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Chemical synthesis of glycosylated spike RBD peptides

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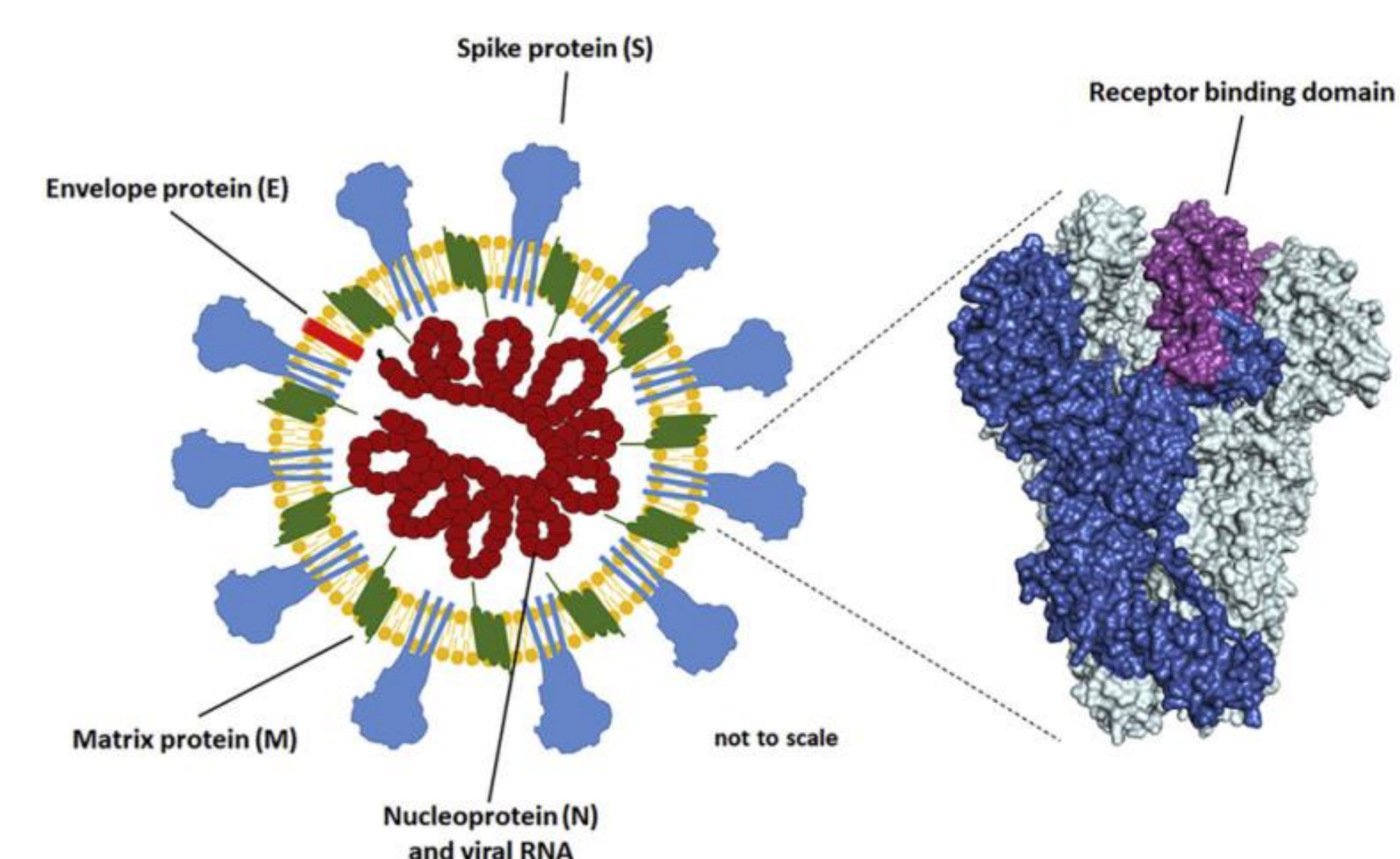
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Introduction

SARS-CoV-2 has a linear single-stranded positive-sense RNA genome, encoding 4 structural proteins [spike (S), envelope (E), membrane (M), and nucleocapsid (N)]. Among these proteins, the S protein (**Figure 1**) is the major focus of current research, as it facilitates viral attachment, entry and membrane fusion. The infection of host cells is initiated by the interaction of the viral S protein with receptors on the cell surface, hence it has a crucial role in the spread of the virus. There is surely an urgent need to develop a vaccine to halt the viral infection and curb the pandemic.

Li, F. *Annu Rev Virol*, **2016**, 3 (1), 237-261; Amanat, F.; Krammer, F. *Immunity* **2020**, 52 (4), 583-589.

Figure 1

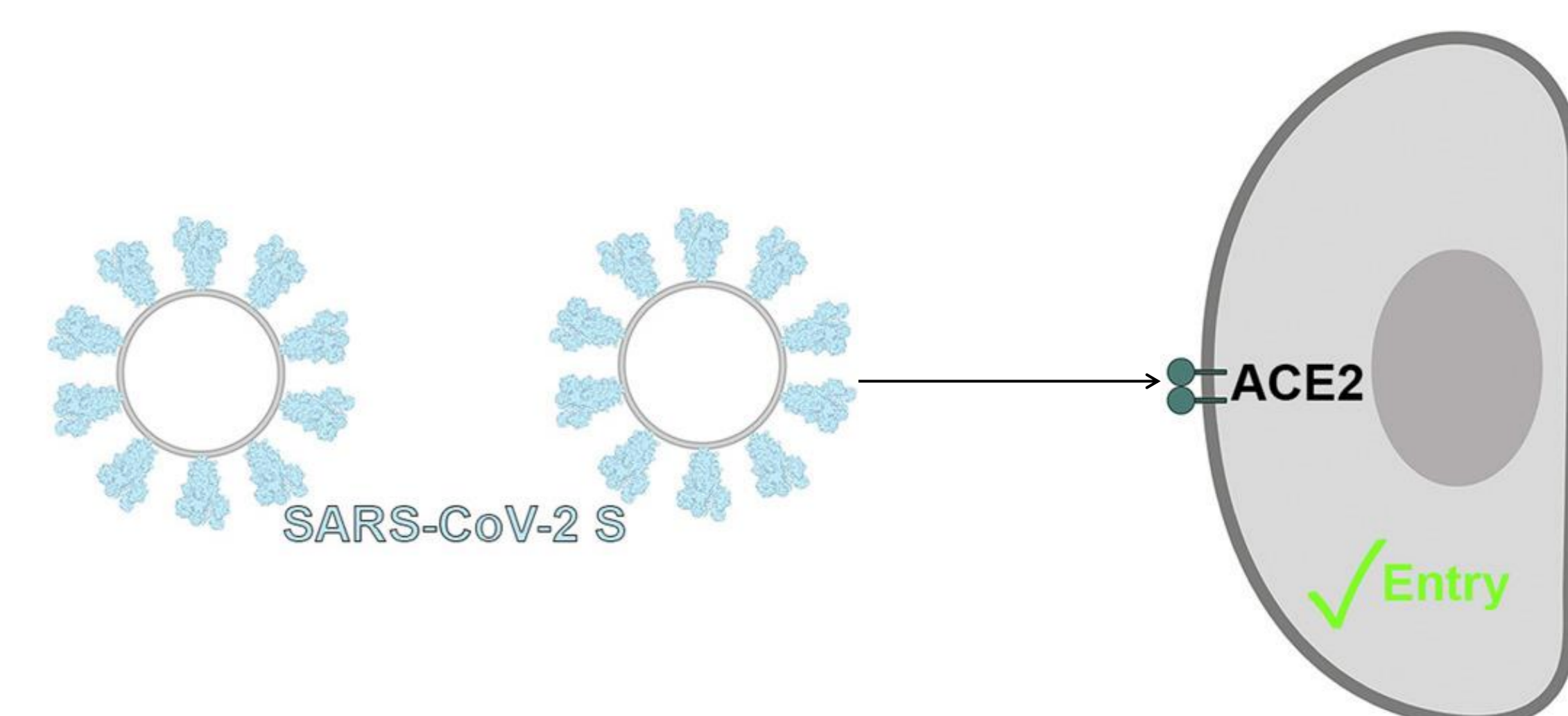


Introduction

The human angiotensin-converting enzyme 2 (hACE2) is the entry receptor for SARS-CoV-2 (**Figure 2**). As vaccine developments aim to elicit adaptive immunity via an antibody response at the sites of viral entry, one main strategy for vaccine development is to use peptides derived from S protein as the antigen to generate the antibody. The S-peptide induced antibody can potentially block the entry of SARS-CoV-2 to the host cell.

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Figure 2



Ser/Thrligation (STL) and Cys/Pen ligation (CPL)

In general, the chemical synthesis of a complex glycopeptide/protein involves two key phases: (1) generation of moderately sized peptides or glycopeptides (~ 20-30 amino acids) by solid-phase peptide synthesis (SPPS) and (2) joining these (glyco)peptide segments together. Towards the goal of chemical protein synthesis, our laboratory has developed in-house chemoselective peptide ligation methods over the past decade, including **Ser/Thrligation (STL)** in 2013 and **Cys/Pen ligation (CPL)** in 2020. STL and CPL could provide many choices for disconnection. The biantennary N-glycan (**Figure 3**) will be obtained by isolation of a sialoglycopeptide (SGP) from egg yolk. After we get glycopeptides by our 'N+1' strategy, we use STL and CPL to ligate then followed by refolding to get the target glycoprotein (**Figure 3**).

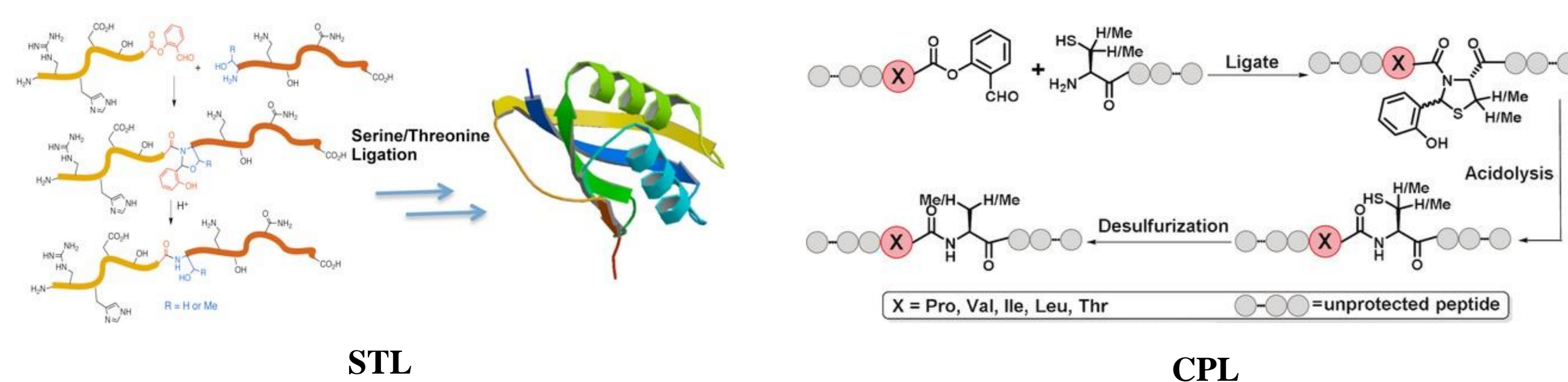
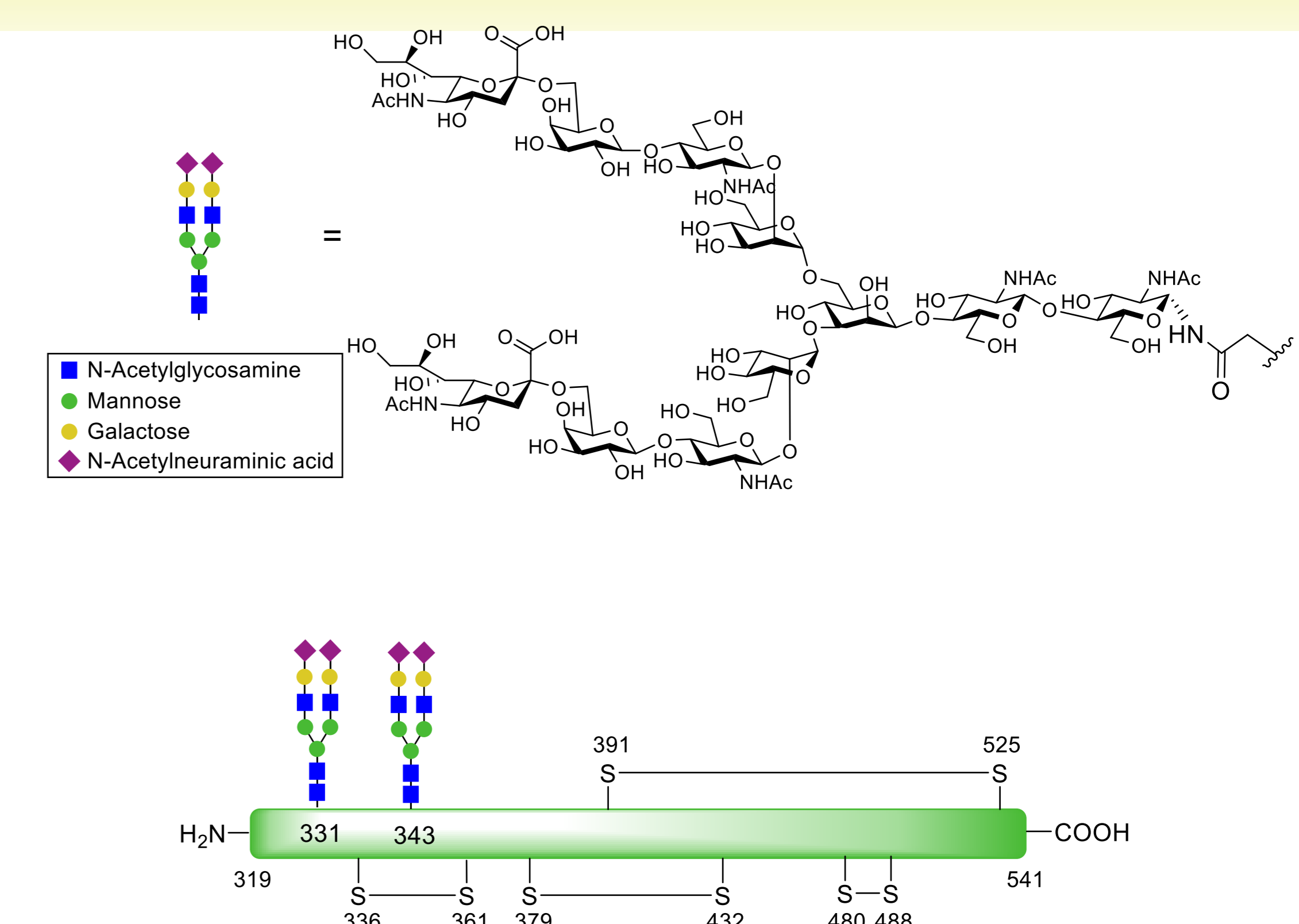


Figure 3



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Zhang, Y.; Xu, C.; Lam, H. Y.; Lee, C. L.; Li, X. *Proc Natl Acad Sci U S A*, **2013**, 110 (17), 6657-62.

Tan, Y.; Li, J.; Jin, K.; Liu, J.; Chen, Z.; Yang, J.; Li, X. *Angew Chem Int Ed Engl*, **2020**, 59 (31), 12741-12745.

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