

Therapeutic antibody glycoforms: structures & function



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The stability of biopharmaceuticals – getting the chemistry right

RSC - MIBio 2014: Cambridge, September 30th 2014

Antibody therapeutics: Glycoform structures & function

Antibody classes & isotypes

Antibody effector activities

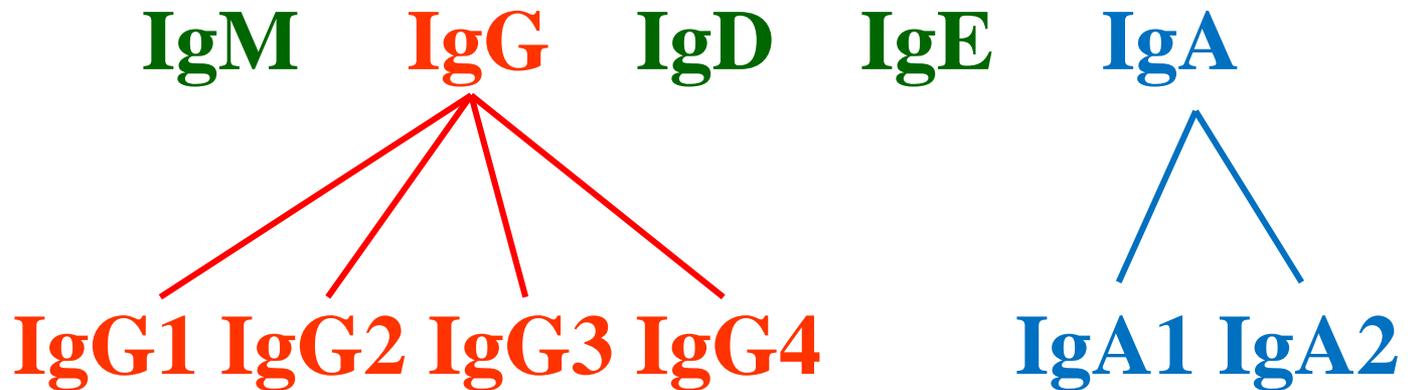
IgG-Fc glycoforms

Ligand binding sites

Engineering “biobetters”

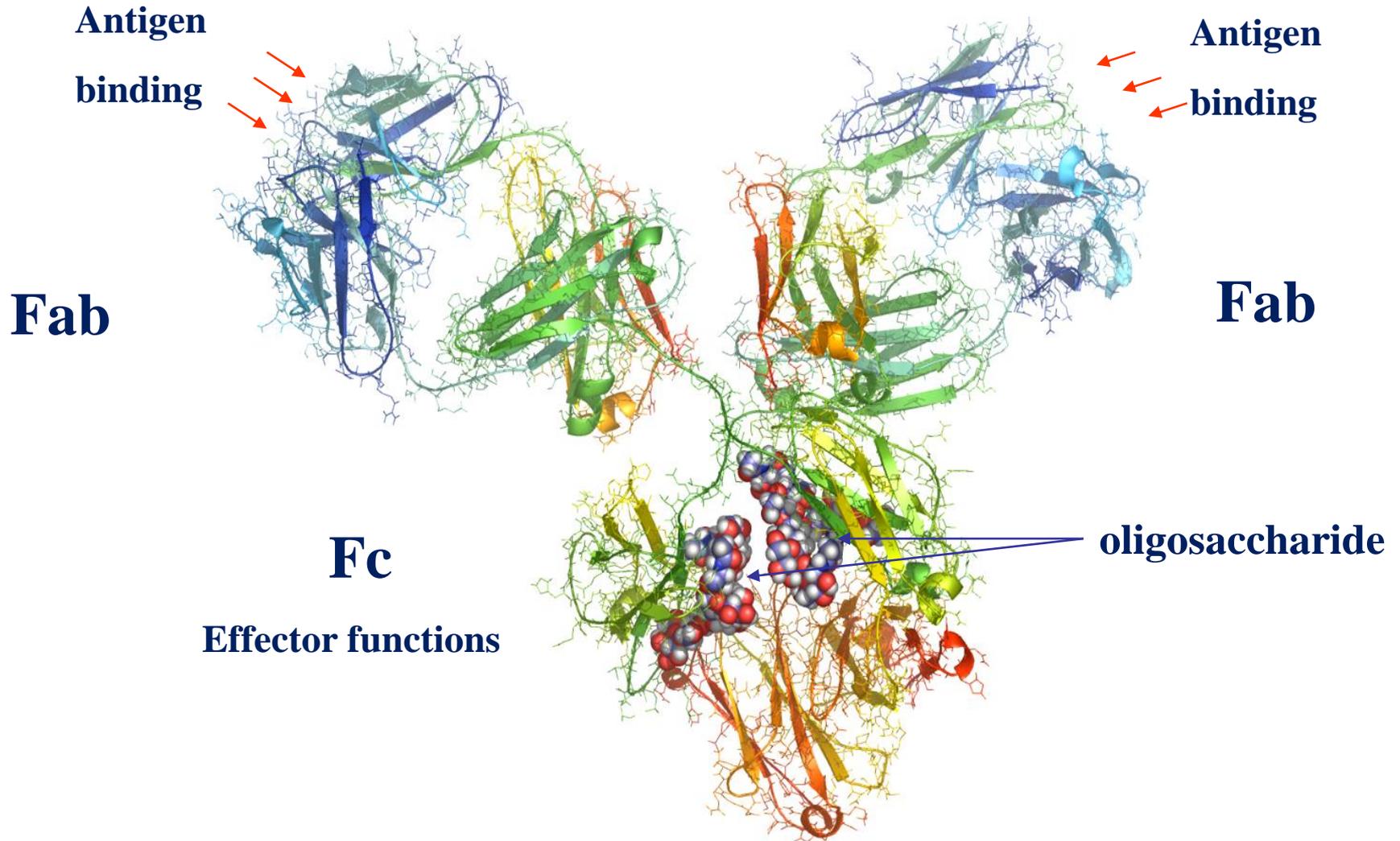
Human immunoglobulin classes & subclasses

The result of gene duplication, mutation & selection



The humoral immune response is “orchestrated” to provide optimal protection to a given “insult”

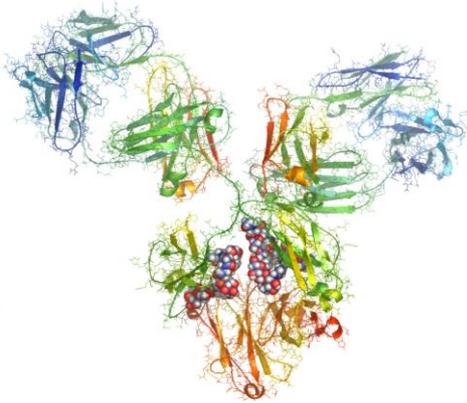
Domain structure of IgG



The human IgG subclasses

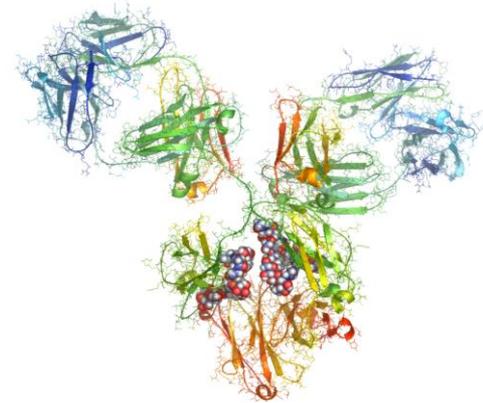
IgG1

~ 60 %



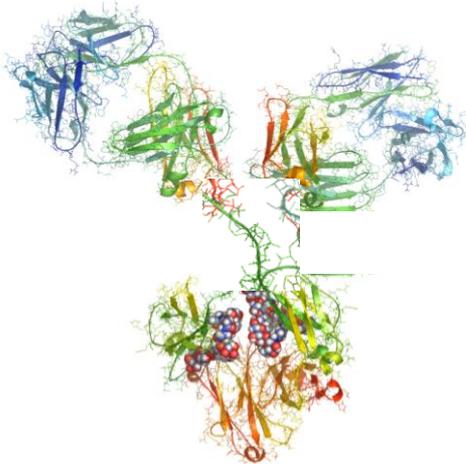
IgG2

~ 25 %



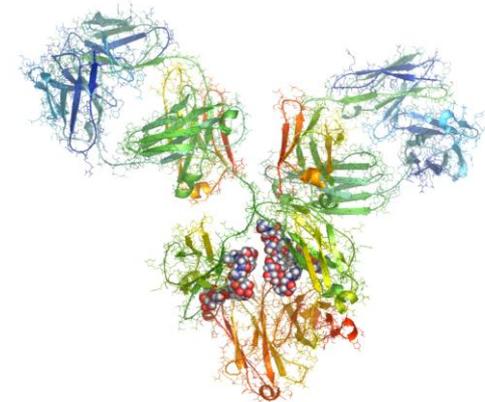
IgG3

~ 10 %



IgG4

~ 5 %



Jefferis R. Arch. Biochem. Biophys. (2012)

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Ligands mediating IgG-Fc effector activities

FcγRI (1,3,4) **FcγRIIa/b**** (1,2,3) **FcγRIIIa/b**** (1,3,4)

phagocytosis, antibody dependent cellular cytotoxicity (ADCC),
apoptosis, generation of superoxide, release of enzymes

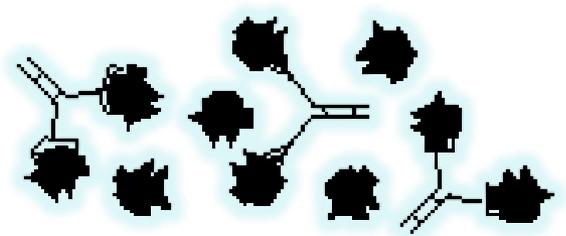
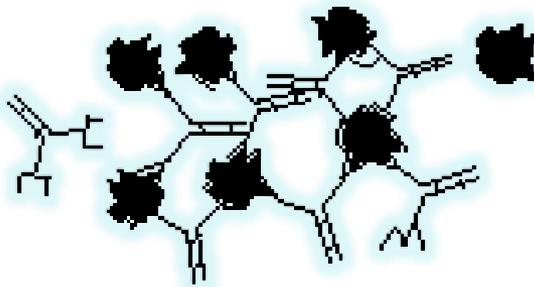
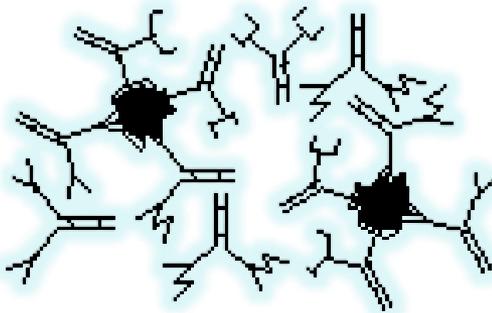
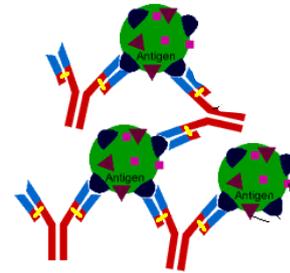
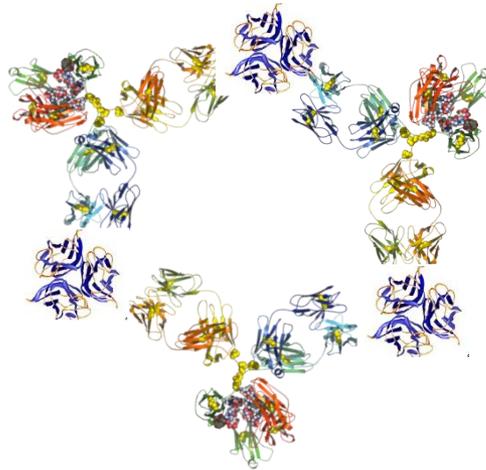
* dependent on FcγR polymorphisms; * dependent on glycoform

C1q: (IgG1,3) complement dependent cytotoxicity (CDC)

MBL (IgG1,2,3,4) Lectin pathway (CDC)

FcRn (1,2,3,4) Catabolism & placental transport

Formation of antigen/antibody complexes



Jefferis R., Steensgaard, J. Immunology. 46:751-60 (1982)

Activities of aglycosylated IgG-Fc

FcγRI	activation	reduced x 100
FcγRII	binding	abolished
FcγRIII	activation	abolished
C1	“ “	abolished
MBL	“ “	abolished
FcRn	binding	unaffected

Aglycosylated antibody products

UCB

Cimzia (Fab-Peg) [TNF– licensed]

Genentech
A Member of the Roche Group

Lucentis (Fab) [VEGFA-licensed]

TOLERX
Immune resolve.

Otelixizumab (CD3) (Asn₂₉₇ – Ala)



Bristol-Myers Squibb

BMS-945429 (IL-6) (Asn₂₉₇ – Ala)

Take home message:

Glycosylation is a Critical quality Attribute (CQA)

**Therapeutic recombinant antibody production processes
must deliver 100 % occupancy or 0 % occupancy**

The glycoform profile of an antibody is achieved as a:

Quality by Design parameter (QbD)

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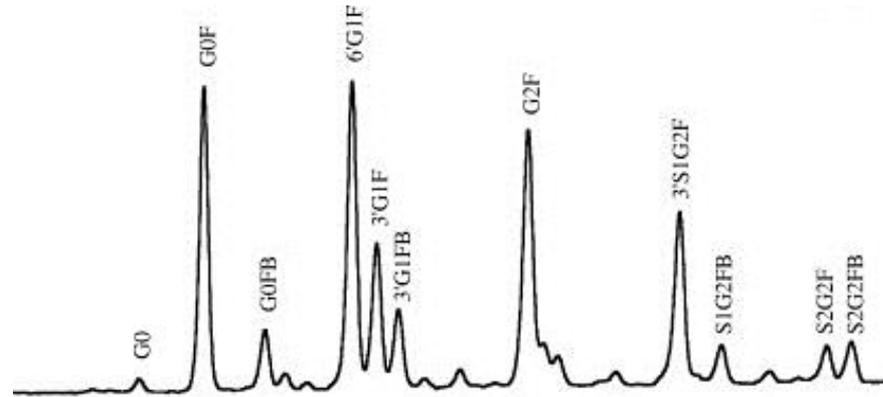
IgG-Fc glycoforms: stability & function

Ligand binding sites

Engineering “biobetters”

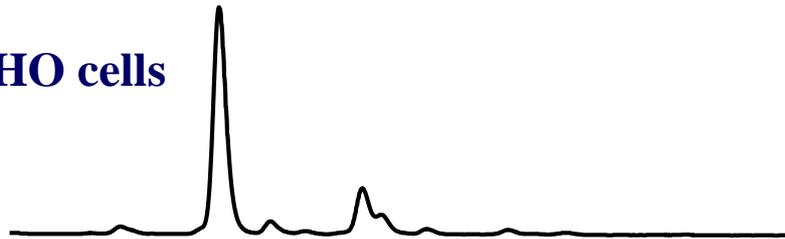
Oligosaccharides released from IgG

Polyclonal IgG



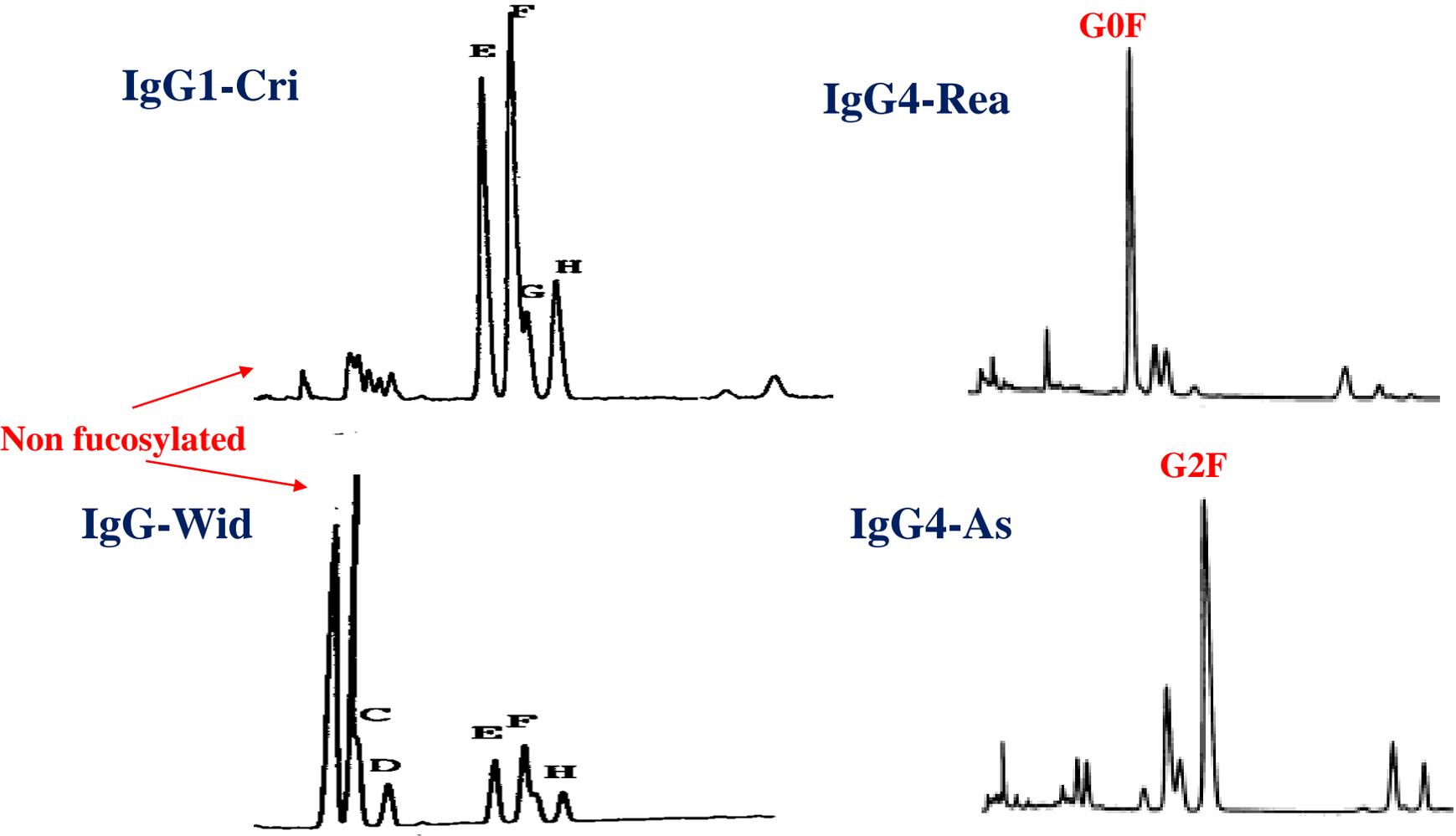
Anumula KR J Immunol Meth 382:167–176 (2012)

Adalimumab: CHO cells



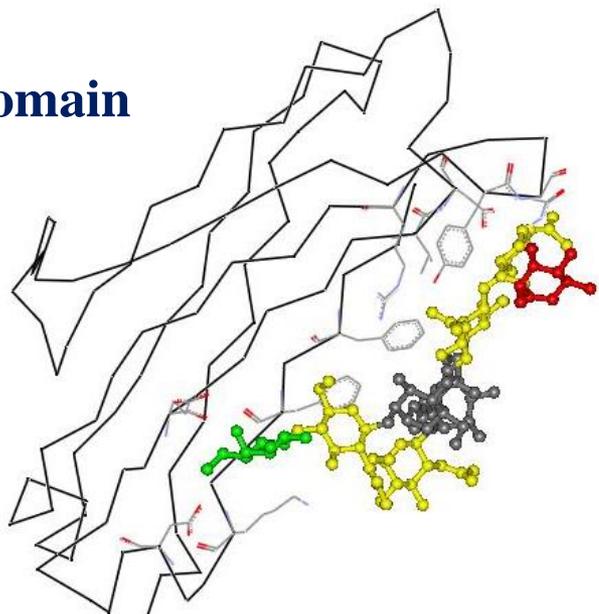
Mimura, Y. et al. Therapeutic antibodies: (Ed) An, Z., Wiley, 2009

Oligosaccharides released from monoclonal IgG1 & IgG4

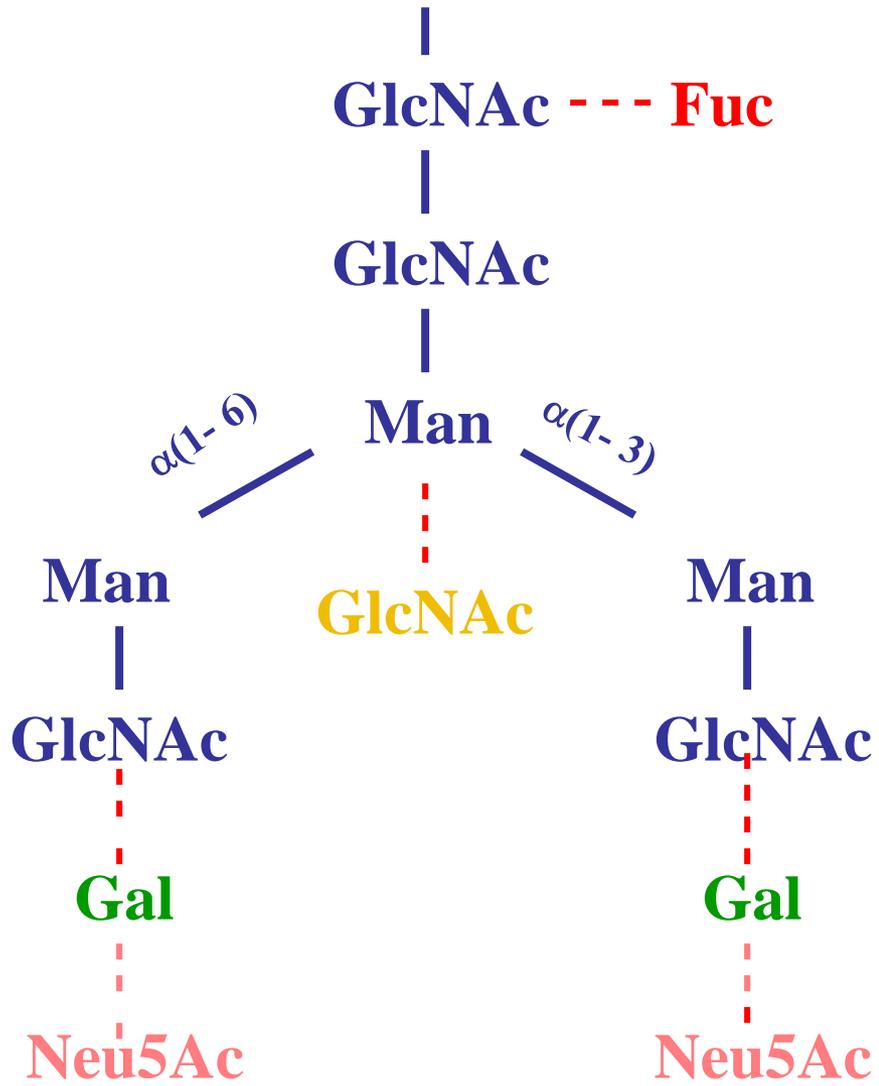


~Gln-Tyr-Asn₂₉₇-Ser-Thr-Tyr-Arg-~

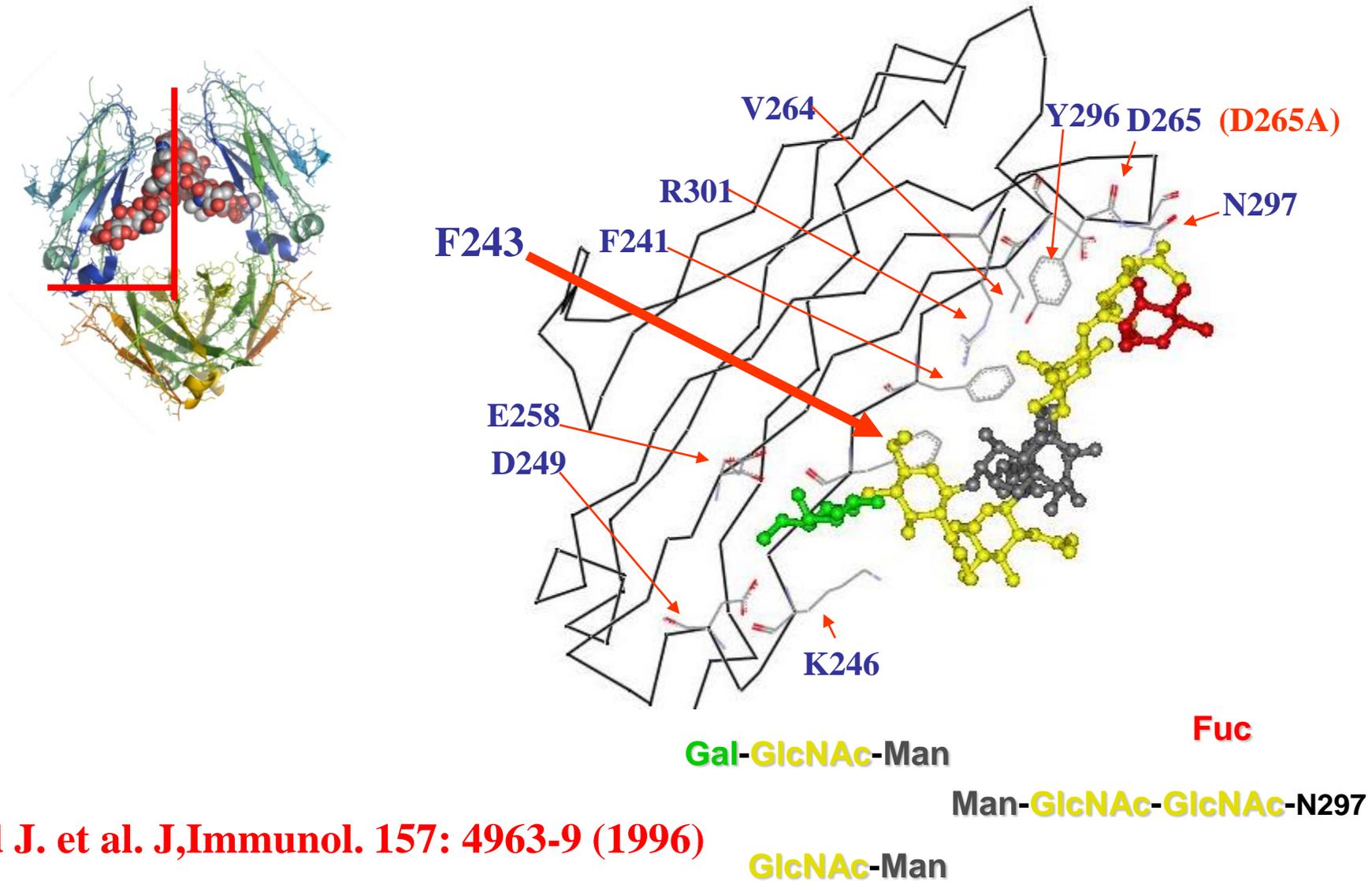
CH2 domain



648/128 possible glycoforms



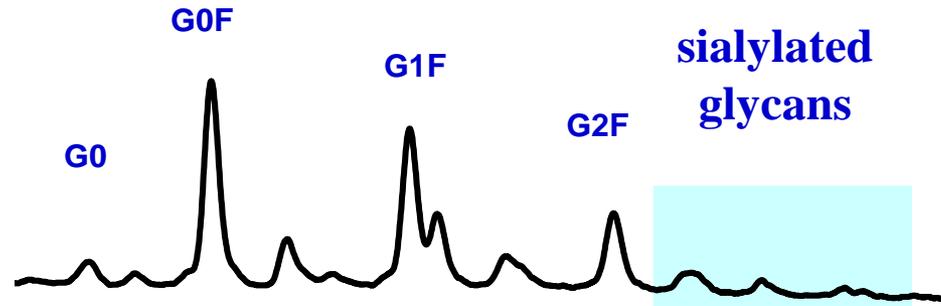
Alanine replacement of amino acids residues forming non-covalent contacts with the oligosaccharide



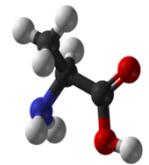
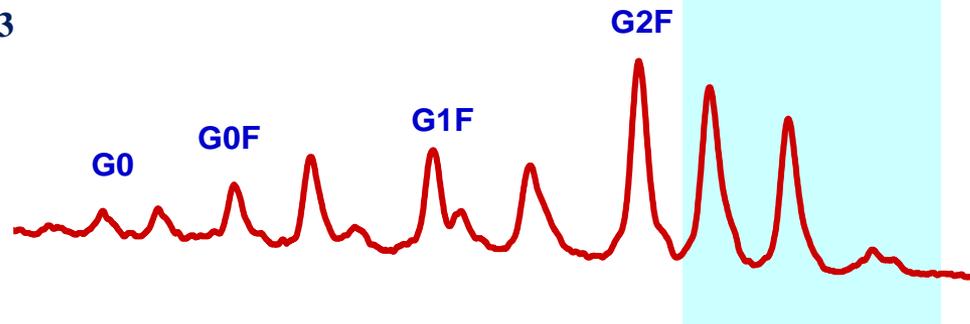
Lund J. et al. J,Immunol. 157: 4963-9 (1996)

The F/A₂₄₃ mutant yields high levels of sialylation

IgG3 WT

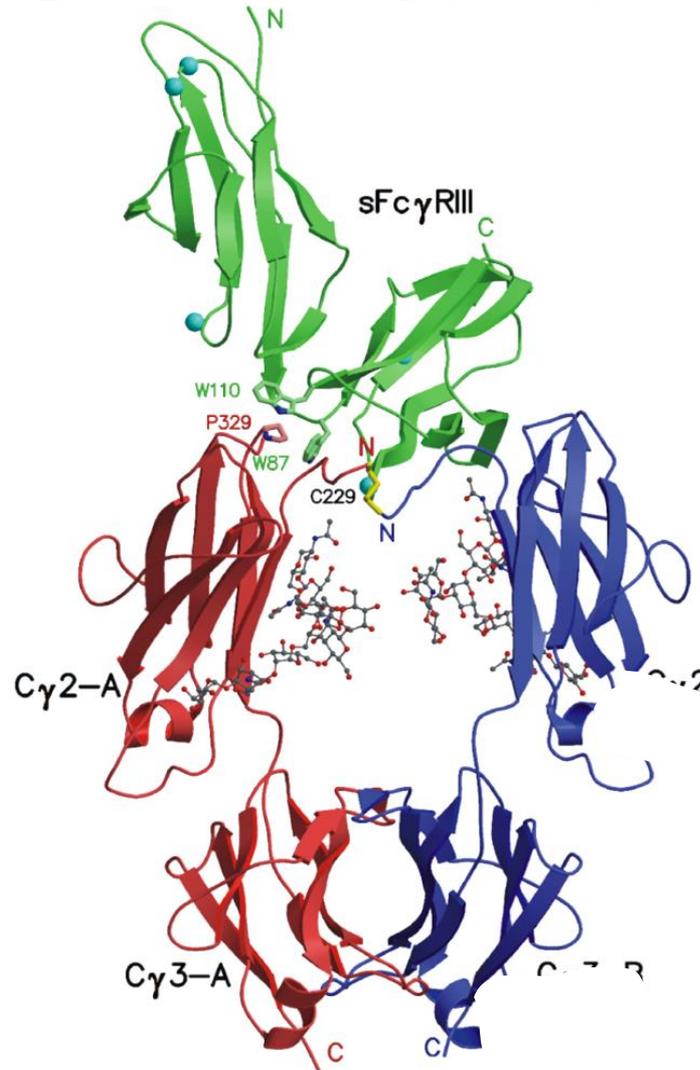


IgG3 F/A₂₄₃



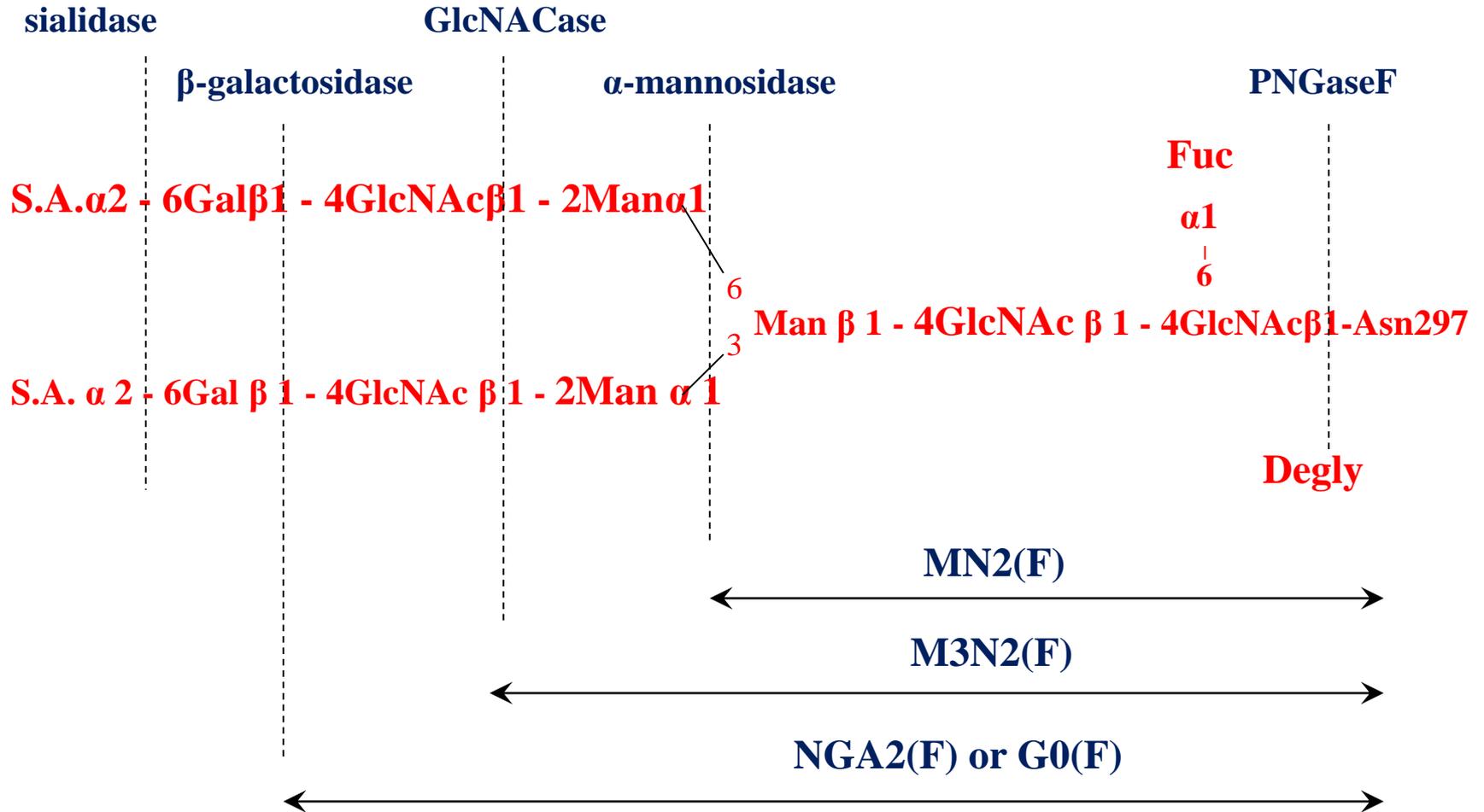
Mimura Y., Jefferis R., Rudd P. et al. unpublished
Lund J, Jefferis R., et al. J Immunol. 157:4963-4969 (1996).

IgG1-Fc in complex with aglycosylated FcγRIII



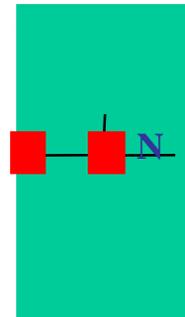
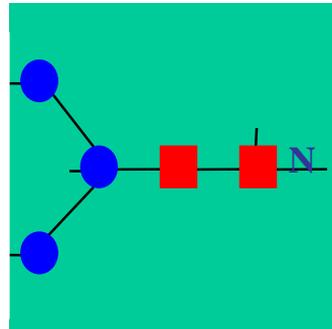
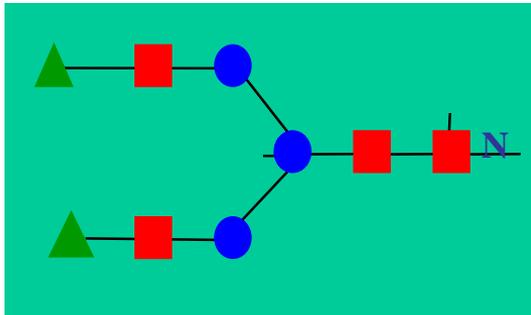
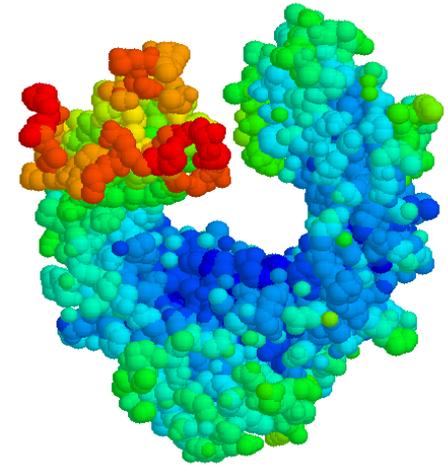
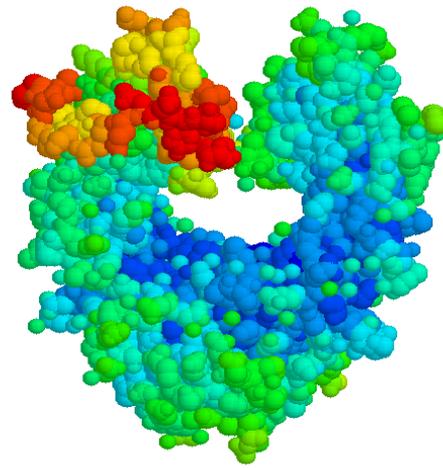
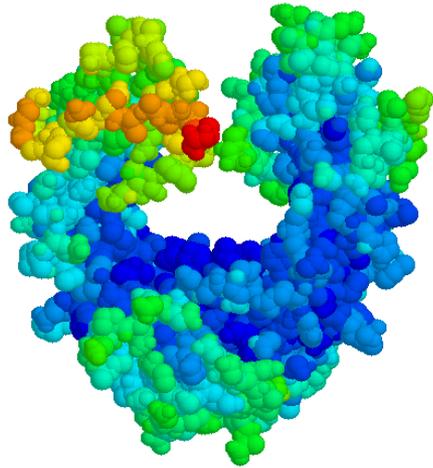
Sondermann P et al. Nature 406:267-273 (2000)
Radaev S et al. Mol Immunol. 38:1073-1083 (2002)

Preparation of homogeneous glycoforms of IgG



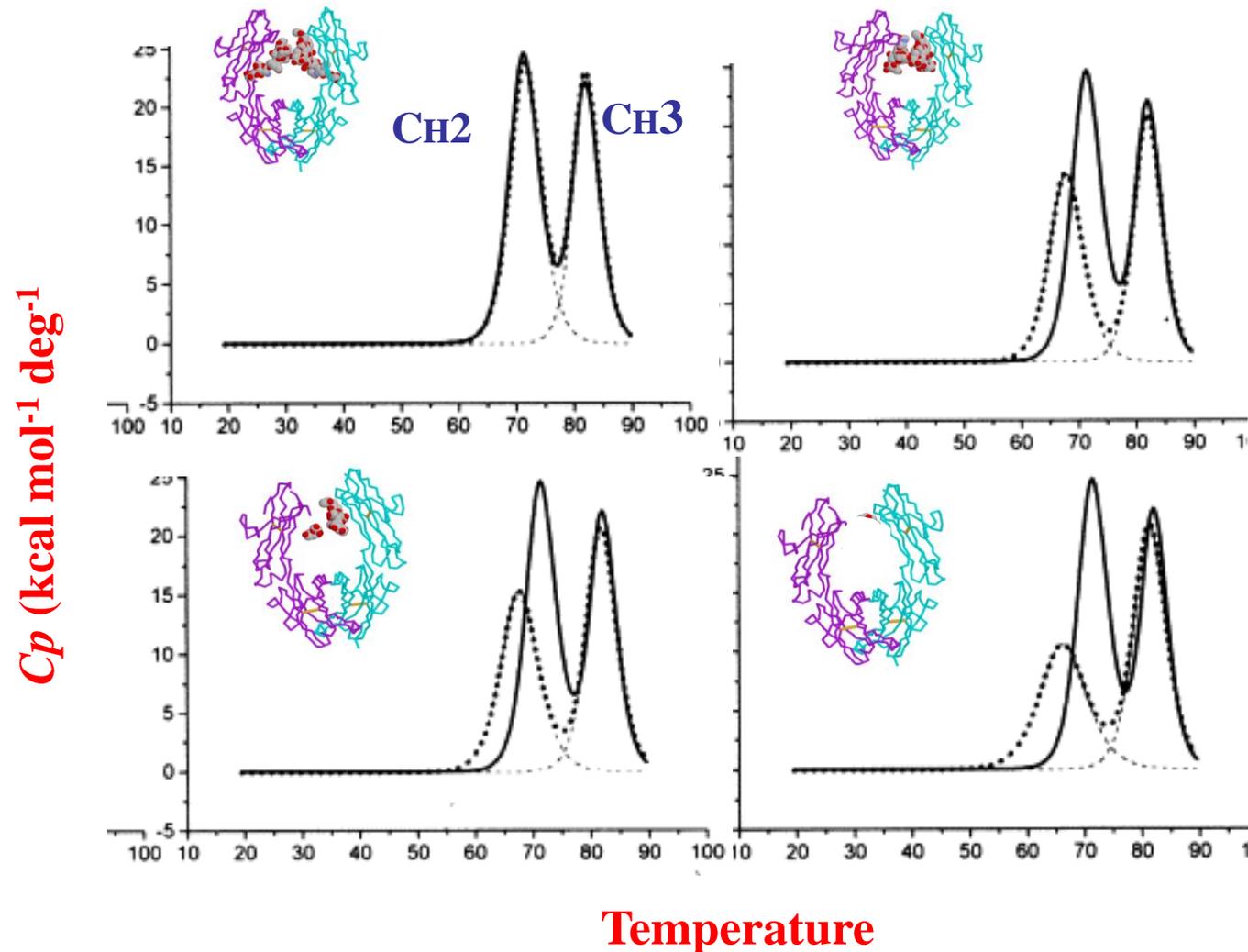
Mimura Y. et al. J Biol Chem. 276:45539-45547.(2001)

Truncated glycoforms of IgG-Fc



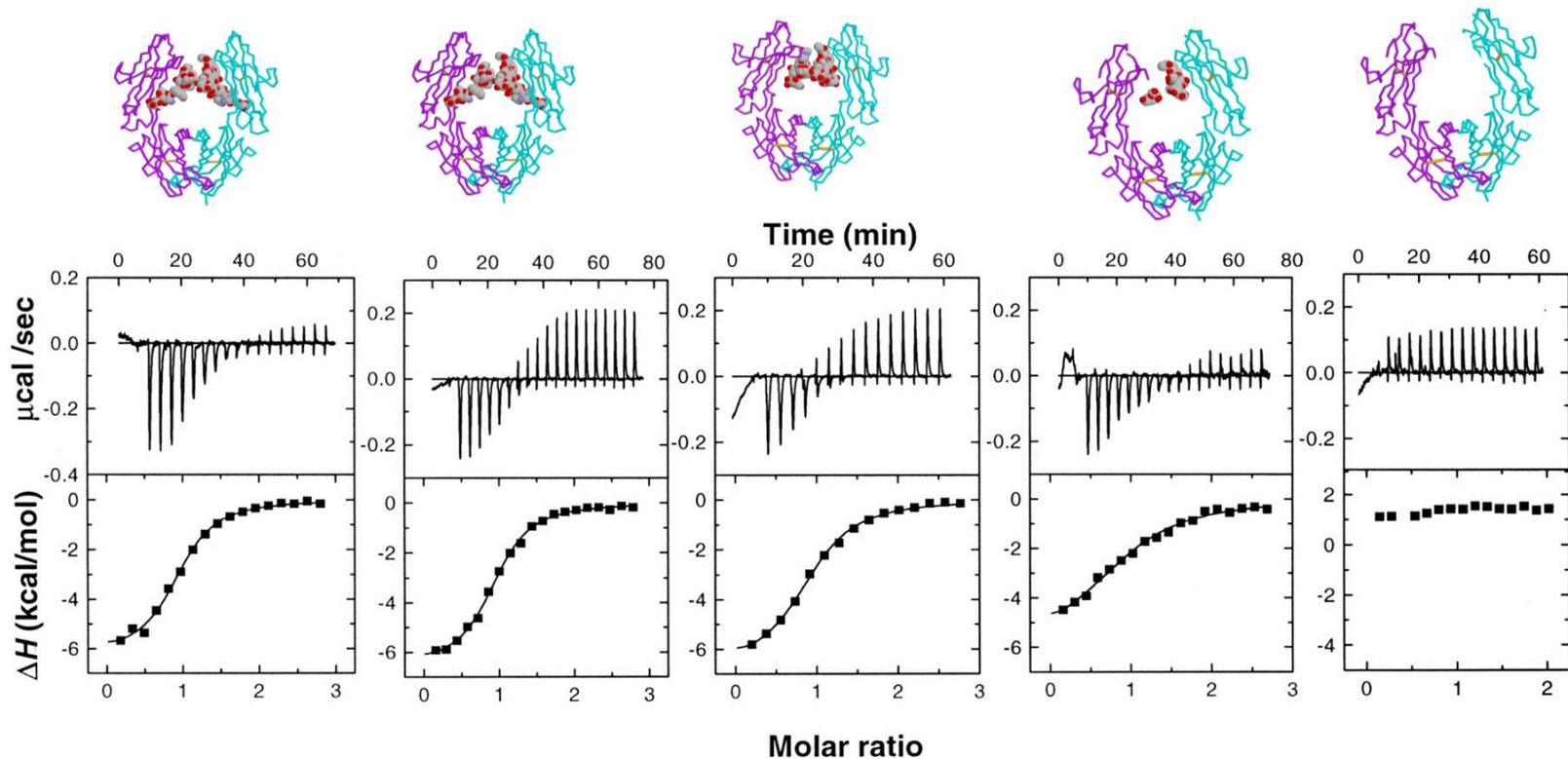
Krapp, S., Mimura, Y. et al. *J.Mol.Biol.* 325:979-989 (2003)

Differential scanning micro calorimetry of IgG-Fc



Mimura Y. et al. Mol Immunol. 37:697-706 (2000)

Binding isotherms for FcγRIIb/IgG1-Fc glycoforms

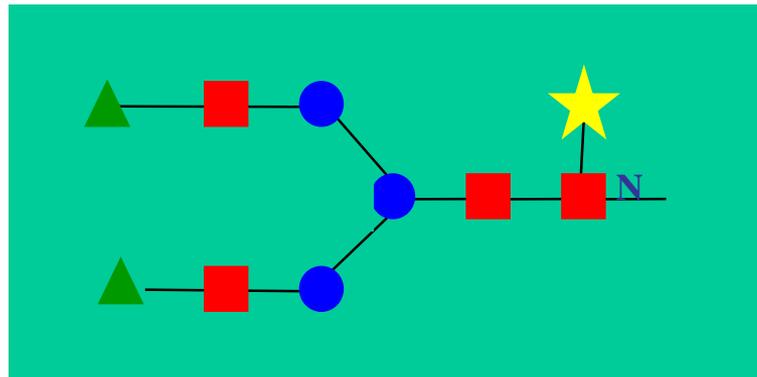


Mimura, Y., Jefferis, R. et al. *J.Biol.Chem.* 276:45539-45547 (2001).

Fully galactosylated glycoforms: G2

exhibit enhanced:

C1 complement activation; FcRn placental passage



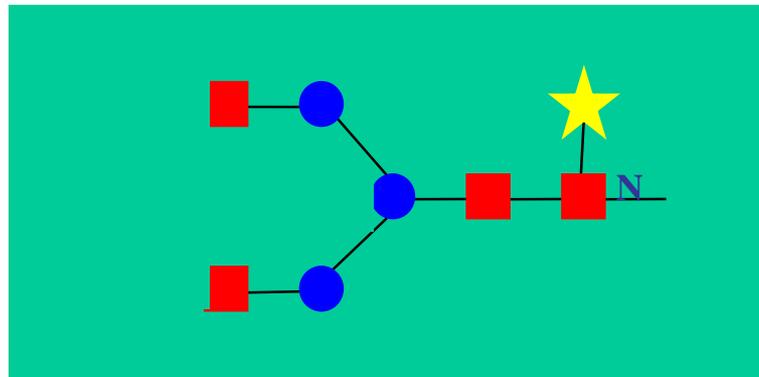
Jefferis, R. *Nature Reviews: Drug Discovery*. 8: 226-234 (2009)

Hodoniczky J, et al.. *Biotechnol Prog*. 21:1644-52 (2005)

Absence of galactose: G0

Terminal N-acetyl glucosamine residues may bind/activate pattern recognition receptors, e.g.

Mannan binding lectin (MBL); Mannose receptor (MR)



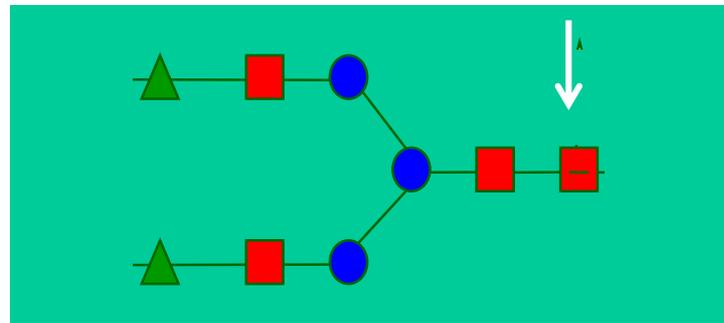
Jefferis, R. *Nature Reviews: Drug Discovery*. 8: 226-234 (2009)

Hodoniczky J, et al.. *Biotechnol Prog*. 21:1644-52 (2005)

Non fucosylated IgG has enhanced NK cell (FcγRIII) mediated ADCC

α 1,6 fucosyltransferase “knockout” CHO cells

Potelligent technology



Lonza

Mogamulizumab anti-CCR4
Anti Ebola virus h-13F



Igeneon AG

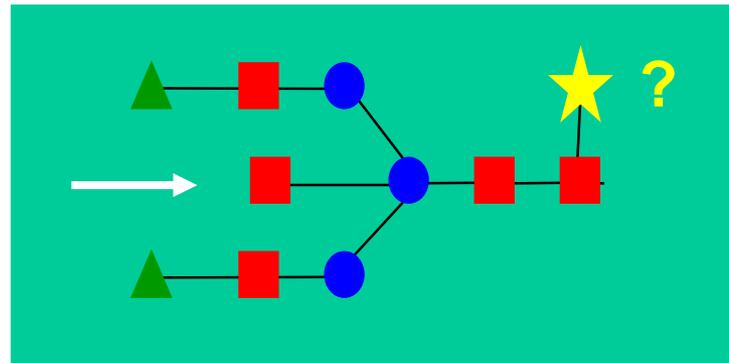
ProBioGen
Supporting Biopharmaceutical Visions

Kanazawa T. et al. Clin Cancer Res. 2014 Aug 12

The presence of bisecting N-acetylglucosamine enhances NK cell mediated ADCC (x 100)

“Knock-in” of GNTIII transferase in CHO cells

GLYCART
biotechnology



GlycoMAb

Obinutuzumab anti-CD20 (Gazyva); GA201 anti-EGFR

<http://www.fda.gov/Drugs/InformationOnDrugs/ApprovedDrugs/ucm373263.htm>

Gerdes CA. & Umana P. Clin Cancer Res. 20:1055 (2014)

Glycoprotein production platforms:

Mammalian: CHO, Sp2/0; NSO; Per.C6 etc



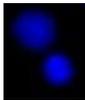
Transgenics: goat; sheep; cows; rabbits; pigs etc



Aves: chickens (eggs)



Yeasts: *Pichia pastoris*; *Saccharomyces cerevisiae*



Insect cells: Sf9 (baculovirus infected)



Plants: tobacco; corn; tomato; potato; moss



Bacteria: *Escherichia coli*; *Bacillus subtilis*

????????????????????????????????

The immune system “orchestrates” the antibody isotype response to be optimal for resolution of an “insult”, i.e. infection by a pathogen.

Can the immune system “orchestrate” the glycoform profile of an IgG response to be optimal for resolution of and insult?

Antibody therapeutics: Glycoform structures & function

Antibody classes & isotypes

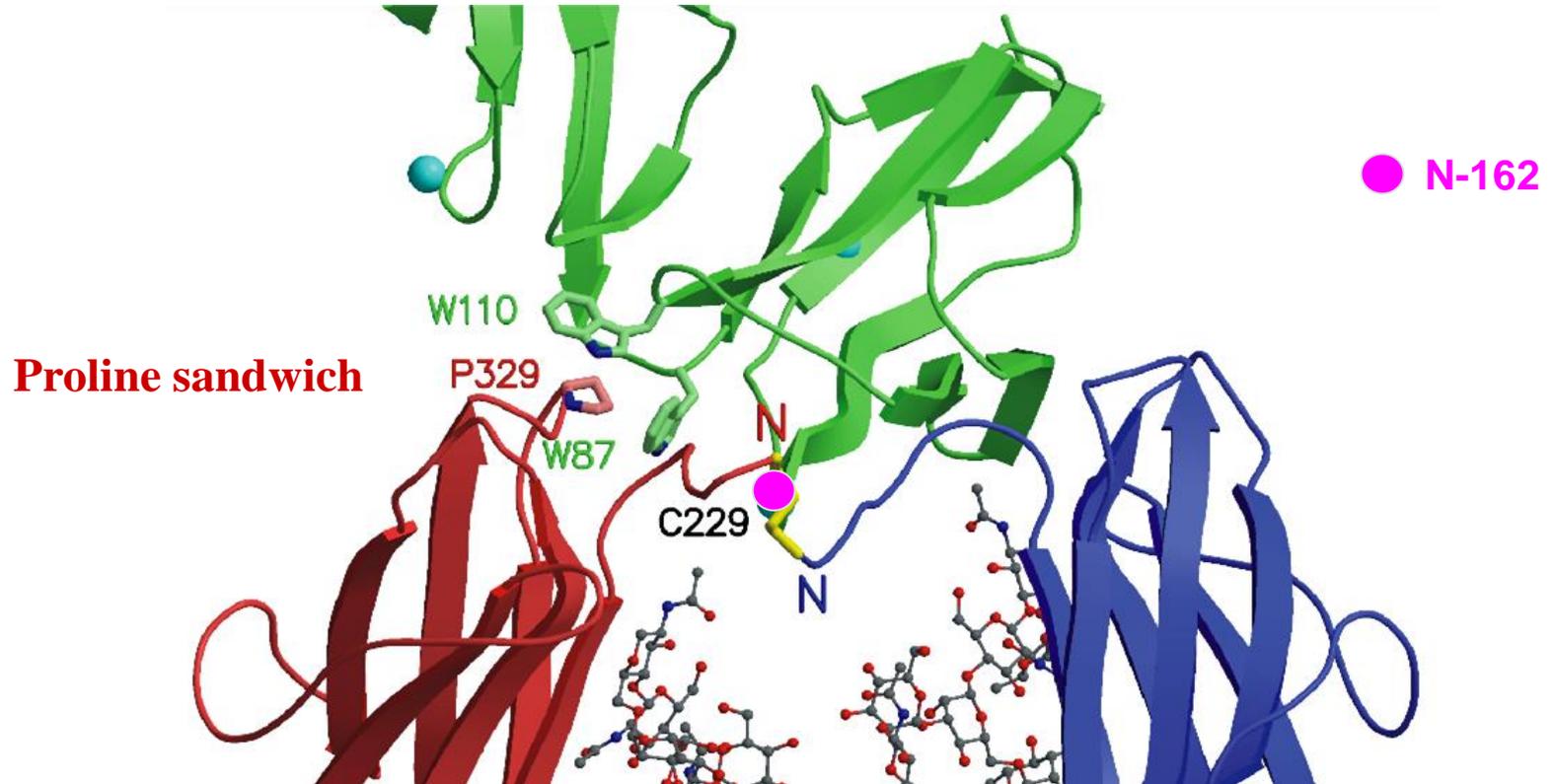
Antibody effector activities

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Ligand binding sites

Engineering “biobetters”

IgG-Fc/FcγRIIIa complex

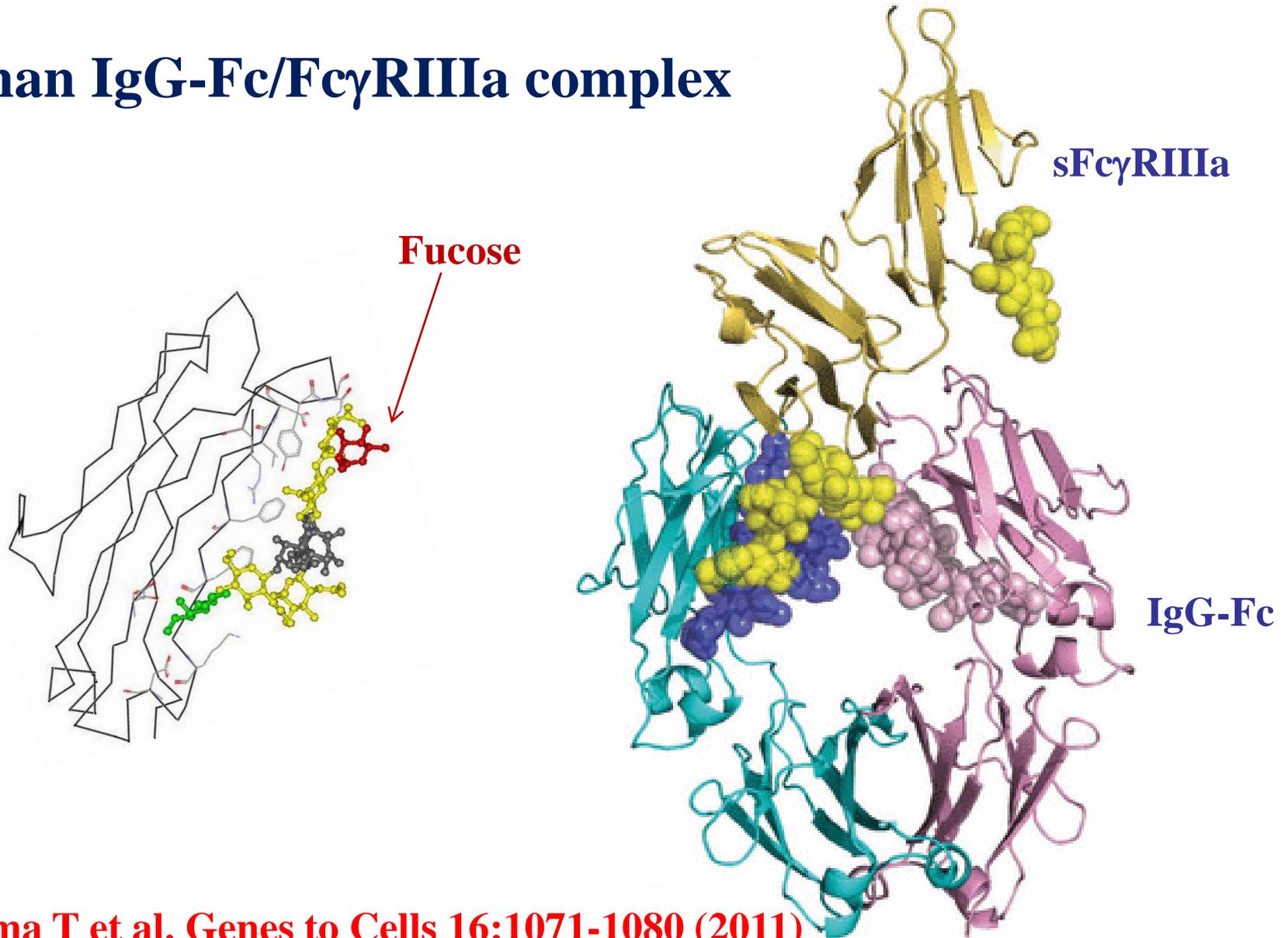


Drescher B. et al., Immunol 110: 335-40 (2003)

Sondermann P et al. Nature 406:267-273 (2000)

Radaev S et al. Mol Immunol. 38:1073-1083 (2002)

Human IgG-Fc/FcγRIIIa complex



Mizushima T et al. Genes to Cells 16:1071-1080 (2011)

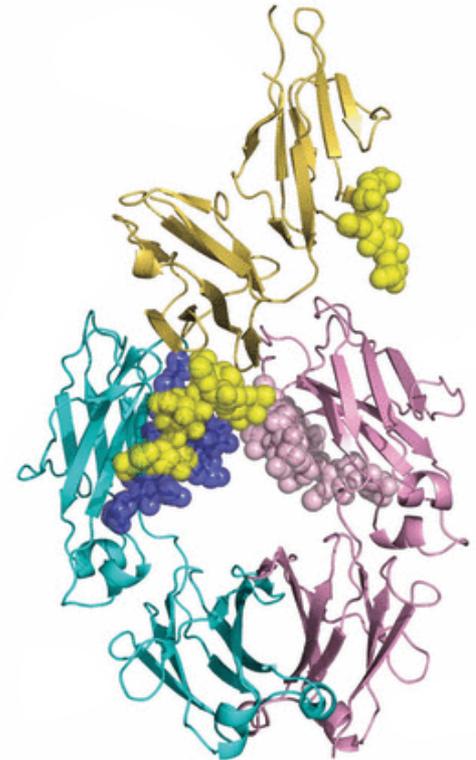
Ferrara C. et al. Proc Natl Acad Sci U S A. 108:12669-12674 (2011)

Features of IgG-Fc/Fc γ RIIIa interactions:

Both heavy chains are involved in the formation of an asymmetric binding site

Affinity of binding is dependent on glycoform profile of the Fc

Affinity of binding is dependent on glycoform profile of the Fc γ RIII



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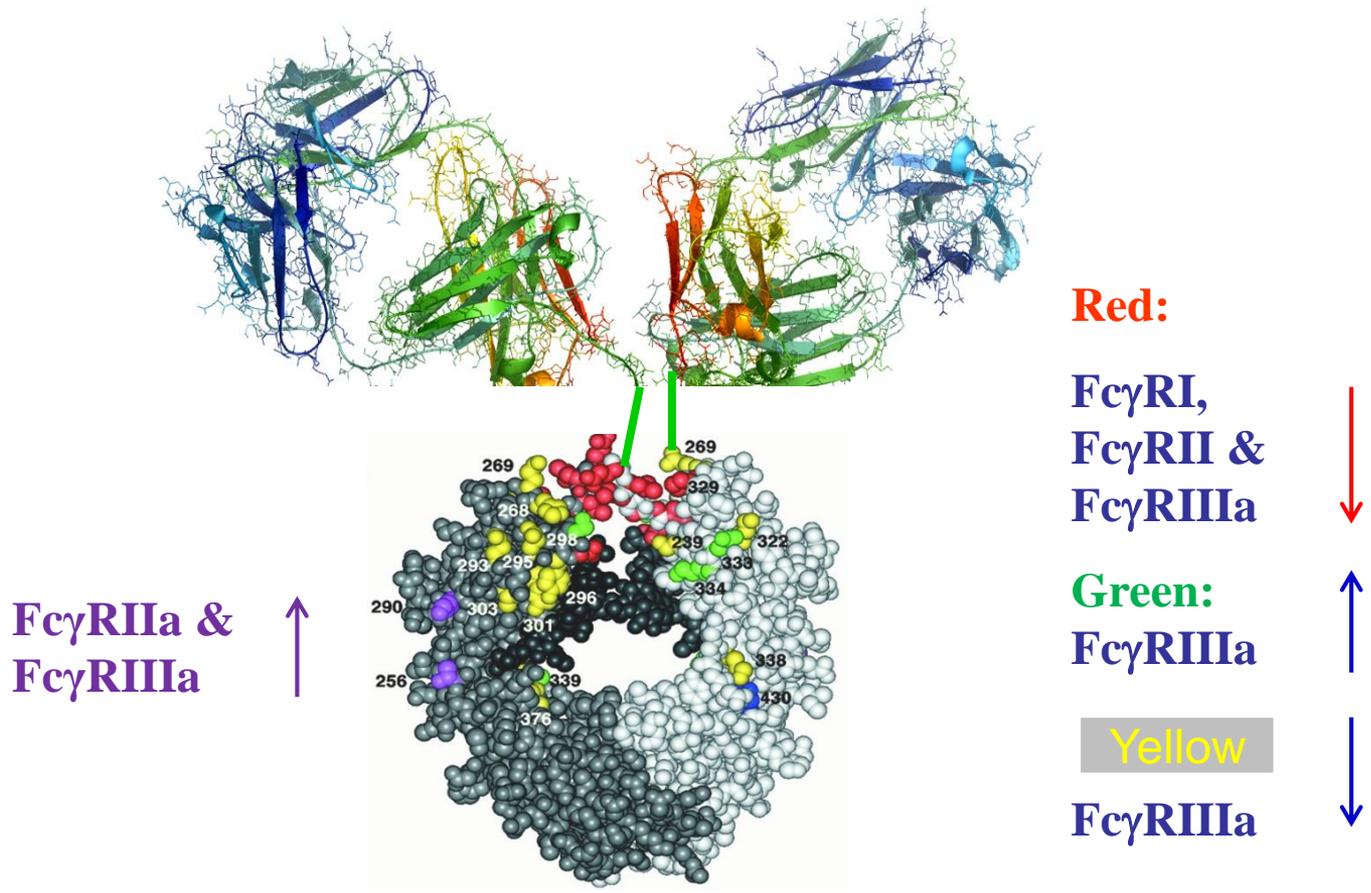
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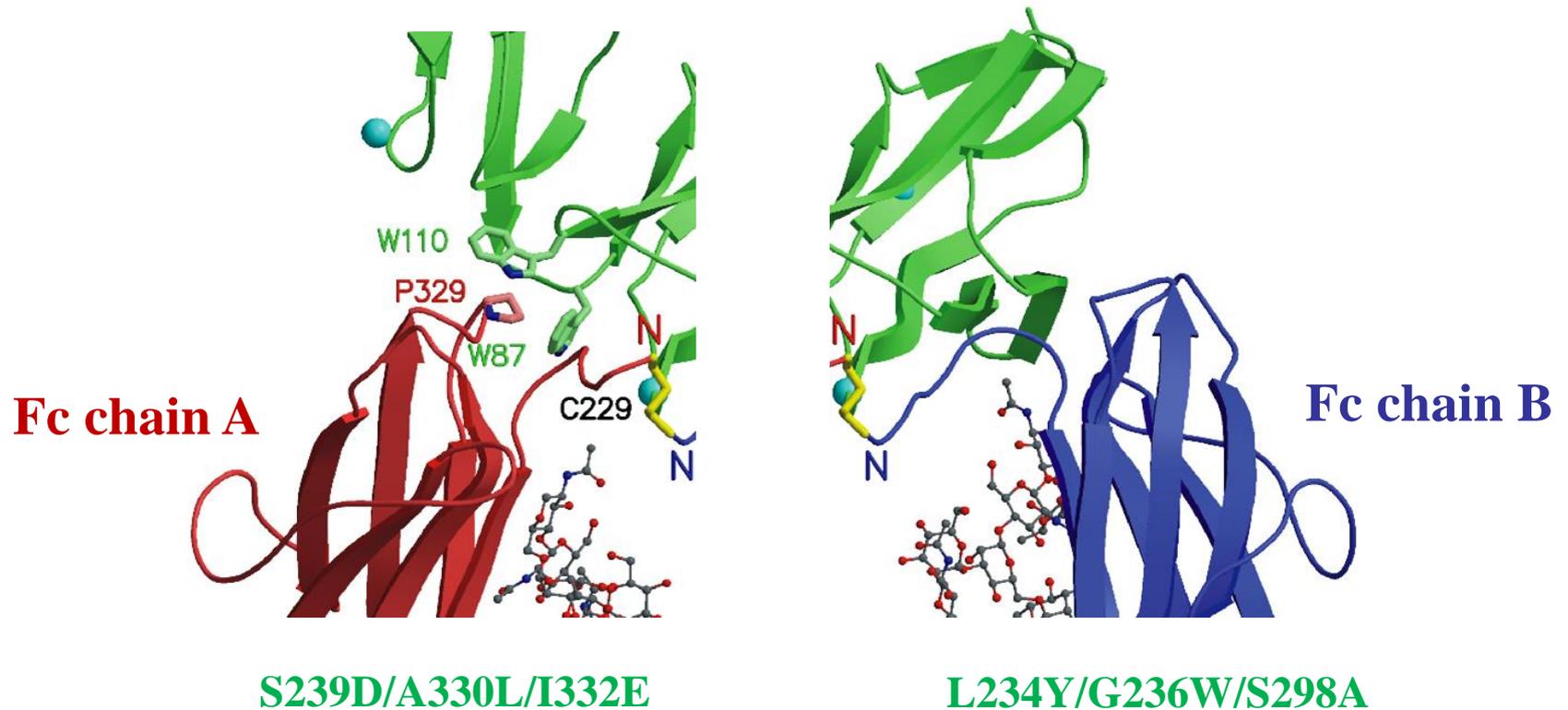
Engineering “biobetters”

Alanine “scanning” identifies residues impacting FcγR & FcRn binding



Lund J. et al. (1991). J.Immunol. 147 2657 - 2662 (1991)
Shields RL. et al., JBC 276:6591-6604 (2001)

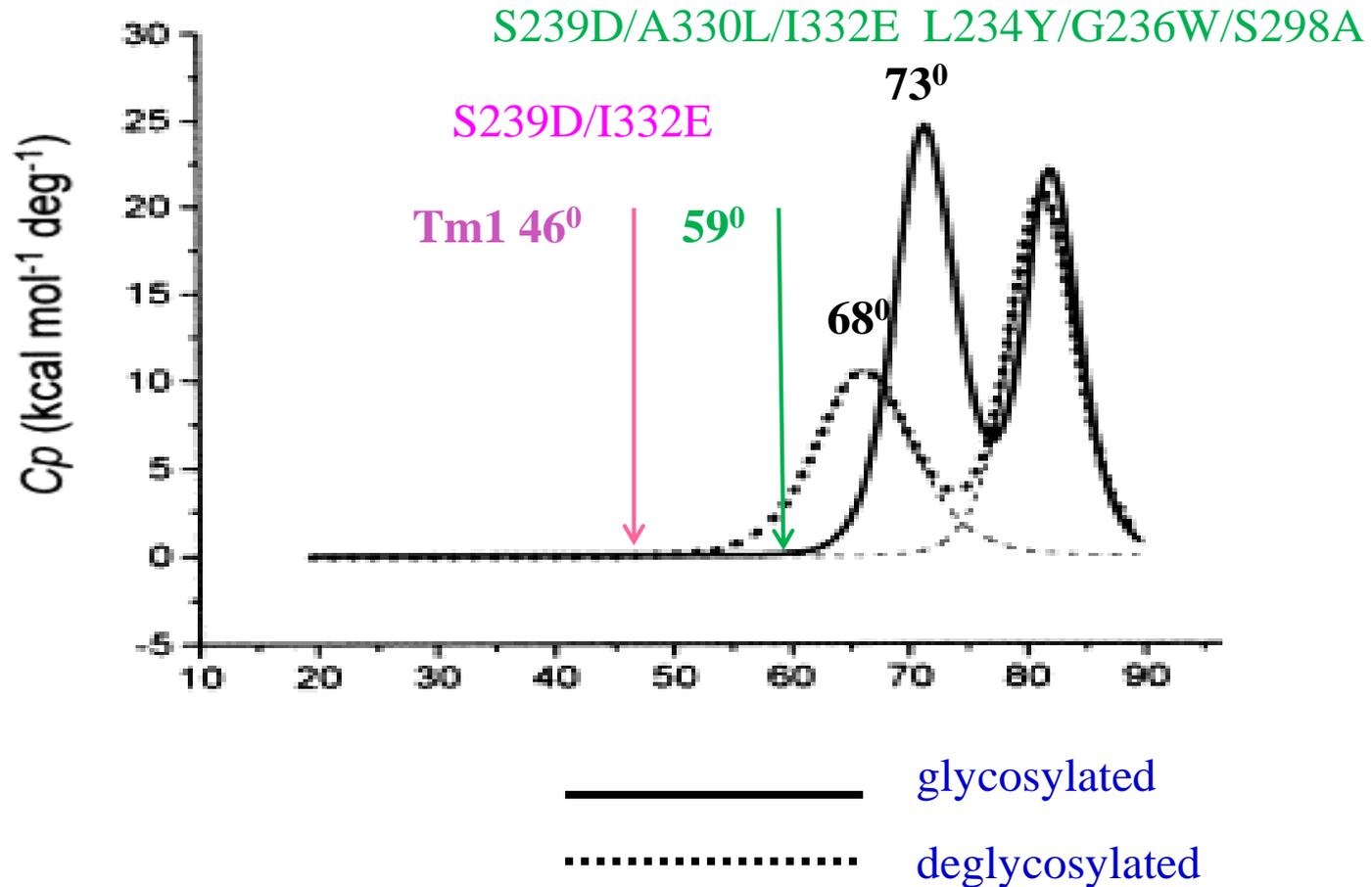
Generation of asymmetrically engineered IgG1-Fc



Mimoto F. et al. MAbs 5:229–236 (2013)

Sondermann P et al. Nature 406:267-273 (2000)

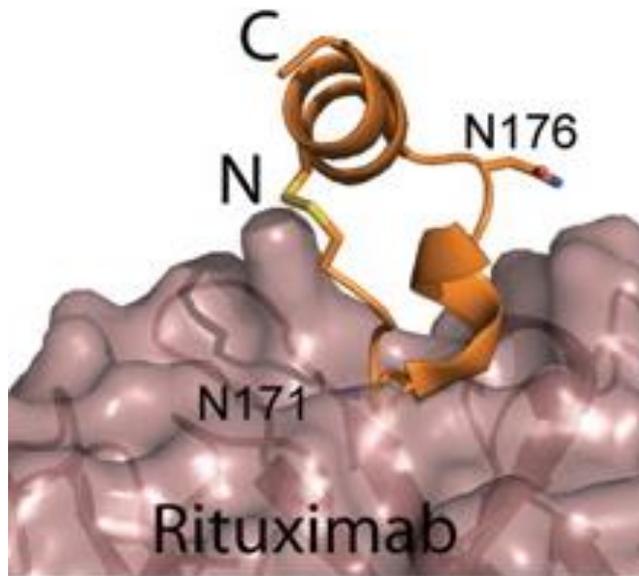
DSC of IgG1 Fc proteins



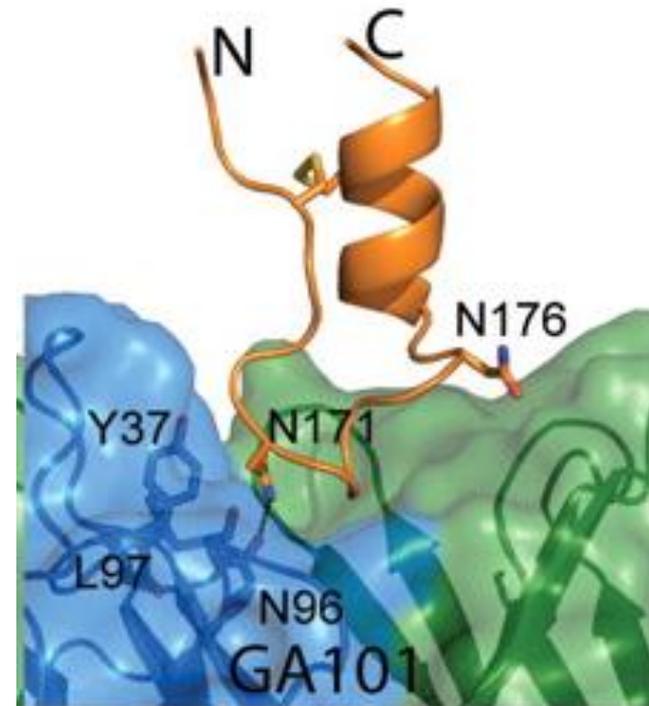
Mimoto F. et al. MAbs 5:229–236 (2013)

Mimura Y., et al., Molec Immunol 37:697 – 706 (2000)

Paratope/epitope orientations for rituximab (Type I) & obinutuzumab (Type II) anti-CD20 antibodies



Rituximab

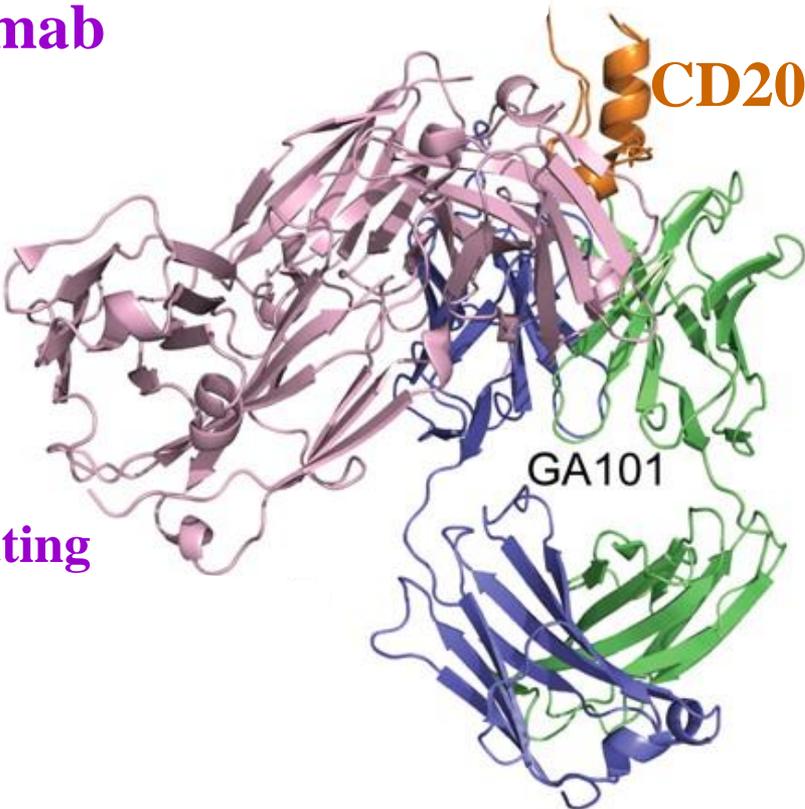


Obinutuzumab
(GA101)

Anti-CD20 Type I (Rituximab) & Type II (GA101) mAbs

Rituximab

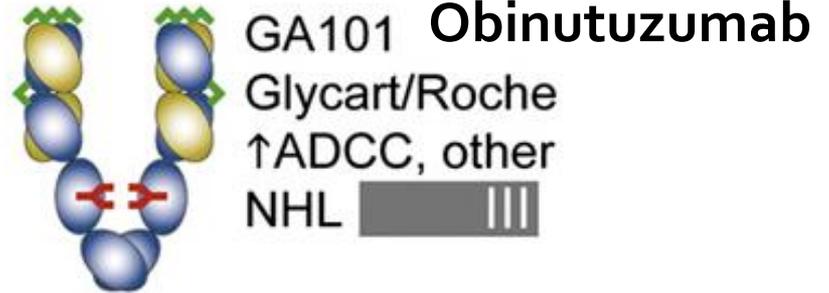
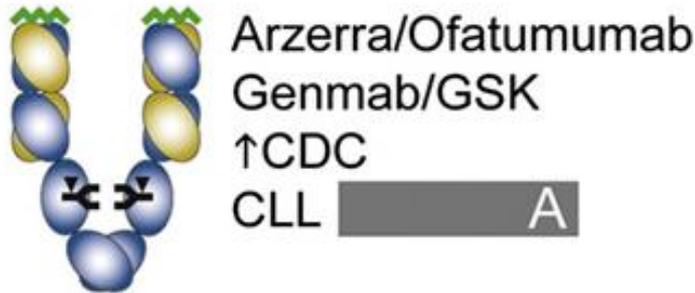
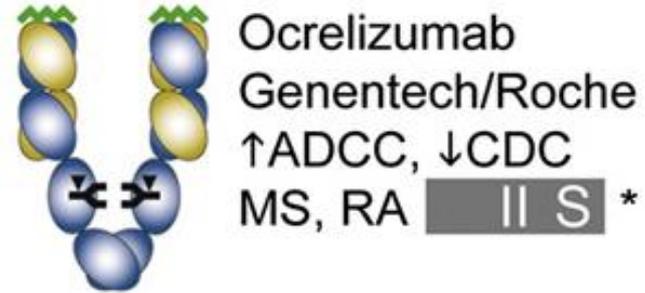
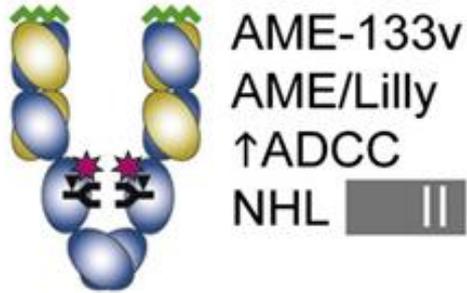
**Obinutuzumab
(GA101)**



Complement activating

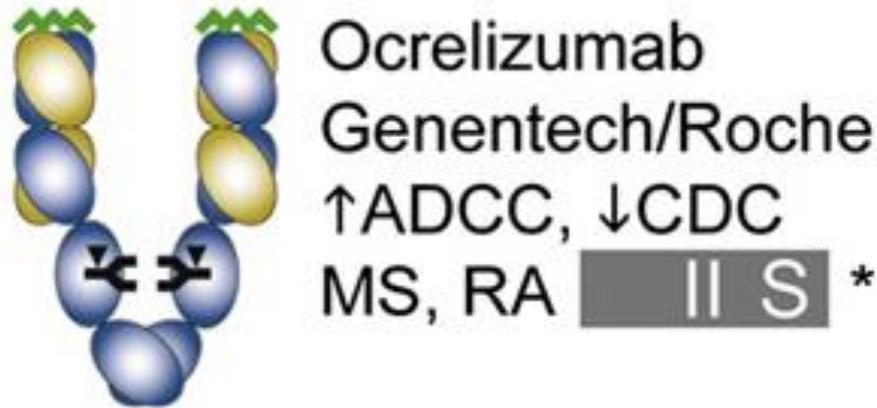
Direct cell killing

Next generation anti-CD20 antibodies



Next generation anti-CD20 antibodies: Adverse effects

Risk/Benefit



In March 2010, Roche announced the suspension of clinical trials in rheumatoid arthritis and lupus erythematosus. This step followed excess deaths due to opportunistic infection

Phase III development for multiple sclerosis continues

Reconstitution of antibody therapeutics for administration

Instructions for Herceptin® states:

- 1) the stream of the diluent should be directed into the lyophilized cake
- 2) the vial should be swirled gently, without shaking
- 3) the vial should stand for five minutes undisturbed
- 4) drawing into a syringe should be performed slowly.

Peters BJM et al. mAbs 5:162–170 (2013)

Arvinte T. et al. MAbs. 5:491-500 (2013)



Davies AM, Jefferis, Sutton BJ. *Structural determinants of unique properties of human IgG4-Fc. J Mol Biol.* 426:630-644 (2014)



Davies AM., Jefferis R., Sutton BJ. *Crystal structure of deglycosylated IgG4-Fc .Mol Immunol.* 62:46-53 (2014)



Jefferis R. *Monoclonal Antibodies: Mechanisms of action. In: Current state of the art for the characterisation of mAbs. Eds: D. Davis, J. Schiel & Borisov O. ACS 2014*



Jefferis R. *Isotype & glycoform selection for antibody therapeutics. Arch Biochem Biophys* 526:159-166 (2012)