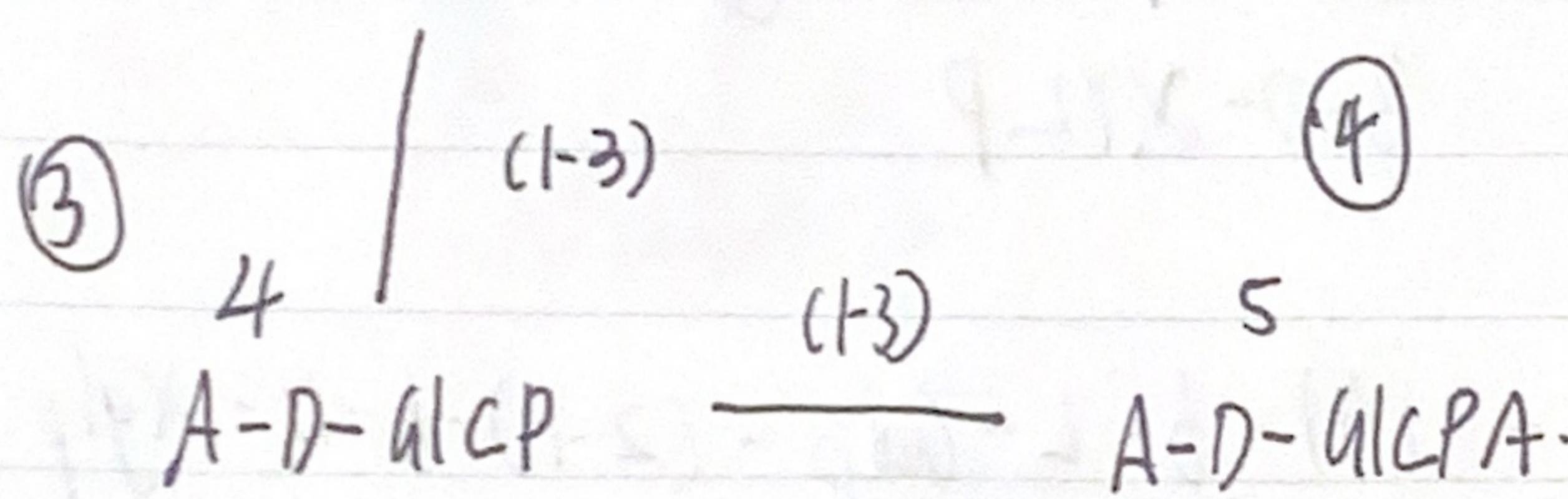
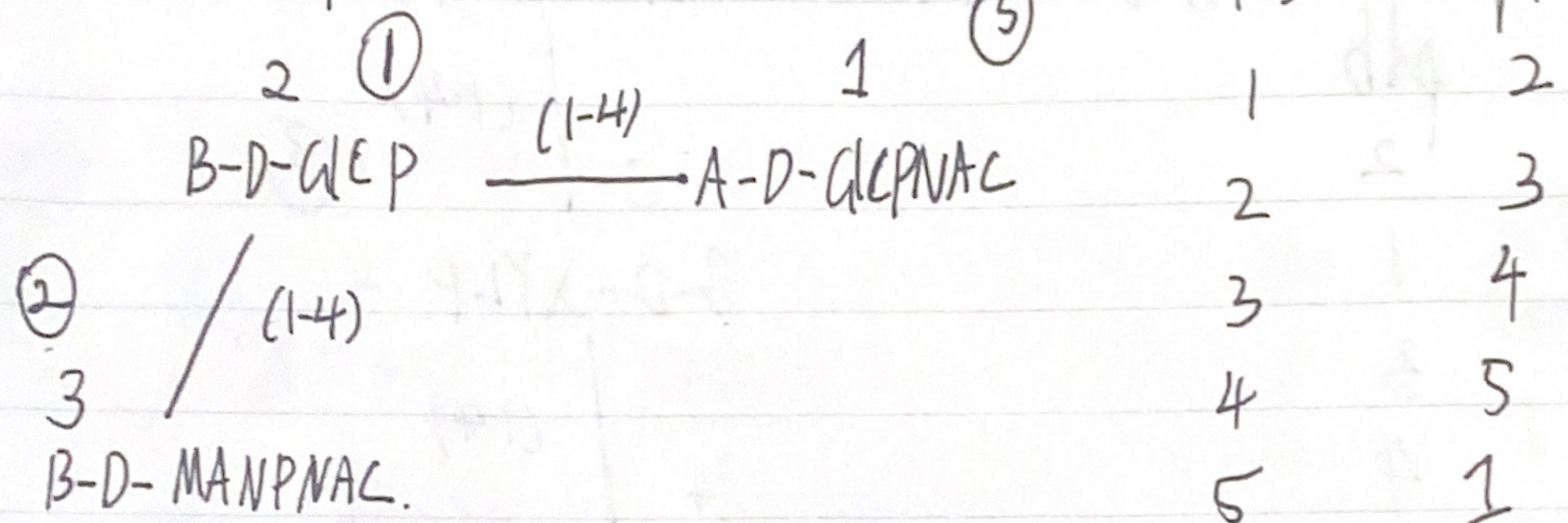


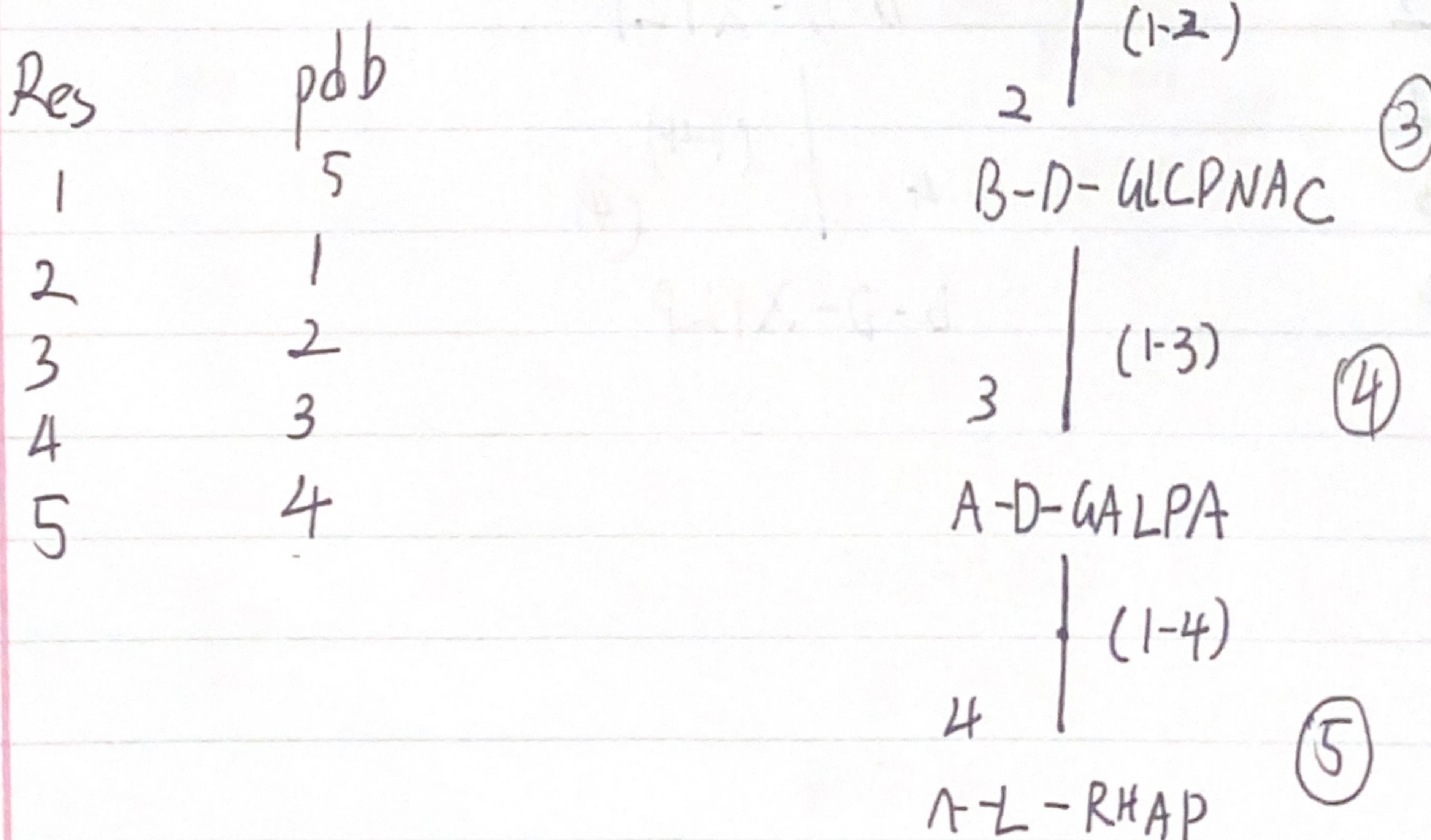
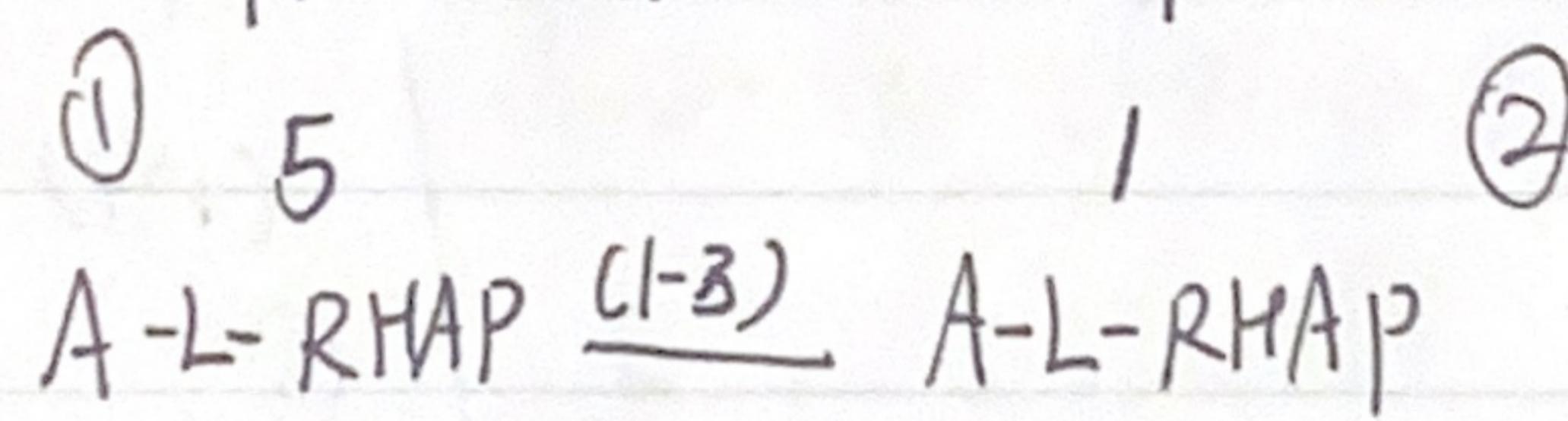
Correct by matching table.

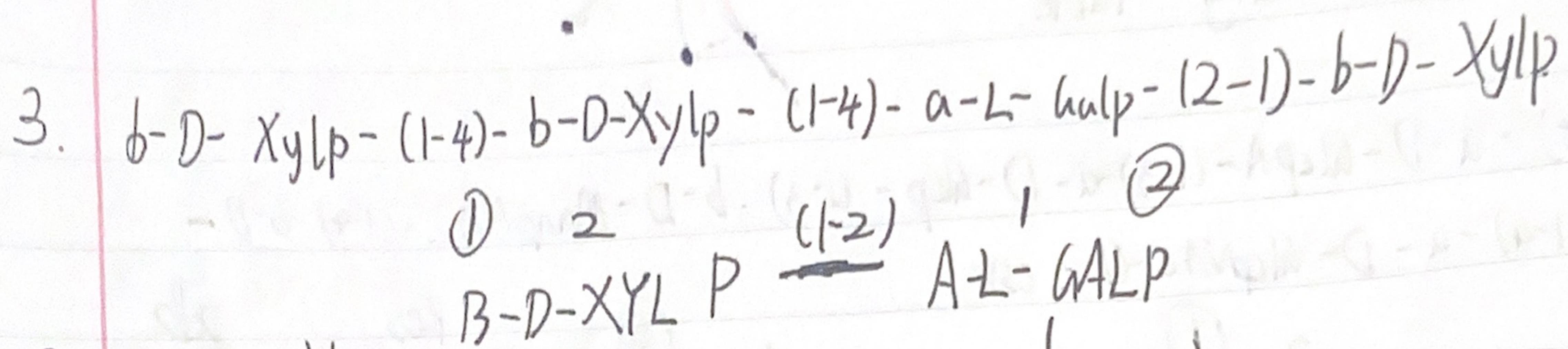


1. Report-4) -  $\alpha$ -D-GlcPA-(1-3)- $\alpha$ -D-GlcP-(1-3)- $\beta$ -D-ManpNAc-(1-4)- $\beta$ -D-GlcP-(1-4)- $\alpha$ -D-GlcPNAc-(1-. csv



2.  $\alpha$ -L-Rhap-(1-4)- $\alpha$ -D-GalpA-(1-3)- $\beta$ -D-GlcPNAc-(1-2)- $\alpha$ -L-Rhap-(3-1)- $\alpha$ -L-Rha,



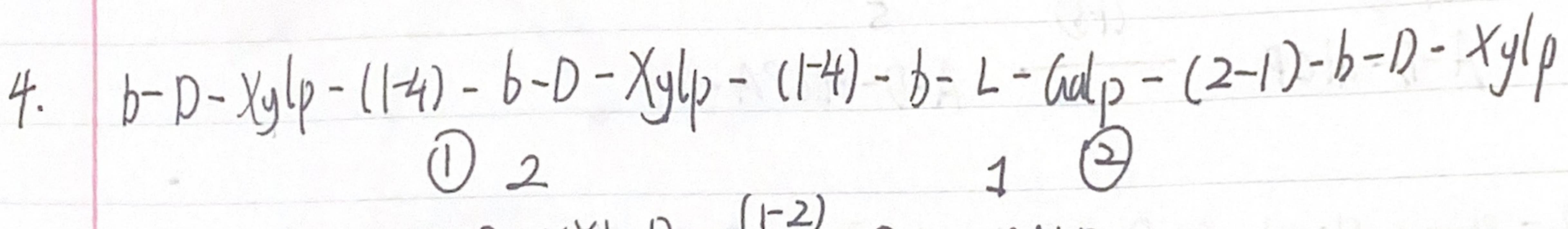


Res	pdb			
1	2			
2	1			$\text{B-D-XYLP}$
3	3			
4	4			

$(1-4)$       ③

$(1-4)$       ④

$\text{B-D-XYLP}$

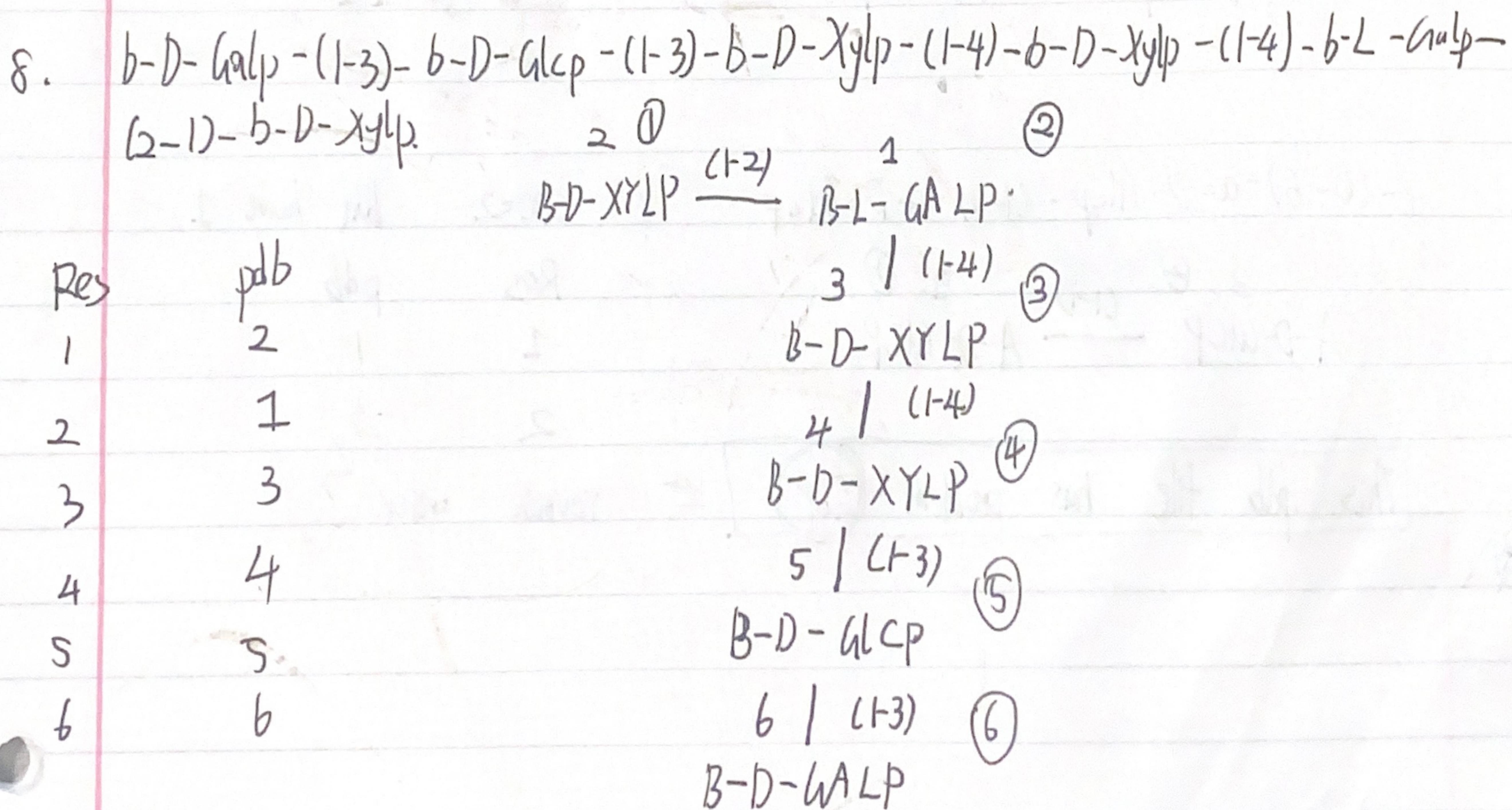
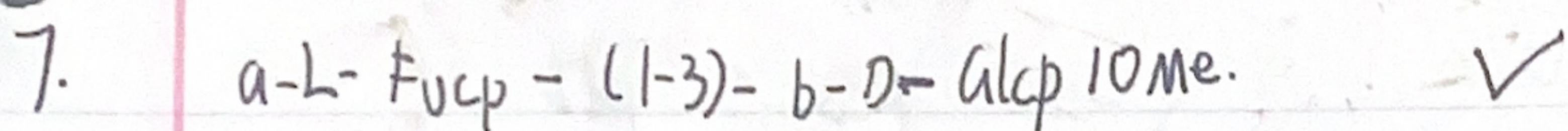
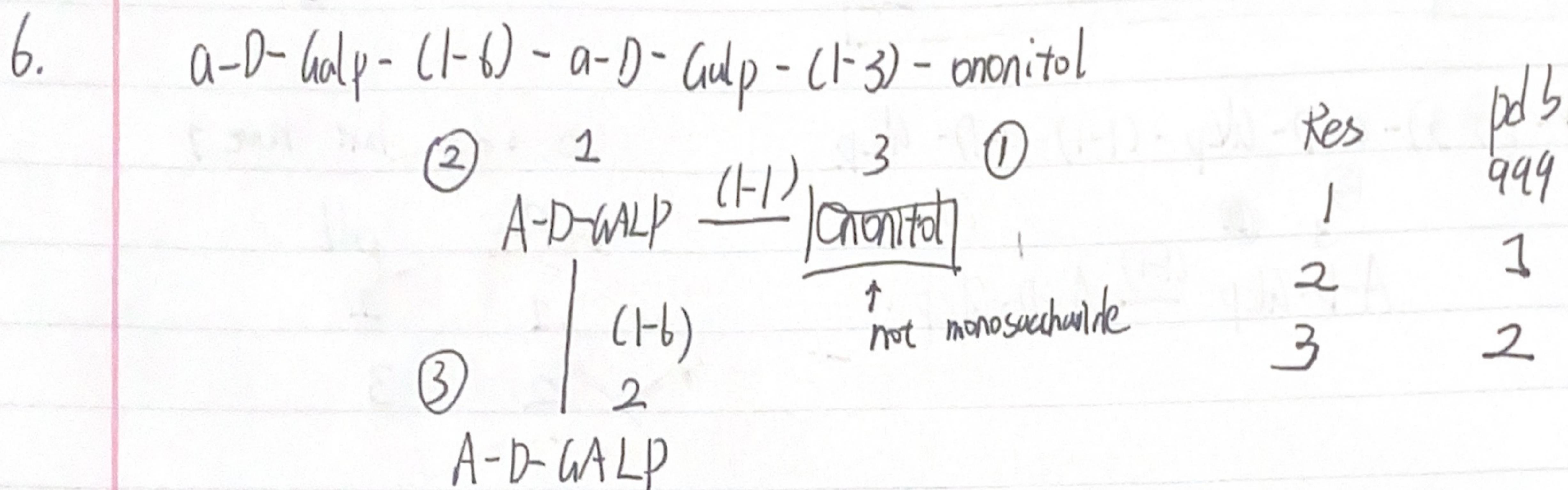
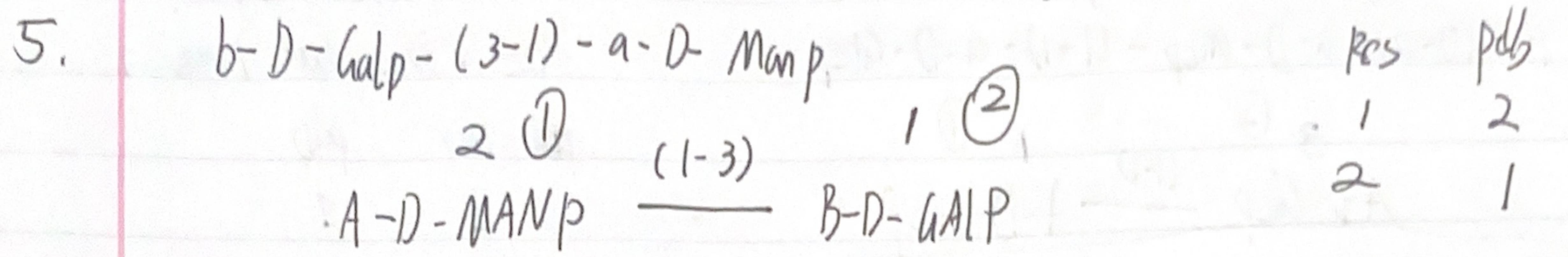


Res	pdb			
1	2			
2	1			$\text{B-D-XYLP}$
3	3			
4	4			

$(1-4)$       ③

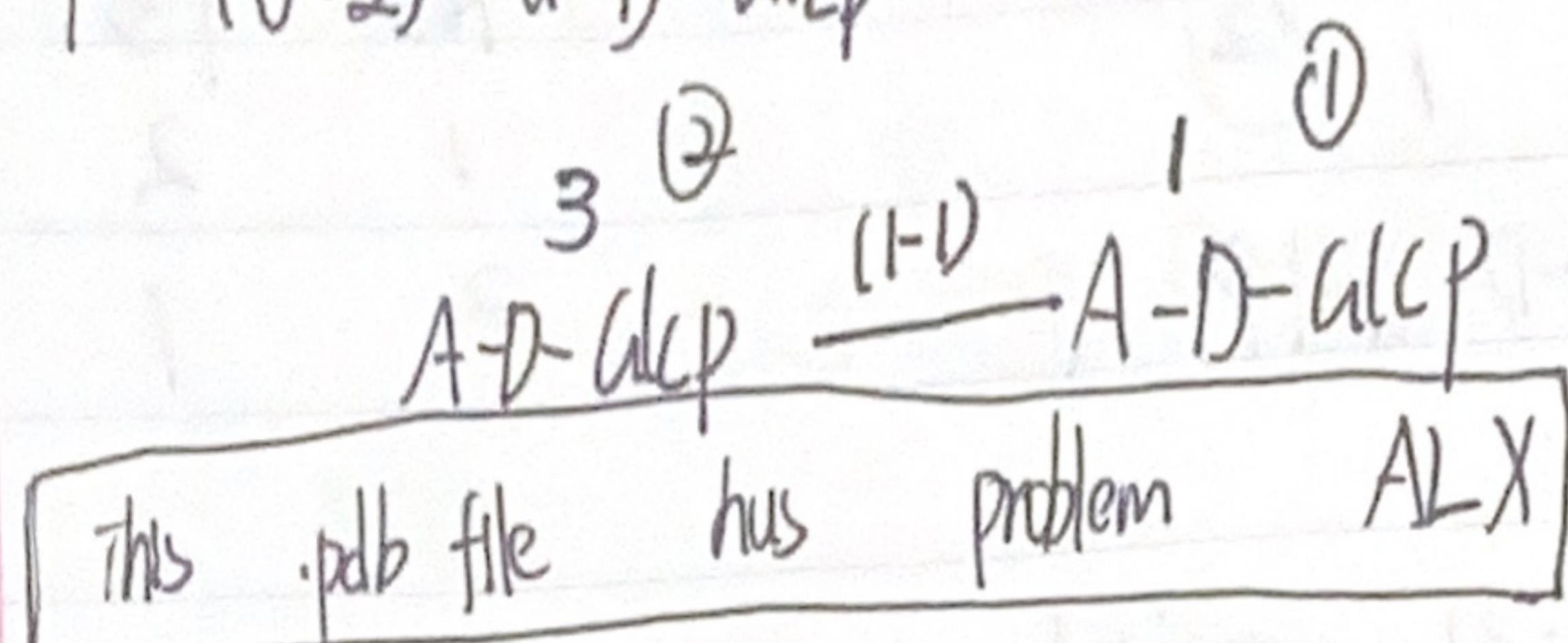
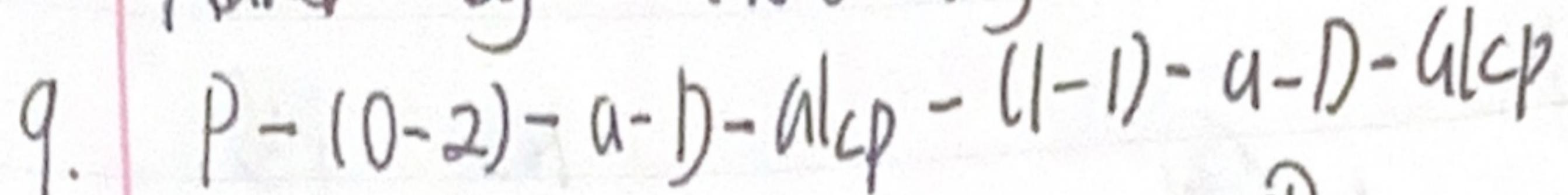
$(1-4)$       ④

$\text{B-D-XYLP}$



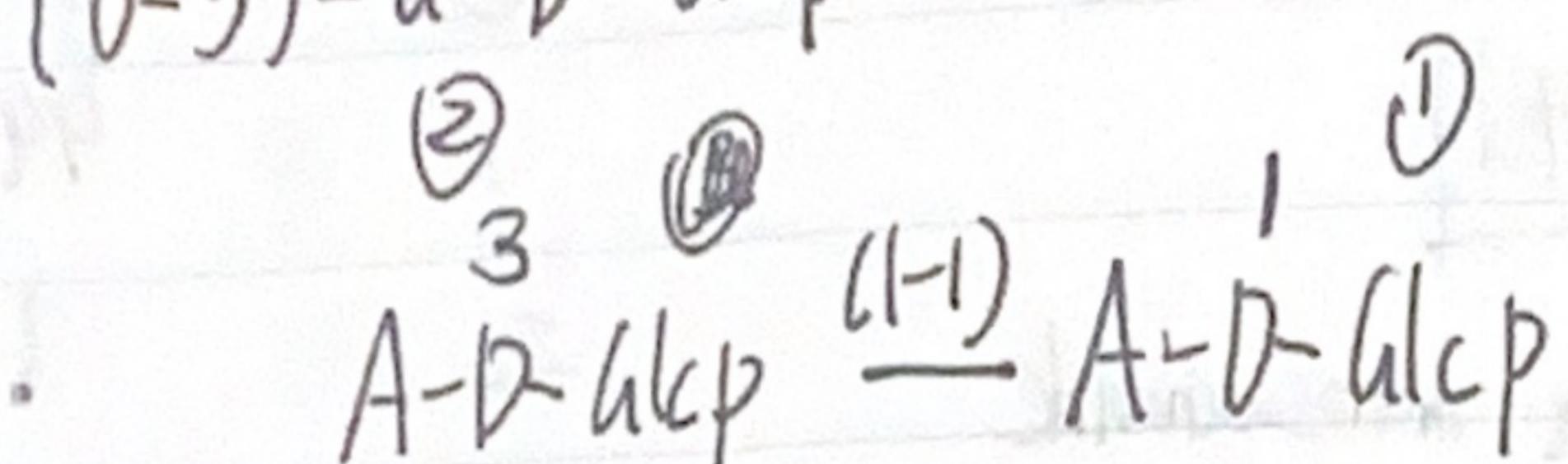
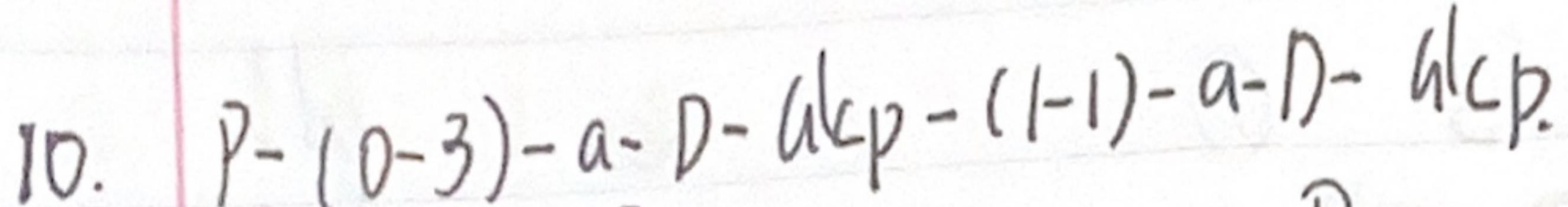
↓ take a look at this, WIA

1. Added by counting number of monosaccharide in a glycan,  
size = 2. but have 1.

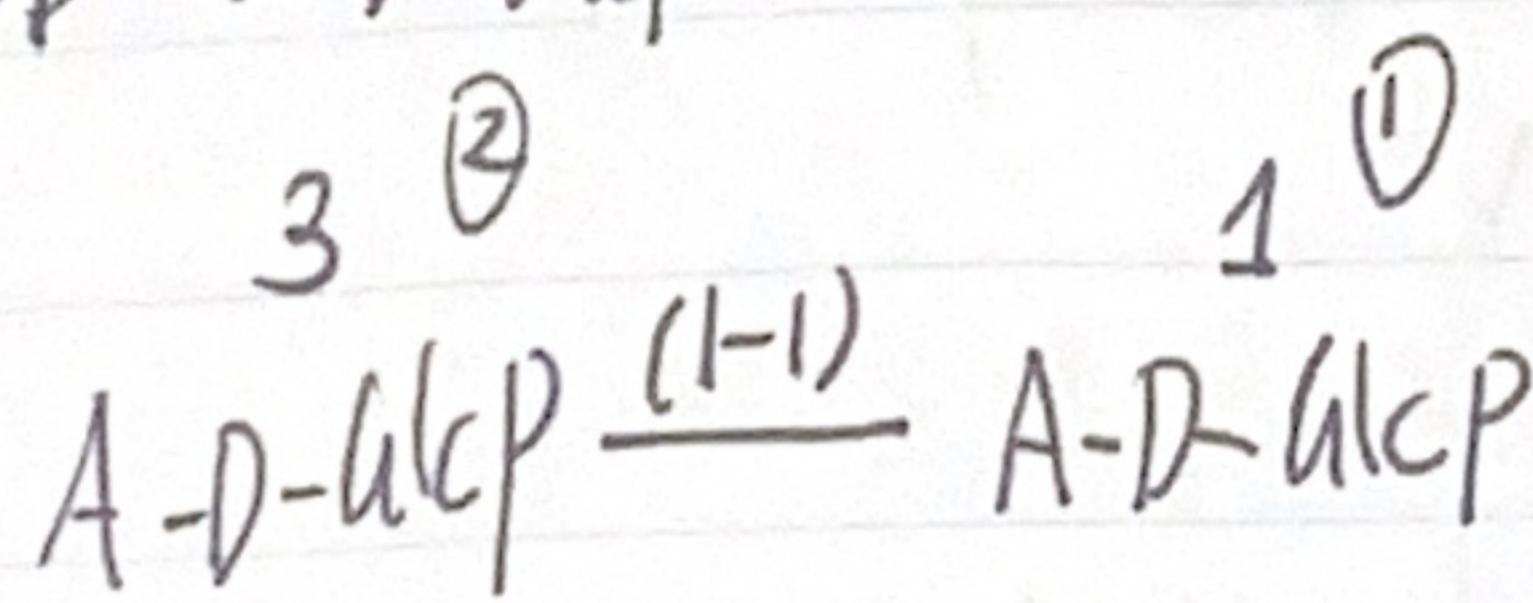
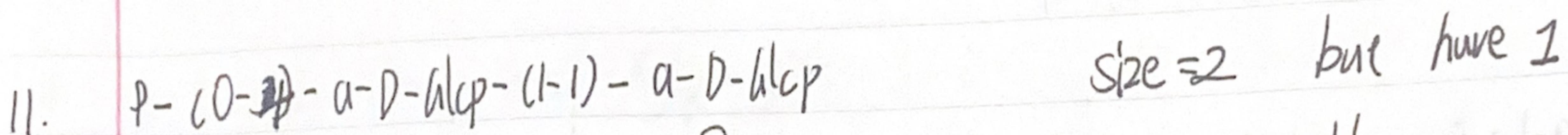


Res	pdb
1	1
2	3.

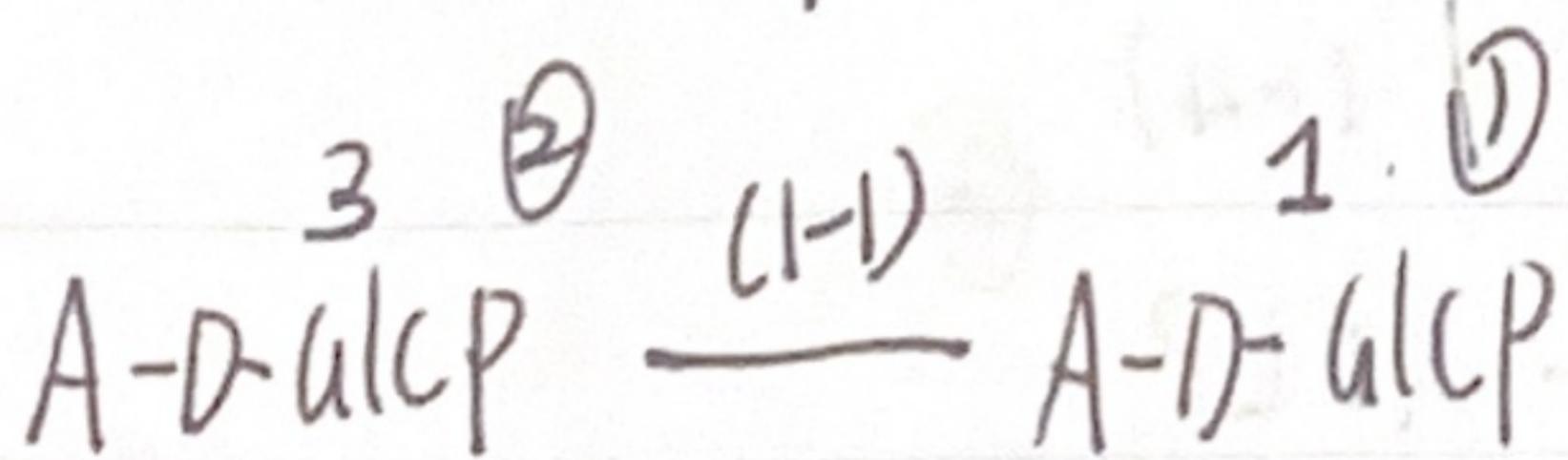
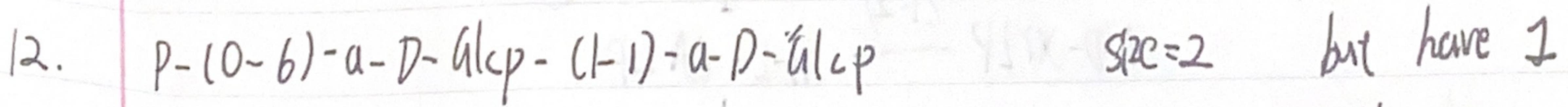
size = 2 but have 1.



Res	pdb
1	1
2	3.



Res	pdb
1	1
2	3.



Res	pdb
1	1
2	3

This .pdb file has problem GlcP ≠ reverse order?

AN

