

# A new approach on targeting aberrant glycans in cancer cells

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## Graphical Abstract

Production and evaluation  
of murine anti-STn mAbs



- Reformatting from murine to chimeric IgG;
- Humanization process

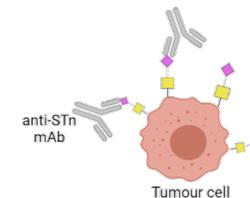


Evaluation of humanized mAbs:

- SPR
- Flow cytometry
- Glycan array



Selection of mAbs with the highest  
affinity and specificity to STn



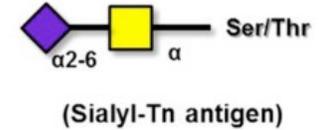
# A – Introduction

**Overexpression of sialyl-Tn (STn), a glycan antigen in Carcinomas** is usually associated with poor prognosis and reduced Overall Survival

**Cancer-Specific target**  
(antigen rarely expressed in normal cells)



Promising target using anti-glycan monoclonal antibodies (mAbs)



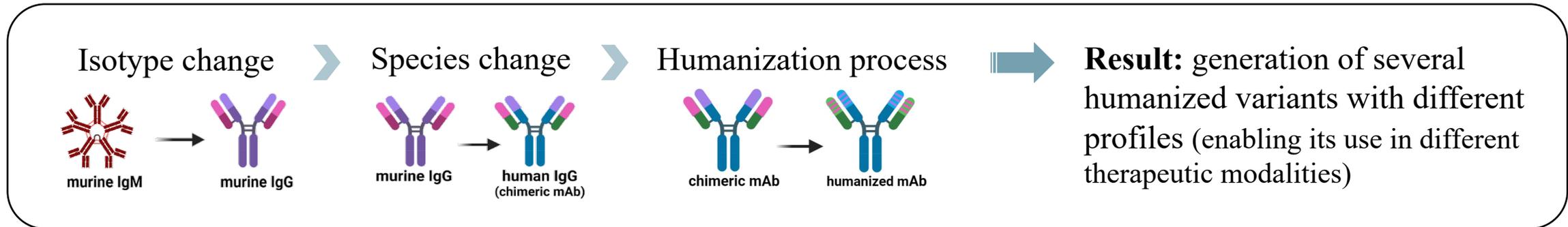
➤ Major challenges in the generation and characterization of highly specific anti-glycan mAbs:

- **Glycan recognition by mAbs** → Need to selectively discriminate between different carbohydrate structures  
→ Compete with the glycome at the cell surface
- **Glycan – mAb interaction** → Binding and affinity depend on:
  - antigen presentation
  - type of carrier (Glycoprotein) expressing the glycan antigen
  - amino acid to which the glycan is attached (Ser/Thr)

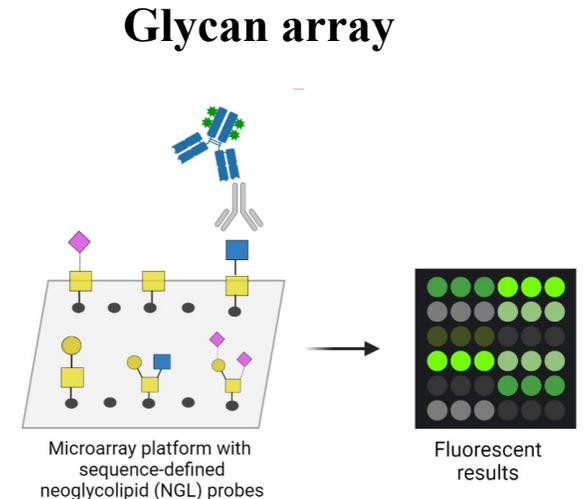
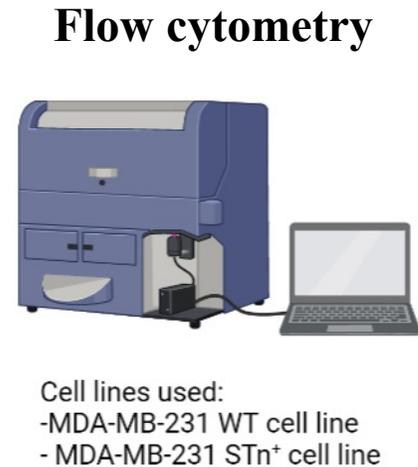
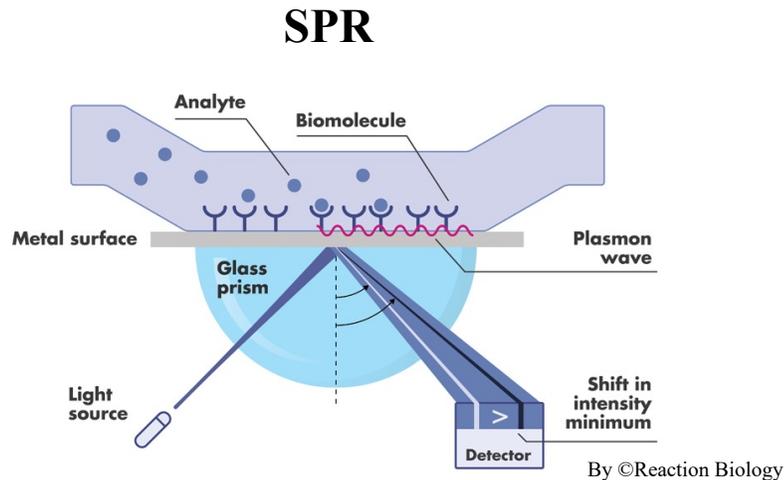
**Objective:** to develop humanized anti-STn antibodies with the most suitable affinity range for addressing solid tumors

# B – Methods

Hybridoma technology → **murine mAb<sup>1</sup>**, that was later reformatted for clinical application:



➤ Humanized mAbs specificity and affinity to STn was assessed by:



<sup>1</sup>Loureiro LR, Sousa DP, Ferreira D, Chai W, Lima L, Pereira C, et al. Novel monoclonal antibody L2A5 specifically targeting sialyl-Tn and short glycans terminated by alpha-2-6 sialic acids. Scientific Reports 2018;8:12196. <https://doi.org/10.1038/s41598-018-30421-w>.

# C – Results & Discussion

## SPR

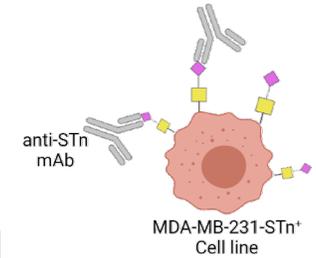
– Immobilized target: BSA-ser-STn conjugated ligand

Affinity measurement of antibodies to STn-serine

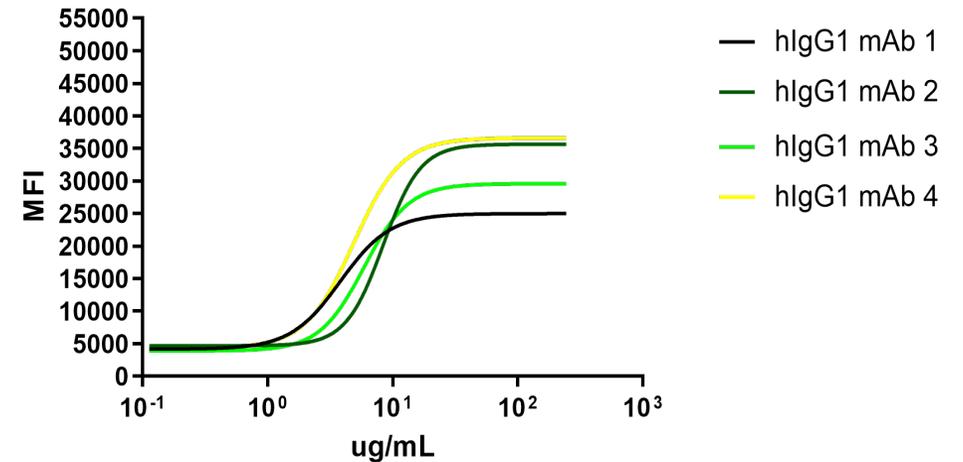
Ligand	Analyte	Chi <sup>2</sup> (RU <sup>2</sup> )	ka (1/Ms)	kd (1/s)	KD (M)	Rmax (RU)
BSA-STn-serine	hIgG1 mAb1	4.13E-01	1.18E+05	3.51E-03	2.97E-08	32.6
BSA-STn-serine	hIgG1 mAb2	2.55E-01	1.74E+05	5.53E-03	3.17E-08	17.8
BSA-STn-serine	hIgG1 mAb3	3.30E-01	1.71E+05	5.68E-03	3.32E-08	21.3
BSA-STn-serine	hIgG1 mAb4	1.35E-01	1.32E+05	5.19E-03	3.94E-08	13.8

↓ KD    ↑ Binding affinity

## Flow cytometry



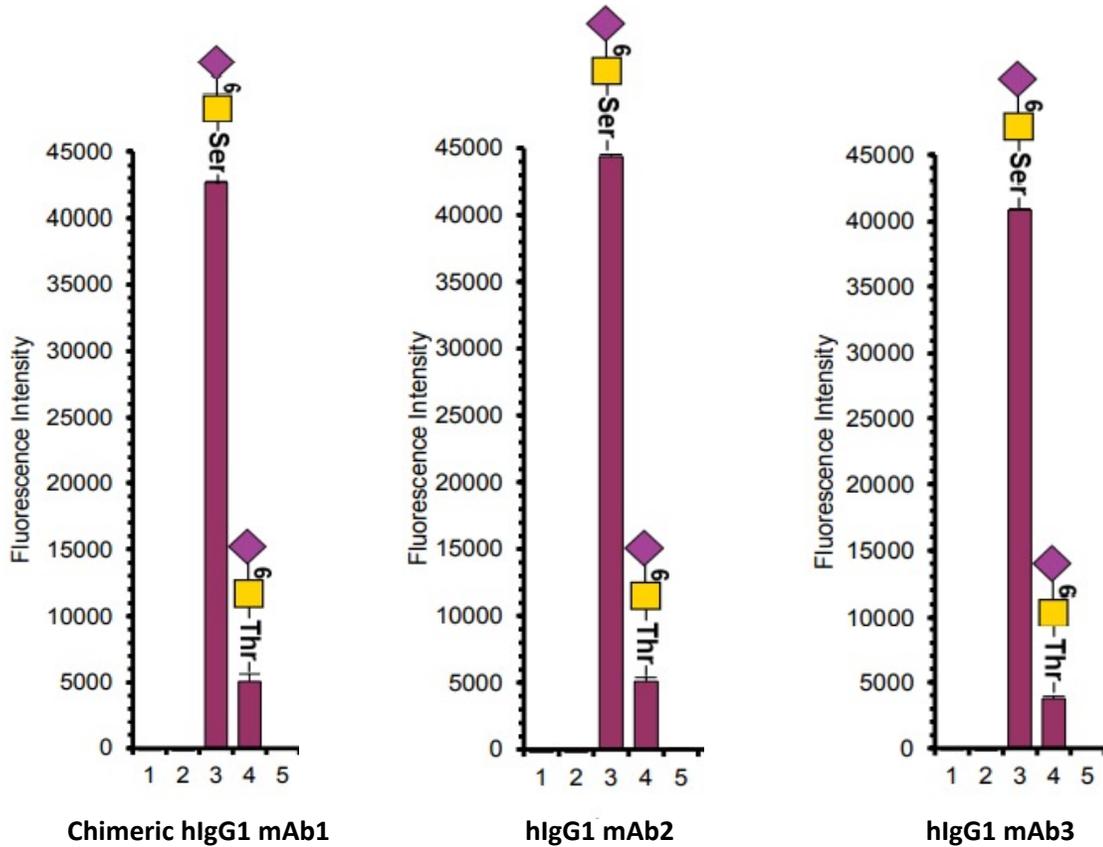
Flow Cytometry - MDA-MD-231 STn+



We screened and characterized some of the different mAbs, evaluating glycan affinity and cellular binding to STn<sup>+</sup> cell lines.

# C – Results & Discussion

## Glycan array



<b>STn(Ac)</b>	NeuAc $\alpha$ -	O-Ser-NH-DA	3
	6GalNAc $\alpha$ -	O-Thr-NH-DA	4

NGL-DHPA microarray platform



32 sialylated and core *O*-glycan related probes

Reformatted antibodies improved selectivity and STn specificity when compared to the murine;

Showed high binding intensity to STn.

# D – Impact

- Several anti-STn mAb variants, were generated, having a broad range of STn affinity;
- By developing mAbs with different profiles, several tumour types at different stages can be targeted by using various therapeutic modalities;
- Due to similarities in the glycosylation pattern of cancer cells and viruses, the antiviral potential of anti-STn mAbs is being evaluated and explored further.

## Acknowledgments

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