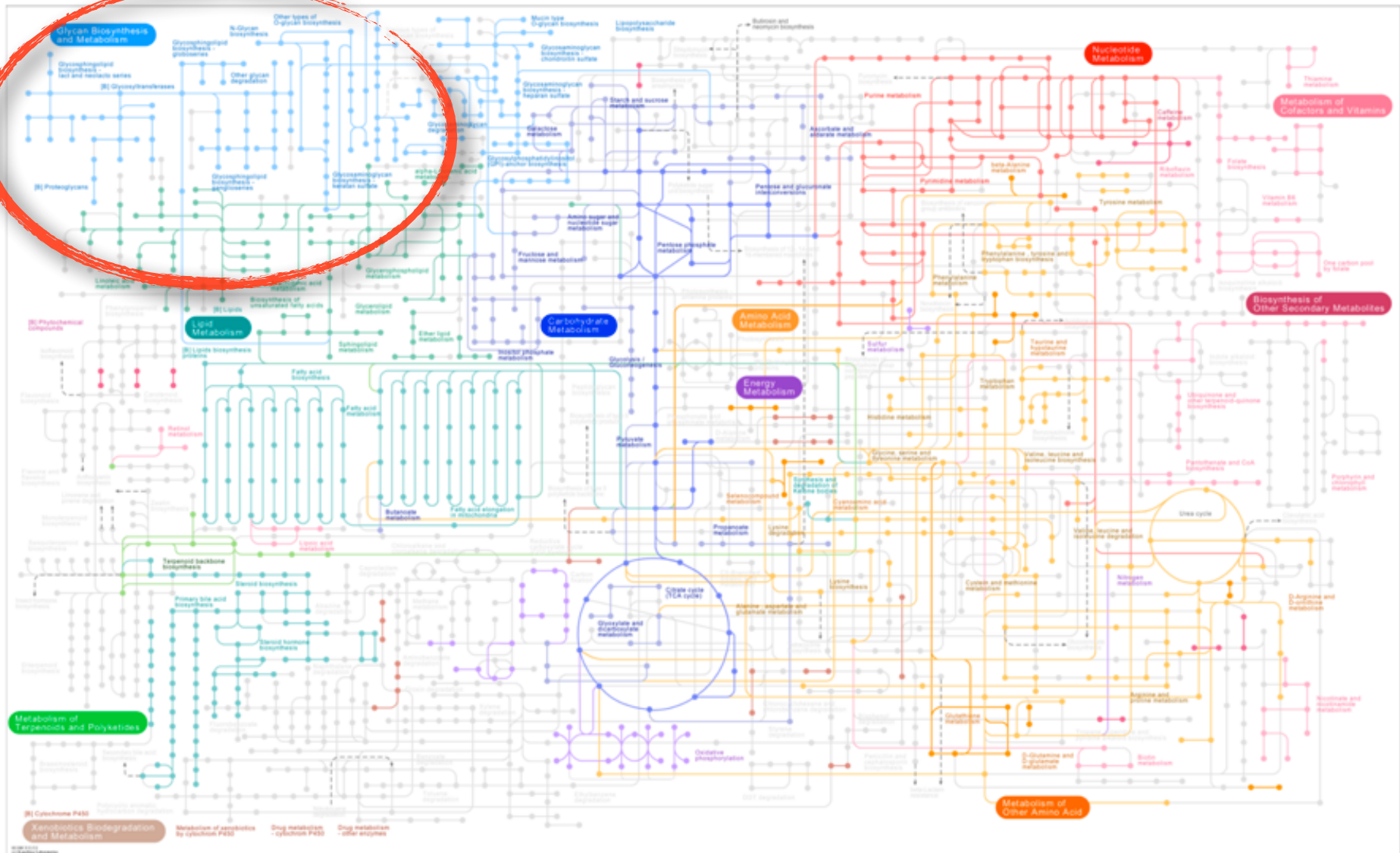


**Imaging beyond the proteome**

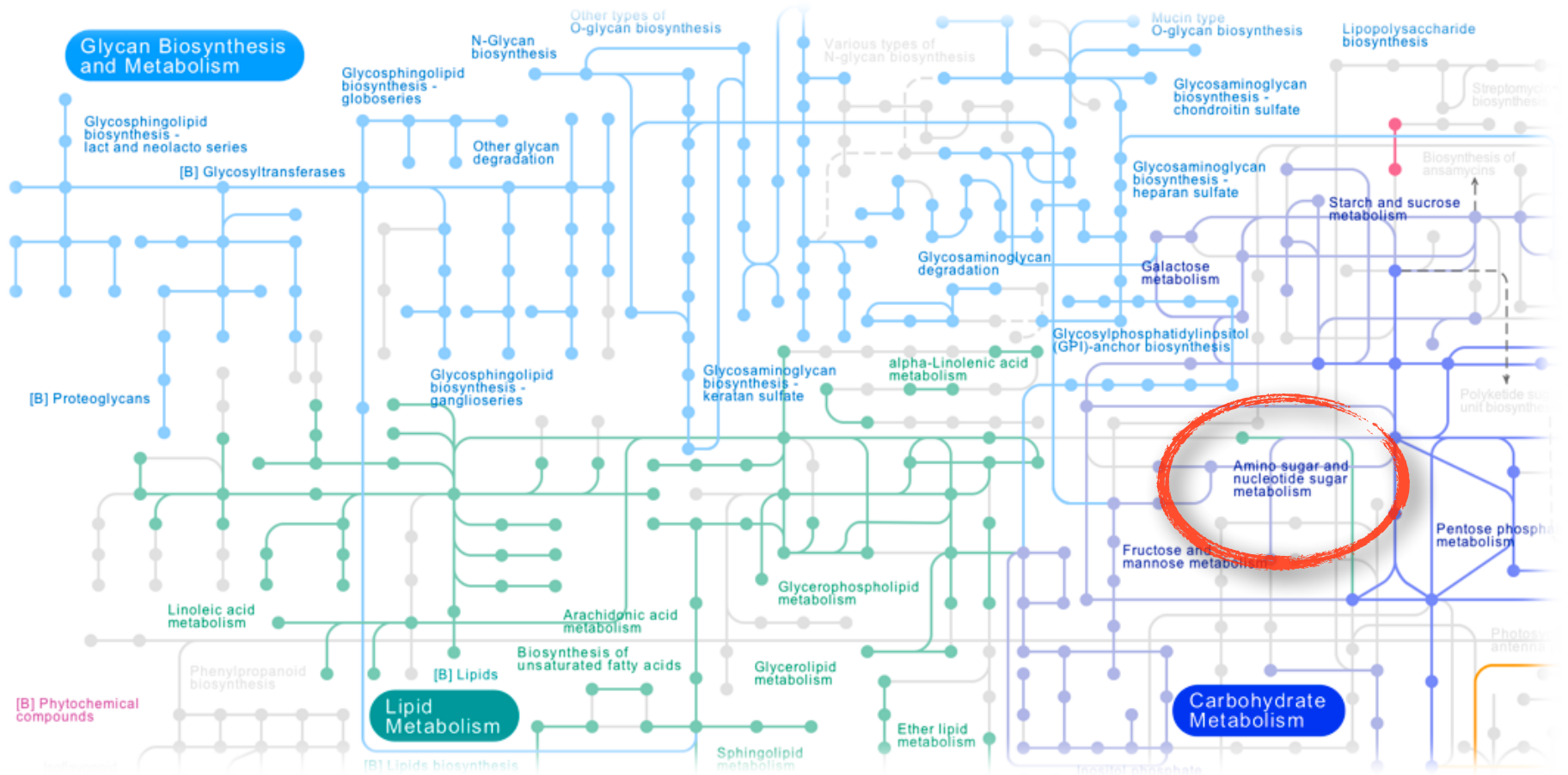
Pamela V. Chang<sup>a</sup> and Carolyn R. Bertozzi<sup>\*abc</sup>

*Chem. Commun.*, 2012, **48**, 8864–8879

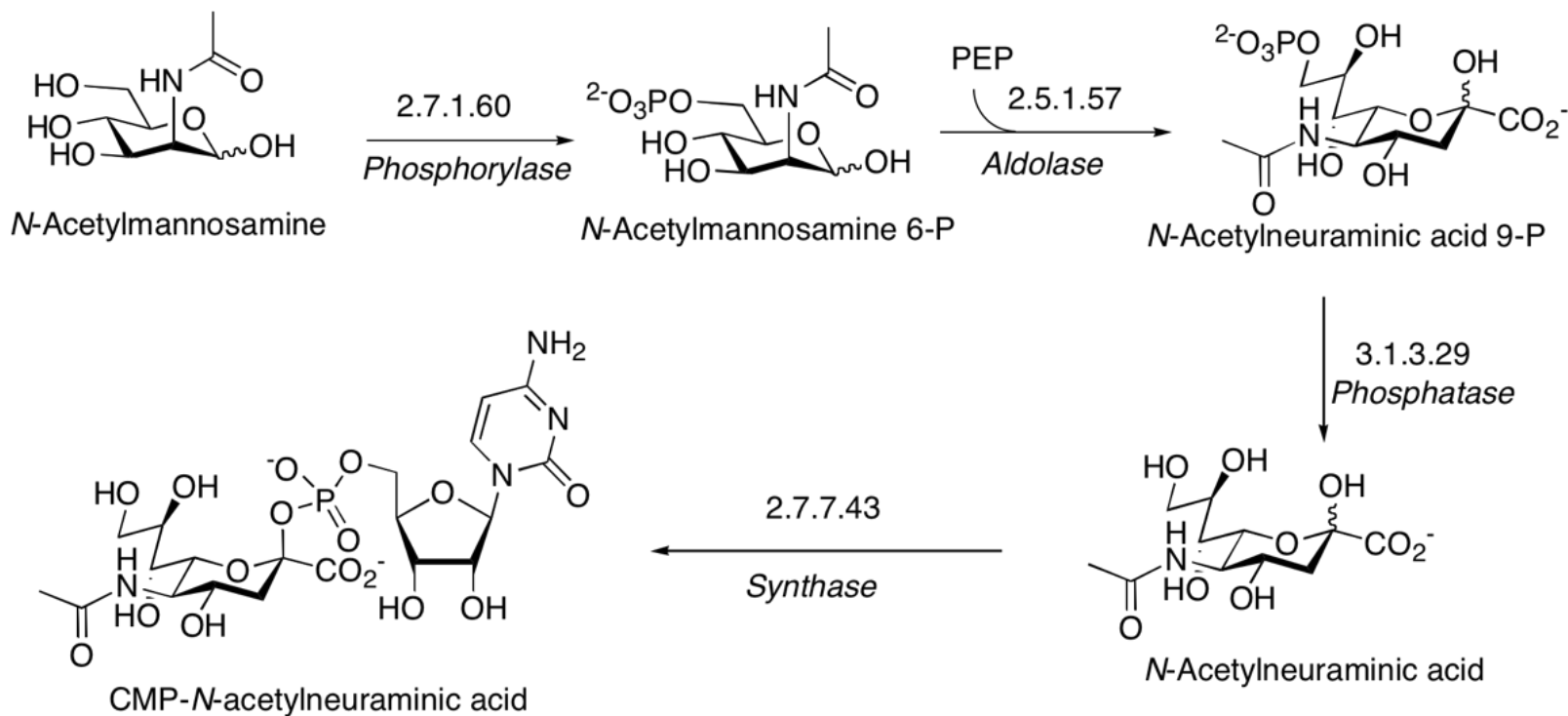
# Metabolic Oligosaccharide Engineering



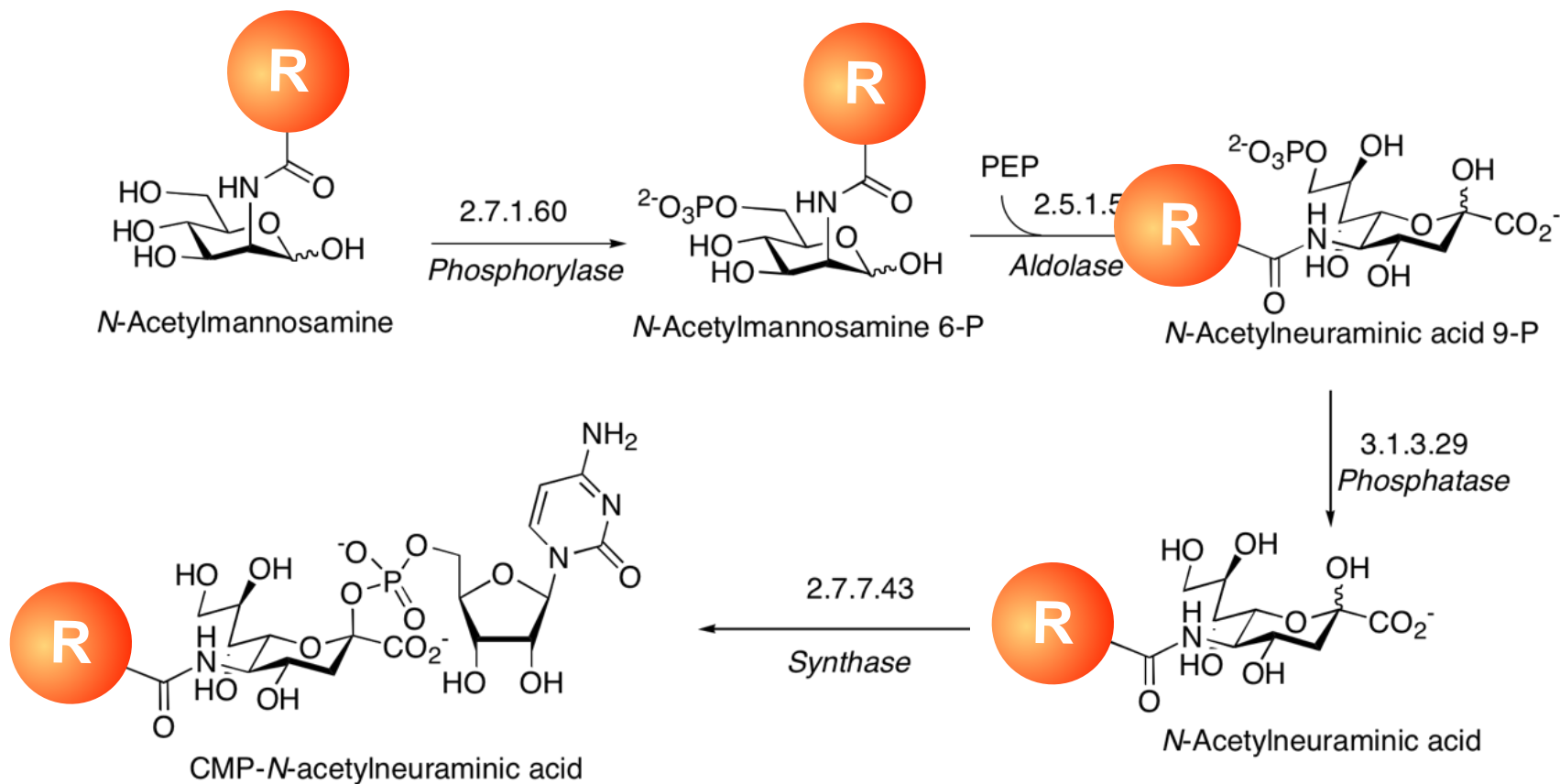
# Metabolic Oligosaccharide Engineering



# N-Acetyl neuraminic acid

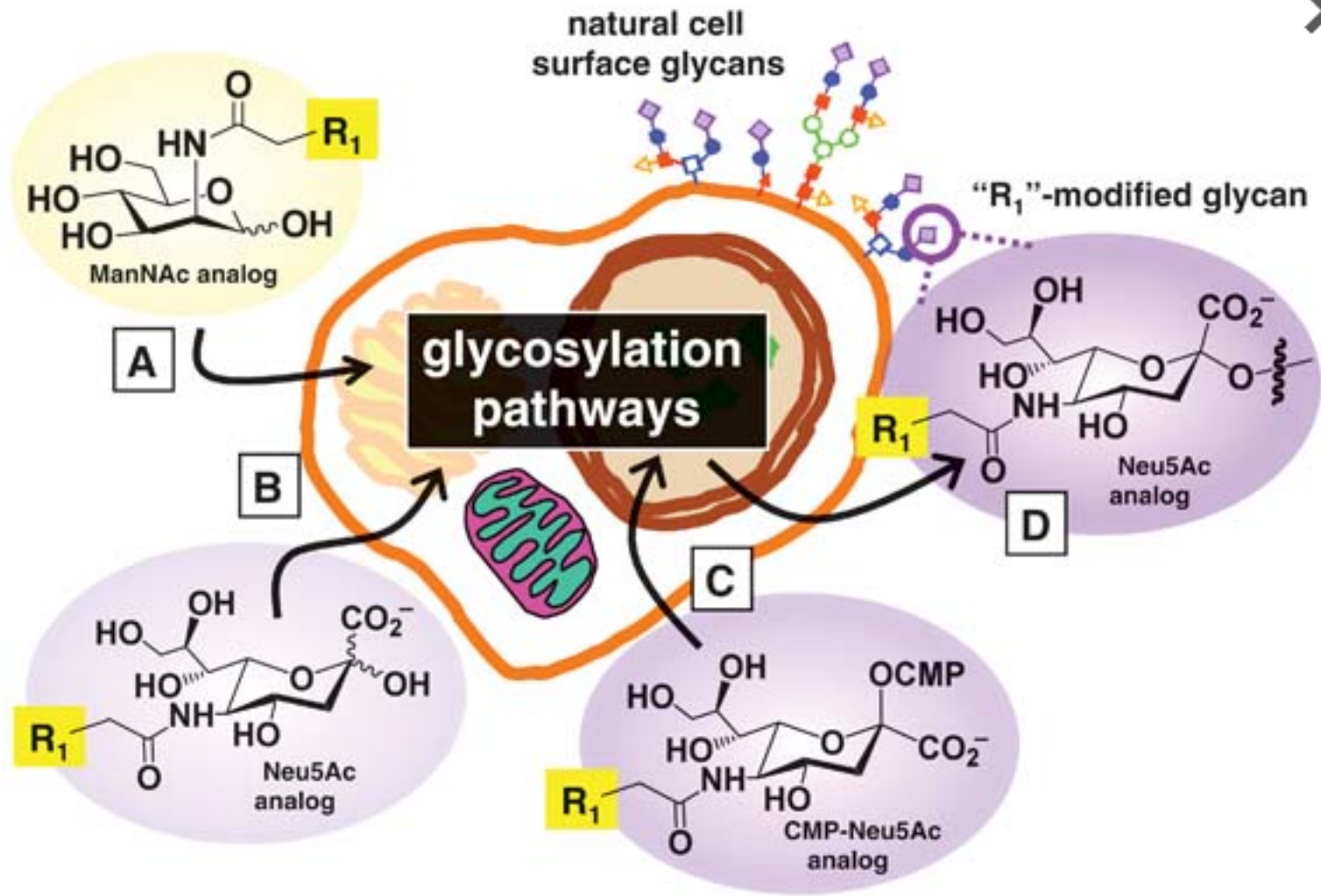


# N-Acetyl neuraminic acid





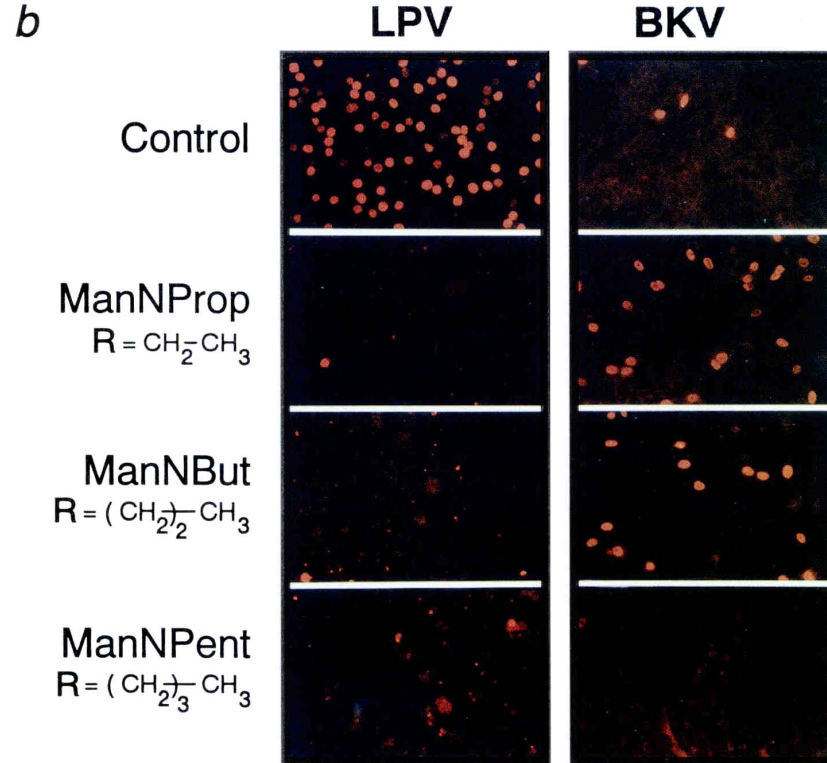
# Cell-surface glycan engineering



X

# Cell surface engineering

FIG. 4. Reduction or enhancement of host cell susceptibility to LPV or BKV infection by pretreatment with sialic acid precursor analogues. *a*, the chair conformation of the applied *N*-substituted D-mannosamines is shown with *R* indicating the modified *N*-acyl group. *b*, host cell lines BJA-B and Vero were cultured for 48 h in the presence of the sialic acid precursor analogues ManNProp, ManNBut, or ManNPent or as control an equivalent volume of PBS and subsequently infected with LPV or BKV, respectively. Virus-infected cells were identified by immunofluorescence staining for LPV and BKV capsid proteins 48 h after infection (this time allows the completion of only one viral replication cycle). Similar numbers of cells are present in microphotographs of each panel as determined by nuclear counterstaining (not shown).



THE JOURNAL OF BIOLOGICAL CHEMISTRY  
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Vol. 270, No. 3, Issue of January 20, pp. 1308-1314, 1995  
Printed in U.S.A.

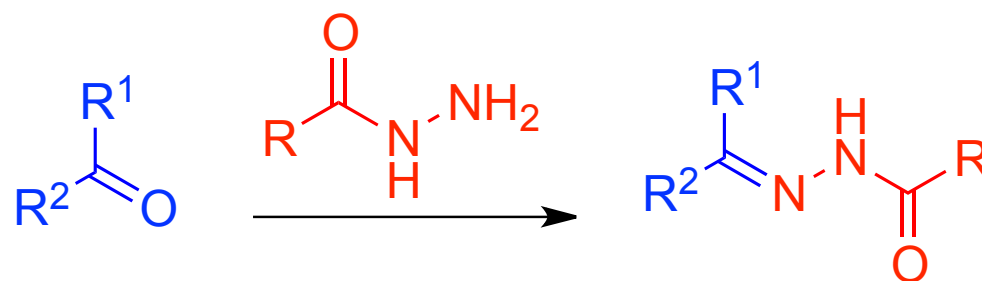
## Biosynthetic Modulation of Sialic Acid-dependent Virus-Receptor Interactions of Two Primate Polyoma Viruses\*

(Received for publication, September 14, 1994, and in revised form, November 2, 1994)

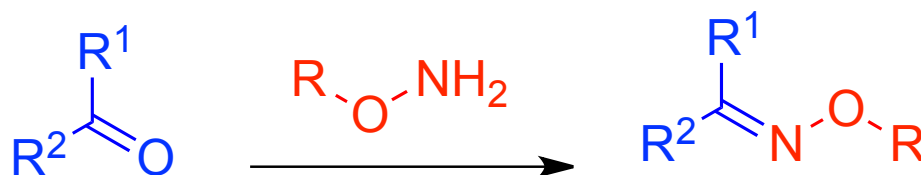
Oliver T. Keppler‡, Peer Stehling§, Markus Herrmann‡, Holger Kayser§, Detlef Grunow§, Werner Reutter§, and Michael Pawlita‡¶

# Bioorthogonal chemistry - polar ligations

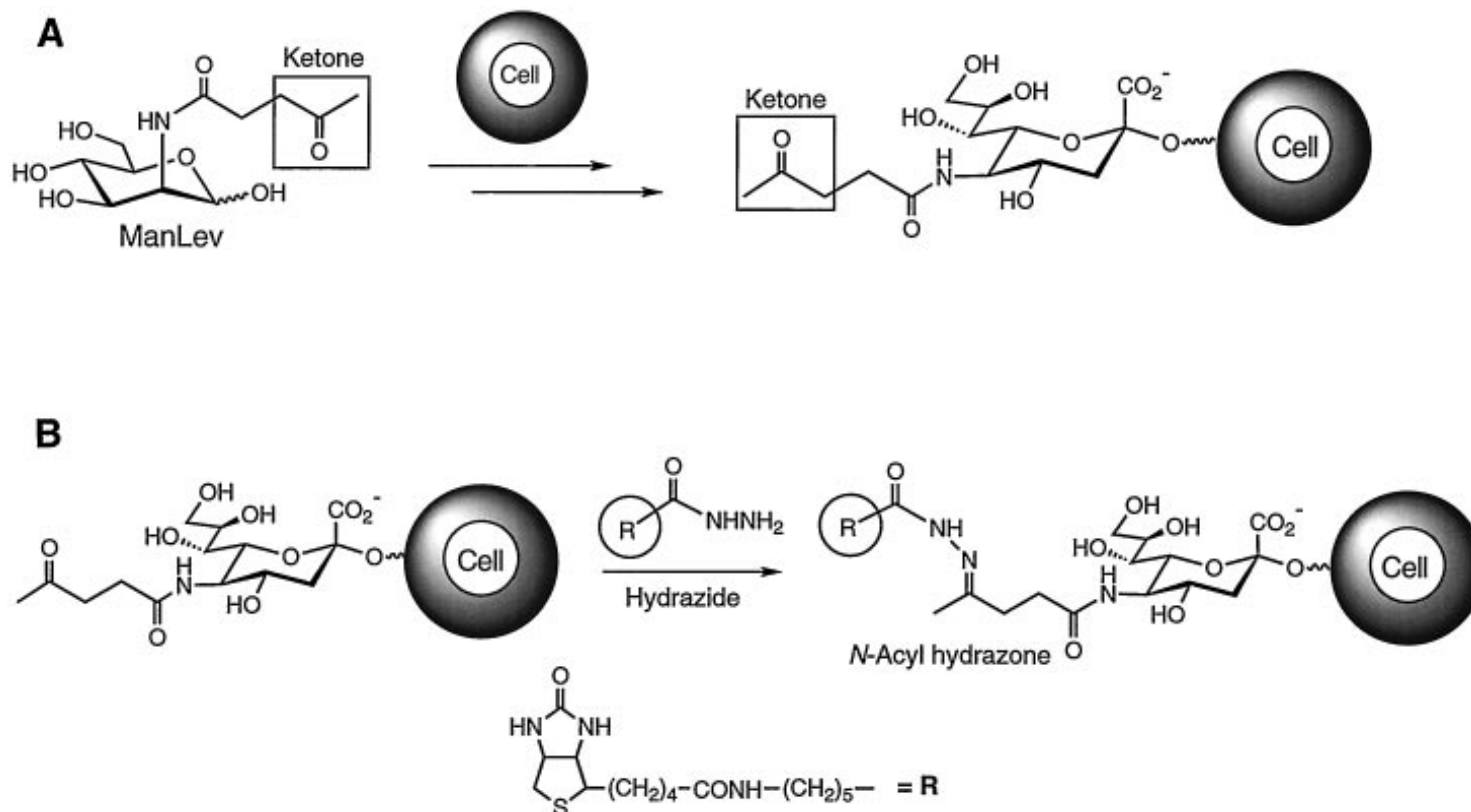
## Hydrazone ligation



## Oxime ligation



# Ketone incorporation on cell surface



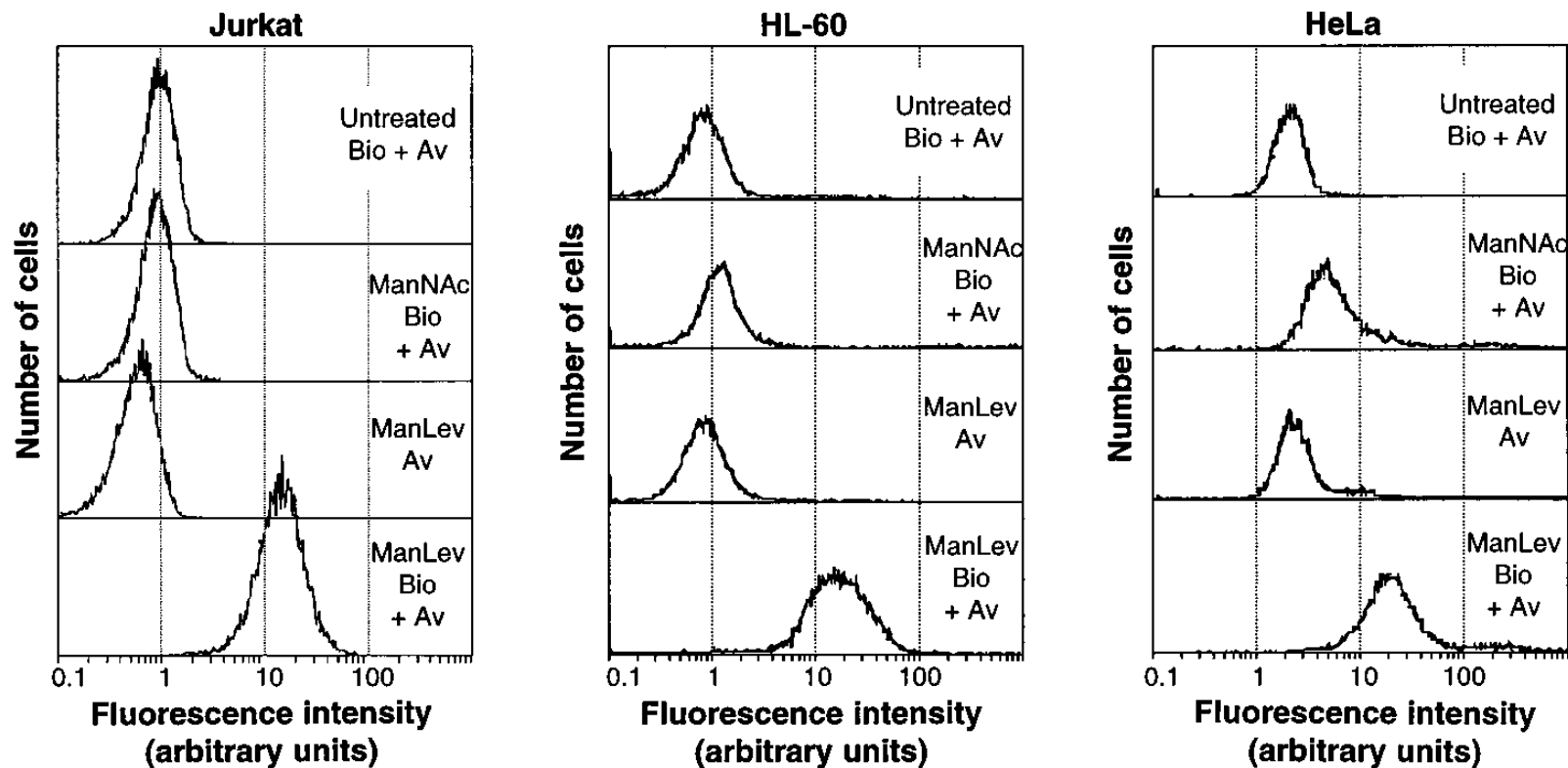
**Engineering Chemical Reactivity on Cell Surfaces Through Oligosaccharide Biosynthesis**

Lara K. Mahal, *et al.*

*Science* **276**, 1125 (1997);

DOI: 10.1126/science.276.5315.1125

# Ketone incorporation on cell surface



FITC-Avidin

Engineering Chemical Reactivity on Cell Surfaces Through Oligosaccharide Biosynthesis

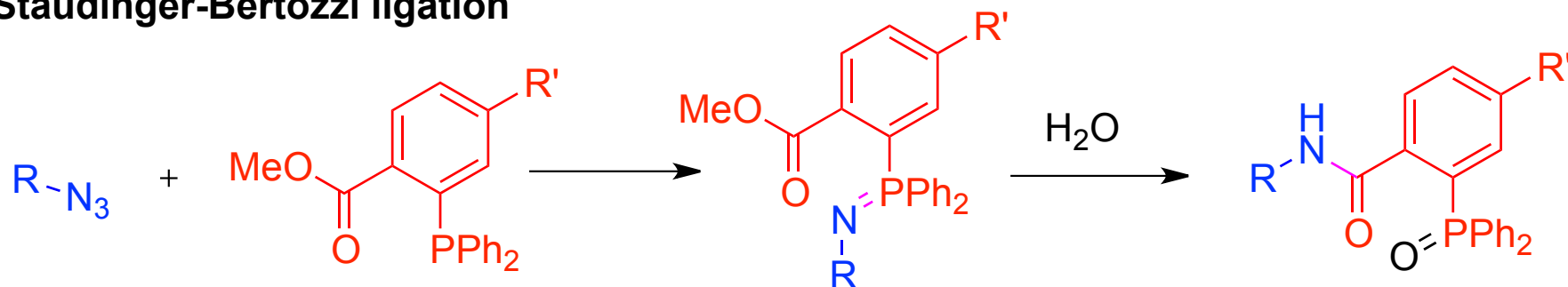
Lara K. Mahal, *et al.*

*Science* **276**, 1125 (1997);

DOI: 10.1126/science.276.5315.1125

# Bioorthogonal chemistry - Staudinger ligation

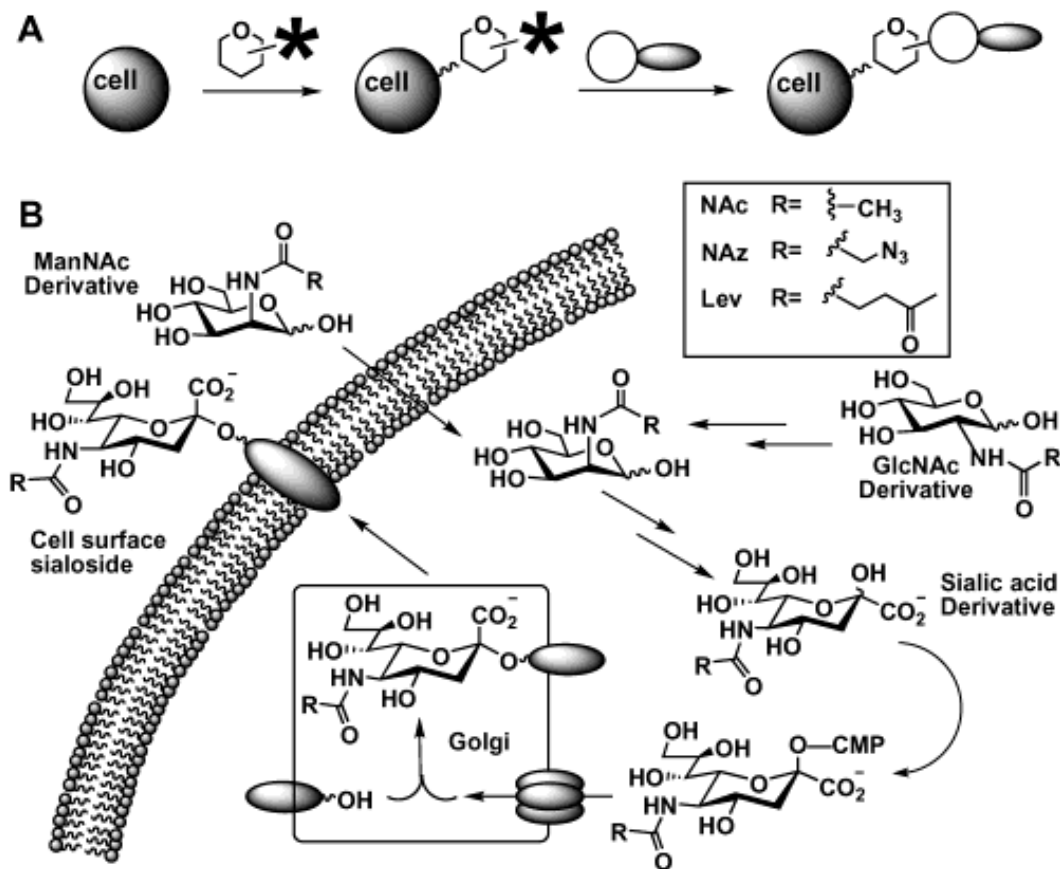
## Staudinger-Bertozzi ligation





Hermann Staudinger

# Azides



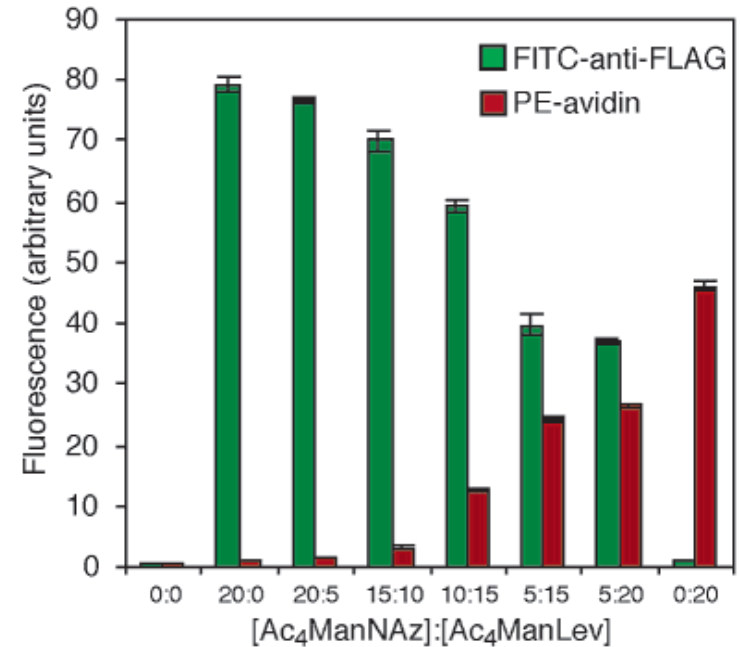
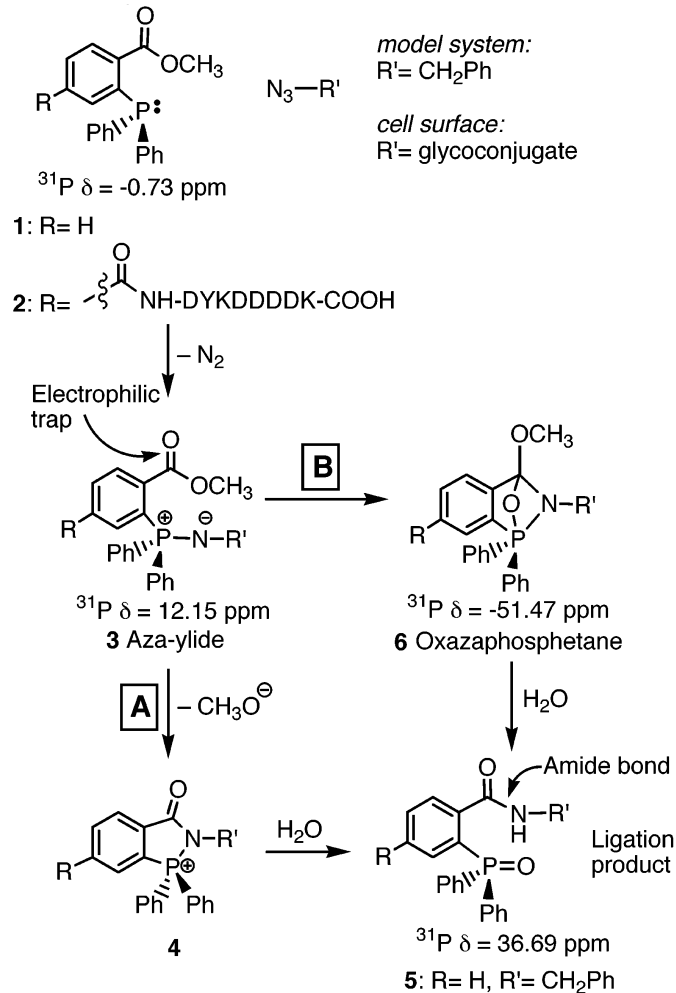
**J|A|C|S**  
ARTICLES  
Published on Web 11/23/2002

## Investigating Cellular Metabolism of Synthetic Azidosugars with the Staudinger Ligation

Eliana Saxon,<sup>†</sup> Sarah J. Luchansky,<sup>†</sup> Howard C. Hang,<sup>†</sup> Chong Yu,  
Sandy C. Lee, and Carolyn R. Bertozzi<sup>\*,†,‡</sup>



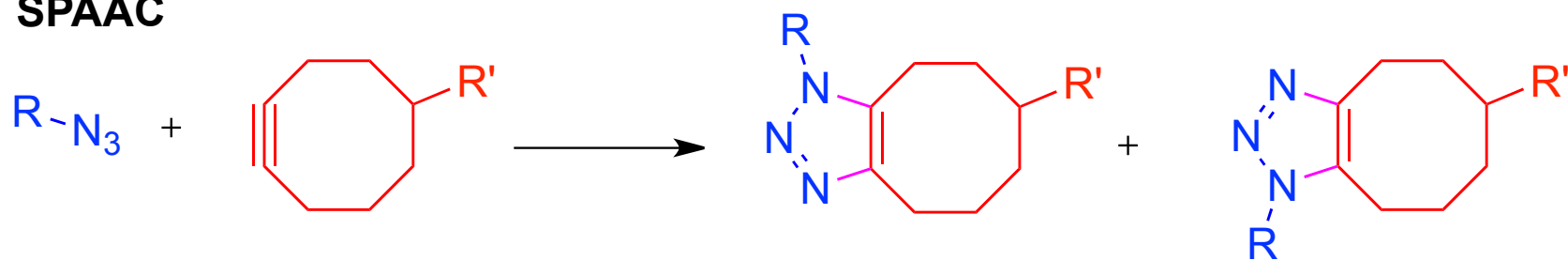
# Staudinger ligation



FLAG peptide, NH<sub>2</sub>-DYKDDDDK-COOH  
PE, phycoerythrin

# Bioorthogonal chemistry - 2+3 Cycloadditions

**SPAAC**



# Copper-free click chemistry

J|A|C|S  
COMMUNICATIONS

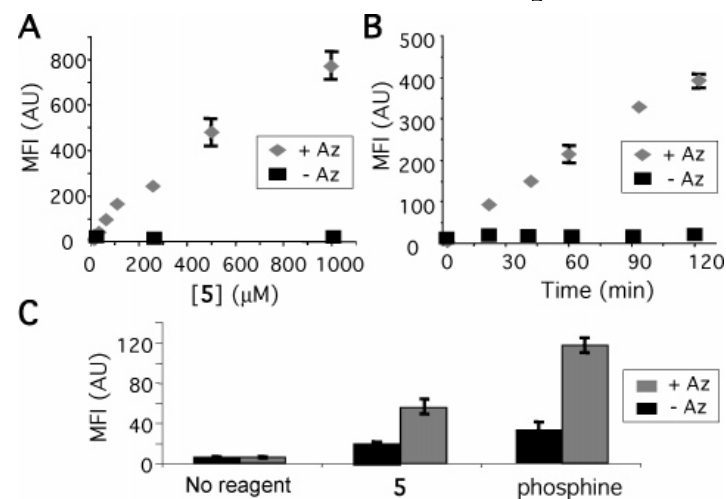
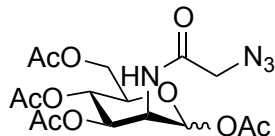
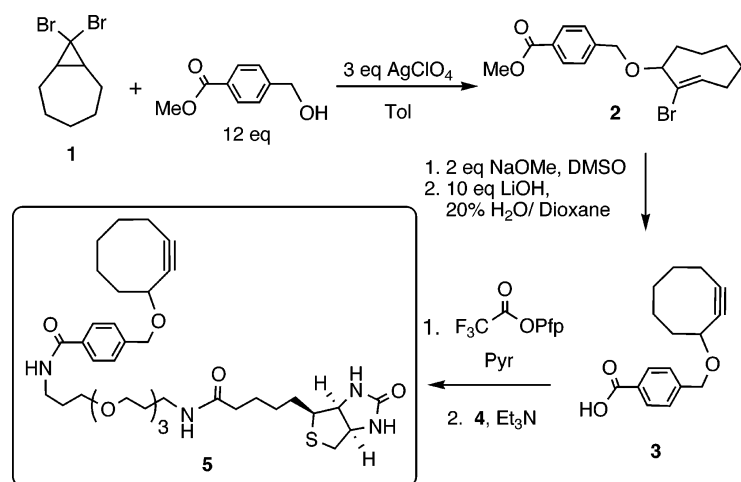
Published on Web 11/02/2004

## A Strain-Promoted [3 + 2] Azide - Alkyne Cycloaddition for Covalent Modification of Biomolecules in Living Systems

Nicholas J. Agard, Jennifer A. Prescher, and Carolyn R. Bertozzi\*

*Departments of Chemistry and Molecular and Cell Biology and Howard Hughes Medical Institute, UniVersity of California, and Materials Sciences Division, Lawrence Berkeley National Laboratory, Berkeley, California 94720*

Received August 19, 2004; E-mail: crb@cchem.berkeley.edu

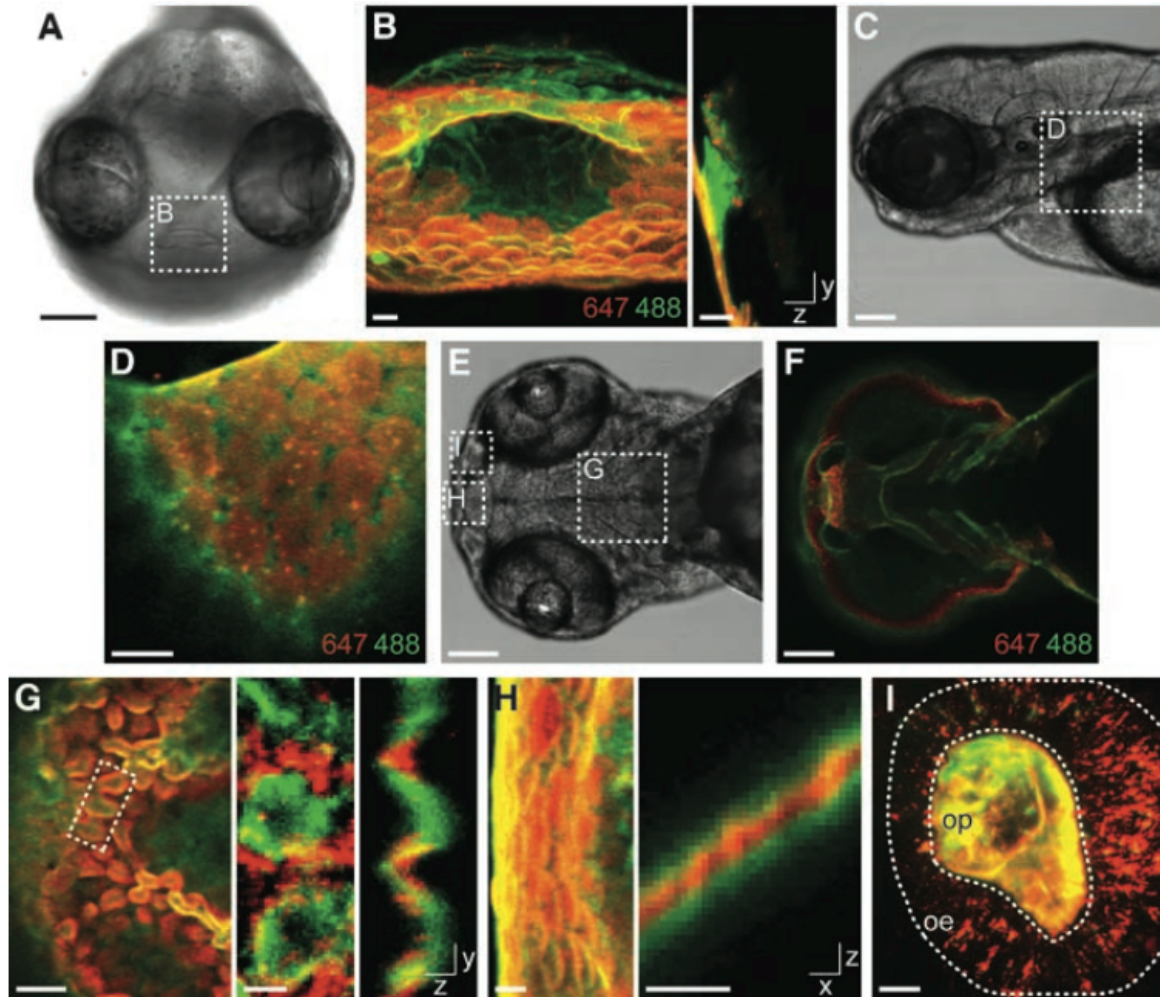


**Figure 3.** Cell-surface labeling with compound **5**. Jurkat cells were incubated in the presence (+Az) or absence (-Az) of 25 μM Ac<sub>4</sub>ManNAz for 3 d. (A) The cells were reacted with various concentrations of **5** for 1 h at room temperature and treated with FITC - avidin; mean fluorescence intensity (MFI) was determined by flow cytometry. (B) Cells were incubated with 250 μM **5** at room temperature and analyzed as in A. (C) Cells were incubated with 100 μM probe for 1 h at room temperature and analyzed as in A. Error bars represent the standard deviation from three replicates. AU ) arbitrary fluorescence units.

# Zebrafish embryos

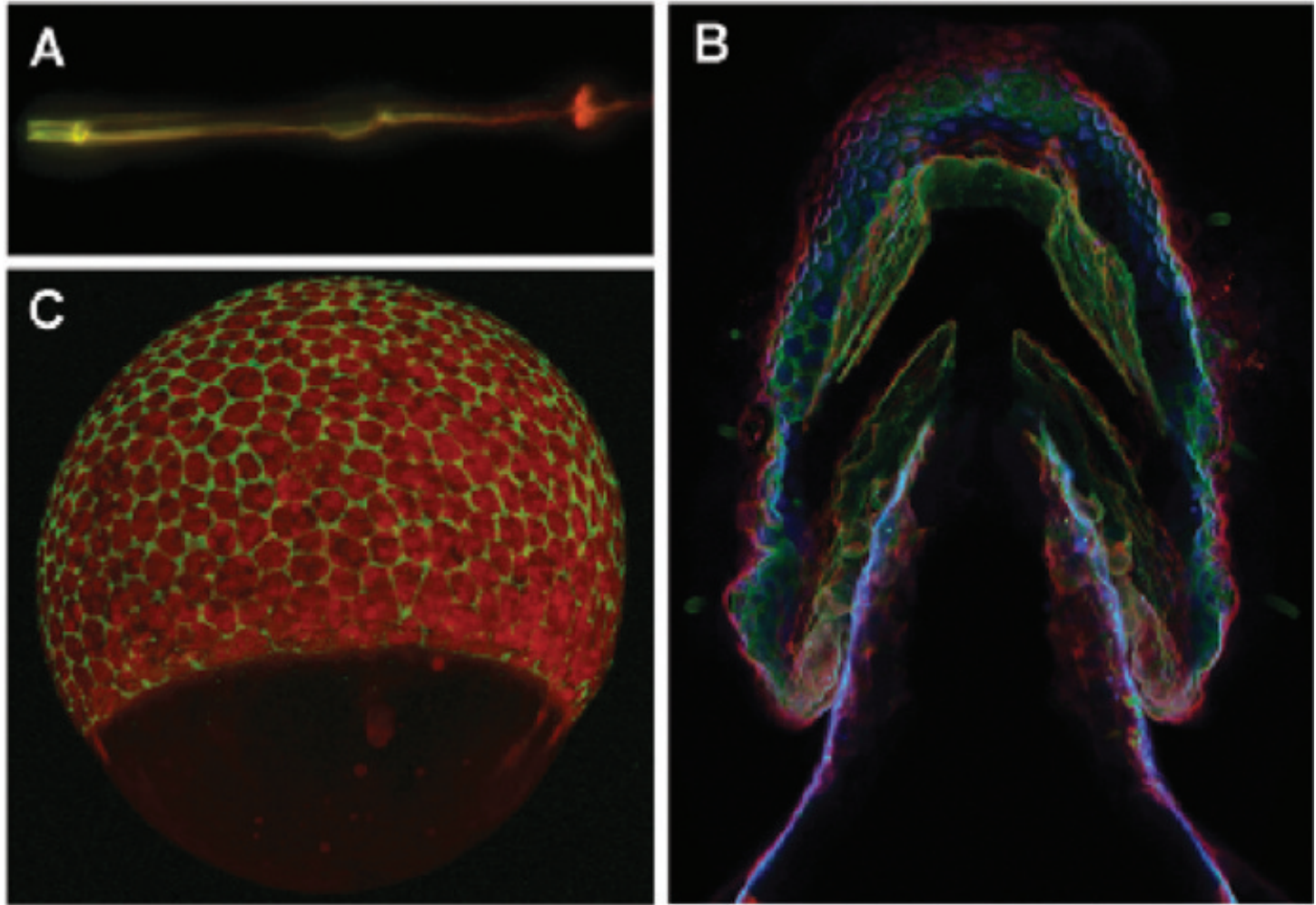
## In Vivo Imaging of Membrane-Associated Glycans in Developing Zebrafish

Scott T. Laughlin,<sup>1\*</sup> Jeremy M. Baskin,<sup>1\*</sup> Sharon L. Amacher,<sup>2</sup> Carolyn R. Bertozzi<sup>1,2,3,4†</sup>



Science, 2008

# C. elegans / Zebrafish

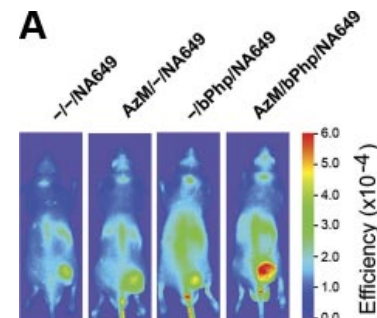
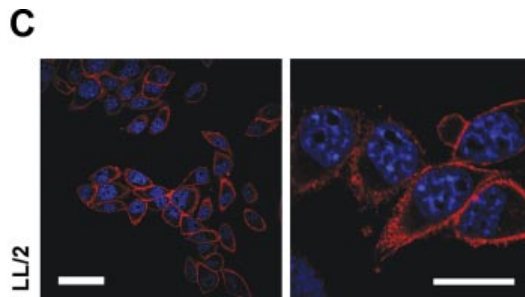
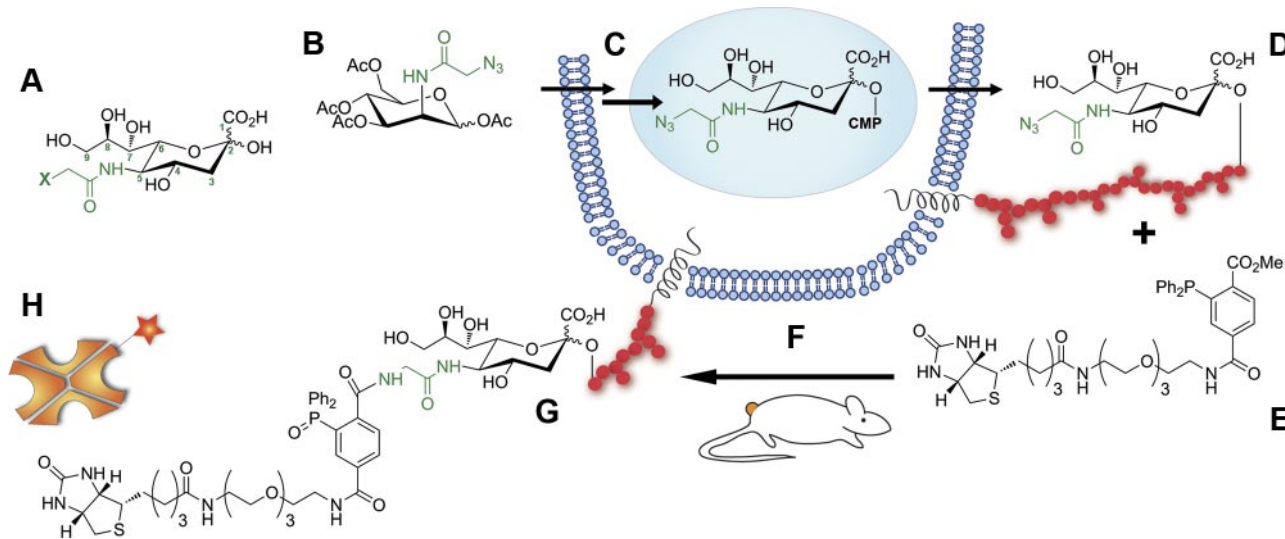


# Sialylated tumor glycan imaging

## Imaging sialylated tumor cell glycans *in vivo*

André A. Neves,\* Henning Stöckmann,\*<sup>†</sup> Rebecca R. Harmston,\* Helen J. Pryor,\*  
Israt S. Alam,\* Heather Ireland-Zecchini,\* David Y. Lewis,\* Scott K. Lyons,\*  
Finian J. Leeper,<sup>†</sup> and Kevin M. Brindle\*<sup>†,1</sup>

*FASEB J* 2011, 25, 2528-2537.

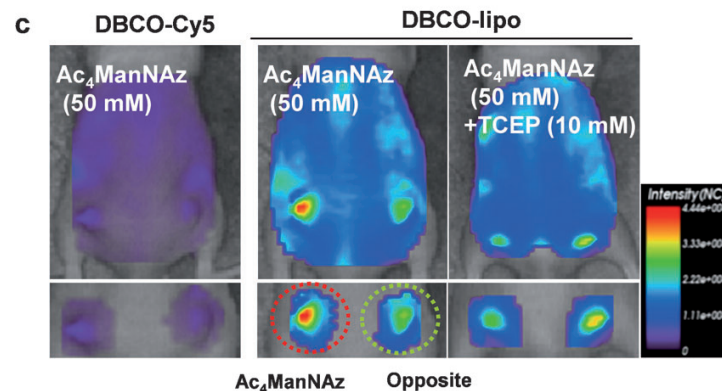
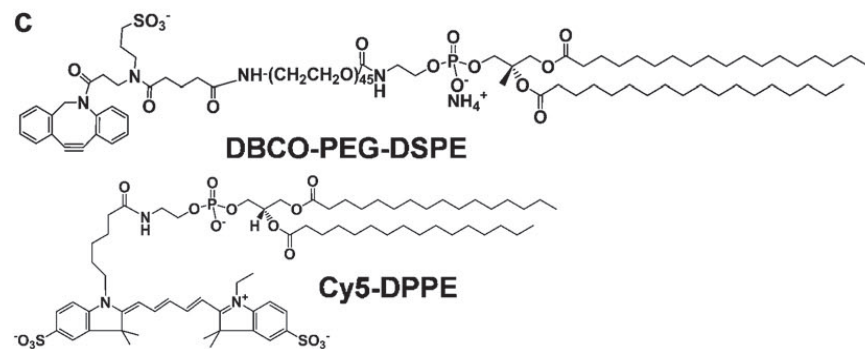
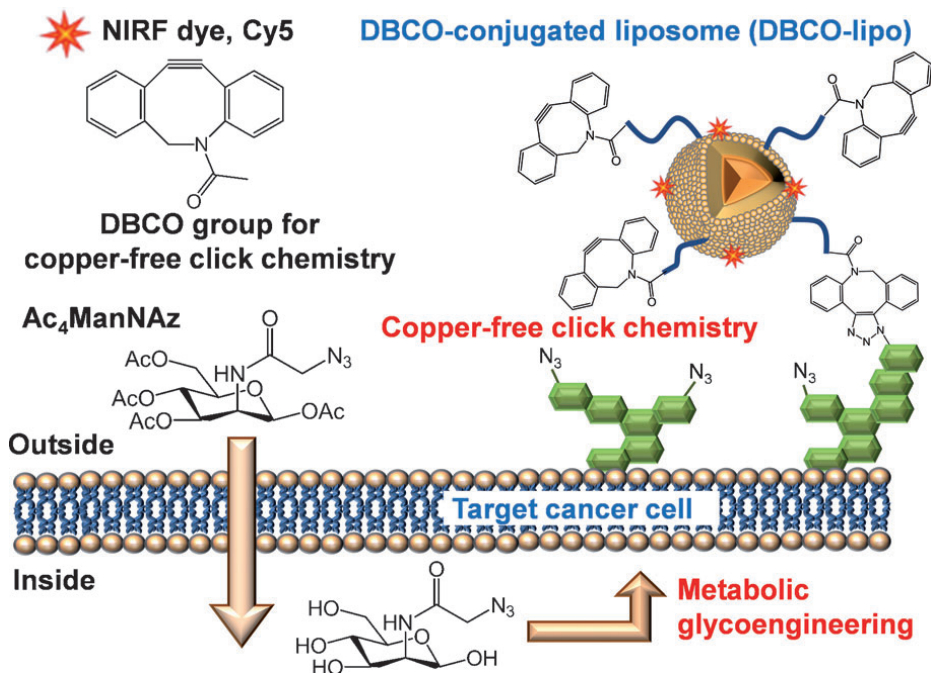


# Tumor-targeting

## Bioorthogonal Copper-Free Click Chemistry In Vivo for Tumor-Targeted Delivery of Nanoparticles\*\*

Heebeom Koo, Sangmin Lee, Jin Hee Na, Sun Hwa Kim, Sei Kwang Hahn, Kuiwon Choi, Ick Chan Kwon, Seo Young Jeong, and Kwangmeyung Kim\*

Angew. Chem. Int. Ed. 2012, 51, 1–6

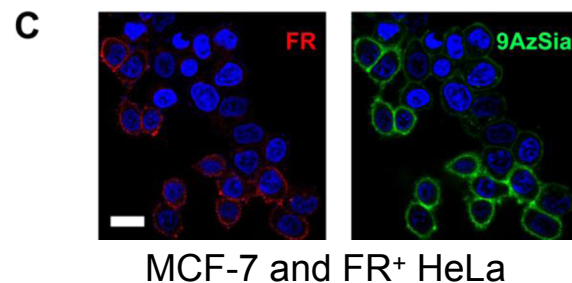
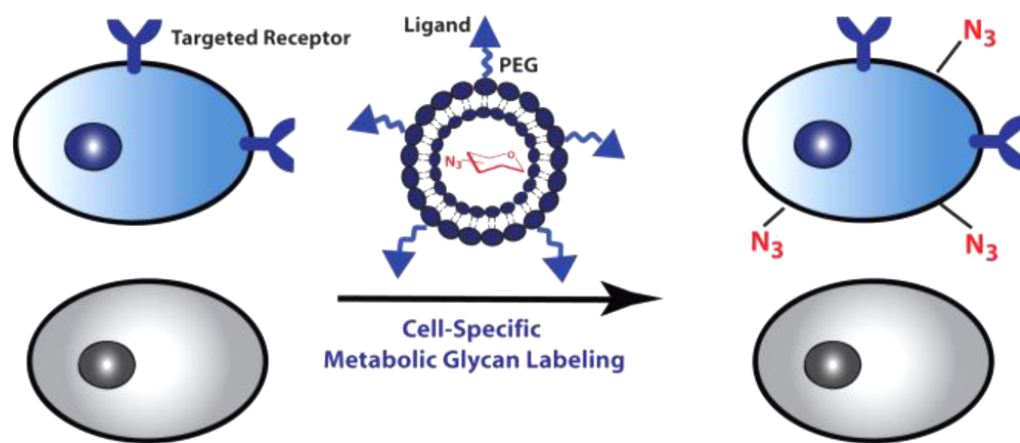
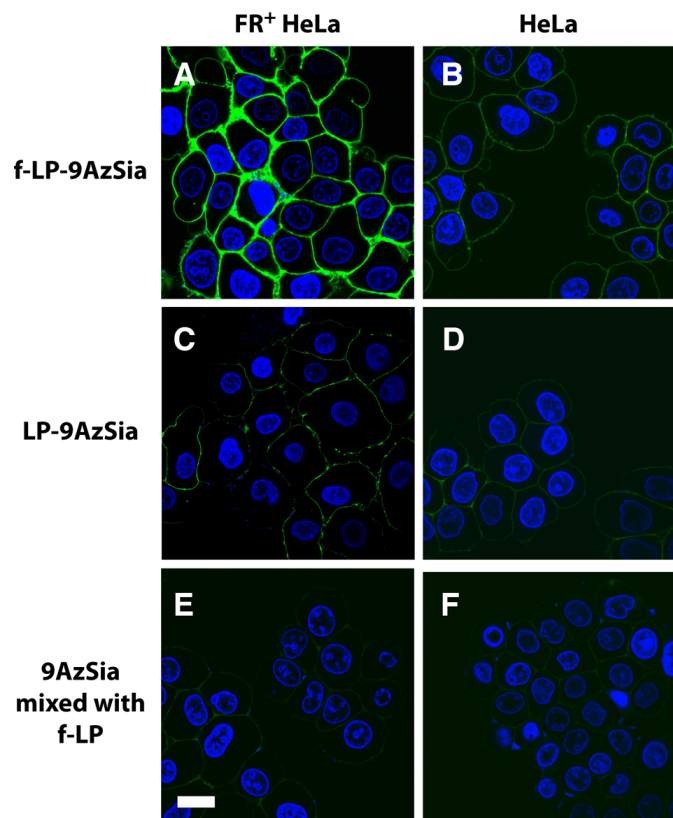


# Cell-selective glycan labeling

## Cell-Selective Metabolic Glycan Labeling Based on Ligand-Targeted Liposomes

Ran Xie,<sup>†,‡</sup> Senlian Hong,<sup>¶,‡</sup> Lianshun Feng,<sup>†</sup> Jie Rong,<sup>†</sup> and Xing Chen<sup>\*,†,§,||</sup>

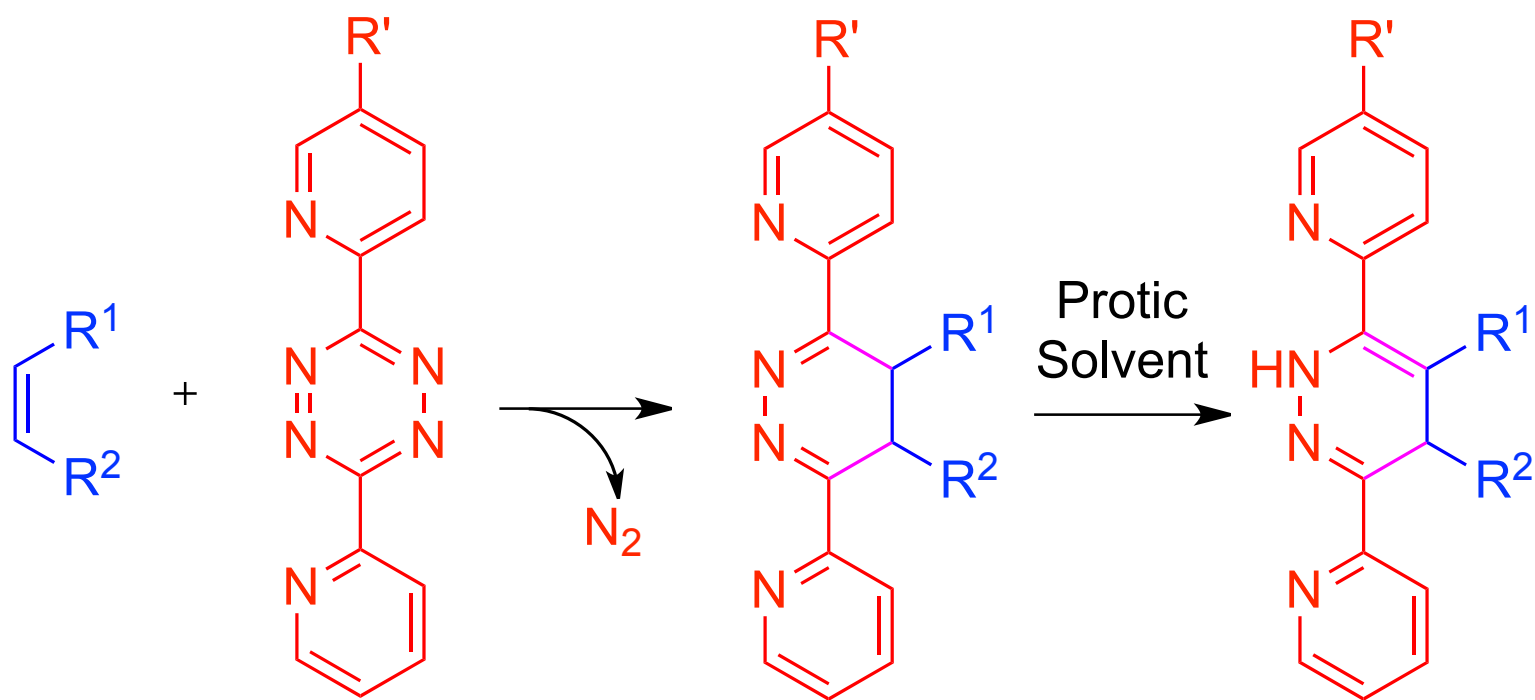
*J. Am. Chem. Soc.* 2012, 134, 9914–9917



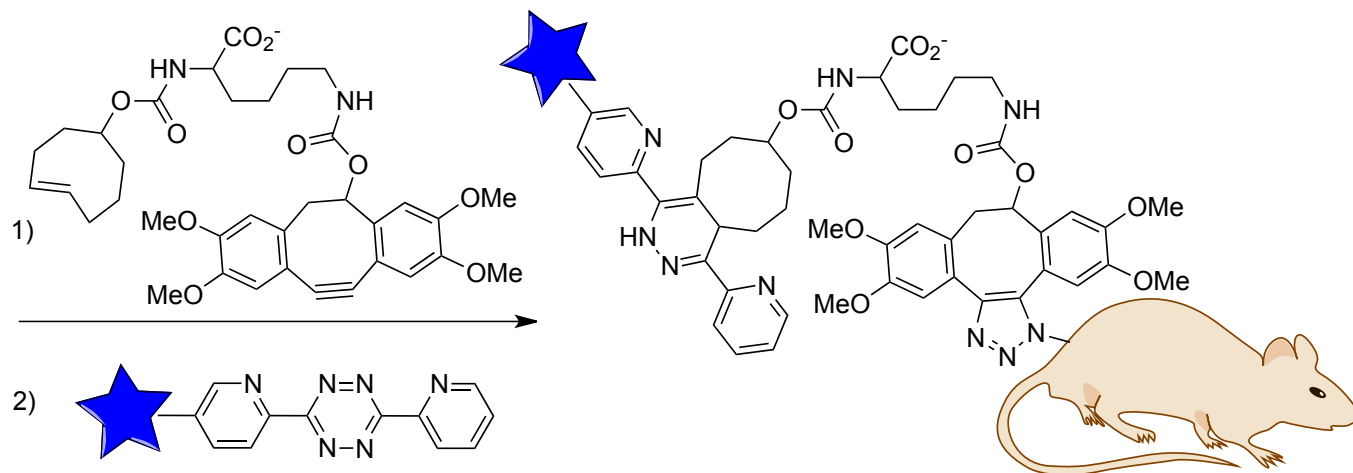
MCF-7 and FR<sup>+</sup> HeLa



# Bioorthogonal Chemistry - Inverse electron demand Diels Alder

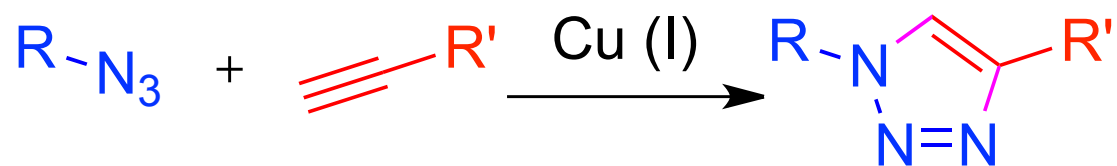


# Sequential double click

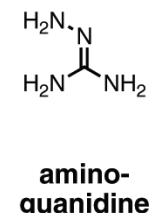
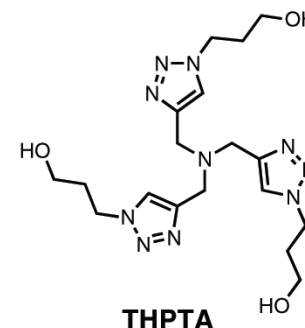
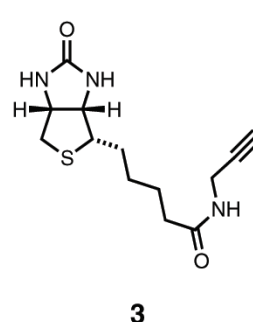
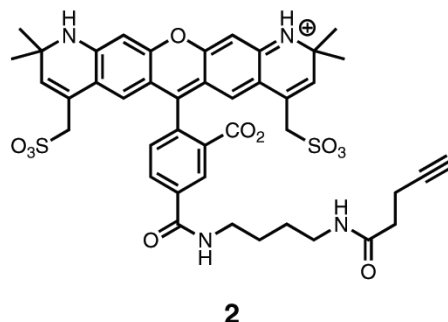
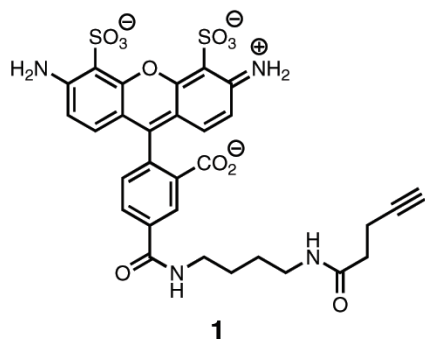
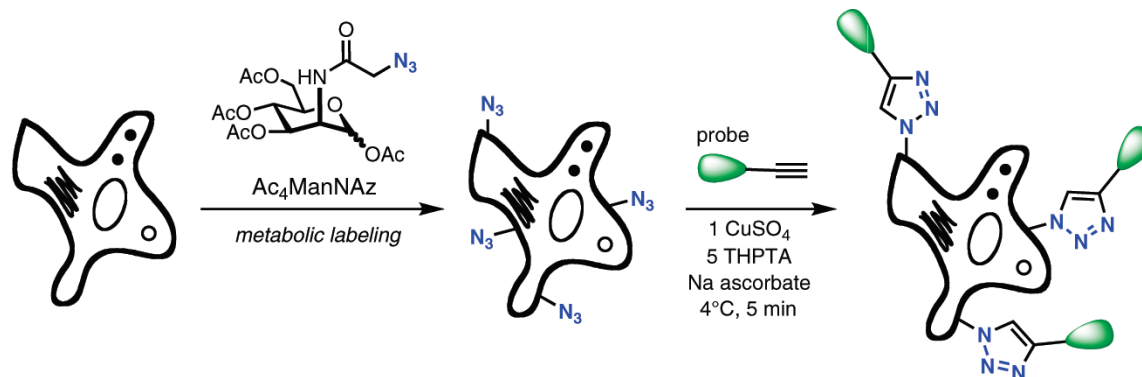


# Bioorthogonal chemistry - 2+3 Cycloadditions

## CuAAC



# Copper-catalyzed click chemistry



1912

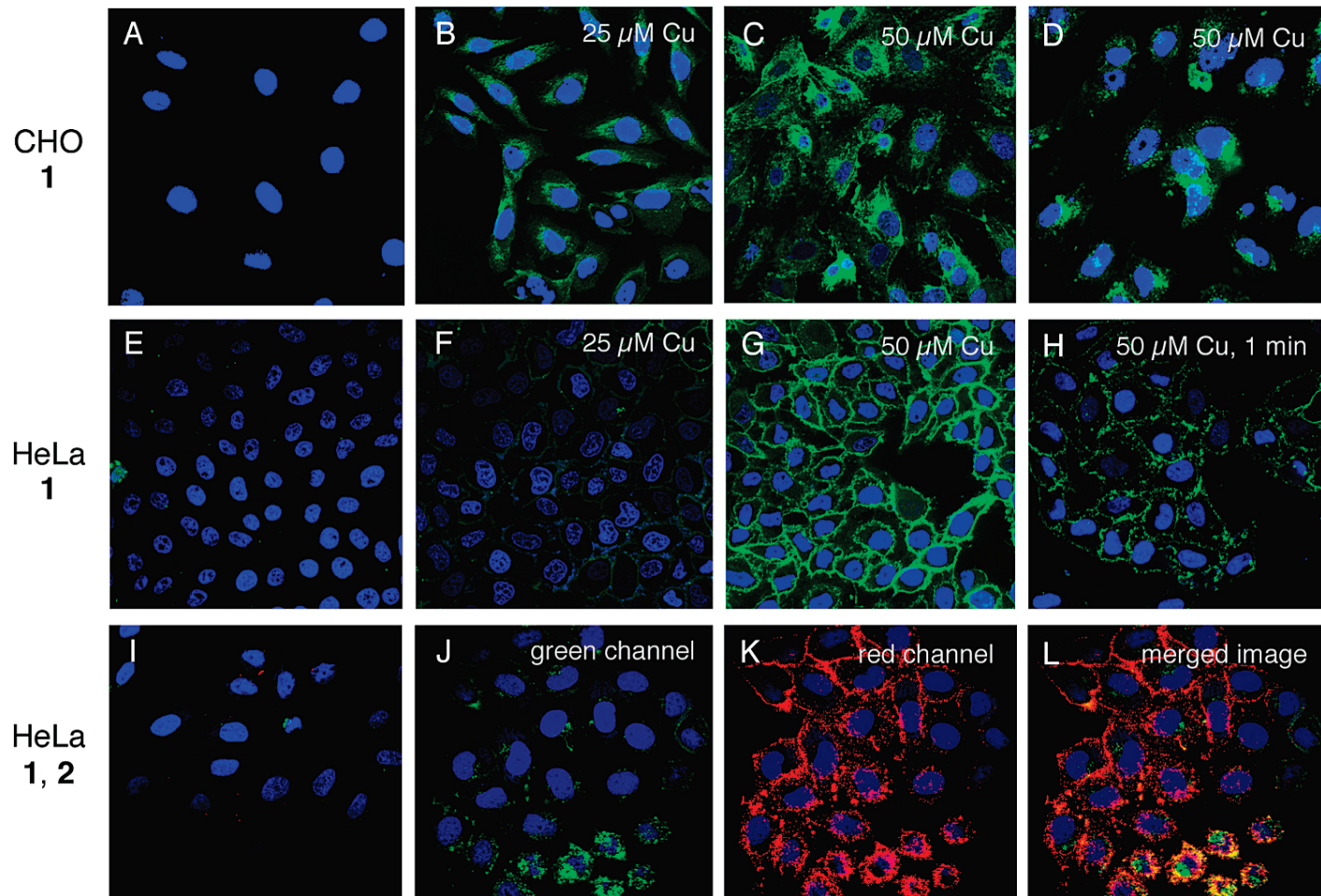
Bioconjugate Chem. 2010, 21, 1912–1916

## Labeling Live Cells by Copper-Catalyzed Alkyne - Azide Click Chemistry

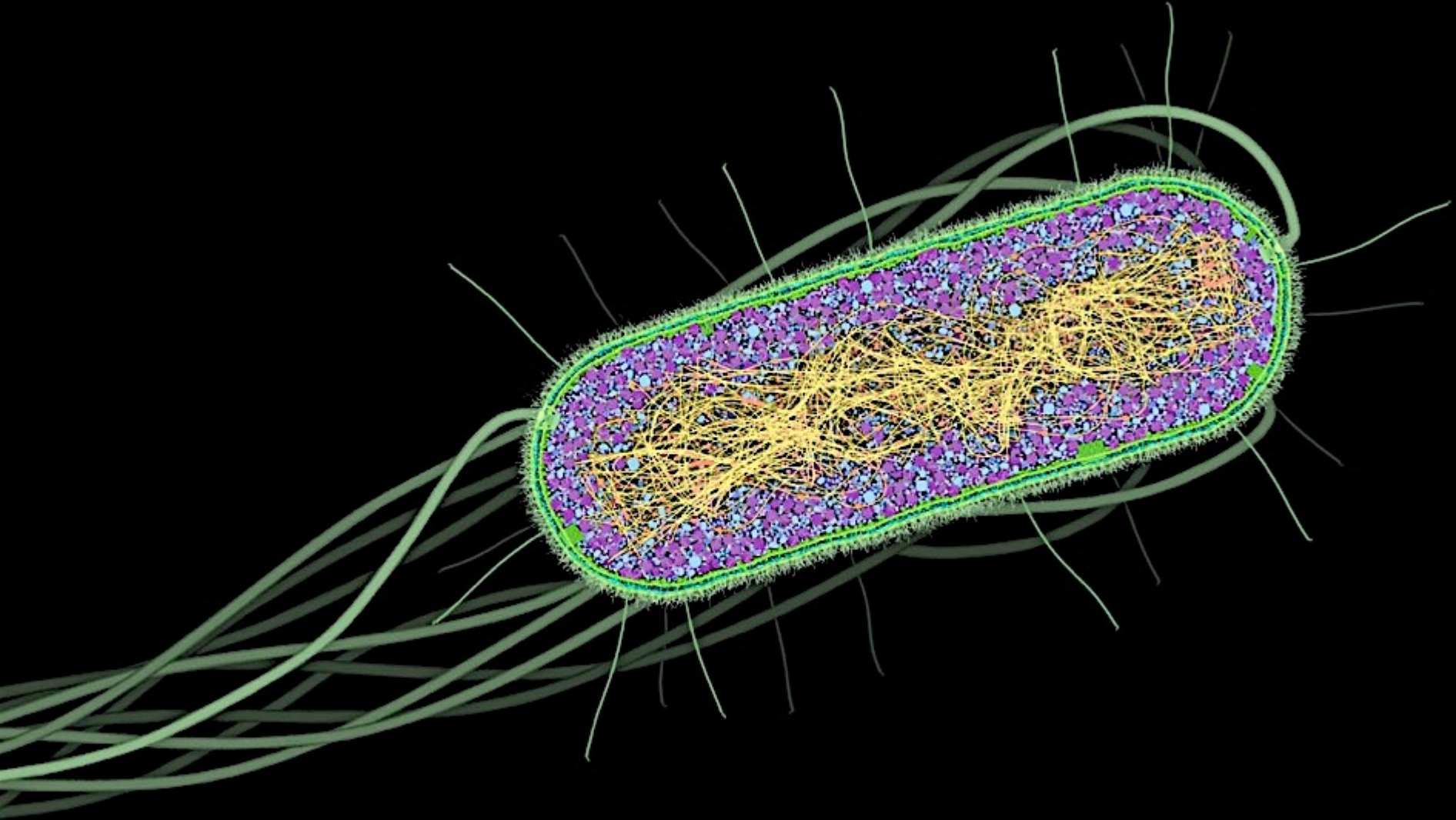
Vu Hong,<sup>S,†</sup> Nicole F. Steinmetz,<sup>S,‡</sup> Marianne Manchester,<sup>l</sup> and M. G. Finn<sup>\*,†</sup>

Department of Chemistry and The Skaggs Institute for Chemical Biology, Department of Cell Biology and Center for Integrative Molecular Biosciences, The Scripps Research Institute, 10550 North Torrey Pines Road, La Jolla, California, Skaggs School of Pharmacy and Pharmaceutical Sciences, University of California San Diego, 9500 Gilman Drive, La Jolla, California. Received June 17, 2010; Revised Manuscript Received August 24, 2010

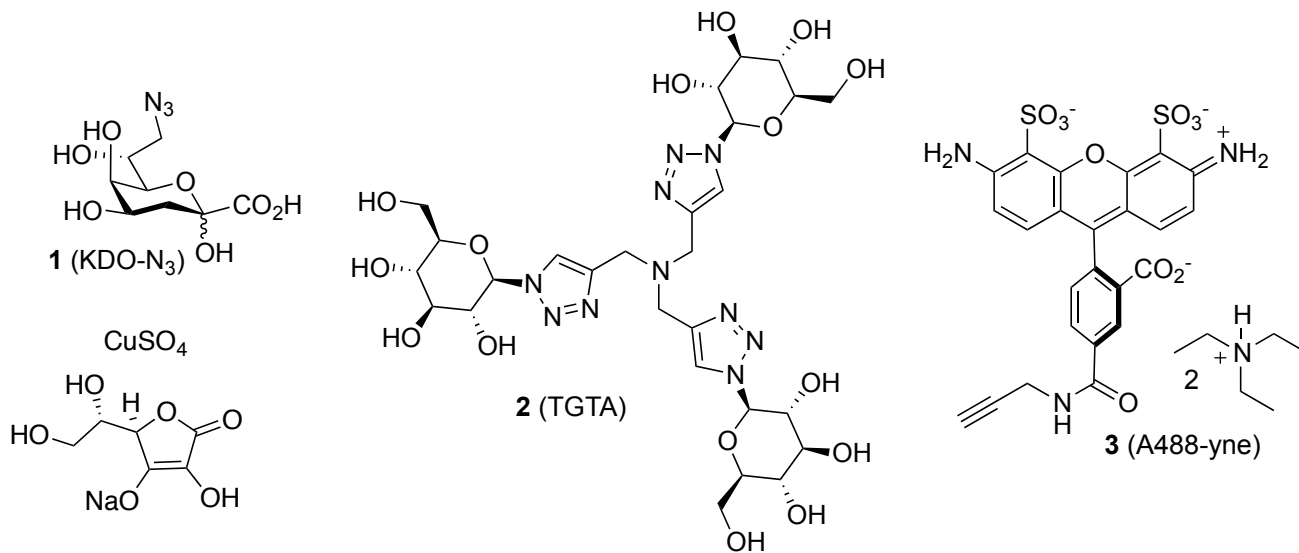
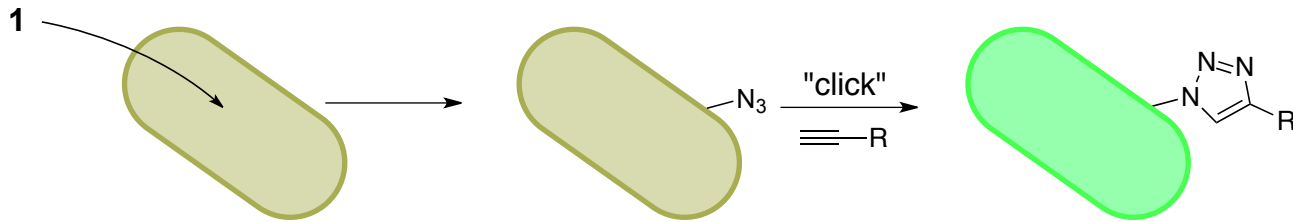
# Copper-catalyzed click chemistry



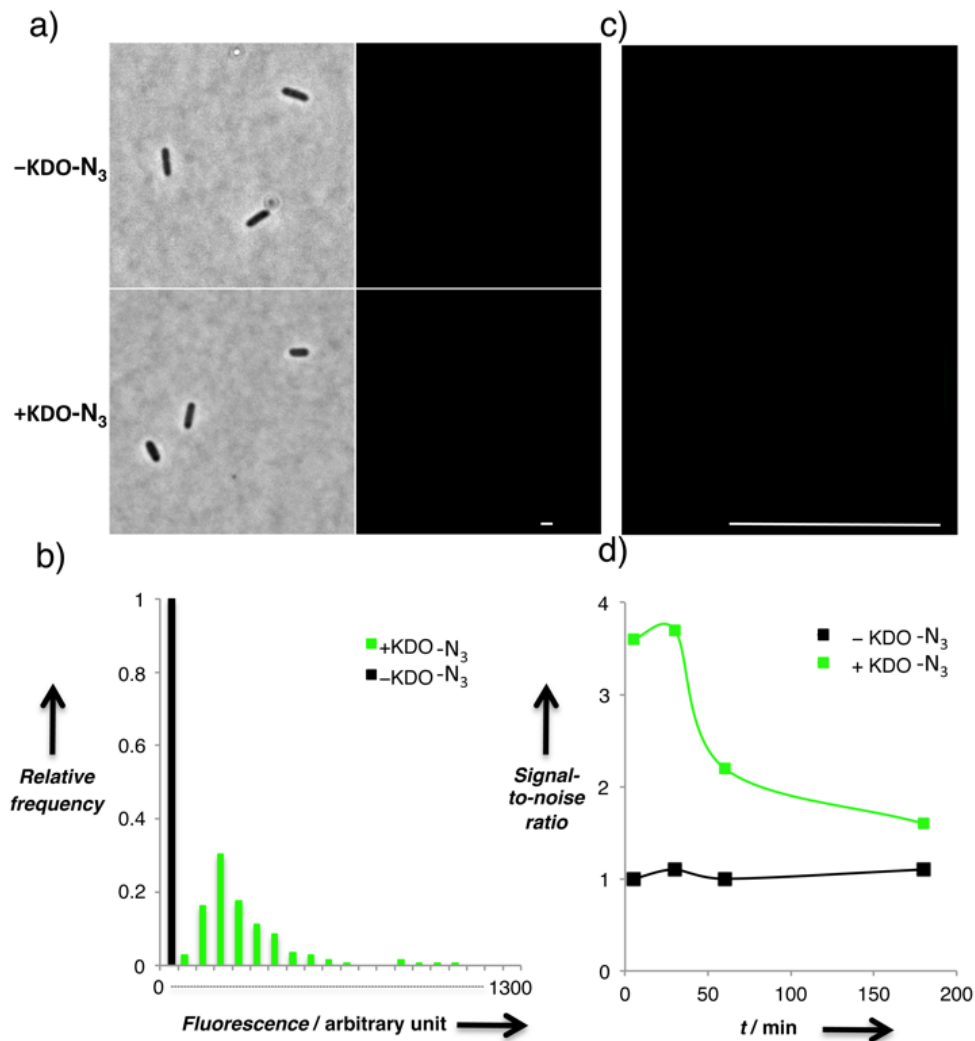
# METABOLIC LIPOPOLYSACCHARIDE LABELING



# Click-chemistry

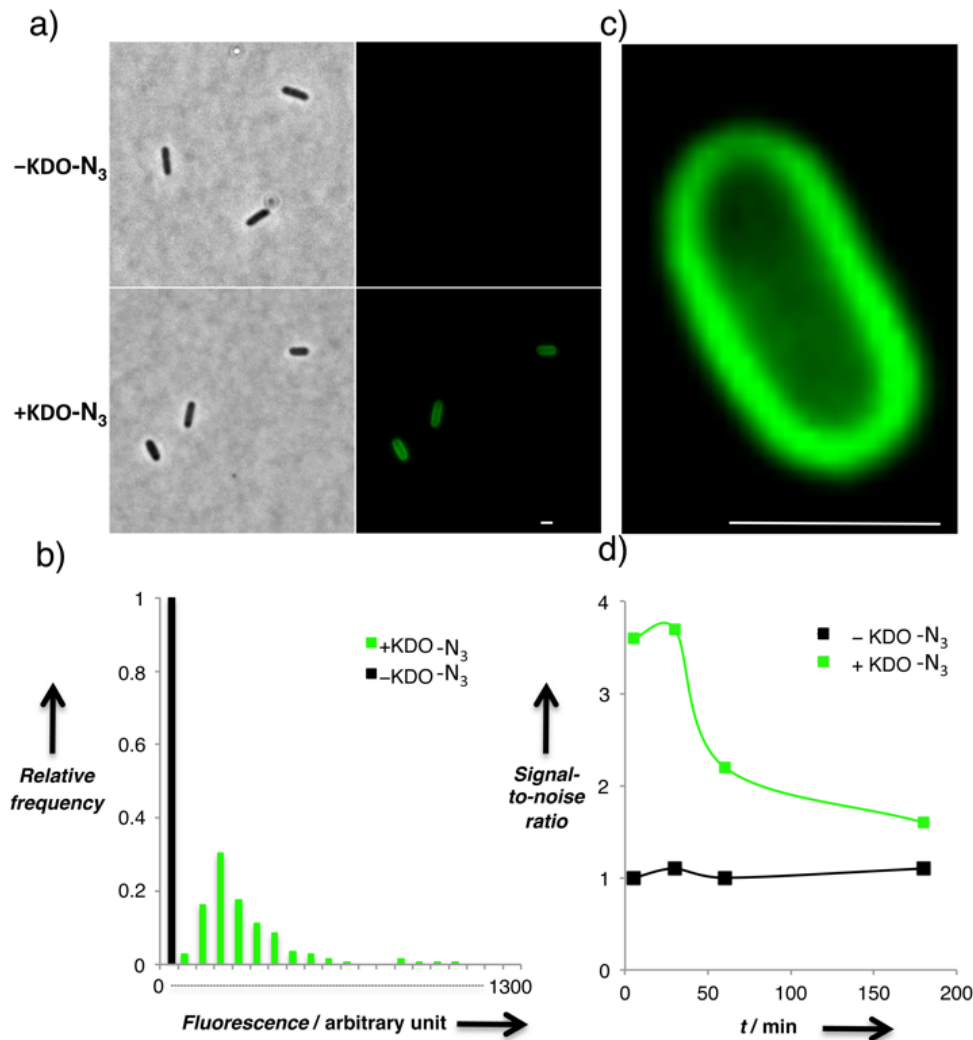


# *E. coli* labeling





# *E. coli* labeling

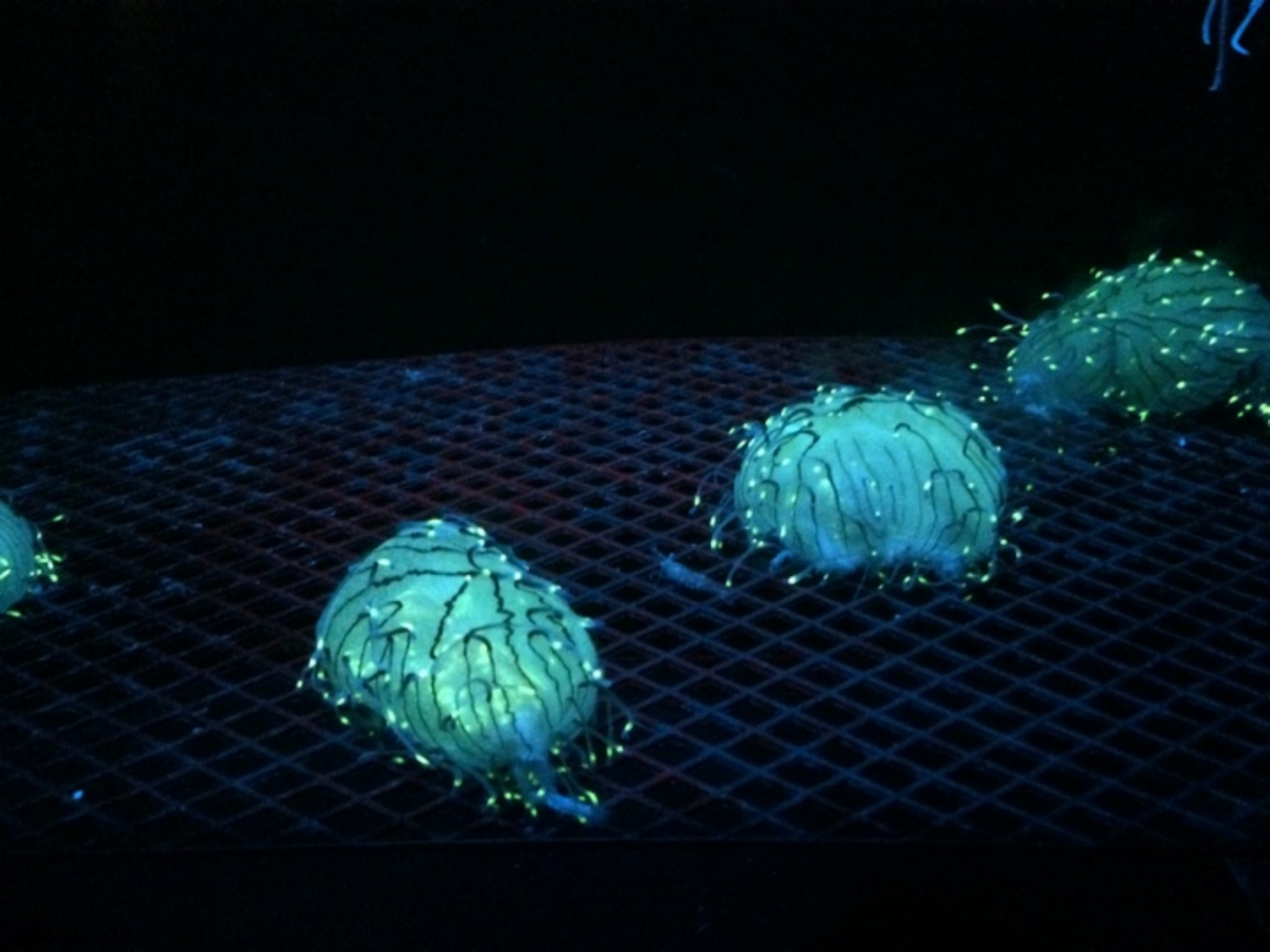


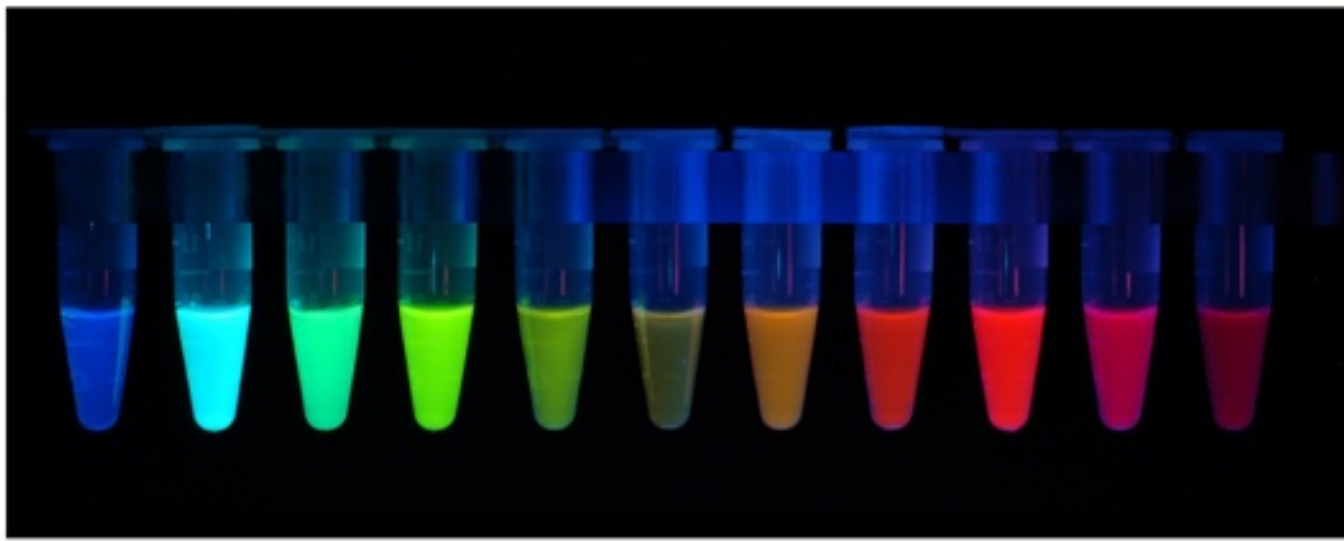
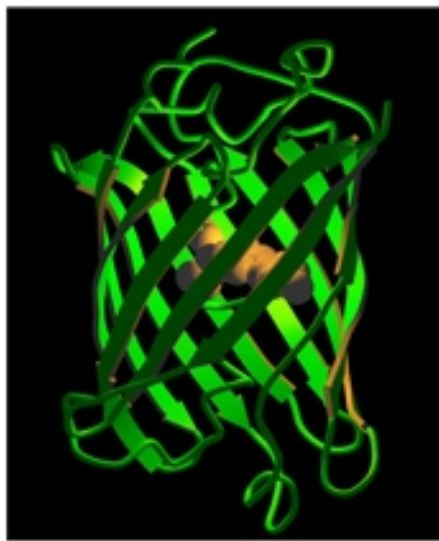
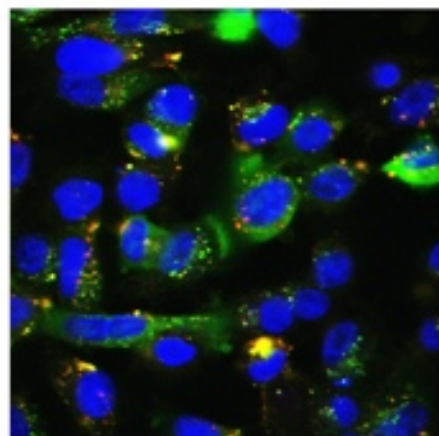
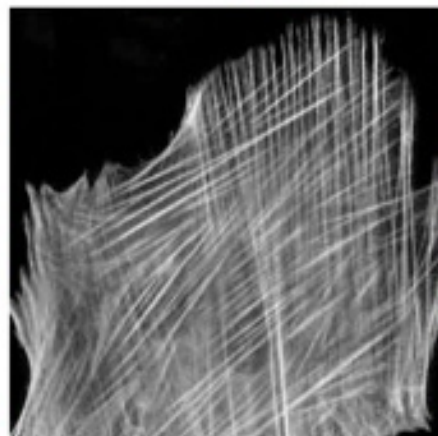
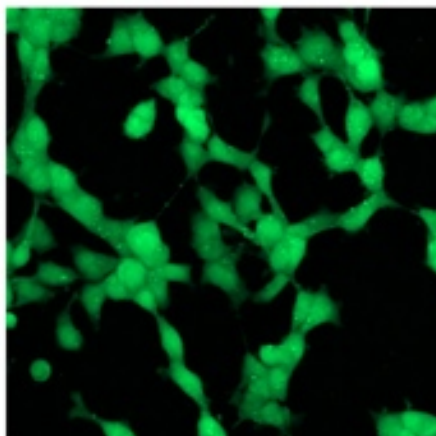
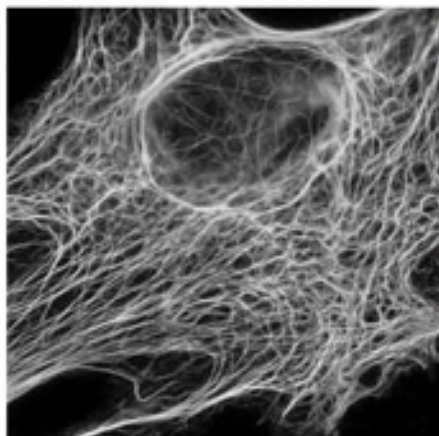
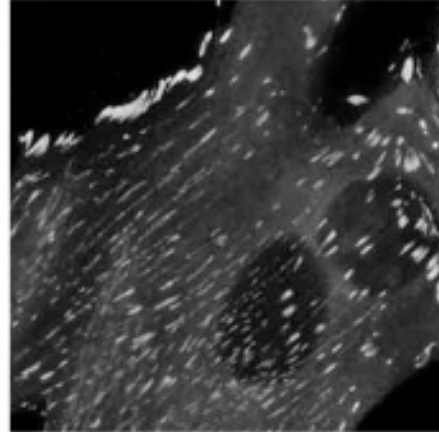
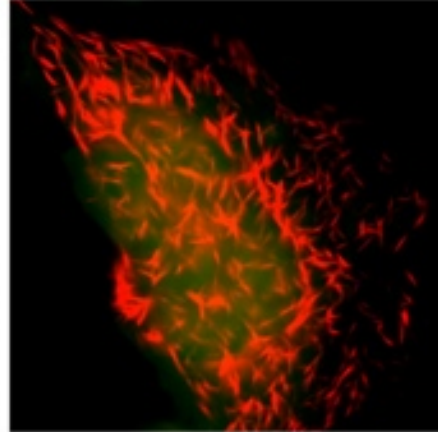
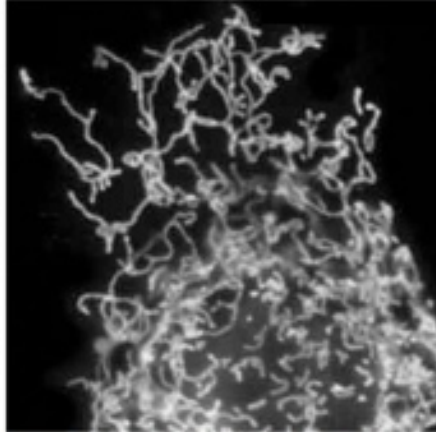
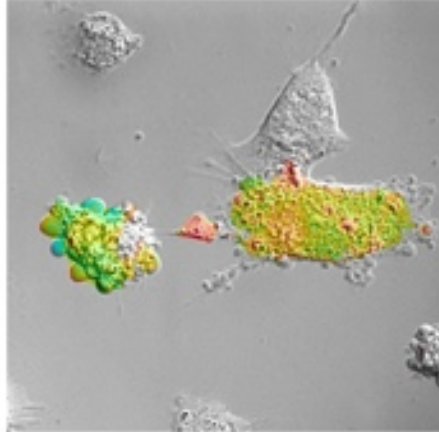
Labeling  
proteins  
*in vivo*

# Fluorescent proteins



Roger Y. Tsien





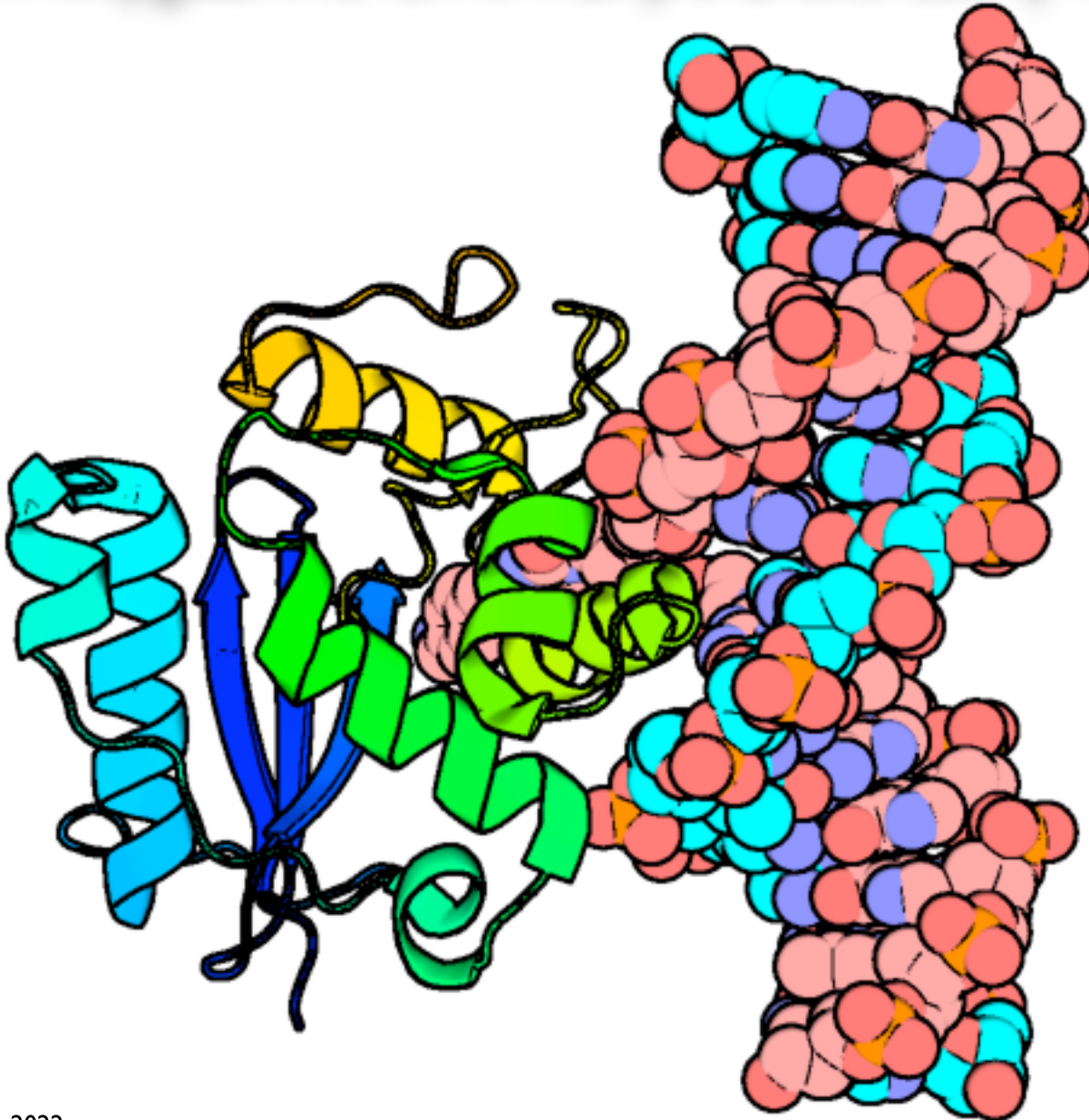
# SNAP-Tag - CLIP-Tag



Kai Johansson

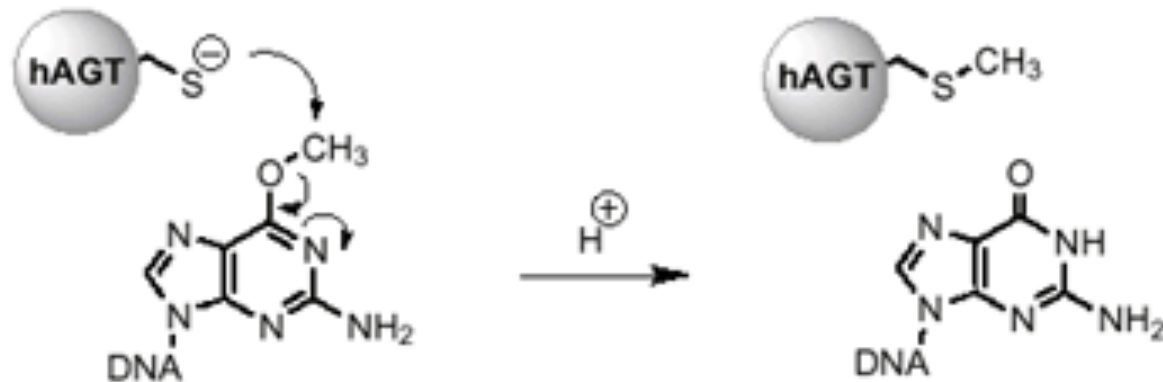


# O6-Alkylguanine-DNA alkyltransferase (AGT)

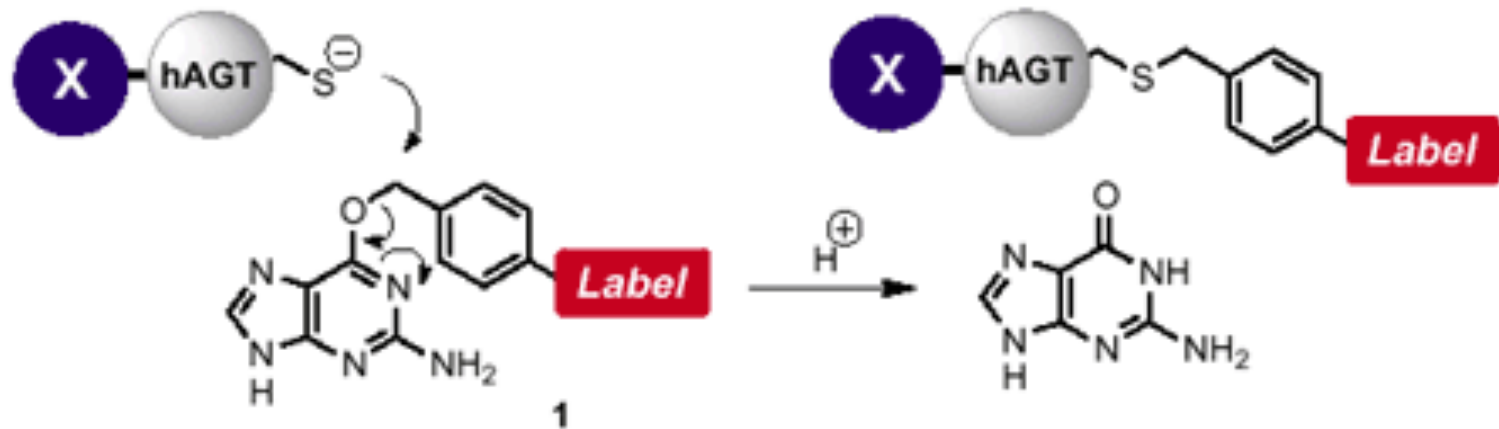


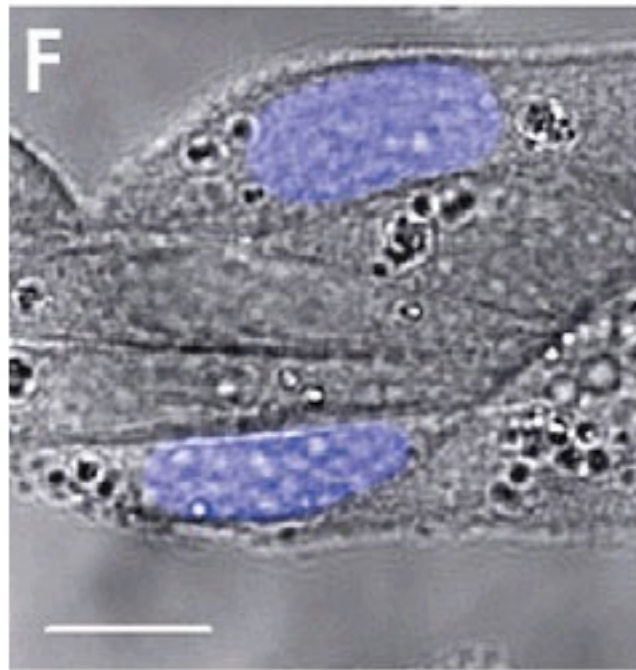
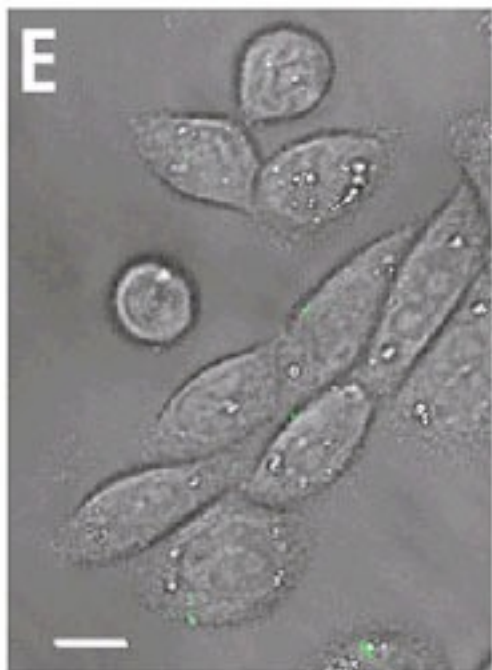
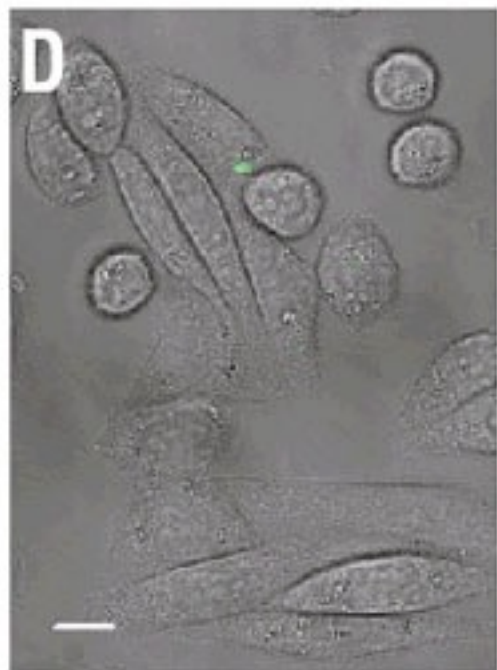
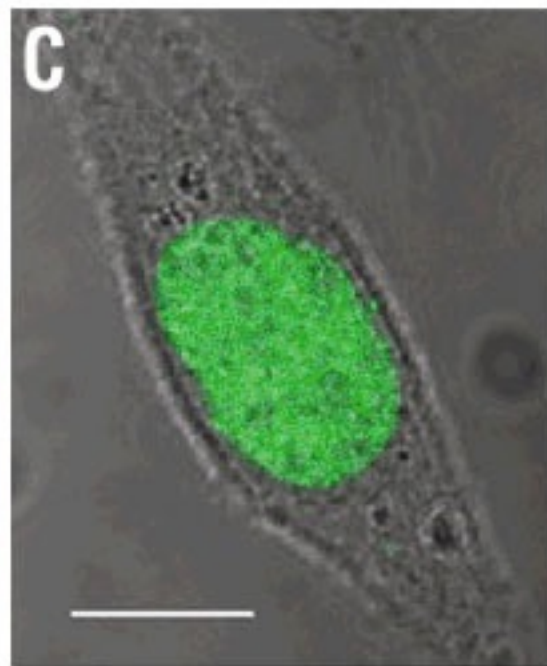
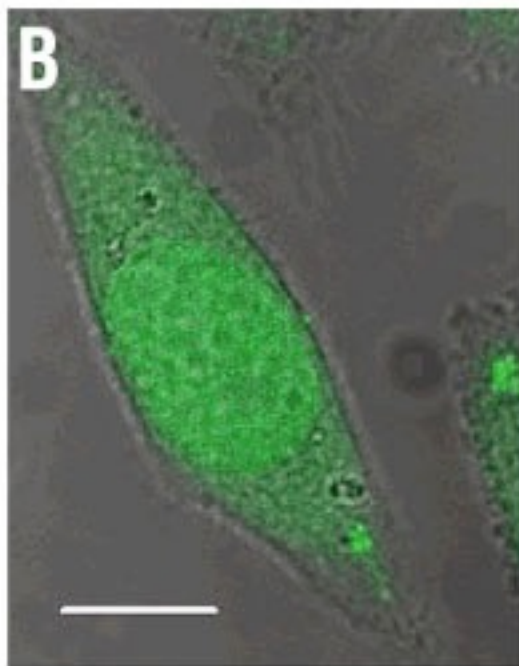
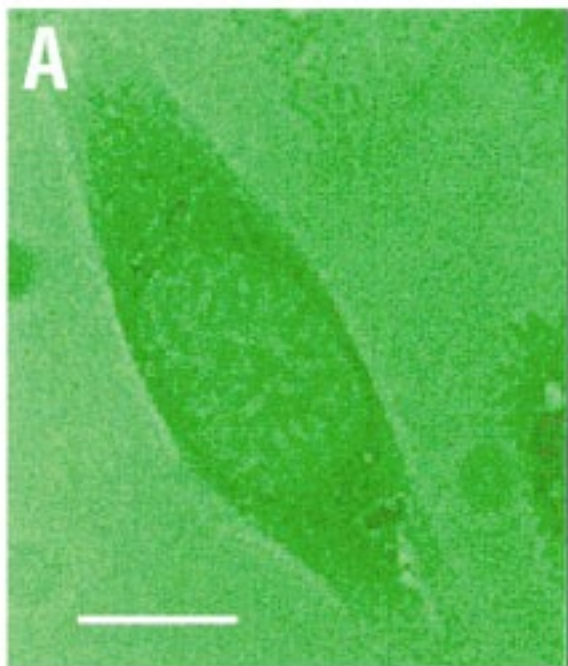
# SNAP-Tag

## A

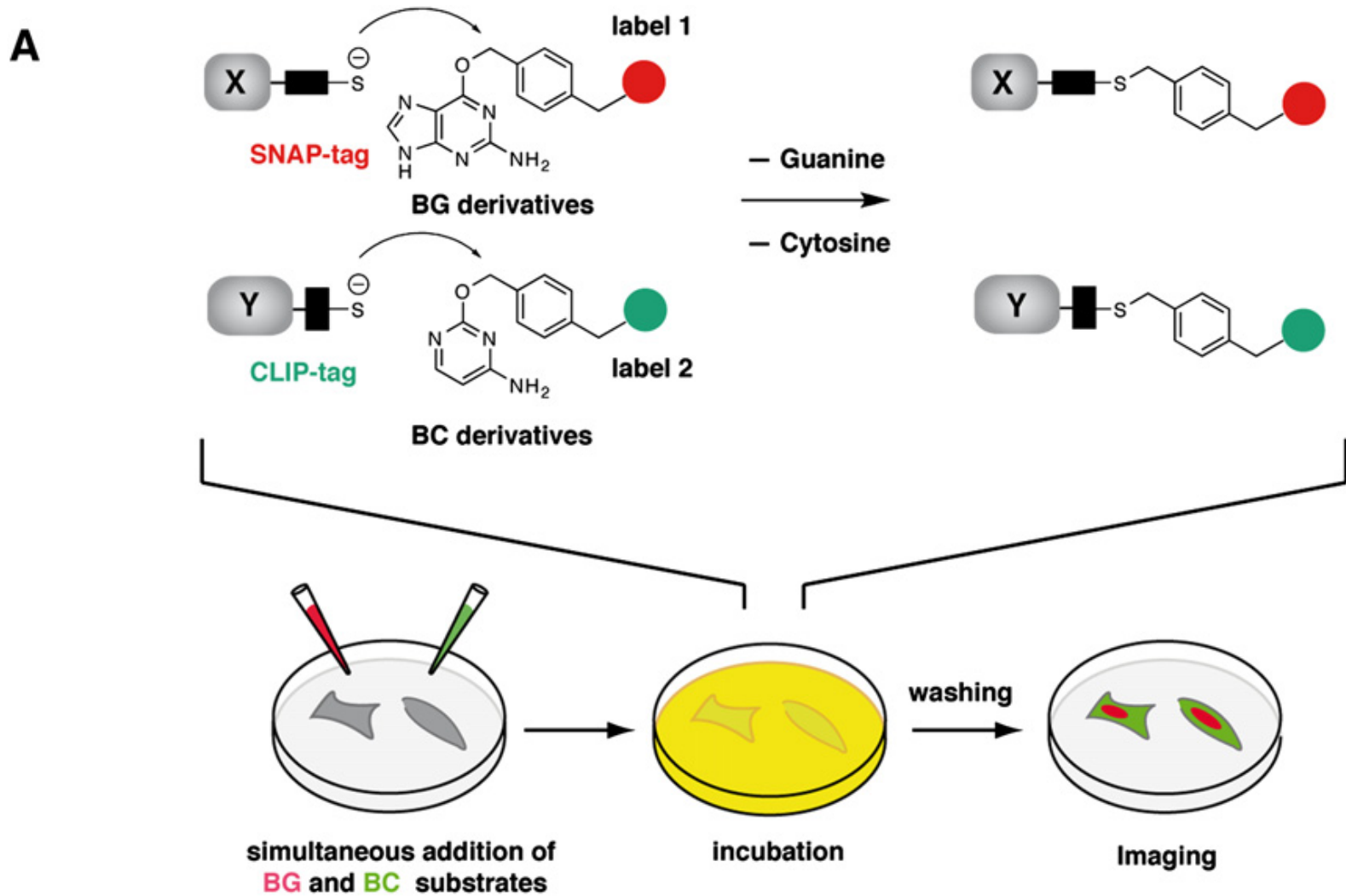


## B

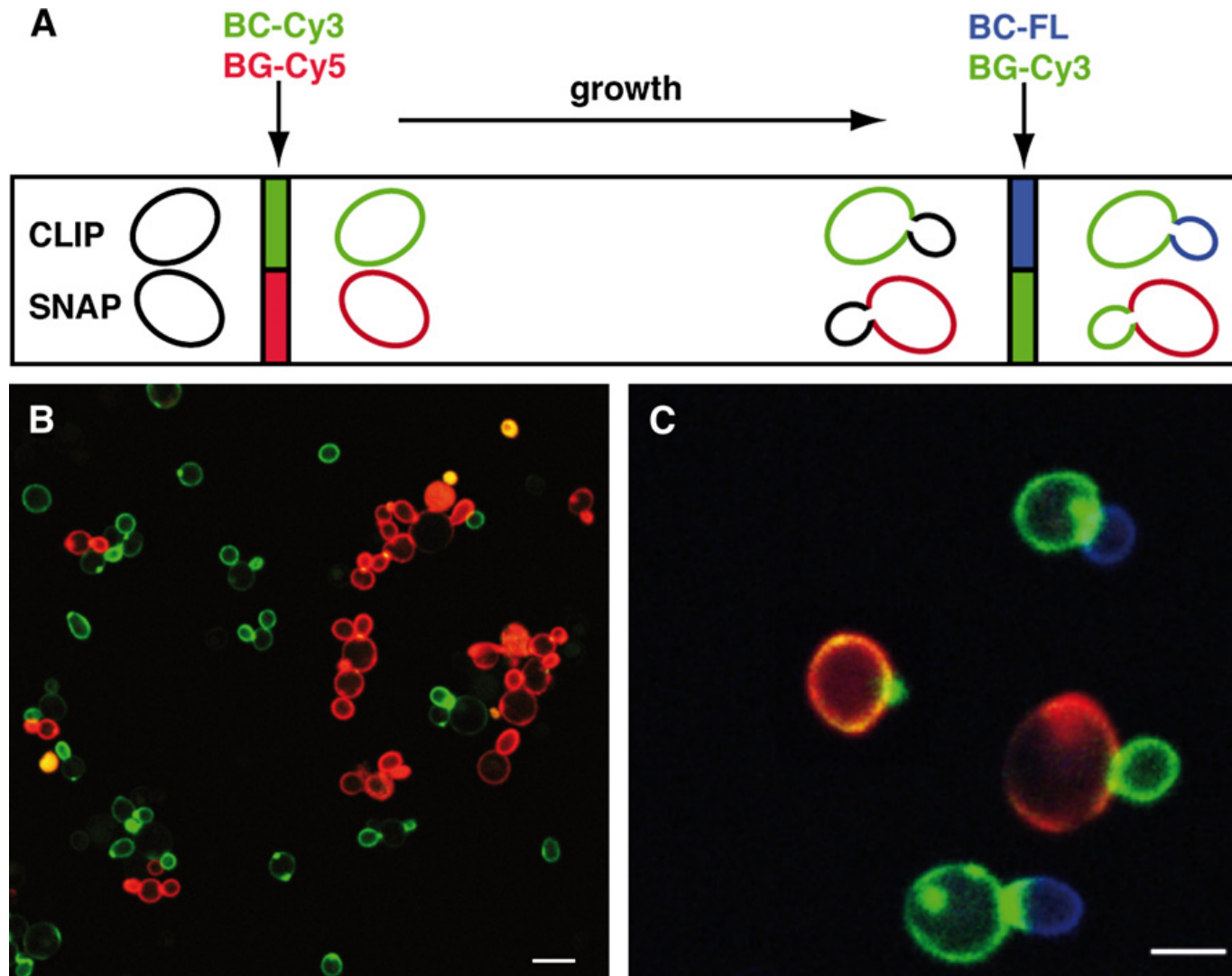




# CLIP-Tag



# Yeast labeling



# Expanding the genetic code



**Peter**  
Schultz  
TSRI: The California Institute  
for Biomedical Research  
**SPEAKER**

# Genetic code... RNA

nonpolar polar basic acidic (stop codon)

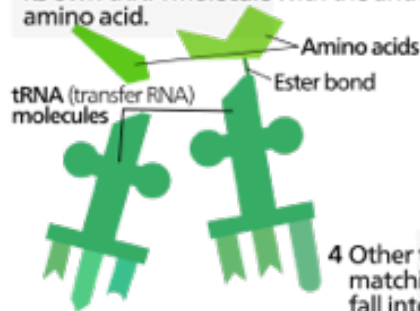
Standard genetic code

1st base	2nd base								3rd base
	U		C		A		G		
U	UUU	(Phe/F) Phenylalanine	UCU	(Ser/S) Serine	UAU	(Tyr/Y) Tyrosine	UGU	(Cys/C) Cysteine	U
	UUC		UCC		UAC		UGC		C
	UUA	(Leu/L) Leucine	UCA		UAA	Stop ( <i>Ochre</i> )	UGA	Stop ( <i>Opal</i> )	A
	UUG		UCG		UAG	Stop ( <i>Amber</i> )	UGG	(Trp/W) Tryptophan	G
C	CUU	(Leu/L) Leucine	CCU	(Pro/P) Proline	CAU	(His/H) Histidine	CGU	(Arg/R) Arginine	U
	CUC		CCC		CAC		CGC		C
	CUA		CCA		CAA	(Gln/Q) Glutamine	CGA		A
	CUG		CCG		CAG		CGG		G
A	AUU	(Ile/I) Isoleucine	ACU	(Thr/T) Threonine	AAU	(Asn/N) Asparagine	AGU	(Ser/S) Serine	U
	AUC		ACC		AAC		AGC		C
	AUA	ACA	AAA		(Lys/K) Lysine	AGA	(Arg/R) Arginine	A	
	AUG <sup>[A]</sup>	(Met/M) Methionine	ACG			AAG		AGG	G
G	GUU	(Val/V) Valine	GCU	(Ala/A) Alanine	GAU	(Asp/D) Aspartic acid	GGU	(Gly/G) Glycine	U
	GUC		GCC		GAC		GGC		C
	GUA		GCA		GAA	(Glu/E) Glutamic acid	GGA		A
	GUG		GCG		GAG		GGG		G

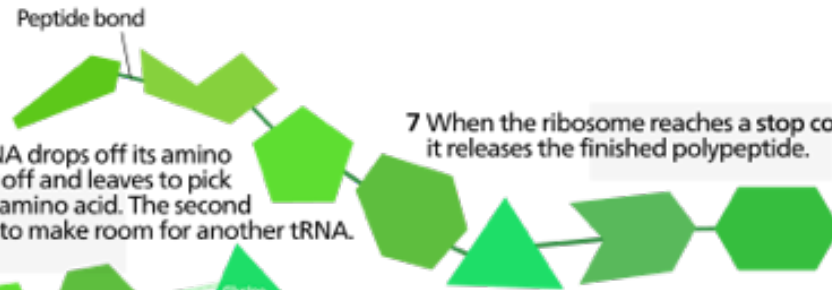


# RNA Translation

1 An enzyme called *aminoacyl tRNA synthetase* (not shown) attaches amino acids to their corresponding tRNA molecules using energy from ATP. Each amino acid has its own tRNA molecule with the anticodon for that amino acid.



5 The first tRNA drops off its amino acid, breaks off and leaves to pick up another amino acid. The second moves over to make room for another tRNA.



7 When the ribosome reaches a stop codon, it releases the finished polypeptide.

4 Other tRNAs with anticodons matching the mRNA codons fall into place in the ribosome.

6 tRNA after tRNA drop off their amino acids. They form a chain of amino acids linked by peptide bonds.

3 It moves along the mRNA until it finds a start codon. There, the first tRNA and the large ribosomal subunit join it.

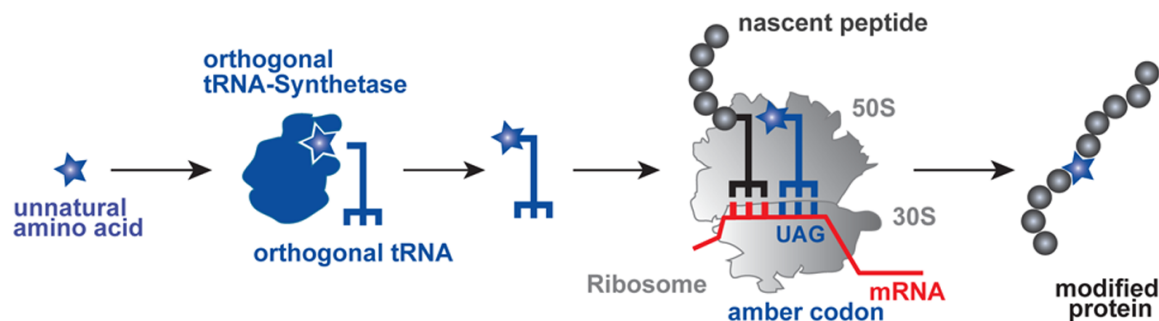


2 A small ribosomal subunit attaches itself to the 5' end of an mRNA strand.



cytoplasm  
**RNA translation**  
a part of **protein synthesis**

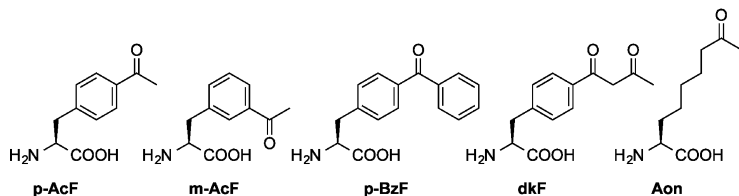
# Expanding the genetic code



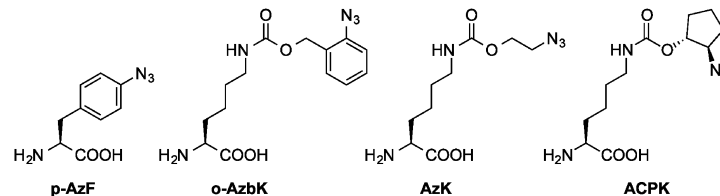
**Figure 16.** Expanding the genetic code. An unnatural amino acid (blue star), added to the cell growth medium, is specifically recognized by an orthogonal aminoacyl tRNA synthetase and attached to an orthogonal amber suppressor tRNA, which is decoded by the ribosome in response to an amber codon (UAG) introduced into the gene of interest, allowing the synthesis of a protein with a site-specifically introduced unnatural amino acid.

# Non natural amino acids...

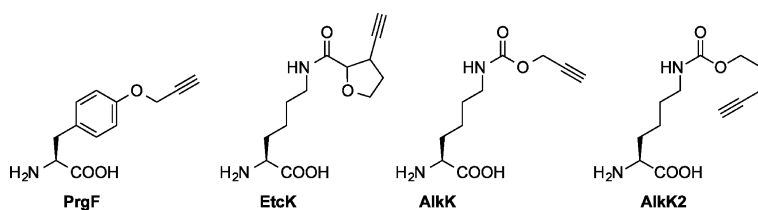
## "Ketones"



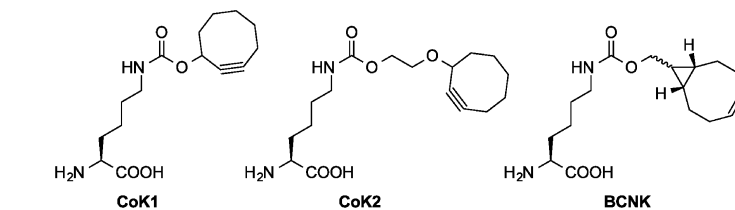
## "Azides"



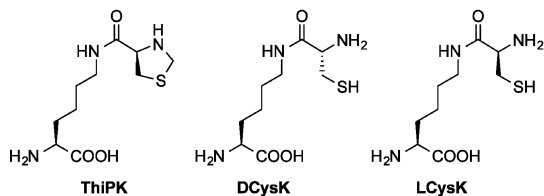
## "Terminal alkynes"



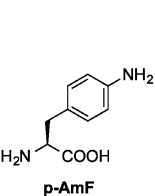
## "Strained alkynes"



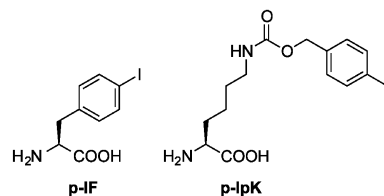
## "1,2-Aminothiols"



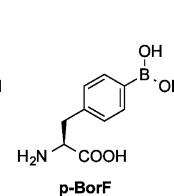
## "Aniline"



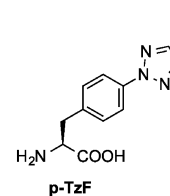
## "Iodides"



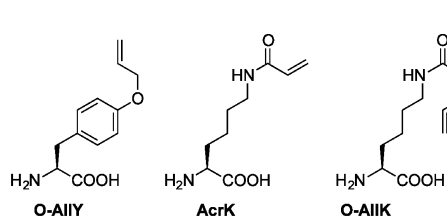
## "Boronate"



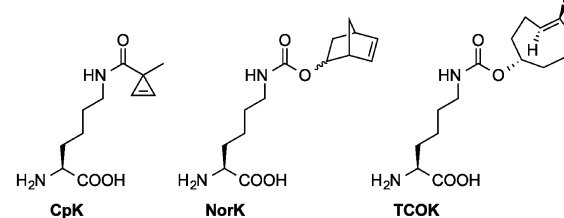
## "Tetrazole"



## "Alkenes"



## "Strained alkenes"



## "Tetrazine"

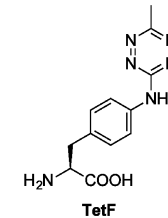
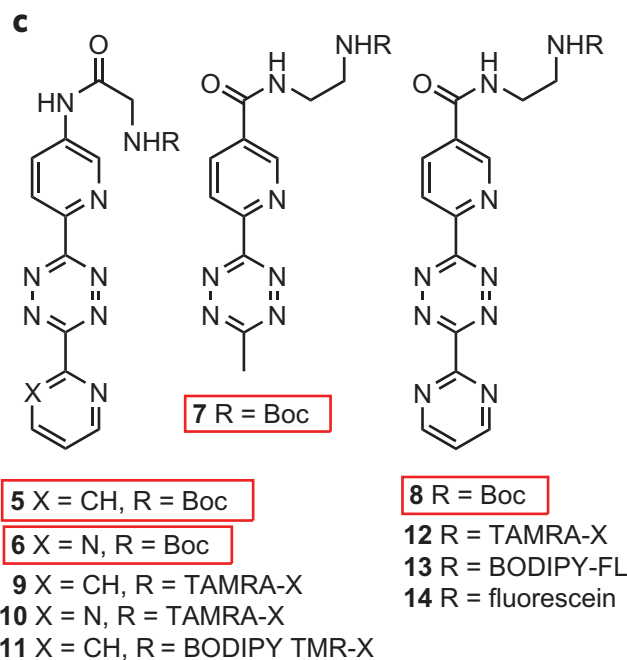
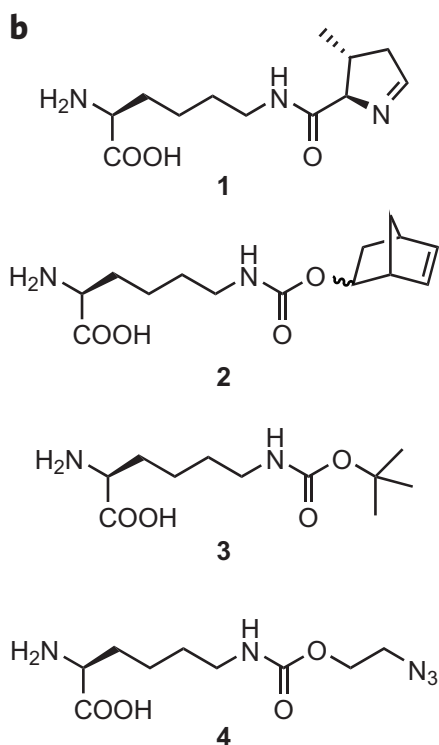
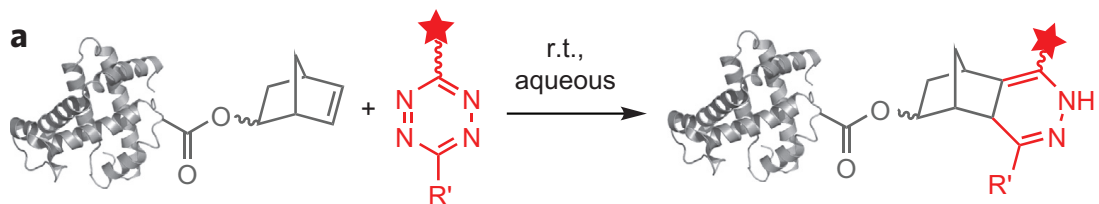


Figure 34. Structural formulas of unnatural amino acid useful for chemoselective labeling that have been incorporated site-specifically into proteins via genetic code expansion.



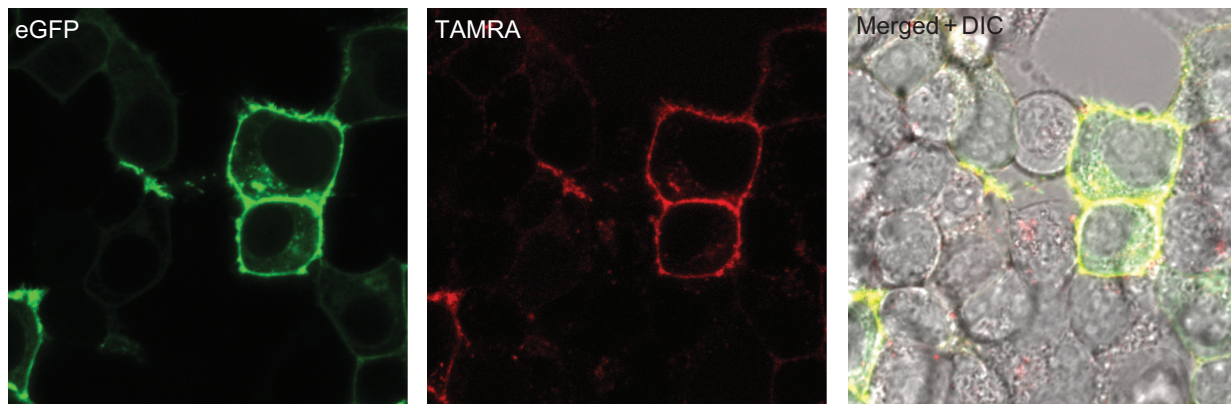
Jason Chin

# Non-natural amino acids...

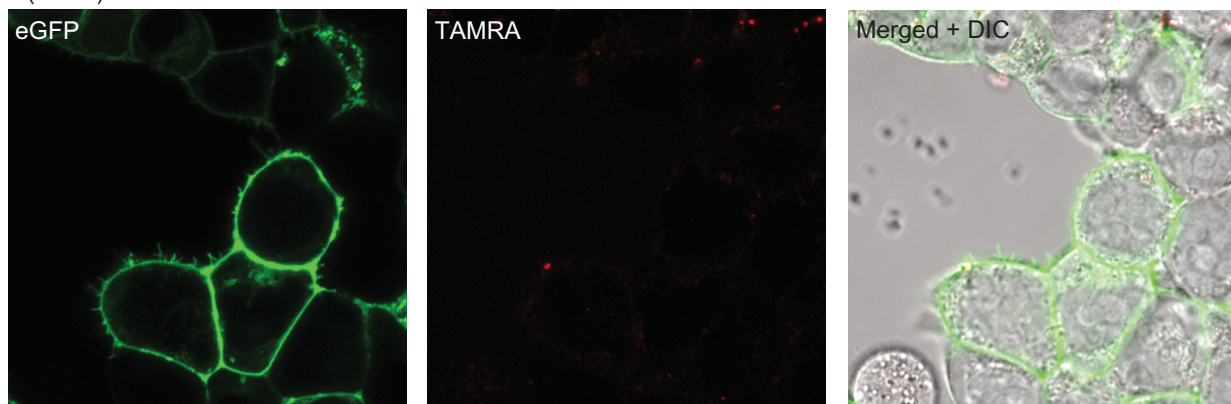


# Non natural amino acids...

**c** 2 (1 mM)

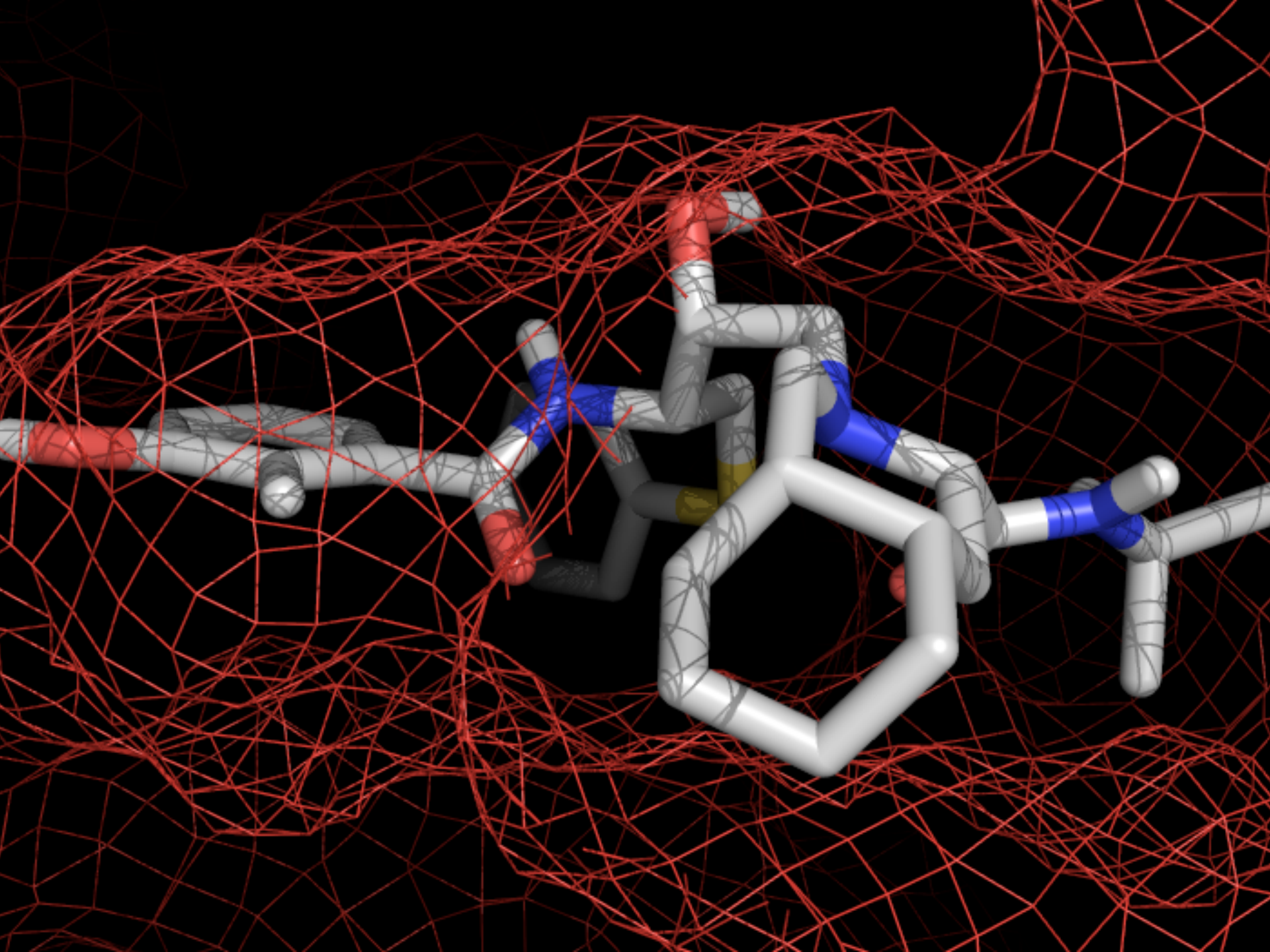


**3** (1 mM)



**Figure 4 | Site-specific incorporation of 2 into proteins in mammalian cells and the specific labelling of EGFR-GFP on the cell surface with 9.** **a**, Cells that contain the PylRS/tRNA<sub>CUA</sub> pair and the mCherry(TAG)eGFP-HA reporter produced GFP only in the presence of **2**. **b**, Western blots confirm that the expression of full length mCherry(TAG)eGFP-HA is dependent on the presence of **2**. **c**, Specific and rapid labelling of a cell surface protein in live mammalian cells. EGFR-GFP that bears **2** or **3** at position 128 is visible as green fluorescence at the membrane of transfected cells (left panels). Treatment of cells with **9** (200 nM) leads to selective labelling of EGFR that contains **2** (middle panels). Right panels show merged green and red fluorescence images, DIC = differential interference contrast. Cells were imaged four hours after the addition of **9**.

Target  
Identification







# Target vs Phenotypic-based Drug Discovery

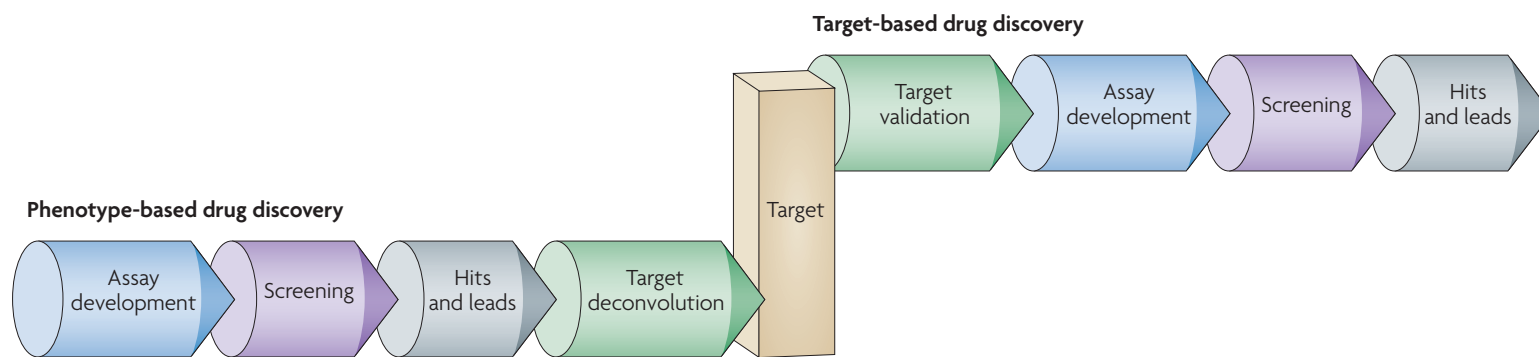
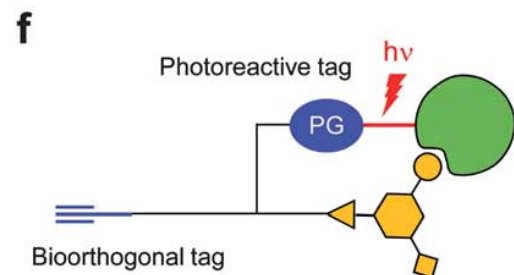
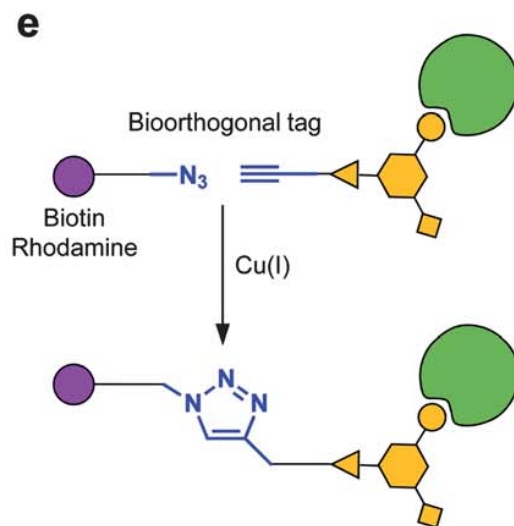
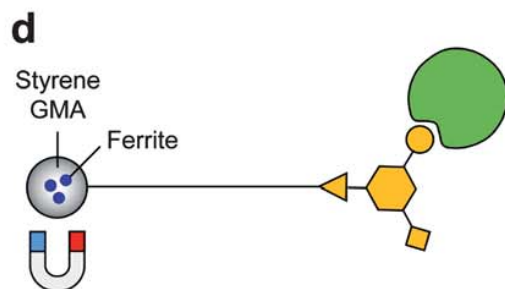
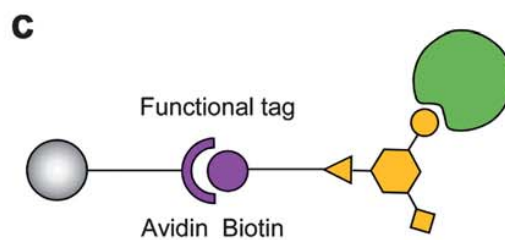
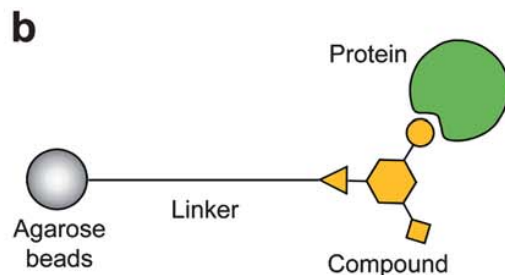
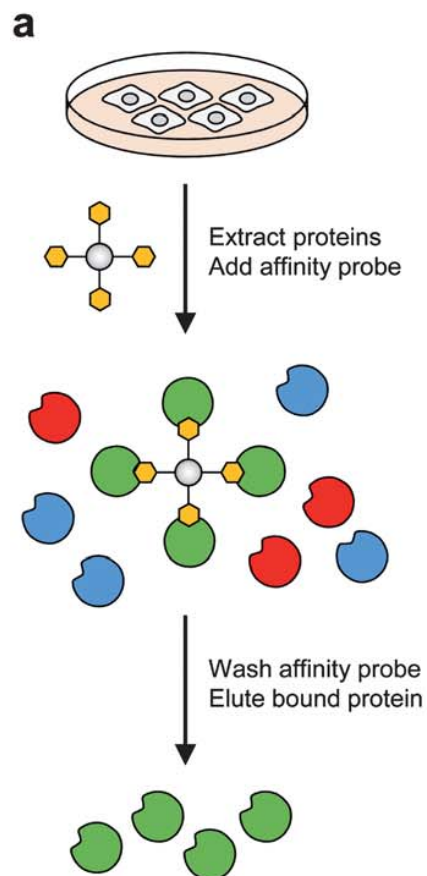
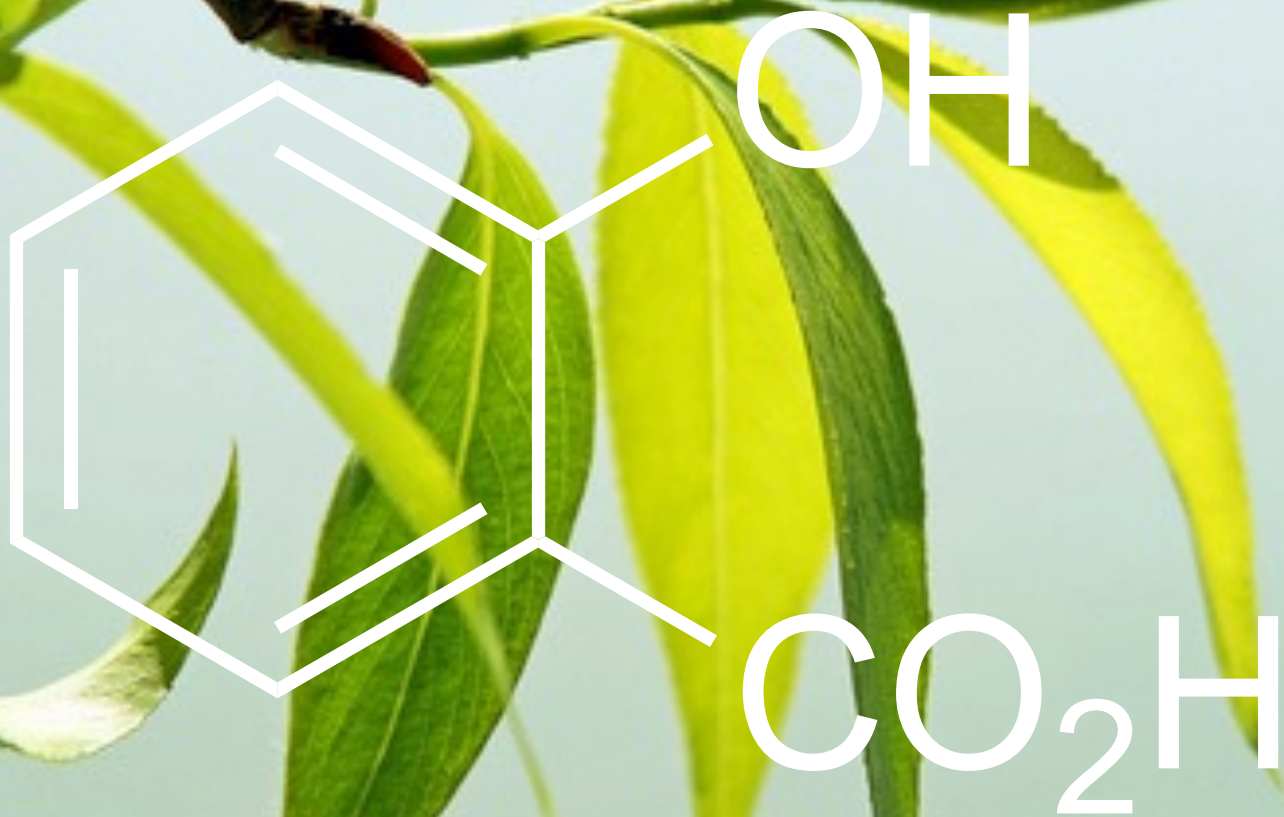


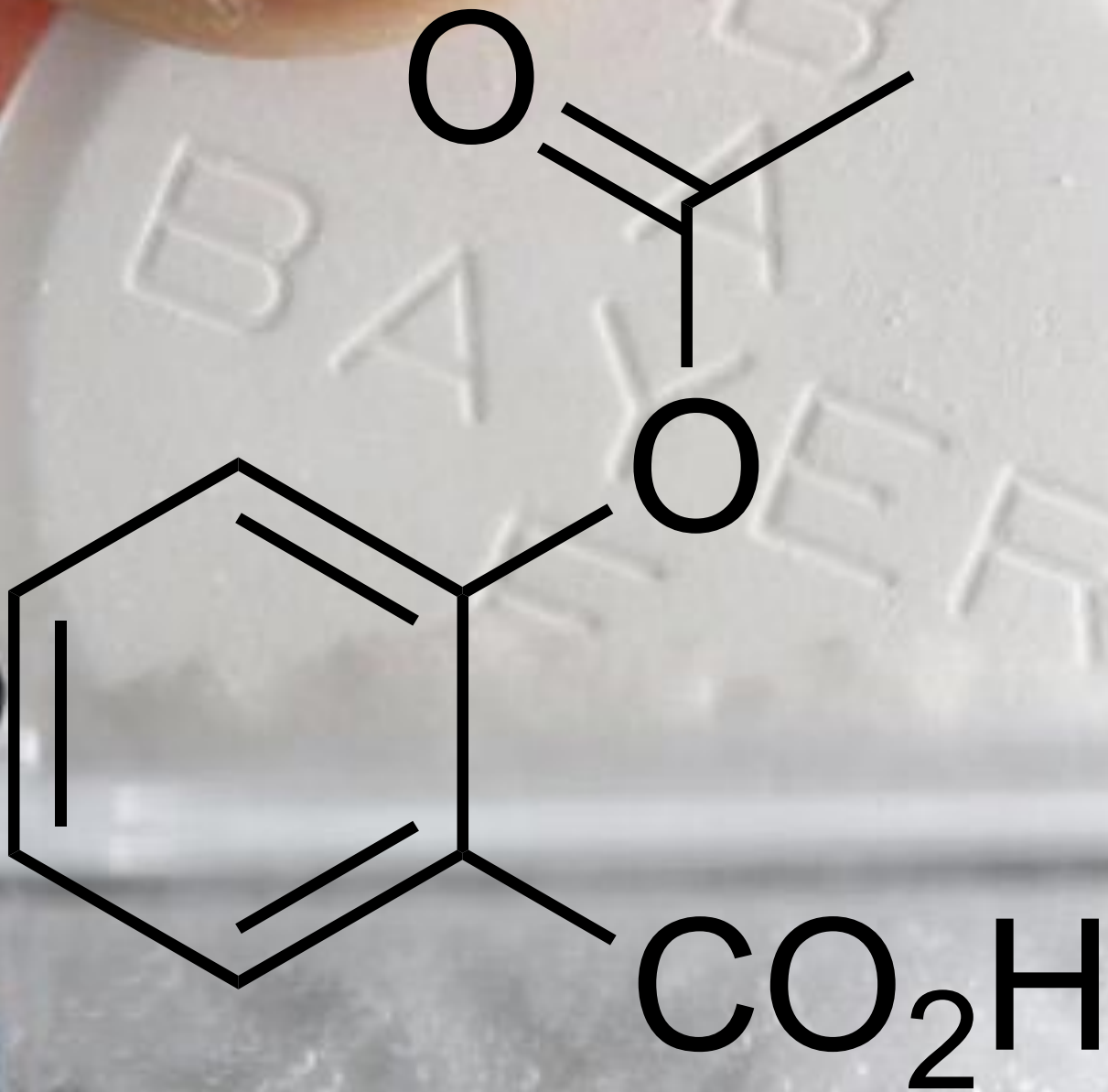
Figure 1 | **Phenotype-based versus target-based drug discovery.** The diagram illustrates the early phase of drug discovery, in which the aim is to identify target and lead molecules. In the phenotype-based approach, lead molecules are obtained first, followed by target deconvolution to

identify the molecular targets that underlie the observed phenotypic effects. In the target-based approach, molecular targets are identified and validated before lead discovery starts; assays and screens are then used to find a lead.

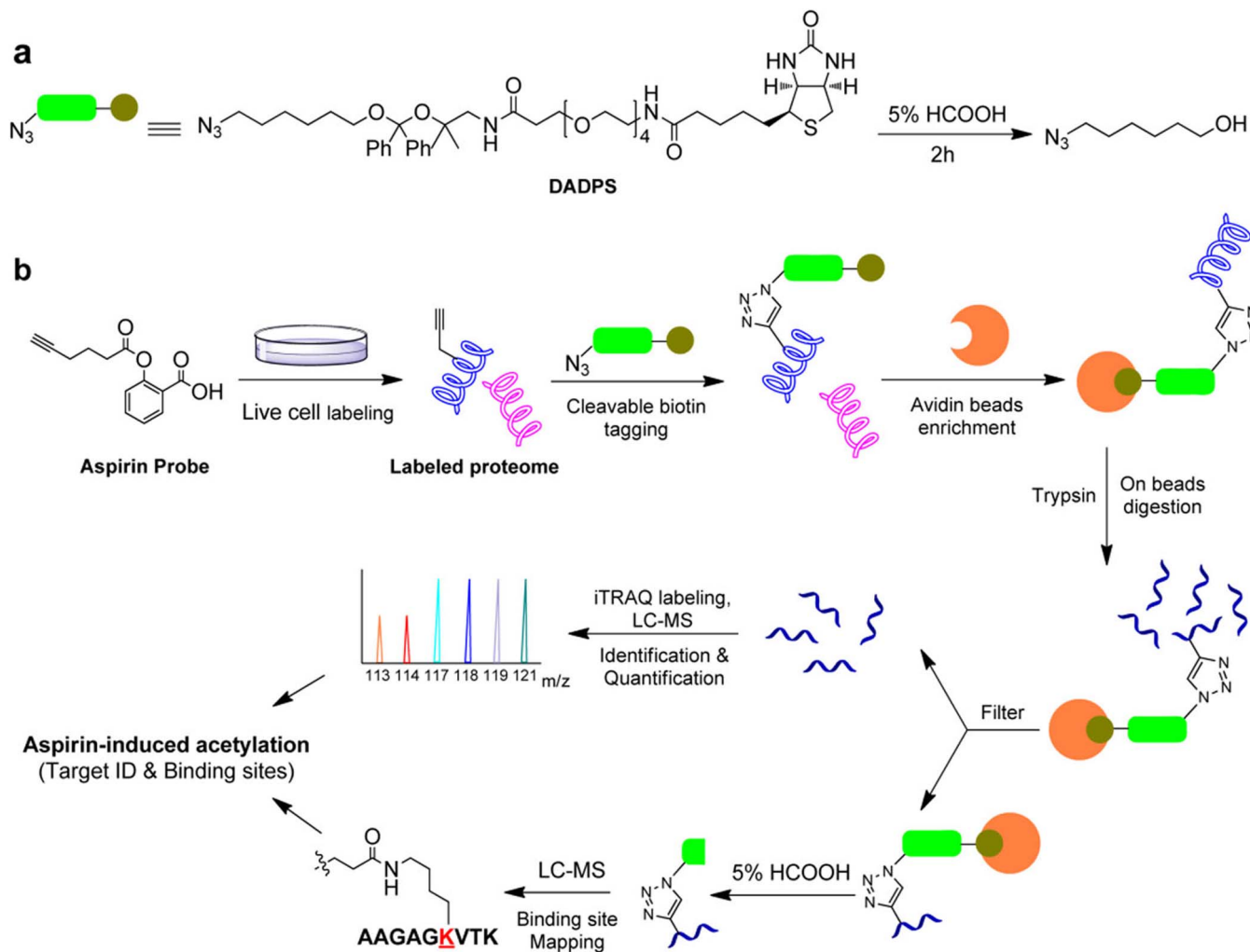
# Affinity purification



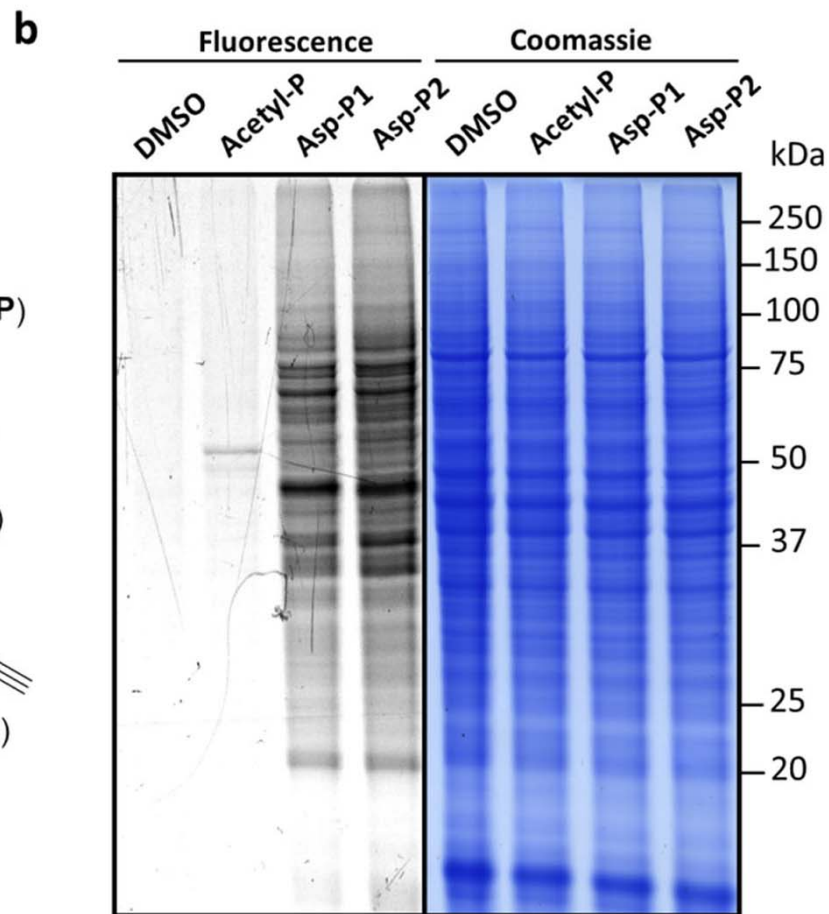
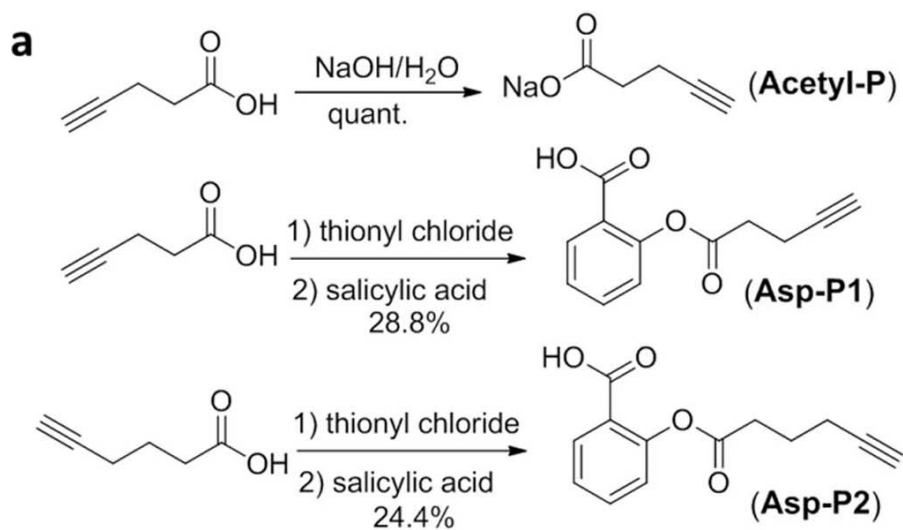




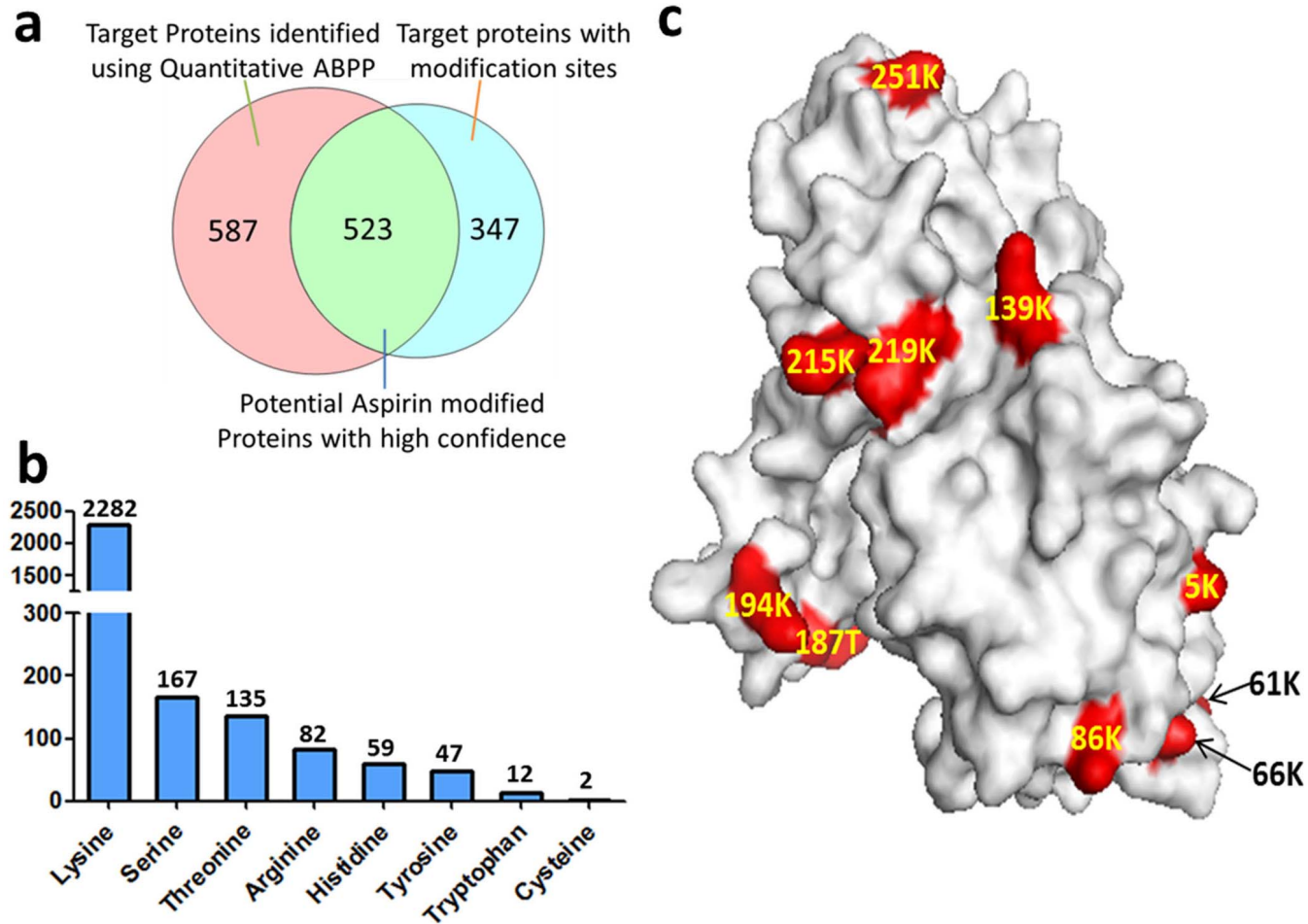
# Target identification strategy



# Targets



# Aspirin Targets



**Figure 3 | Summary of aspirin modified proteins and amino acid residues.** (a) Numbers of proteins identified using quantitative ABPP, proteins with modification sites identified and proteins confirmed with high confidence; (b) Numbers of the aspirin-modified amino acid residues. Numbers on top of the columns are the numbers of peptides modified by Asp-P2; (c) Locations of aspirin-modified residues in the protein GAPDH.



Chimie

Combinatoire

Dynamique



Jorge Luis Borges (1899 – 1986)

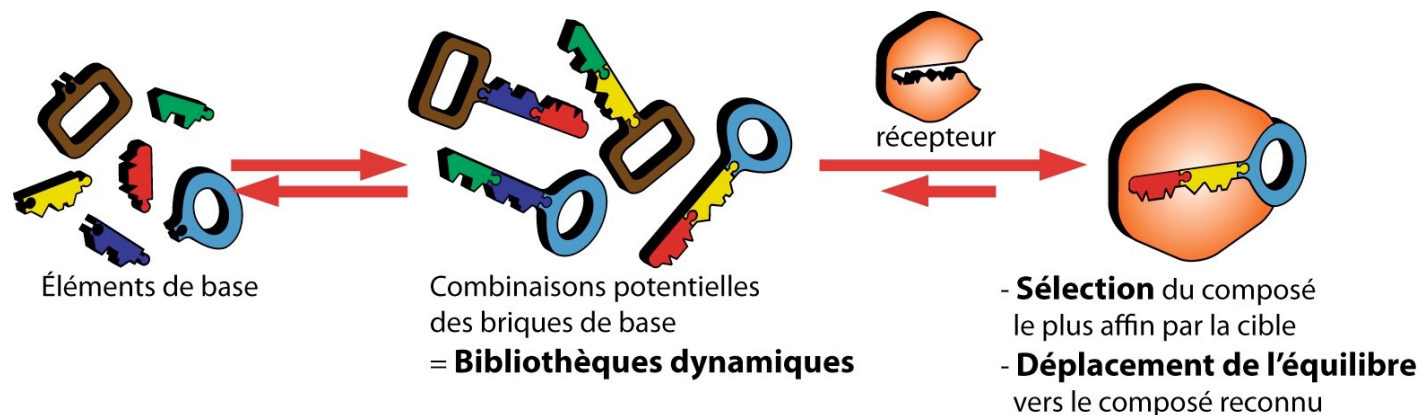
tout ce que je  
vais vous dire  
est déjà écrit  
dans la  
bibliothèque...

rien de ce que  
je vous  
raconte n est  
ecrit dans la  
bibliotheque...

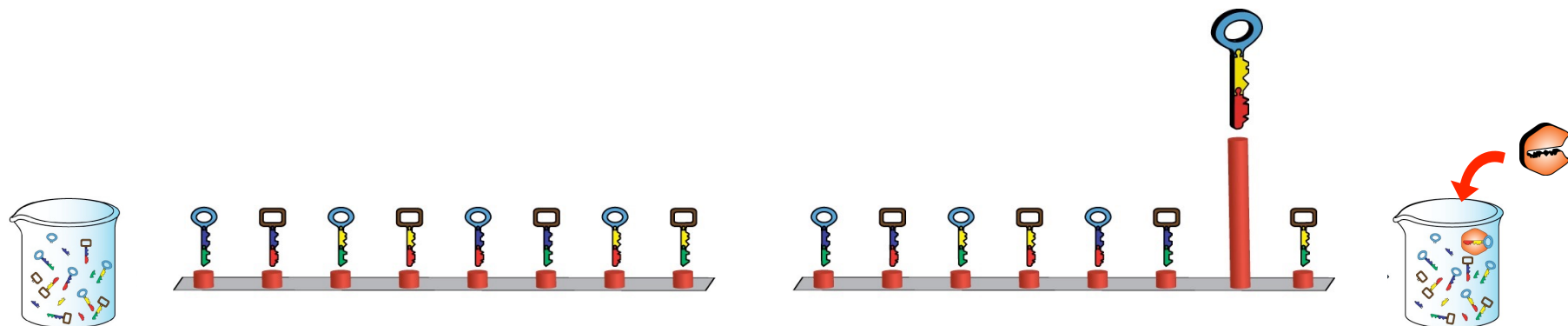
$$K_a = \frac{[A \cdot B]}{[A][B]}$$



# Chimie Combinatoire Dynamique

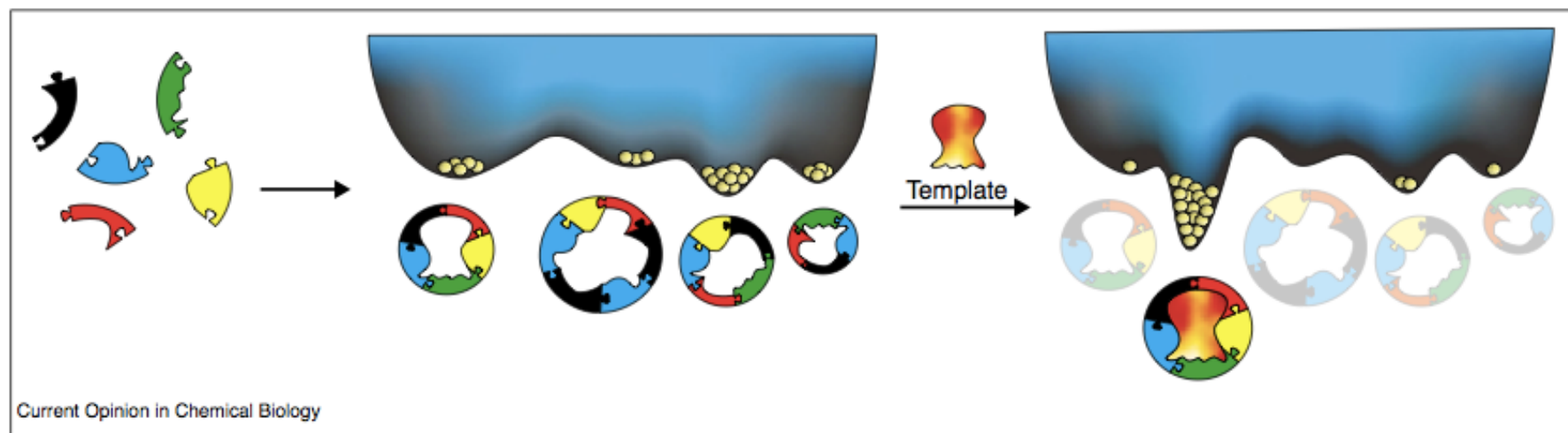


I. Huc, J.-M. Lehn, *Proc. Natl. Acad. Sci. USA* **1997**, 94, 2106



Reviews: I. Huc, R. Nguyen, *Comb. Chem. High Throughput Screening*, **2001**, 4, 109  
S. Otto, R. L.E. Furlan, J. K.M. Sanders, *Curr. Opin. Chem. Biol.*, **2002**, 6, 321  
O. Ramström, T. Bunyapaiboonsri, S. Lohmann, J.M. Lehn, *Biochim. Biophys. Acta*, **2002**, 1572, 178

# Chimie Combinatoire Dynamique



A small dynamic combinatorial library and its free energy landscape showing the effect of adding a template that strongly and selectively binds to one of the equilibrating species.

# Bibliothèques Combinatoires Virtuelles

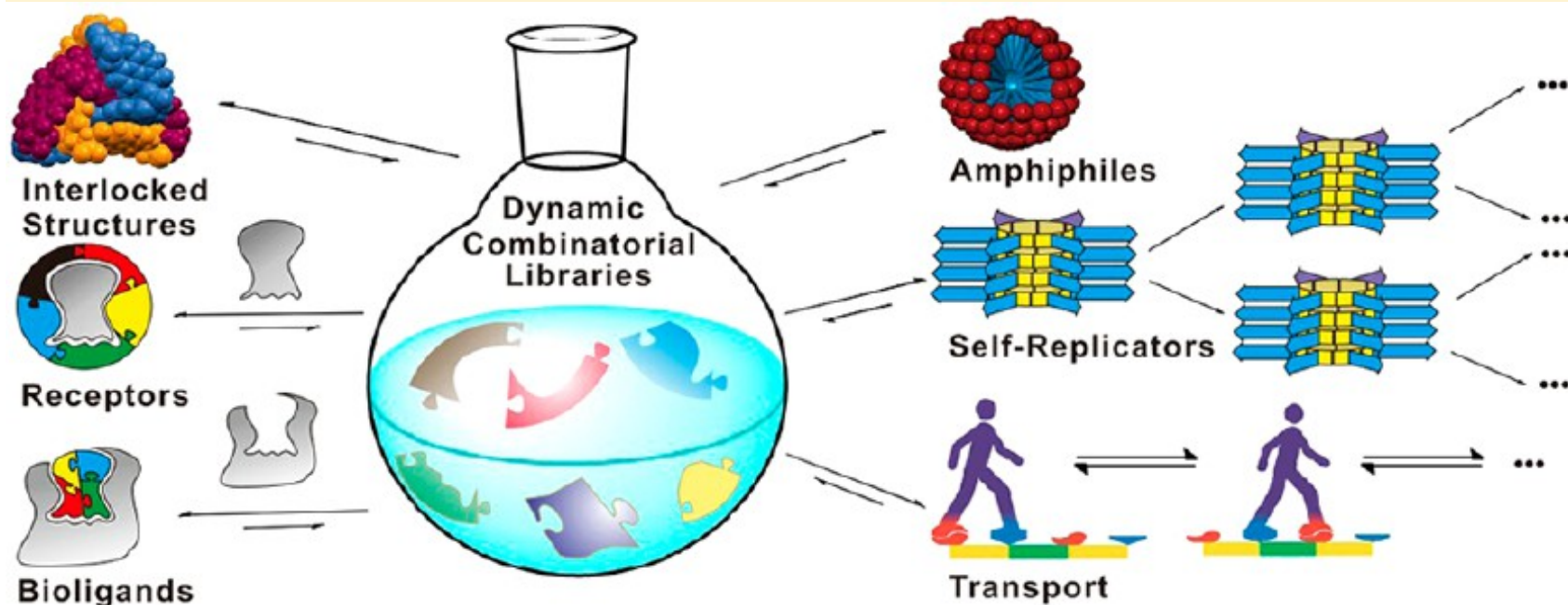
**Dynamic combinatorial chemistry** is a conceptually different approach that rests on supramolecular chemistry

It relies on a **reversible connection process** for the spontaneous and continuous generation of all possible combinations of a set of basic components, thus **making virtually available all structural and interactional features** that these combinations may present.

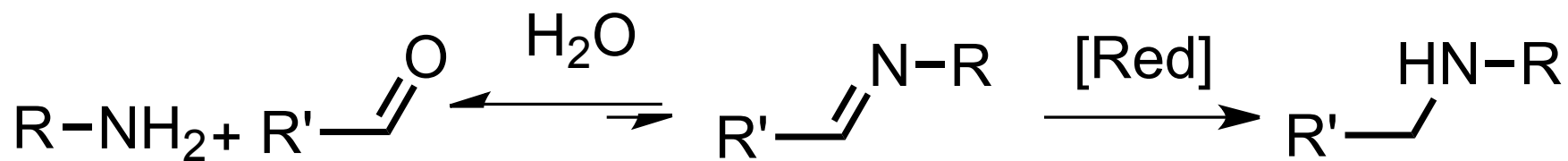


# Cibler des protéines

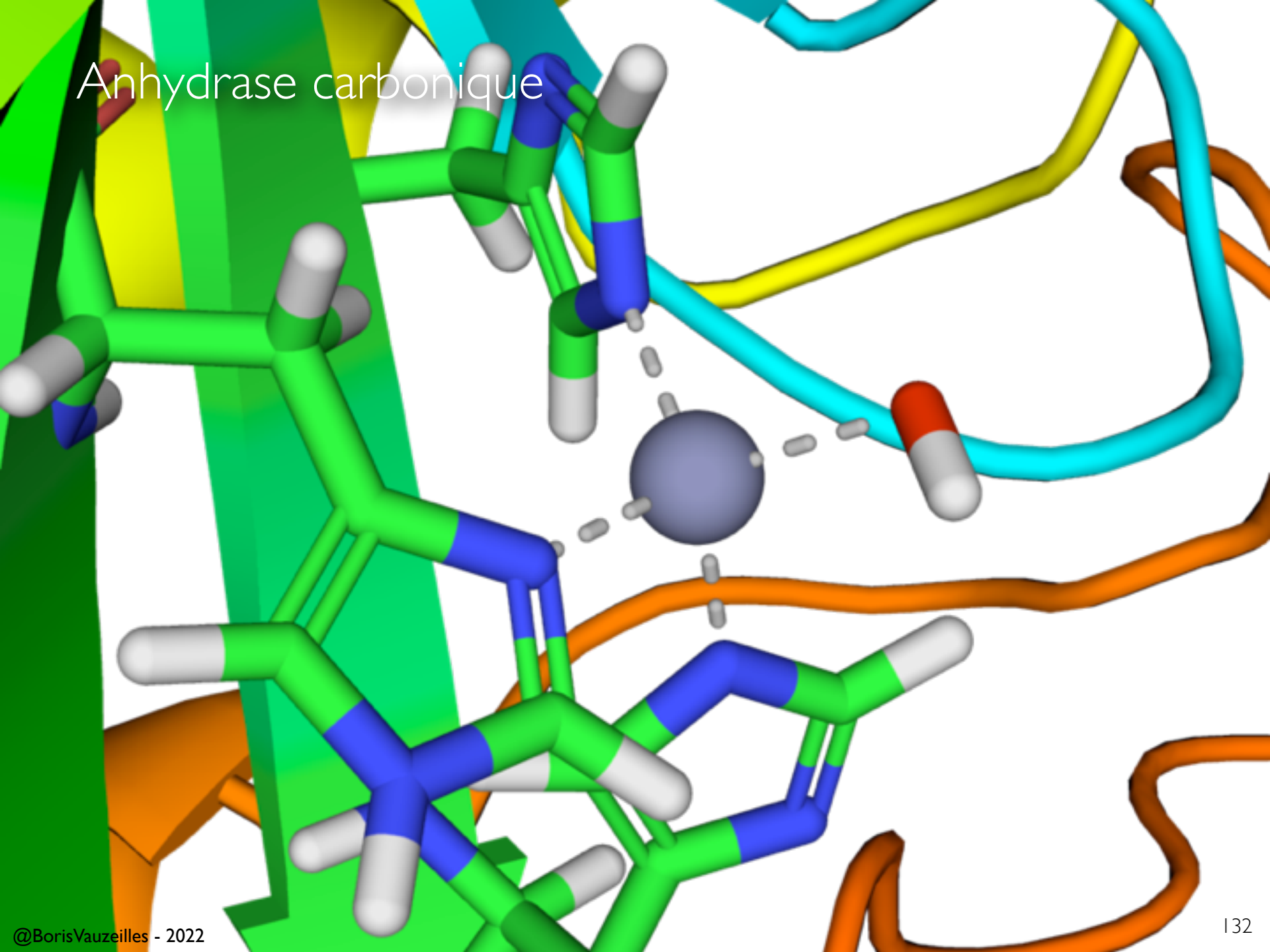
# Bibliothèques combinatoires dynamiques



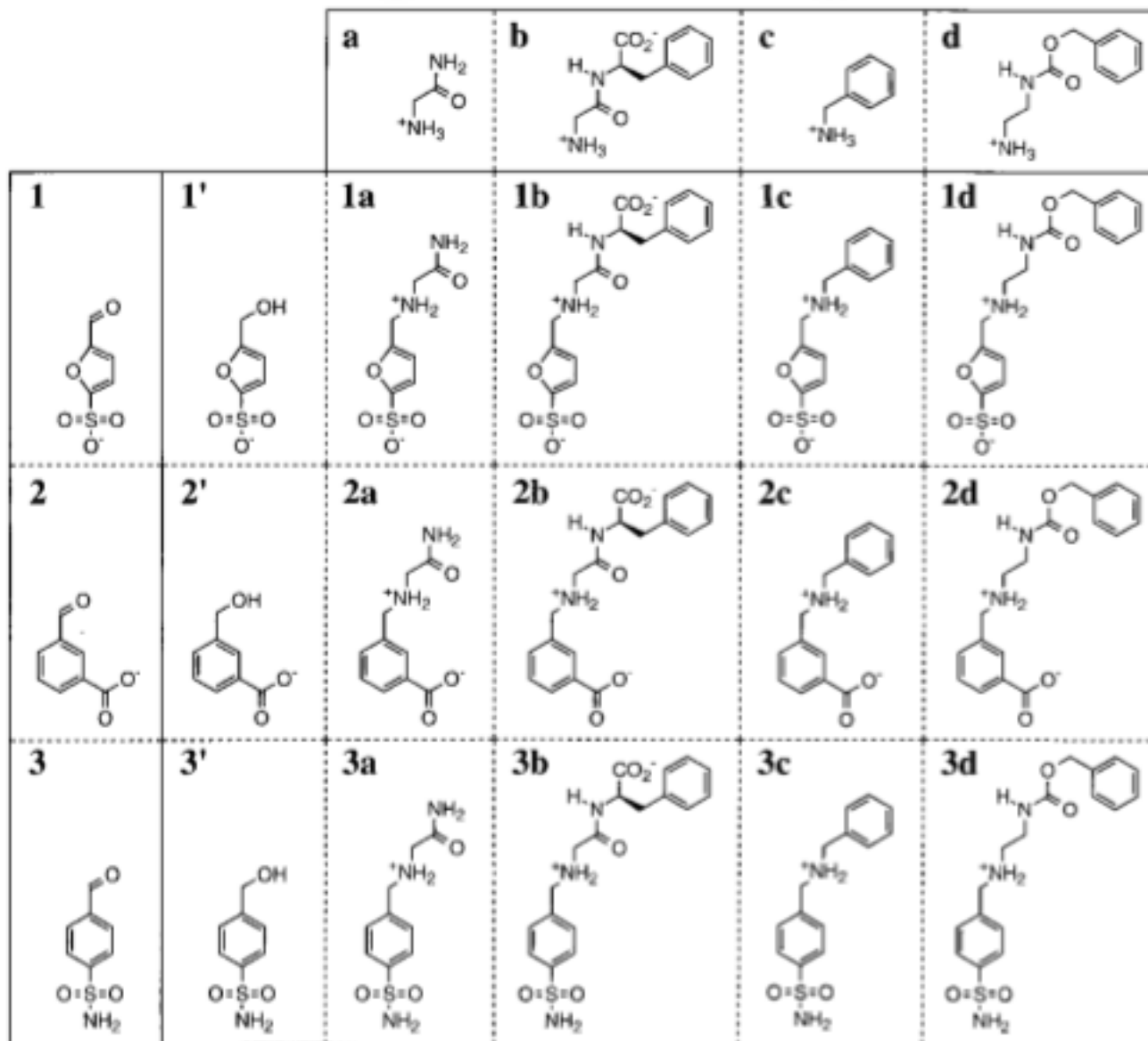
# Réactions réversibles - Formation d'imines



Anhydrase carbonique



# Inhibiteurs de l'anhydrase carbonique II



# Profil HPLC

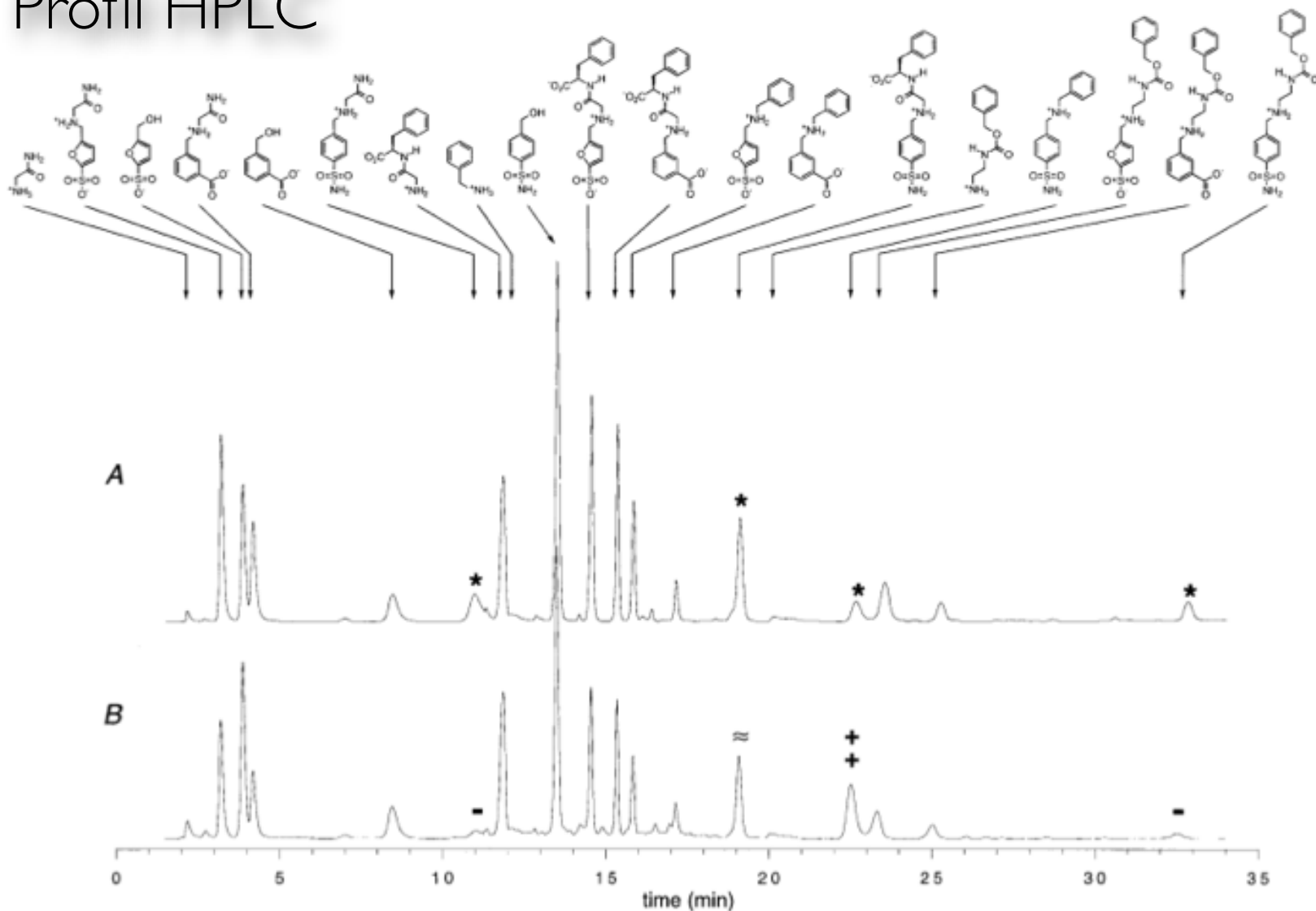


FIG. 3. HPLC traces of the final reaction mixtures of **a** + **b** + **c** + **d** (each at 4 mM) with **1** + **2** + **3** (each at 0.4 mM) and NaBH<sub>3</sub>CN (1.2 mM) in aqueous phosphate (20 mM, pH 6). (A) Trace corresponds to the reaction without additive. (B) Trace corresponds to the reaction in the presence of 1 equivalent (0.4 mM) of CA.

# Product amplification

Table 1. Relative proportions between the products **3c** and **3a**, **3b**, or **3d**, obtained with and without enzyme in reactions involving the aldehyde **3** and two competing amines **c** and **a**, **b**, or **d**

Starting materials	Normalized proportion of the products	Normalized value
<b>3</b> , <b>c</b> , <b>a</b>	$(3c/3a)_{rel}$	15
<b>3</b> , <b>c</b> , <b>b</b>	$(3c/3b)_{rel}$	4.5
<b>3</b> , <b>c</b> , <b>d</b>	$(3c/3d)_{rel}$	21
<b>3</b> , <b>c</b> , <b>d</b> , <b>I</b>	$(3c/3d)_{rel}$	2

Value of  $x$  means that the ratio between the two products is  $x$  times higher with CA than without CA. **I** is the inhibitor hexyl 4-sulfamoylbenzoate.

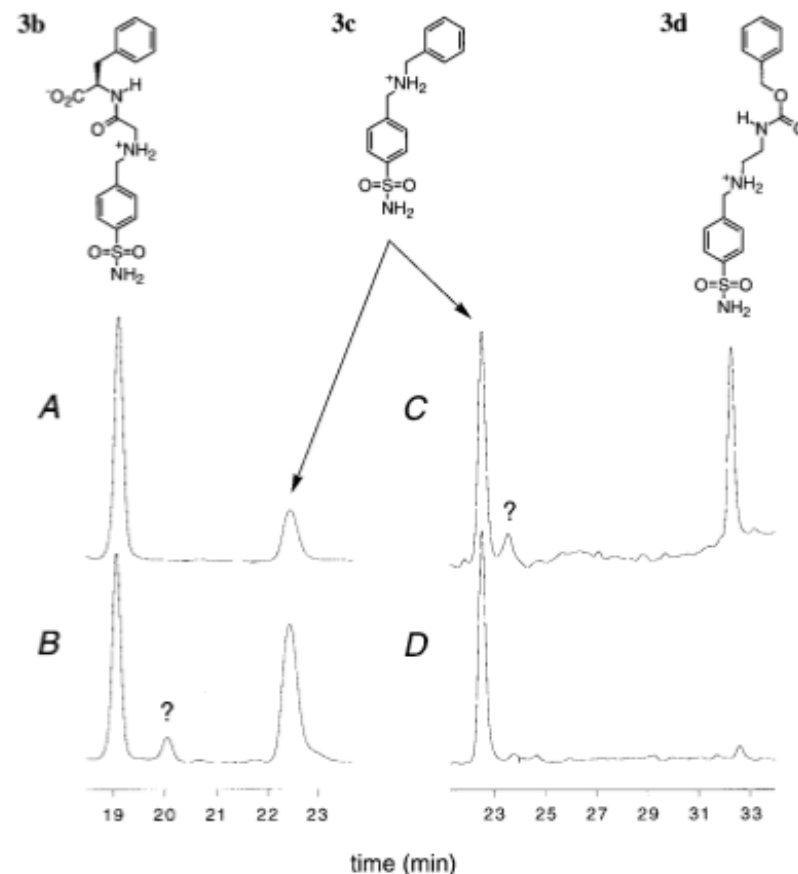
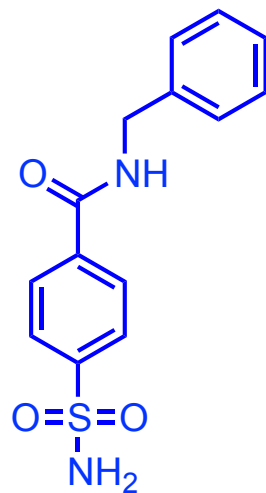
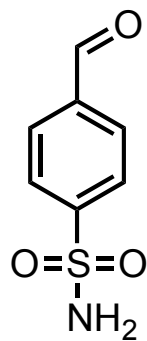


FIG. 4. HPLC traces of the final reaction mixtures of **3** (0.4 mM) with **b** + **c** (each at 2 mM) (*A* and *B*) or **c** + **d** (each at 2 mM) (*C* and *D*) and NaBH<sub>3</sub>CN (1.2 mM) in aqueous phosphate (20 mM, pH 6). (*A* and *C*) Traces correspond to the reaction without additive. (*B* and *D*) Traces correspond to the reaction in the presence of 1 equivalent (0.4 mM) of CA. Question marks indicate unidentified impurities.

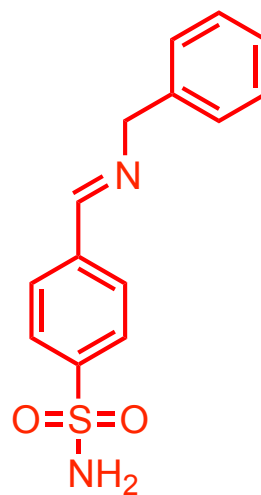
# Bilan



Inhibiteur connu  
1.1 nM



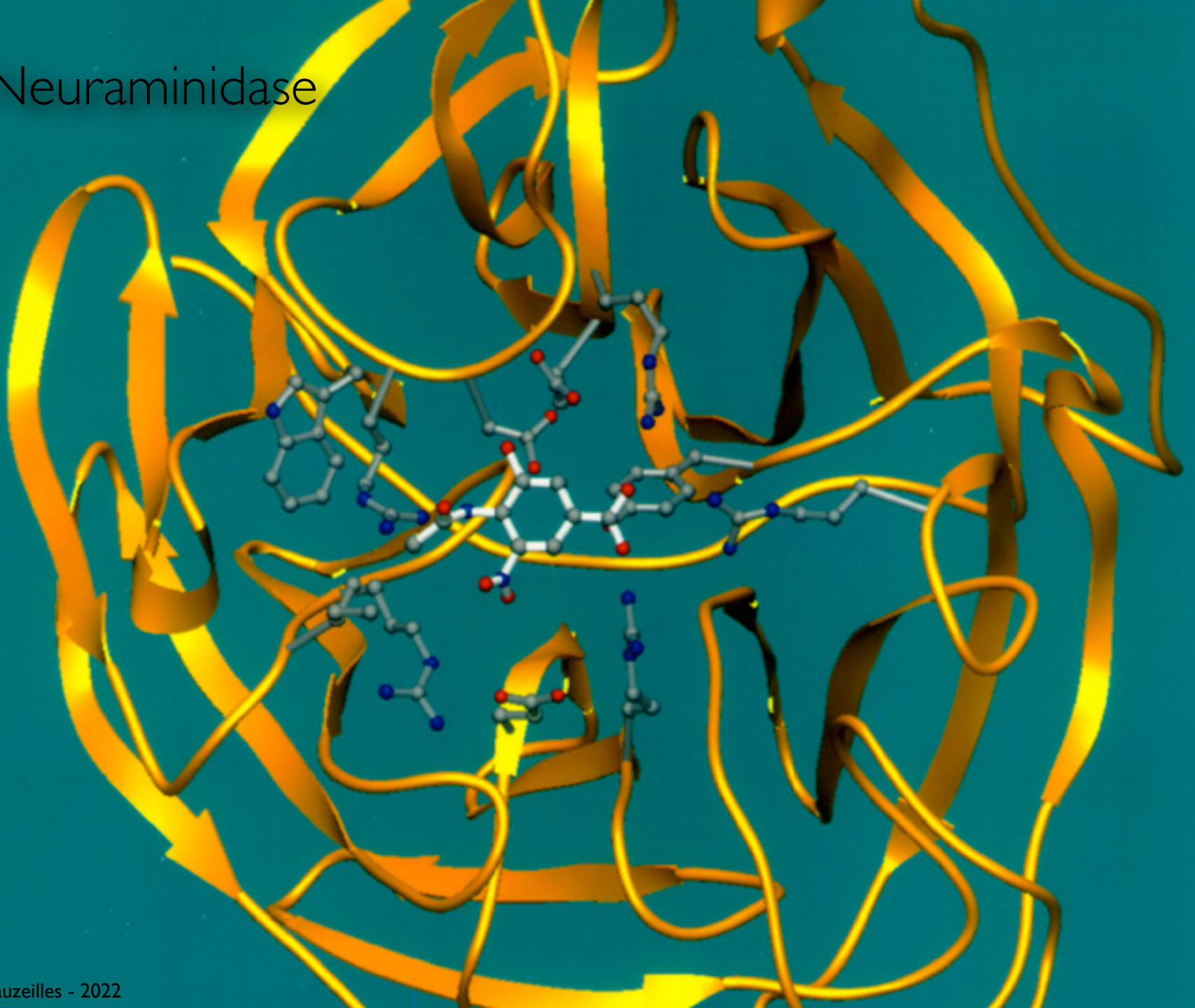
Plateforme  
submicromolaire



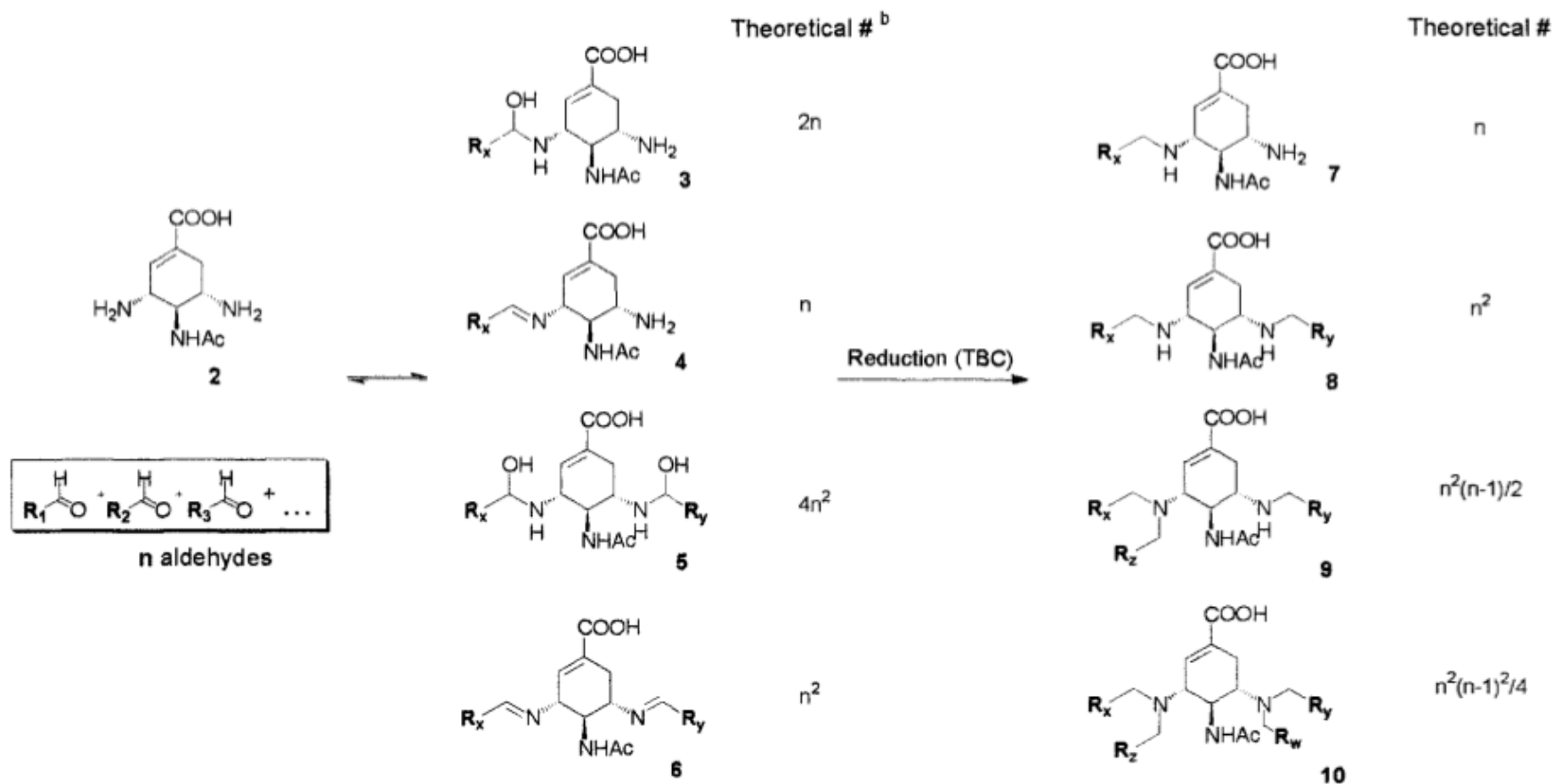
Hit



# Neuraminidase

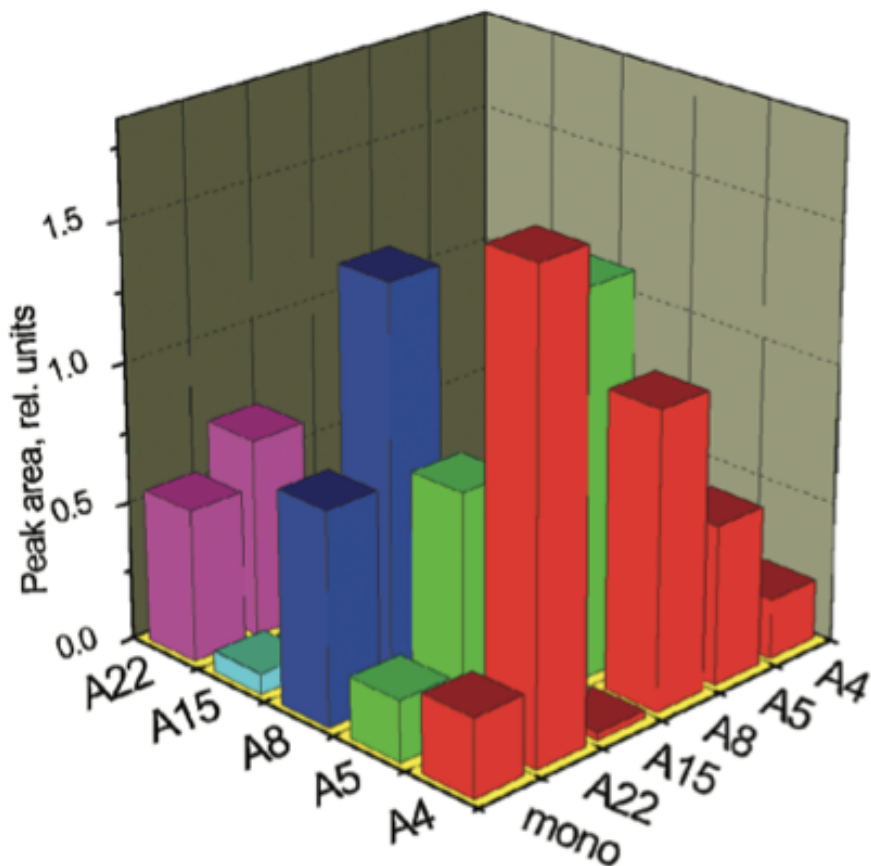


# Neuraminidase

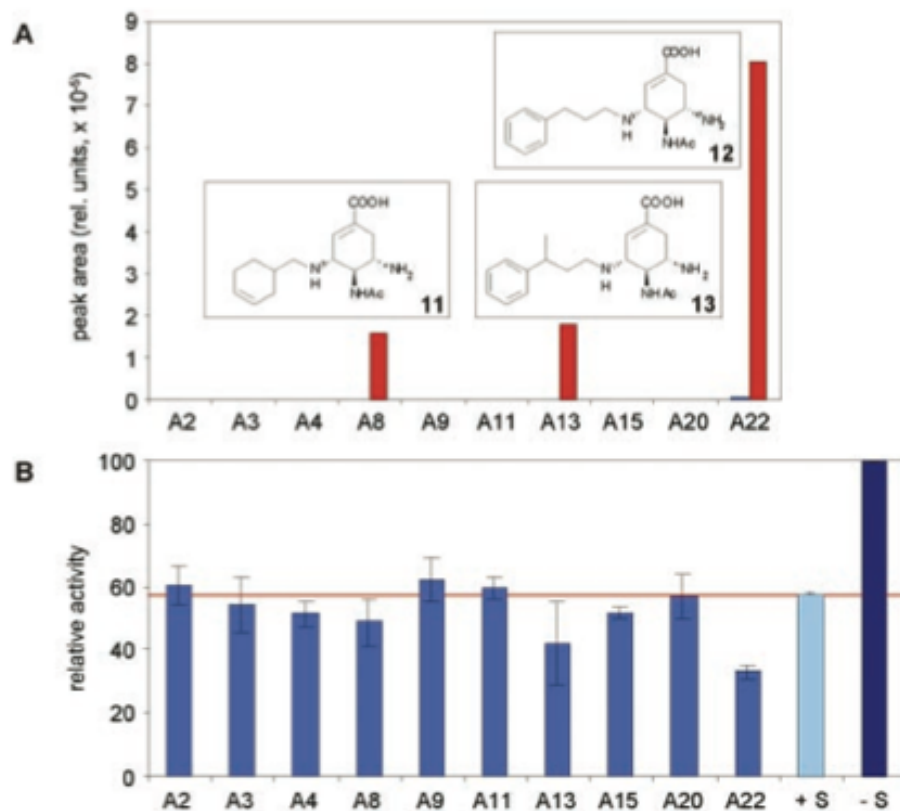


**Scheme 1.** Structures of the scaffold and library composition. <sup>a</sup>, Hemiaminal species are restricted to those detected by NMR. <sup>b</sup>, Numbers of compounds **3** and **5** account for diastereomers.

# Bibliothèque combinatoire dynamique

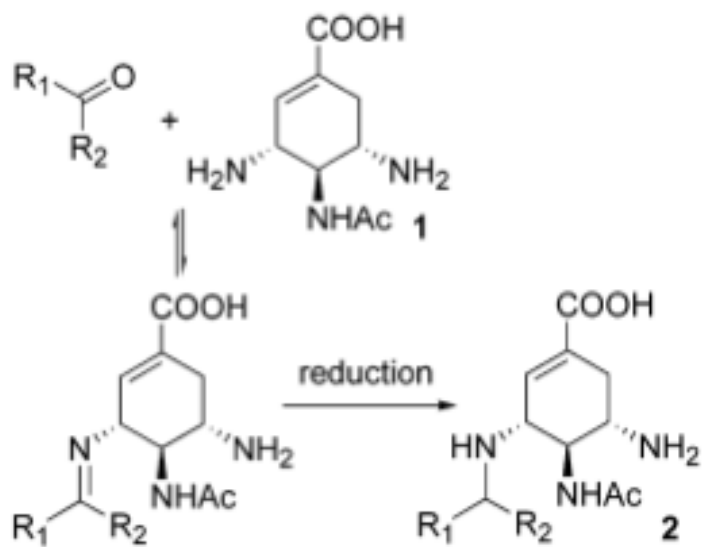


**Fig. 1.** Mono- and disubstituted components (structures 7 and 8, respectively) detected in a static library of potential neuraminidase inhibitors (see *Materials and Methods* for synthesis protocol) after 24 h of incubation. Heights of the bars correspond to peak areas of the compound masses in the extracted ion chromatograms.

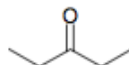


**Fig. 2.** (A) Library components of general structure 7 formed in the presence (red bars) and in the absence (blue bars, where applicable) of 10  $\mu\text{M}$  neuraminidase on incubation of the mixture of 2 (10  $\mu\text{M}$ ) with A2–A4, A8, A9, A11, A13, A15, A20, and A22 (200  $\mu\text{M}$  each) and 100  $\mu\text{M}$  tetrabutylammonium cyanoborohydride (TBC) for 12 h. See *Materials and Methods* for analytical protocol. (B) Inhibition of neuraminidase by mixtures of 2 (10  $\mu\text{M}$ ) with individual aldehydes, as marked on the axis (2 mM each). Control bars correspond to the enzyme activity without inhibitors (–S) and with the scaffold 2 alone (+S).

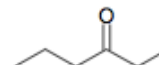
# Cétimines



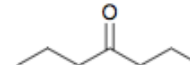
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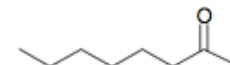
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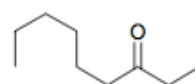
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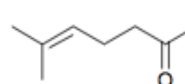
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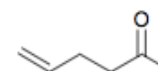
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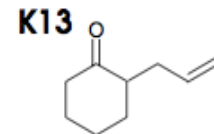
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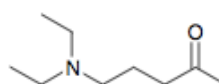
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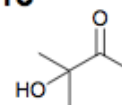
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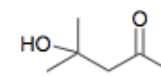
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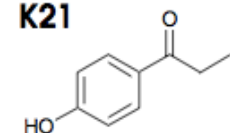
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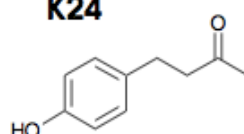
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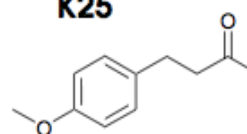
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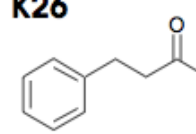
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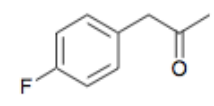
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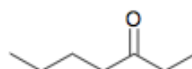
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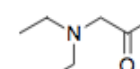
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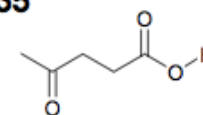
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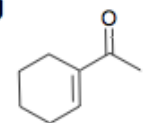
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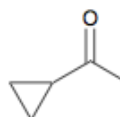
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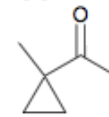
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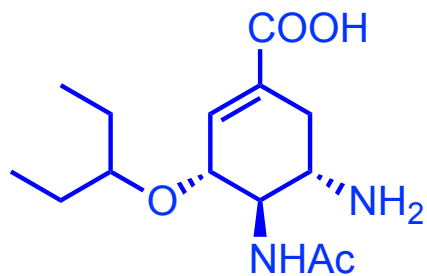
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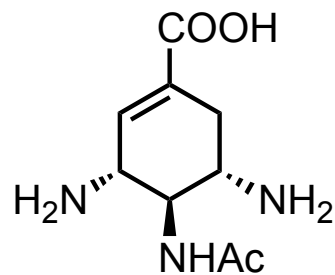
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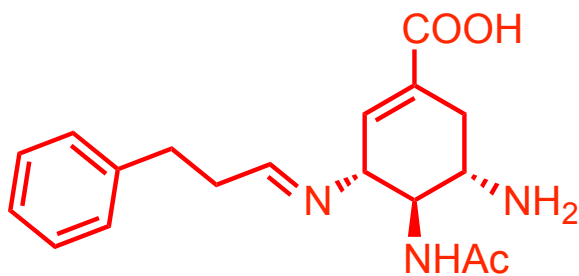
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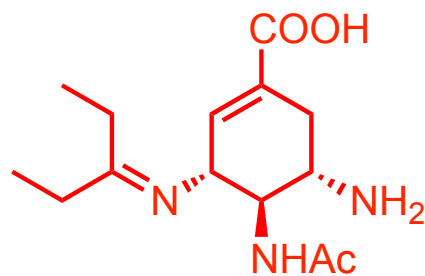
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1.3 nM



Plateforme  
31 μM



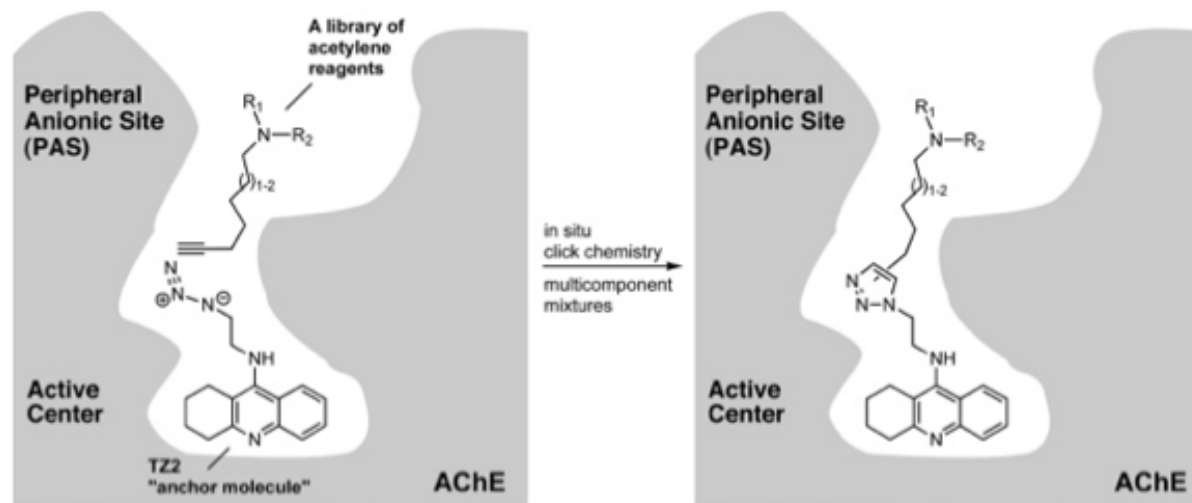
Hit 1  
1.6 μM



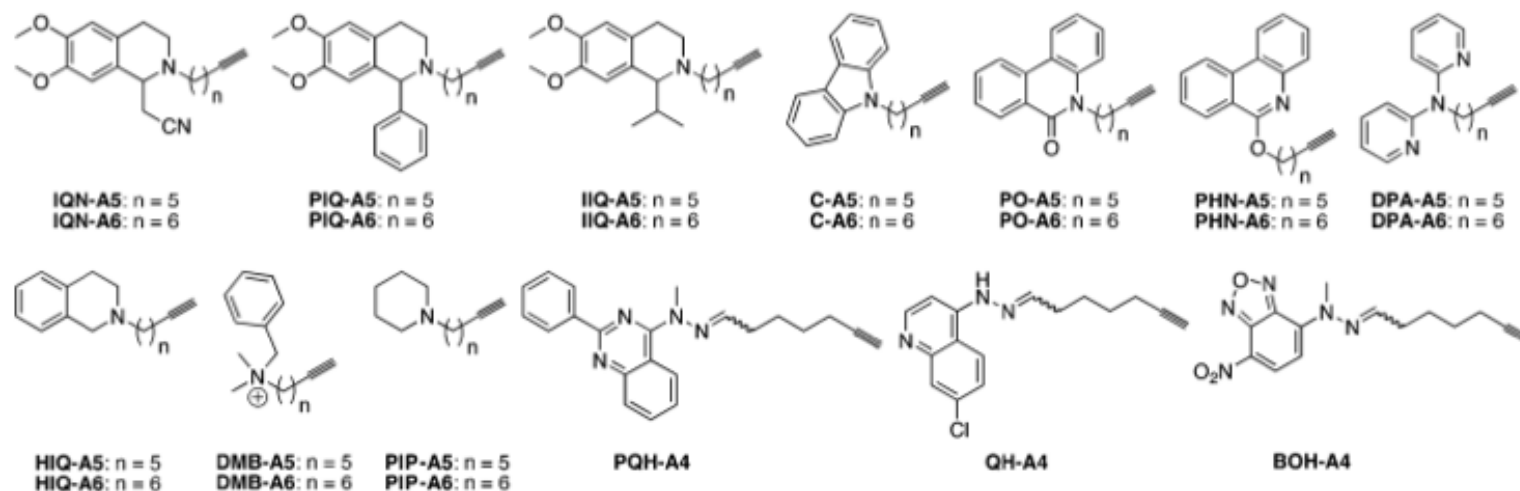
Hit 2  
85 nM

# Synthèse accélérée par la cible

# Chimie click *in situ*

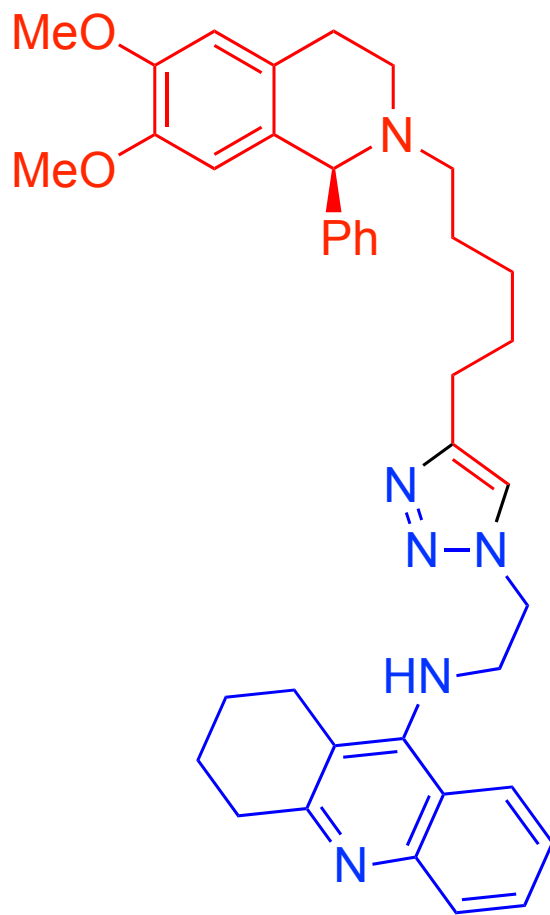


**Figure 2.** In situ click chemistry screening for AChE inhibitors, containing novel peripheral site ligands.



**Figure 3.** A library of 23 acetylene reagents for in situ click chemistry screening. All chiral compounds are racemic.

# Inhibiteur femtomolaire



Ki = 33 fM (AChE anguille)



The  
End