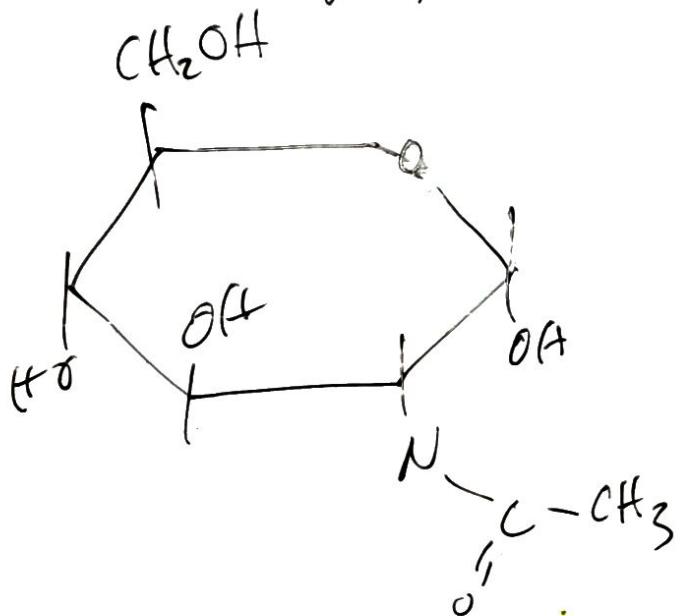


## 11 November 2019 ##

Amino derivatization

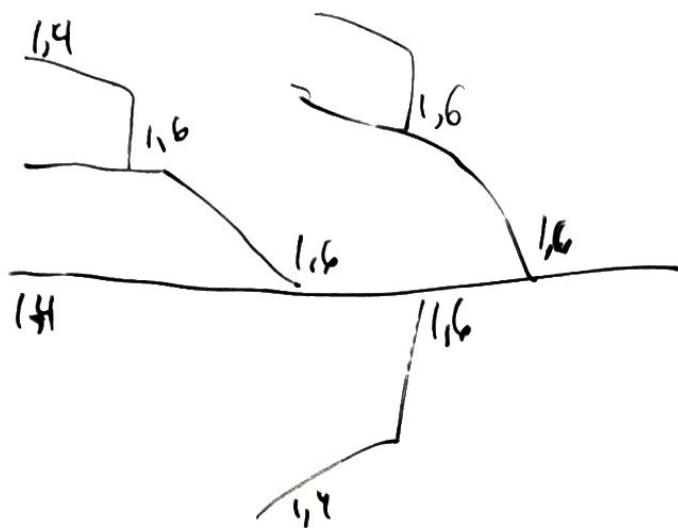
Always @ carbon 2 of aldoses

$\alpha$ -D-n-acetyl-glucosamine



GlcNAc

amylopectin: Every 30  $\alpha$ 1,4 monomers  
gets an  $\alpha$ 1,6 branch



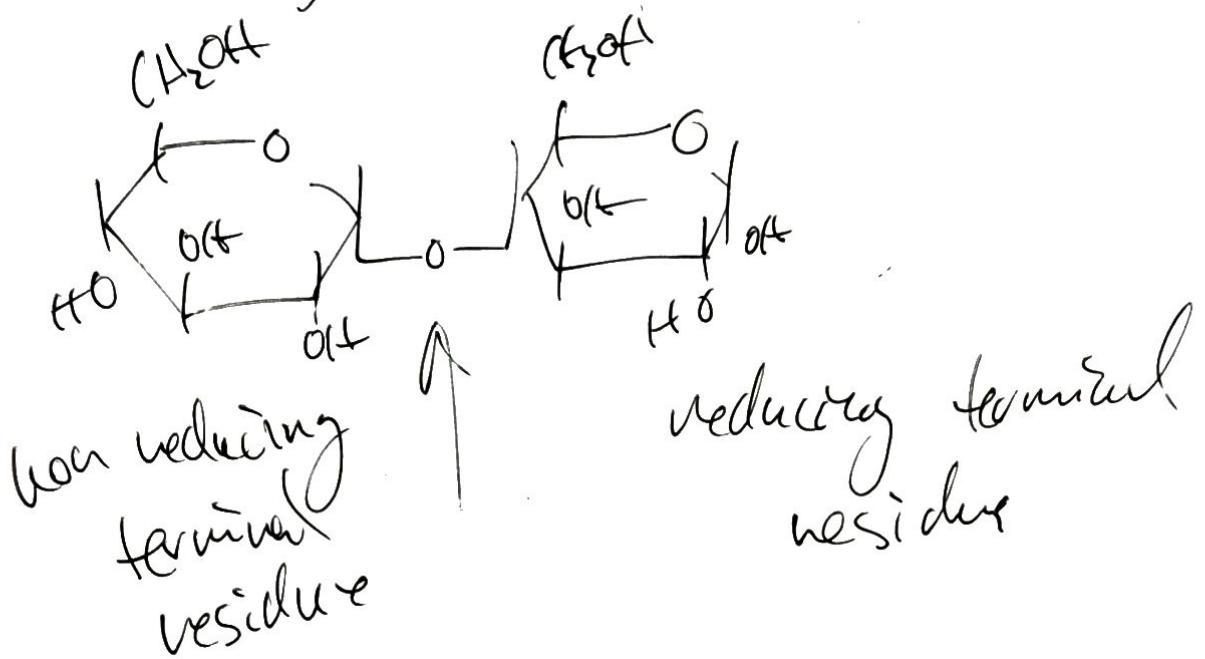
glycogen :  $\alpha(1 \rightarrow 6)$  branches every  
 $8 \alpha(1 \rightarrow 4)$  residues

Reducing sugars have free  
 aldehyde carbons

Nonreducing sugars have aldehyde  
 carbons involved in a glycosidic

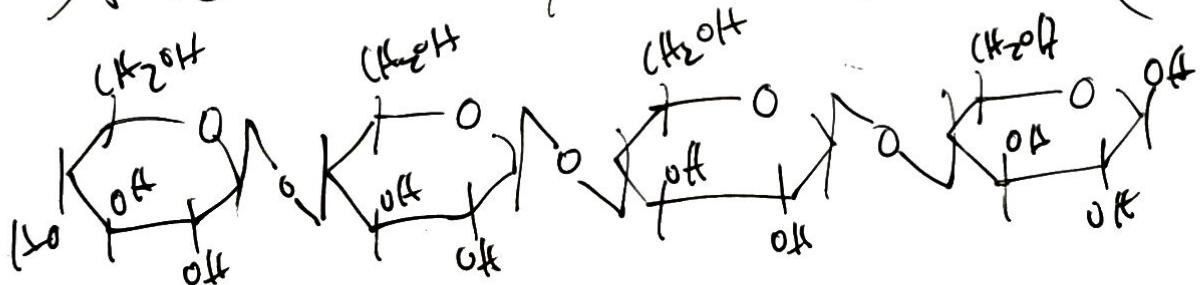
Maltose

$\alpha$ -D-glucose (1 $\rightarrow$ 4) -  $\alpha$ -D-glucose



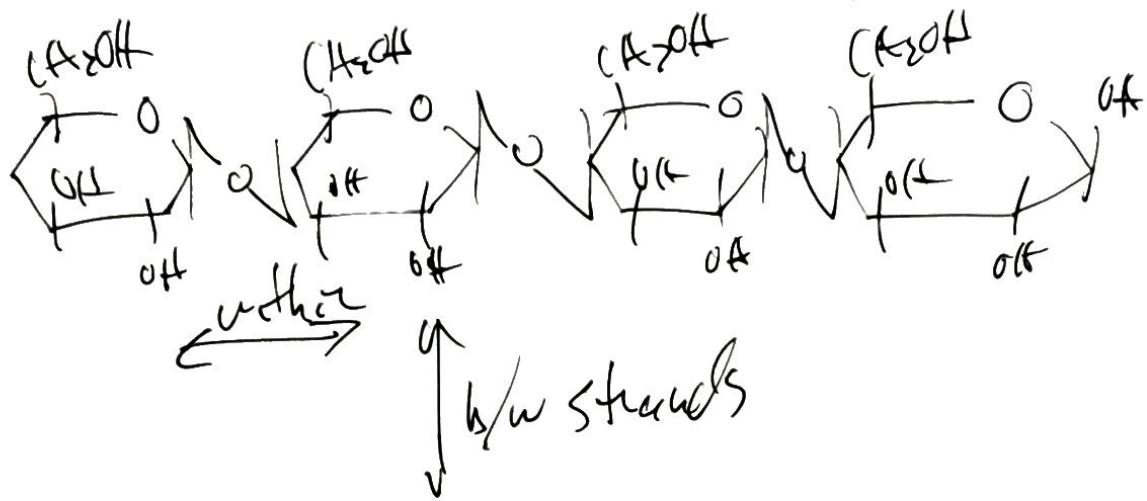
Every Polysaccharides =  $\alpha$  linked

Structural Polysaccharides =  $\beta$  - linked  
Cellulose ( $\beta$  1 $\rightarrow$ 4)



$\beta$ -linked polysaccharides have:

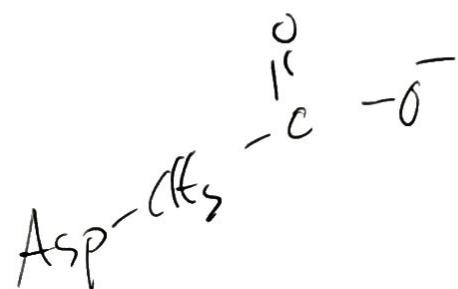
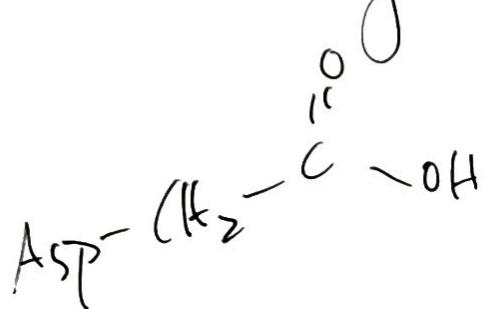
Inter  
molecular hydrogen bonds



modify cellulose @  
positions 2 and 3 to get  
different properties

N-acetyl group, arabinose

The more exposed to solvent you are  
the more likely you are to behave  
normally.



Predominant in  $\text{H}_2\text{O}$   
at  $\text{pK}_a \approx 7.2$

Burying an ionizable amino acid  
side chain inside an enzyme  
can shift its  $\text{pK}_a$  up or  
down.

KEY TO ENZYME FUNCTION