Engineering of Antibodies, a Knowledge Based Approach

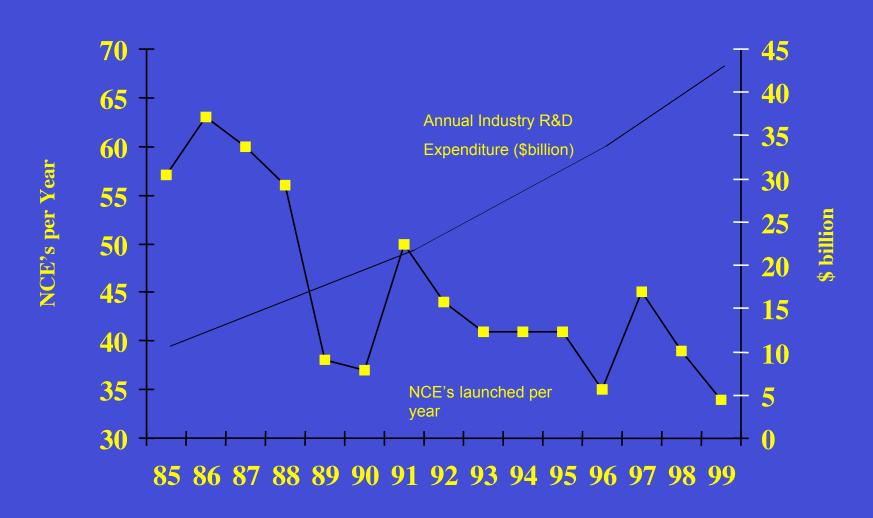
Nottingham: 21st March 2007 dee@moksha8.com

Introduction

- Introduction
- Antibody structure
- Engineering Antibodies: Humanisation
- Expression
 - Mammalian
 - E.coli
- Clinical Utility
- Alternatives

Commercial value of Antibodies

Discovery Spending vs Output



Licensed mAb Products

<u>Product</u>	Indication Year Lie	cenced
Reopro	Coronary artery disease	1994
Synagis	Prevent RSV infection	1998
Rituxan	Lymphoma therapy	1997
Herceptin	Breast Cancer	1998
Zenapax	Kidney Transplantation	1997
Remicade	Arthritis/Crohn's disease	1998
Orthoclone	Organ Transplantation	1986
Simulect	Kidney Transplantation	1988
Mylotarg	AML	2000

Total Global 1999 Sales ≈ \$1.5 Billion

Value of Antibody Technology

- Four marketed humanised antibodies had combined total sales in excess of \$700M
 - Mylotarg
 - Herceptin
 - Synagis
 - Zenapax
- Total global sales in 1999 of therapeutic Mab products ≈ \$1.5 Billion

Avastin had approx sales of \$1.1Billion in 2005

Value of Antibody Technology..contd

- one of the fastest growing and most lucrative sectors of the pharmaceutical industry, with exceptional 48.1% growth between 2003 and 2004.
- In 2005, the antibodies market was valued at an estimated US\$14 billion, accounting for over 24 percent of the total protein therapeutics market.
- Potential to triple in value over the next few years and reach \$30.3 billion in 2010
- A wave of fully human products are expected to launch from 2007 onwards, accounting for 12 of the 20 launches between 2007 and 2010.

Immunogenicity of Therapeutic Antibodies versus other Biologics

- For naked, humanised antibodies or fragments, there is a high probability that the response will be directed towards the idiotype
- Important issues are
 - Hypersensitivity reactions (safety)
 - However it is unlikely that anti-idiotype antibodies will cross-react with native proteins
 - Blocking/ neutralising/ clearing of drug (efficacy)
 - Anti-idiotype antibodies will have the potential to neutralise if they are sufficiently high titre or affinity

Immunogenicity of Therapeutic Antibodies – conjugates or chimeric Mabs

- Attachments have the potential to be immunogenic – even weak immunogens may provoke a response when attached to a large protein
- Linkers, neo-epitopes at joining regions, chimeric V-region framework sequences all have potential to be immunogenic

Factors affecting Antibody Immunogenicity

- Molecule
 - Species homology (rat, monkey, man)
 - Expression of antigen
 - Location of antigen e.g. soluble vs surface of immune system cell;
 Affinity of binding;
 - Attachments (e.g. cytotoxic drug, PEG)
- Manufacturing
 - Glycosylation
 - Aggregates
- Administration
 - Route of administration (SC vs IV)
 - Size of dose; Number and frequency of doses
- Recipient
 - Patient population
 - Concomitant medication

Technologies which have been developed to reduce Immunogenicity

- **Biovation:**For DeImmunisation[™] of antibodies and proteins, helper T cell epitopes are identified within the primary sequence of the biopharmaceutical and these sequences are modified, principally by amino acid substitution, to avoid recognition by T cells.
- **AlgoNomicsl:**EpibaseTM is the proprietary platform for T-cell epitope identification.
- Genencor: The i-mune Assay (Epitope Mapping Technology)
- **PEG:** Roche PEGASYS[®] (IFN -α2a)

Human/Humanised Antibodies

- 1st Generation Chimeric Antibodies
- 2nd Generation CDR Grafting
- Fully Human Antibodies:
 - Human Ab Phage Libraries
 - CAT
 - Hu Mice
 - Abgenix
 - Medarex
 - XTL
 - Hu Rabbits

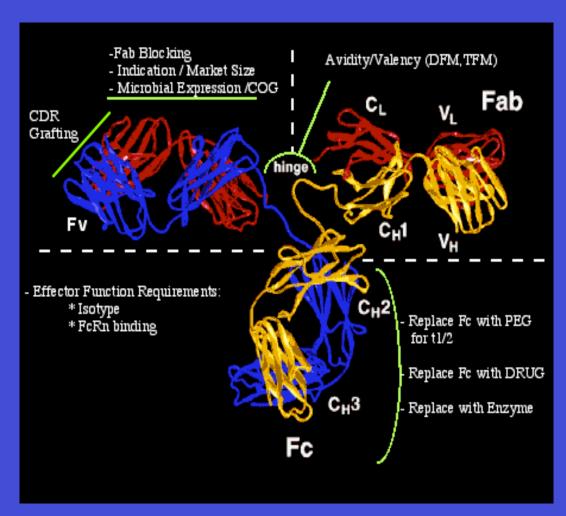
HUMIRA TM

- 1st approved "fully human" Mab (RA)
- Q Is HUMIRATM immunogenic?
- A Yes
 - Overall rate 5.5% in pivotal trials
 - Included 1gG2, IgG3, IgG4 & IgM (No igE)
 - Primarily anti Id's
 - Modest reduction in efficacy in patients on monotherapy
 - No effect on safety

Abbott Lab's presented above data in Sept 2003 (IBC)

SAR of an Antibody

- Properties to be considered:
 - Half Life (t 1/2)
 - Number of binding sites (Avidity / Valency)
 - Immunogenicity
 - Blocking antibody ?
 - Internalising (TFM > DFM))
 - Deliver drug / effector functions / enzyme





Antibodies

• Estimated that the humans can produce as many as 10 million different antibodies in the primary repertoire and this can be further expanded by several orders of magnitude by somatic mutation

• This diversity and specificity of antibodies can be understood by examination of antibody structure

• IgG Structure

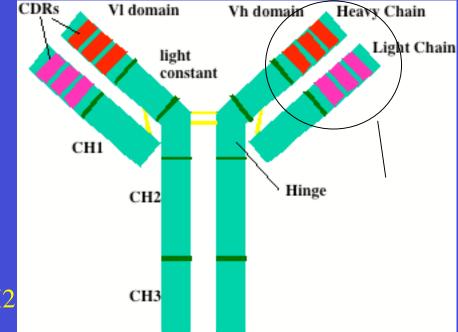
- Consists of 4 polypeptide chains
 - two heavy chains, approx. 50kDa
 - two light chains, approx. 25kDa

• Each is divided into domains of approx. 110aa stabilised by intra-

disulphide bonds.

• Light chain composed of:

- variable domain (VL)
- constant domain (CL)
- Heavy chain composed of:
 - variable domain (Vh)
 - three constant domain (CH1,CH2

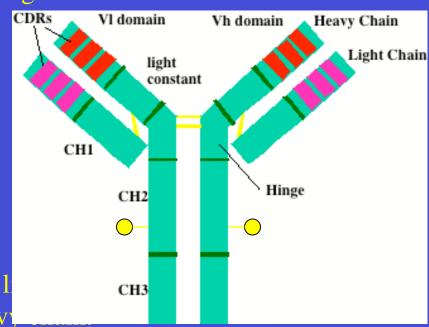


 Heavy and light chain variable domains associate to form the antigen binding site

• IgG Structure Contd.

• The IgG molecule has two antigen binding sites.

 The heavy and light chains are linked by a disulphide bond between the CL and CH1 domains.



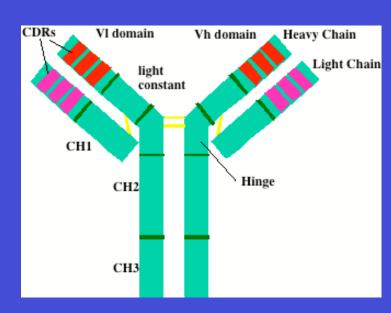
- A flexible region known as the "hinge" l
 the CH1 and CH2 domains of each heavy
- The two heavy chains are linked together at the hinge by disulphide bonds.
- The CH2 domain is normally glycosylated.

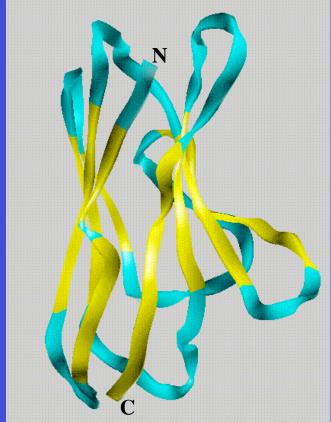
Antibody Architecture

• Variety of immunoglobulin fragments from both human and mouse have been characterised by X-ray Crystallography

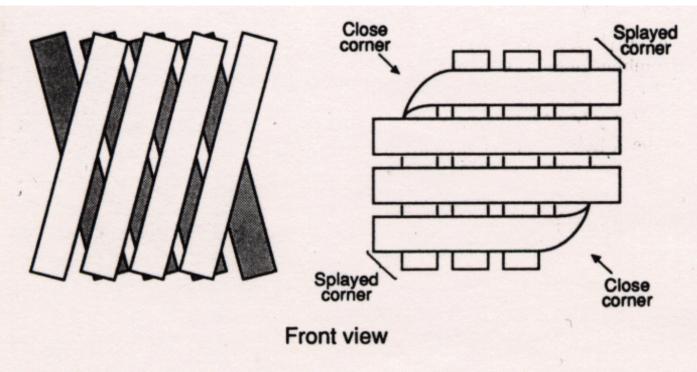
• The individual domains have similar conformation,

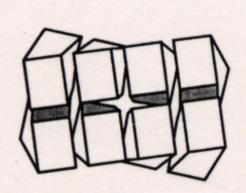
"Immunoglobulin Fold"

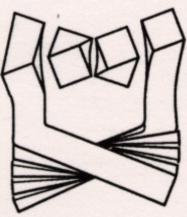




• The Ig Fold consists of 2 layers of anti parallel beta sheet





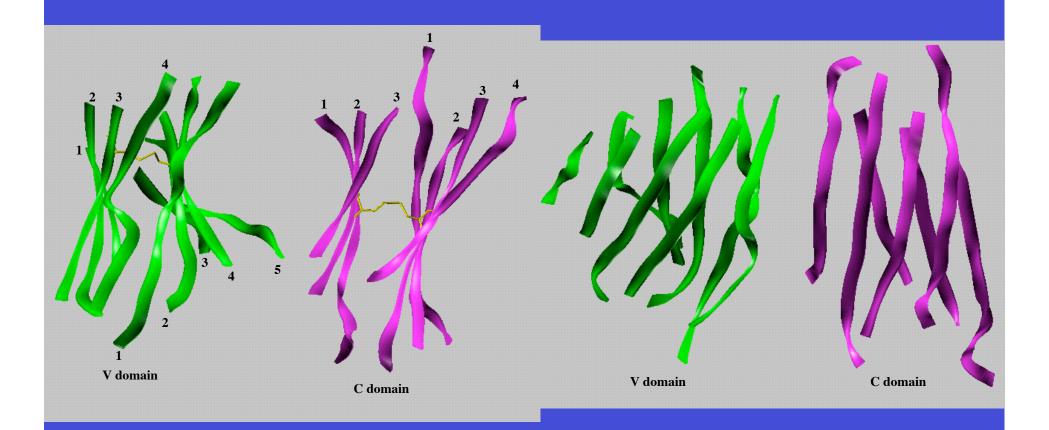


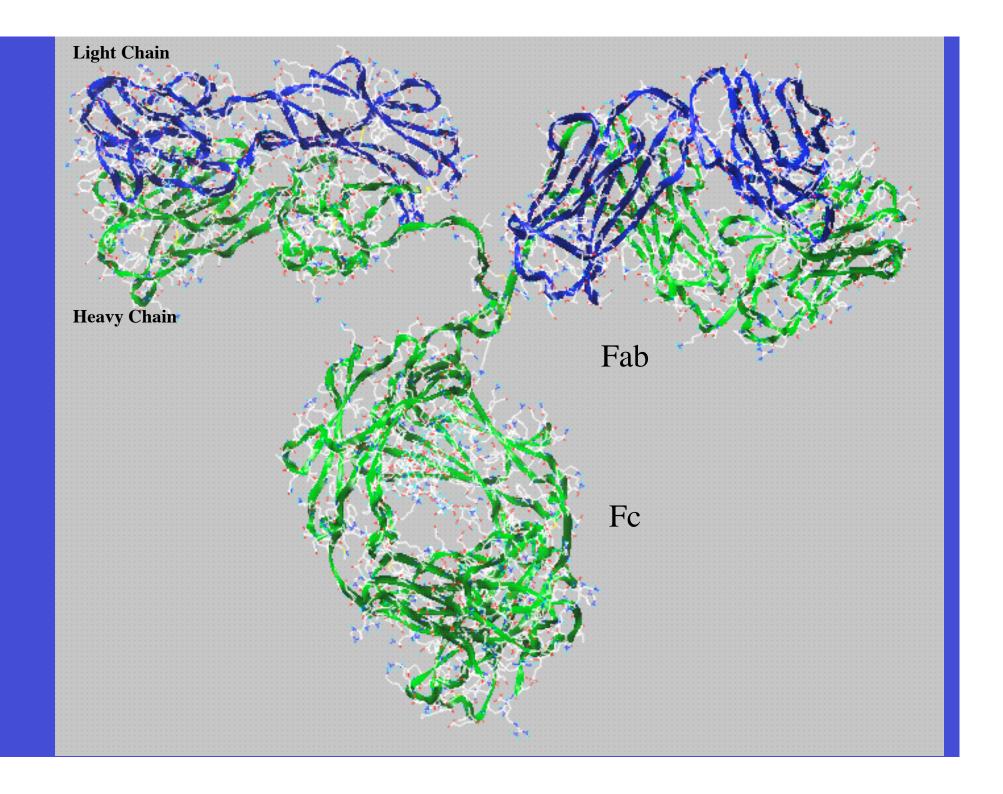
End view

(a) Aligned packing (b) Orthogonal packing

Ig domain structure

- The N-terminal variable domains differs from that of the C-proximal constant domains:
 - V domains β sheets consist of 5 & 4 strands
 - C domains β sheets consist of 4 & 3 strands





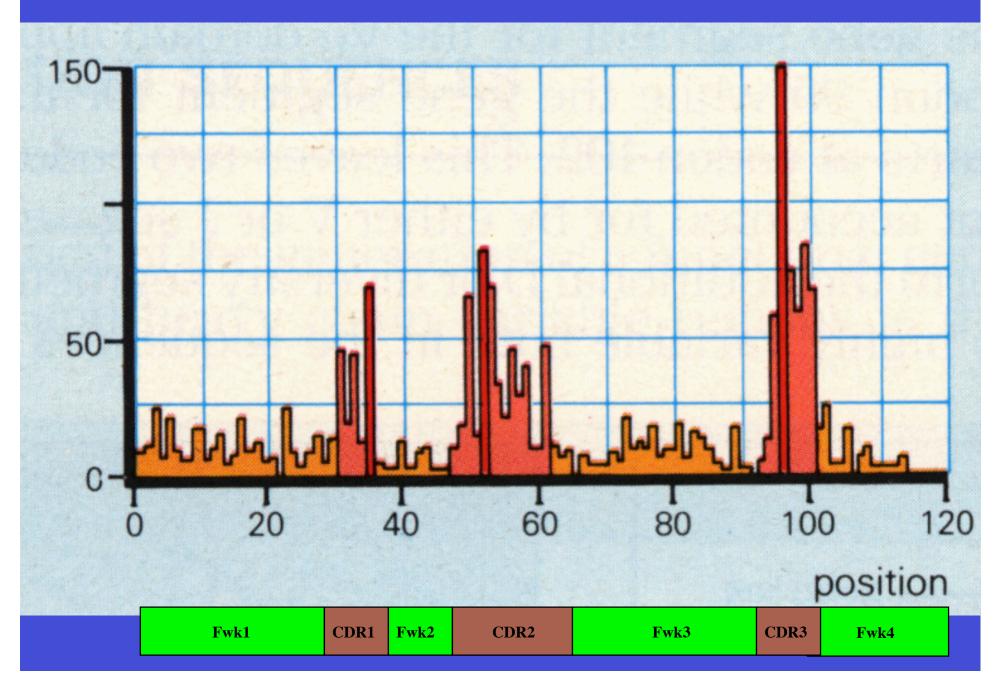
Bio-informatics

• The primary sequence of several thousand antibody molecules are known (largest number of known sequences in one protein family).

• Analysis of these sequences revealed that the variability of the variable domains is largely restricted to three "hypervariable" regions, in each of the light and heavy variable domains.

 Comparatively little variation in the intervening "framework" regions"

Kabat Plot

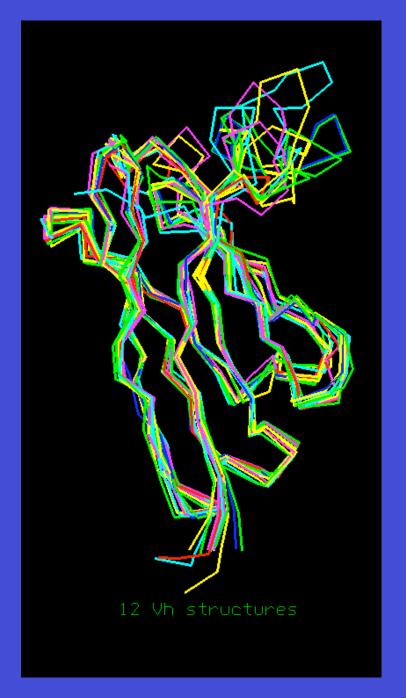




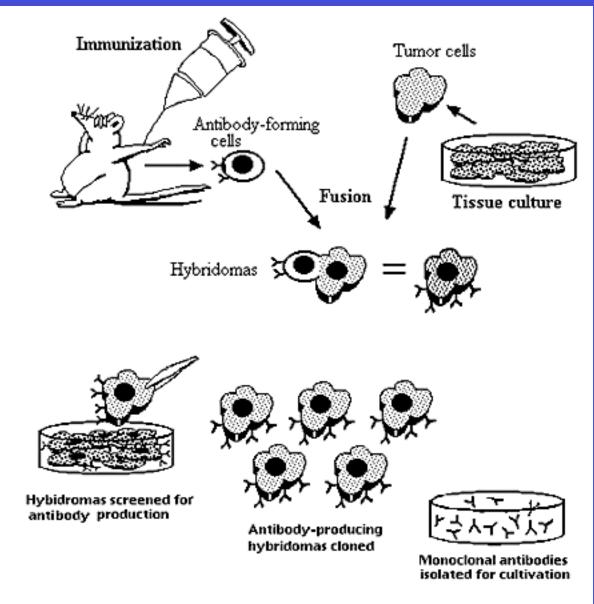
CDRs

• The combining site of antibodies is almost entirely made up of 6 loops, 3 each from the light and heavy variable domains.

- Variability in these loops arises from:
 - sequence
 - Length
- Loops are commonly referred to as
 - Complementarity determining regions (CDRs)
 - Hypervariable loops

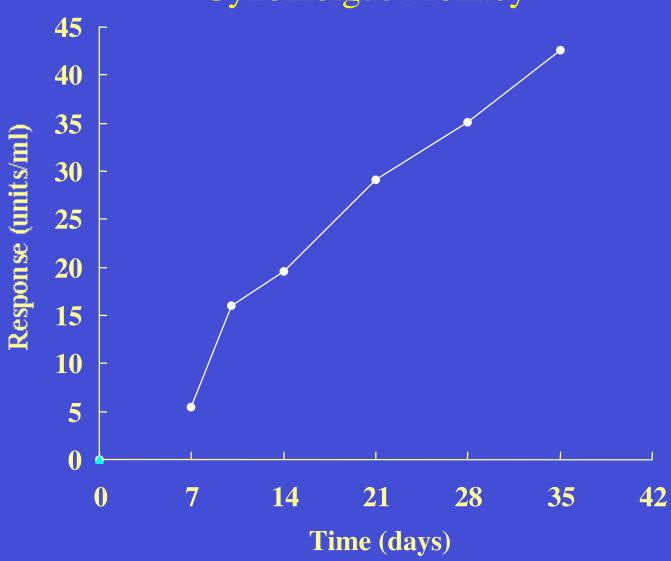






Monoclonal Antibody Production

Antibodies to Murine anti TNF-antibody in Cynomolgus Monkey



Potential Disadvantages of a Human Anti - Mouse Antibody (HAMA) Response

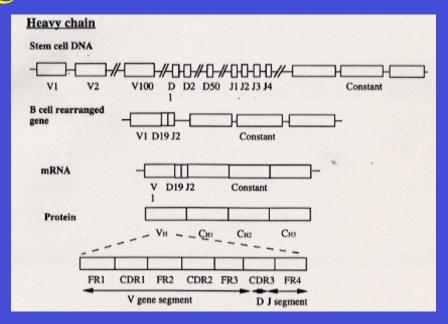
- Acute or Delayed Hypersensitivity Reactions
- Pharmacokinetic Changes
 - » High MW Aggregates
 - » Increased rates of clearance
- Reduced Targeting Efficiency / Efficacy

Strategies to Avoid HAMA

- Co- therapy with immunosuppresent
- Sequential IgG type therapy regime
- Co- administration with anti-CD4 antibody
- Use of Antibody fragments
- Use of human or primate monoclonals
- Use of recombinant chimeric or humanised antibodies

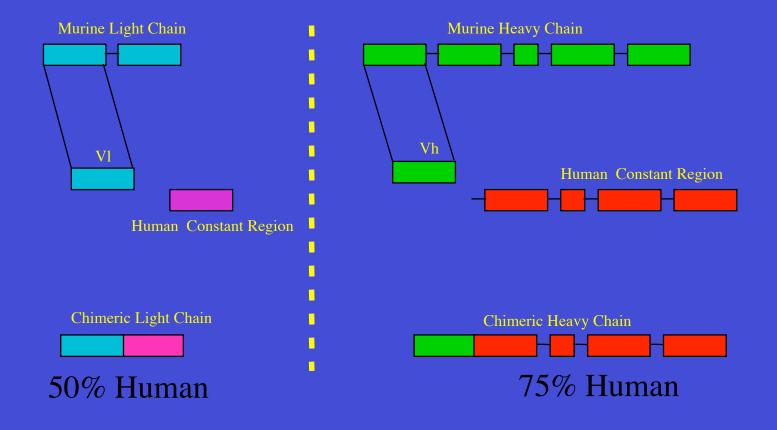
Cloning V regions

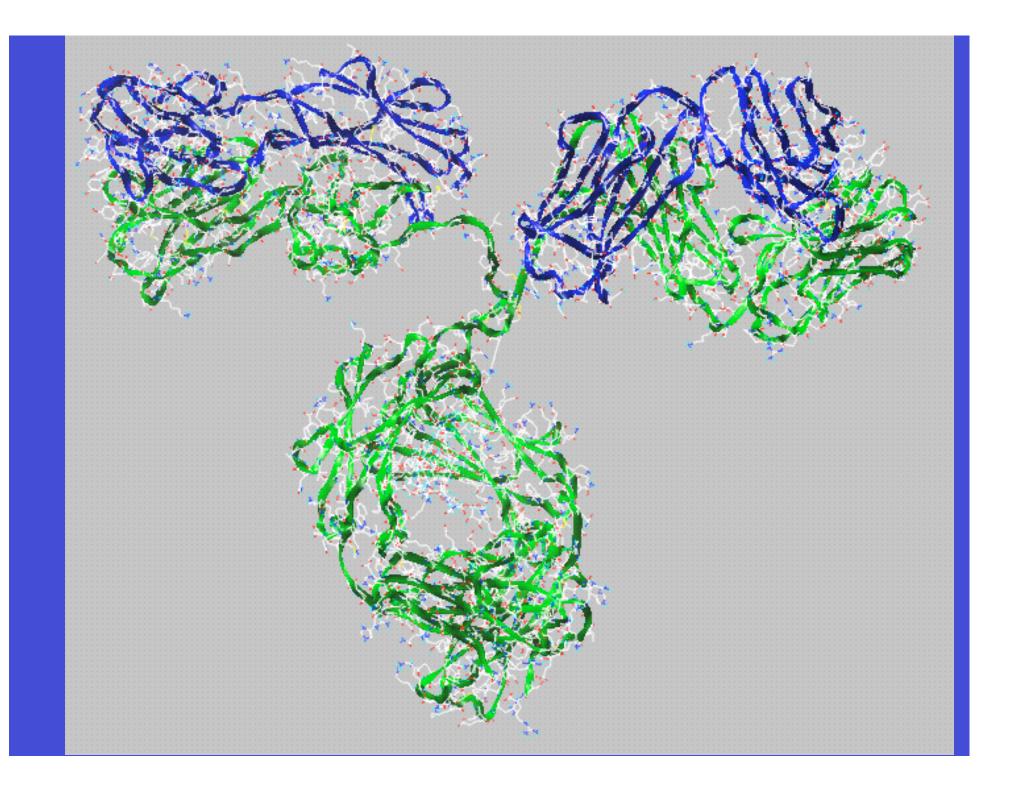
- Clone rodent variable domains using RT-PCR
- Primers designed based on leader sequence and constant regions

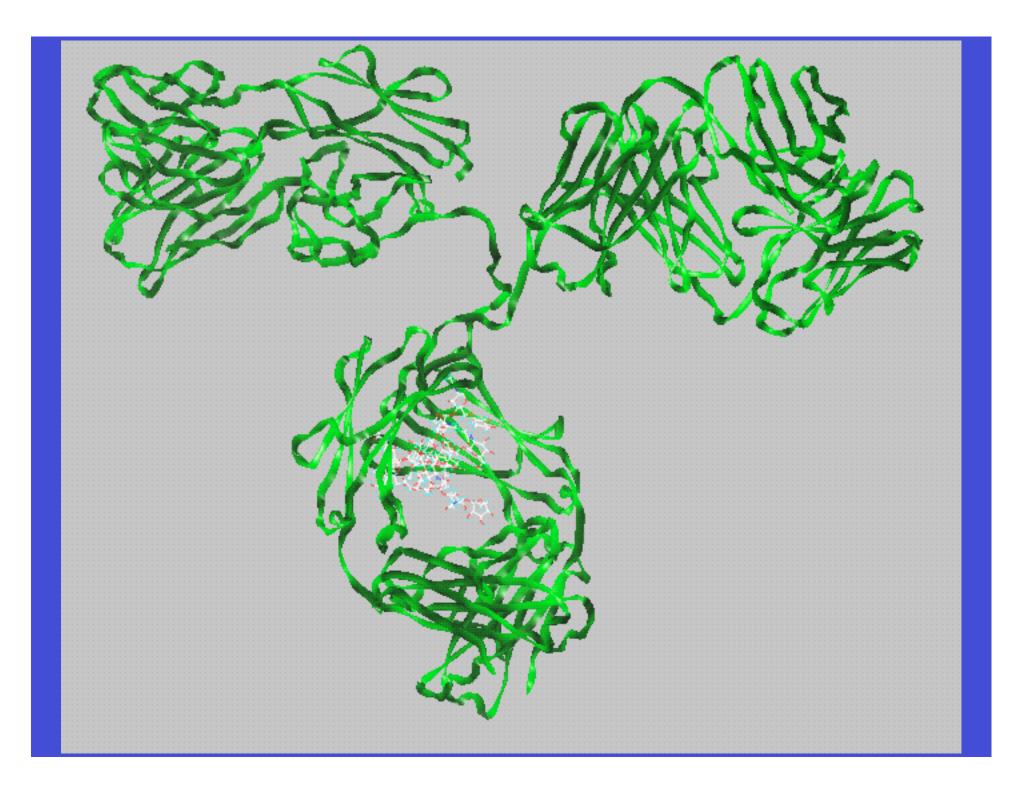


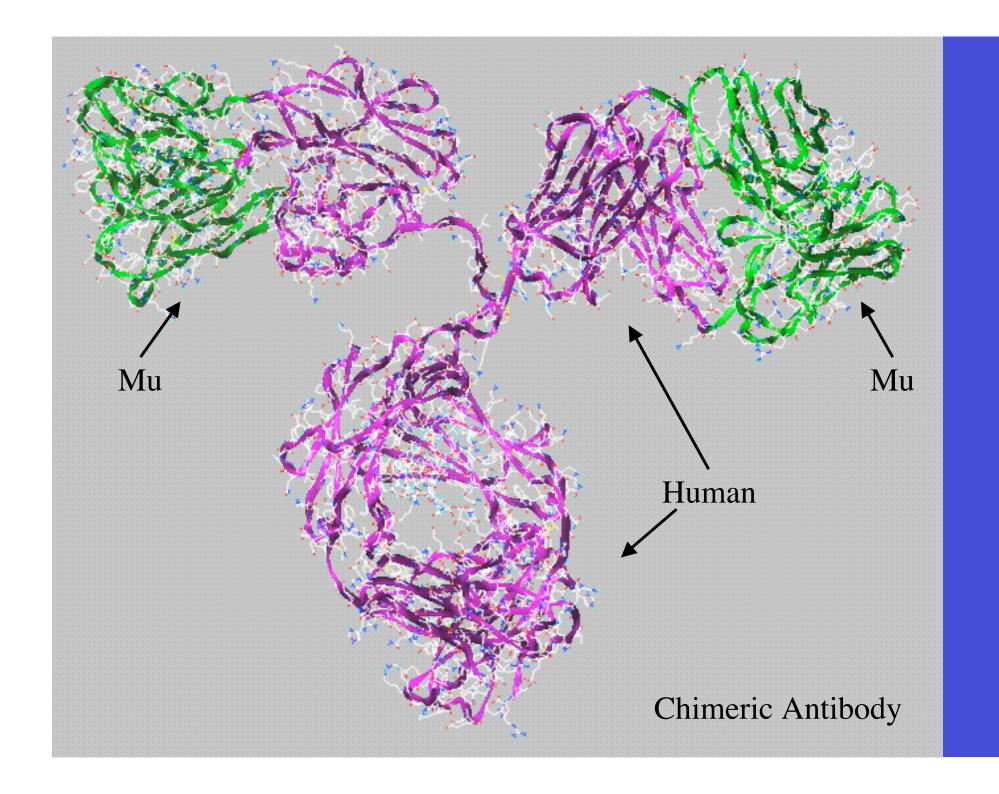
• Cloned directly into expression vectors containing human constant regions.

Chimeric Construction







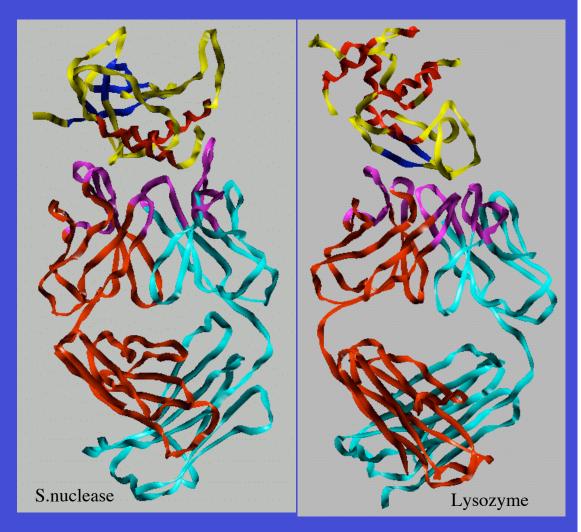


Antibody-antigen structures

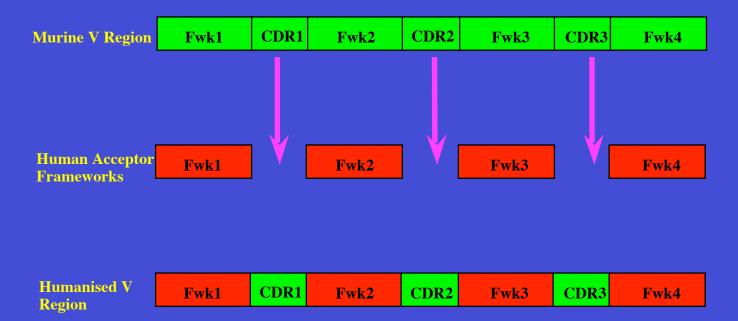
- Fab structures complexed with several antigens have been deposited in PDB:
 - Haptens
 - Phosphocholine
 - Vitamin K₁OH
 - Fluorescein
 - Proteins:
 - Lysozyme
 - Influenza virus neuraminidase
 - DNA
 - Carbohydrate

Humanisation by CDR grafting

- Antibody Antigen interactions occur at the tips of the Fab arms.
- CDR grafting transfers
 the binding specificity of
 the rodent antibody into
 a human antibody
- Transfer CDRs from mouse to human environment.

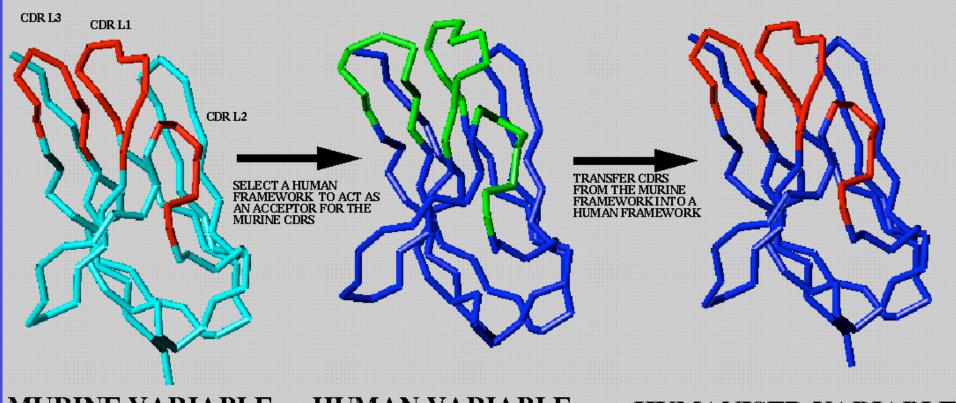


Humanisation of V Region



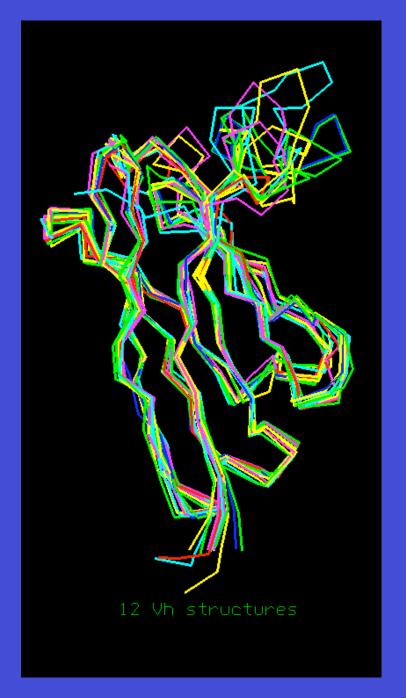
Kabat CDR grafting

PRINCIPLE OF ANTIBODY HUMANISATION



MURINE VARIABLE DOMAIN HUMAN VARIABLE DOMAIN

HUMANISED VARIABLE DOMAIN





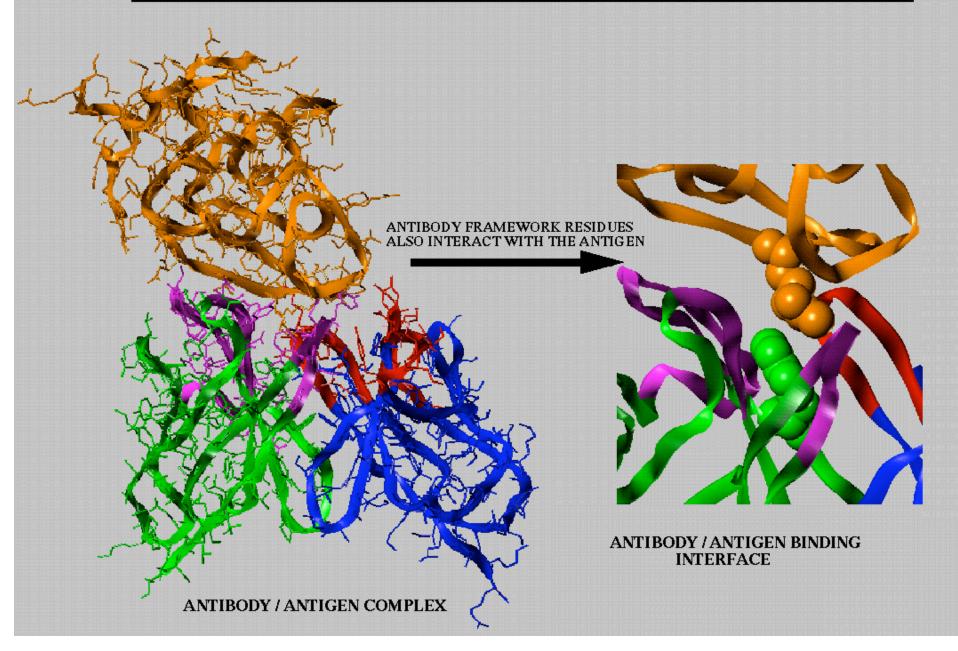
Initial Results of CDR grafting

• The affinities of Kabat grafts, transferring only the CDRs from rodent to human frameworks is generally poor.

 Most if not all of the original affinity is normally lost

WHY?

PRINCIPLE OF ANTIBODY HUMANISATION



Framework Residues

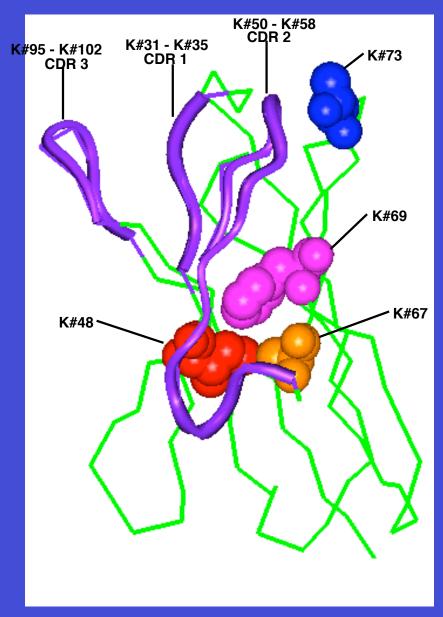
• Selection of framework residues is acheived by visual inspection of antibody models:

 Exposed and spatially adjacent residues to CDRs

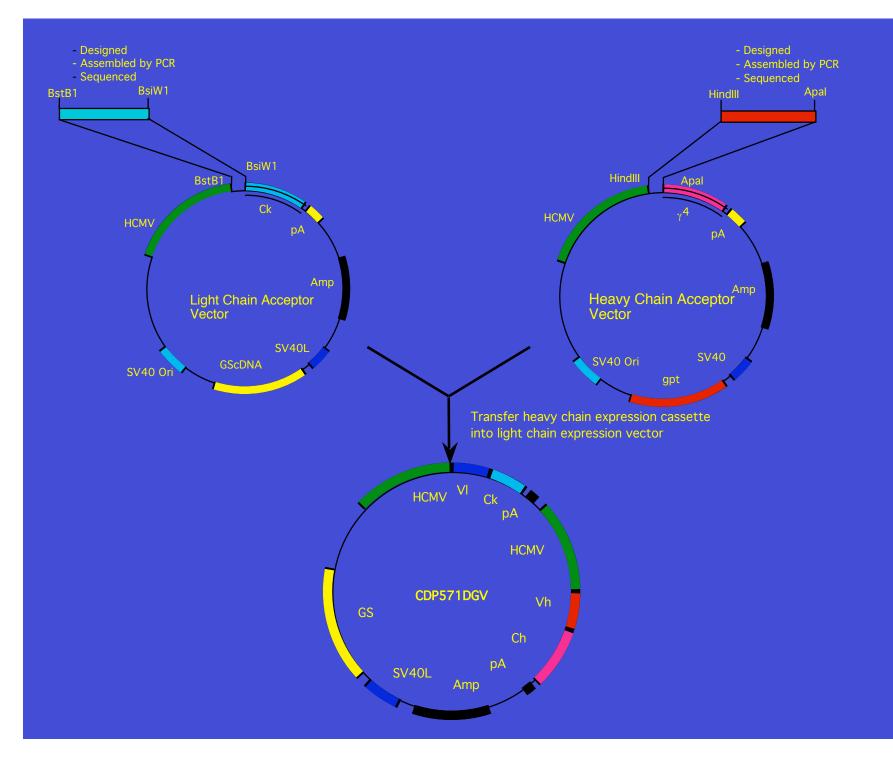
Buried residues which form Ig fold core

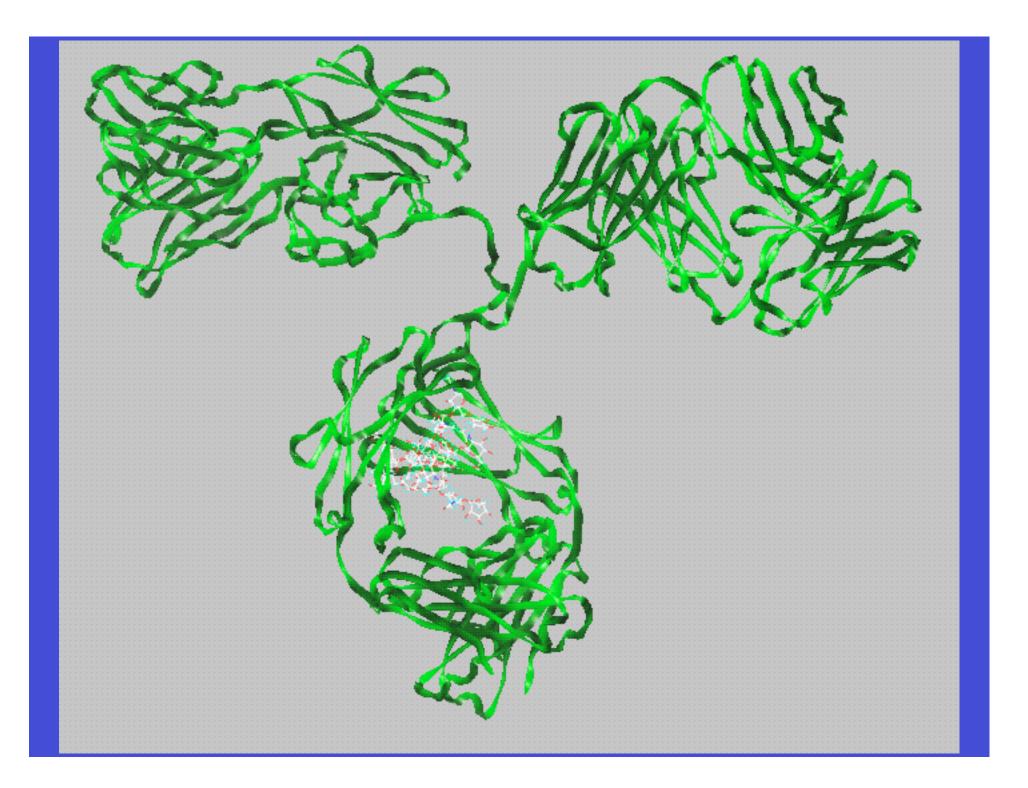
 Buried residues which are capable of forming contacts with the CDRs

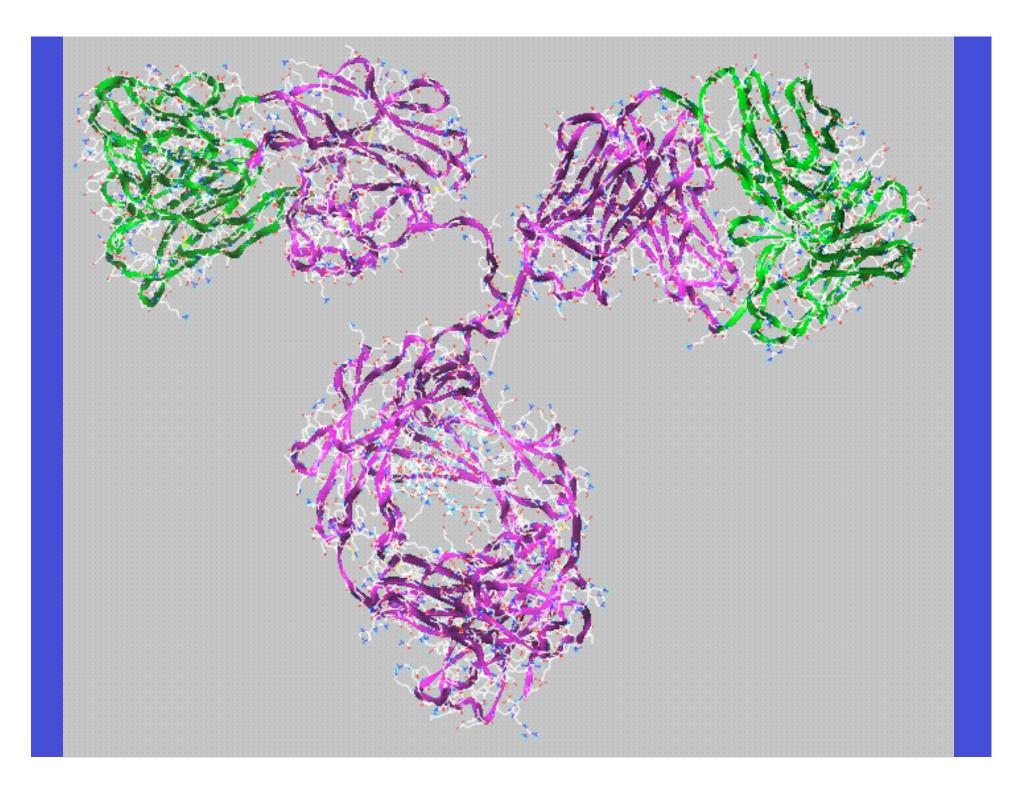
Model of the variable heavy domain of CDP571

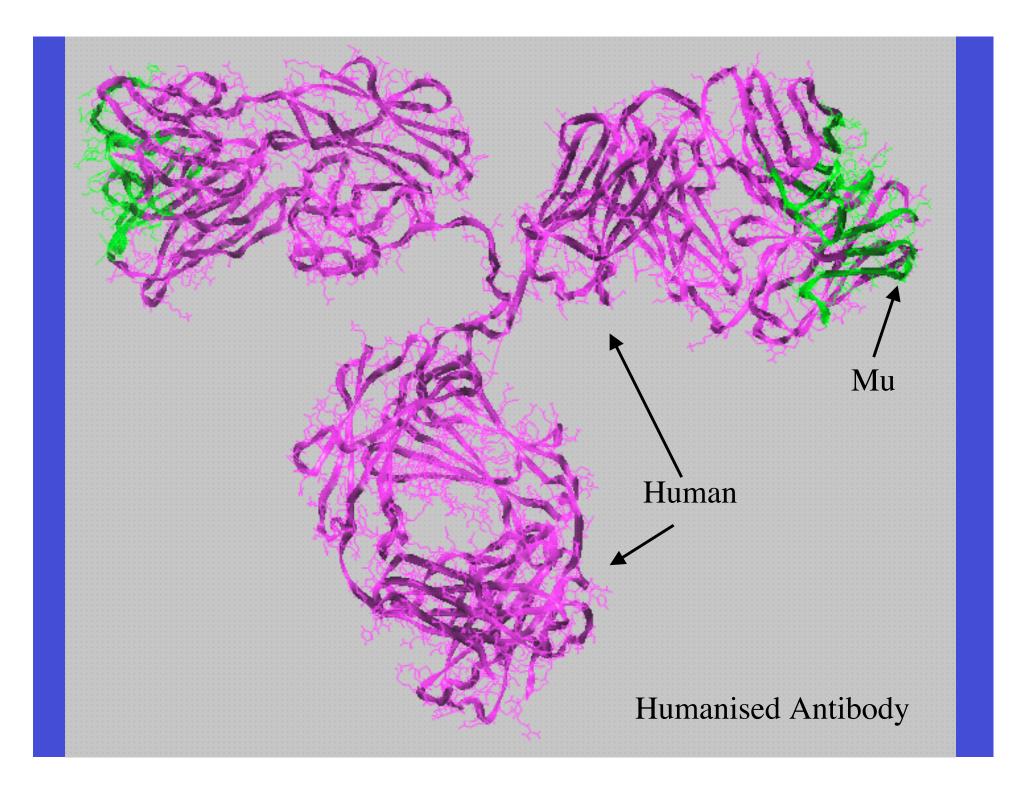


Model showing CDR's and 4 key amino acid substitution positions.









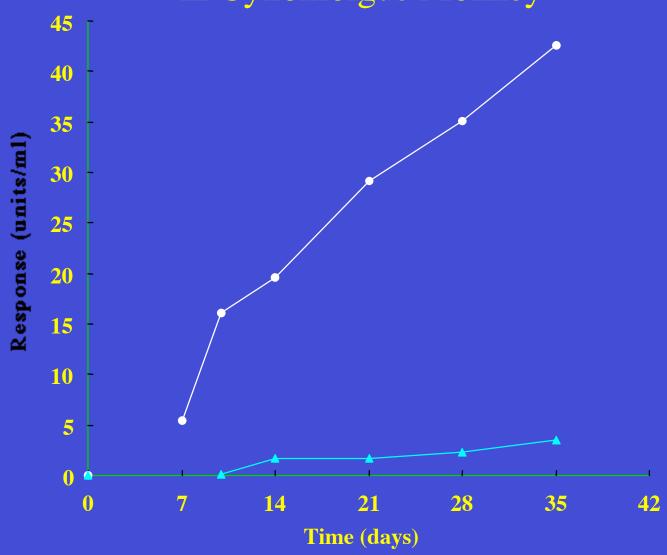
Generation of High Affinity Antibodies

OTypical Affinities:

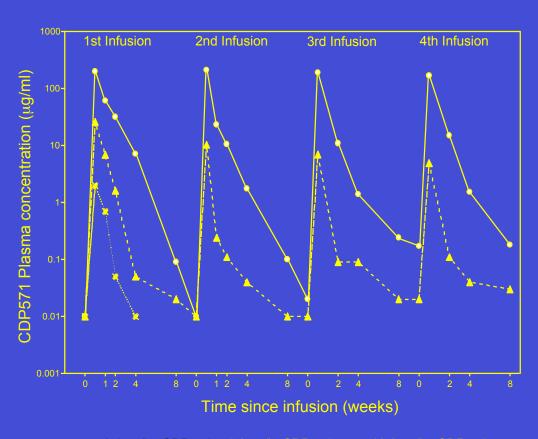
<u>Mab</u>	Mu Affinity	Hu Affinity	<u>Antigen</u>
O Ab1	3. 8 x 10 ⁻¹⁰	3.2 x10 ⁻¹⁰	TNF
O Ab2	0.86 x10 ⁻¹⁰	0.5×10^{-10}	Cytokine
O Ab3	2.5 x10 ⁻¹⁰	4.5 x 10 ⁻¹⁰	Receptor
O Ab4	4.2 x 10 ⁻¹⁰	4.6×10^{-10}	CD33
O Ab5	12.8 x10 ⁻¹⁰	12.7x10 ⁻¹⁰	gp43



Antibodies to Murine and Humanised TNF-antibody in Cynomolgus Monkey



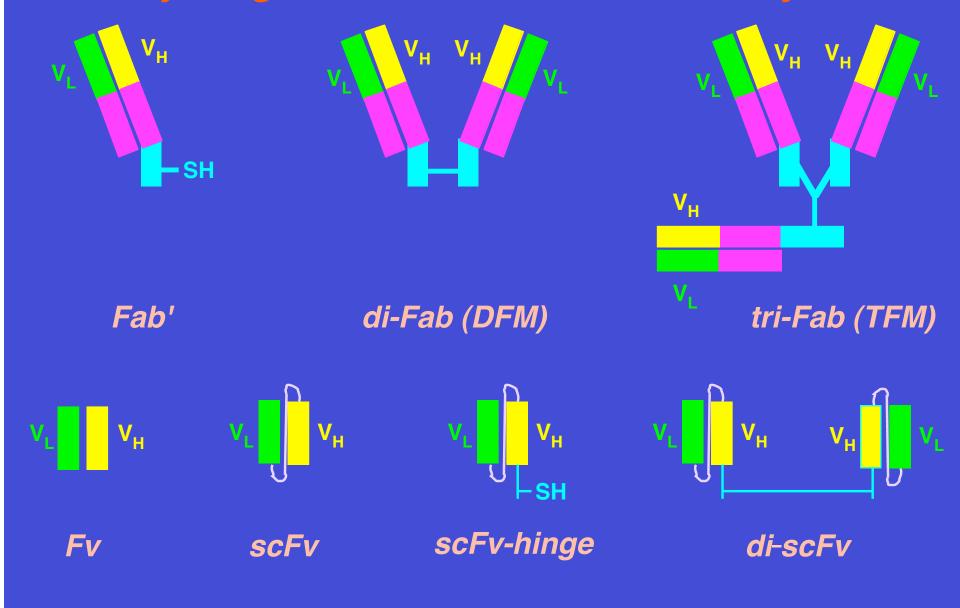
Repeat administration of CDP571 in Patients with Rheumatoid Arthritis



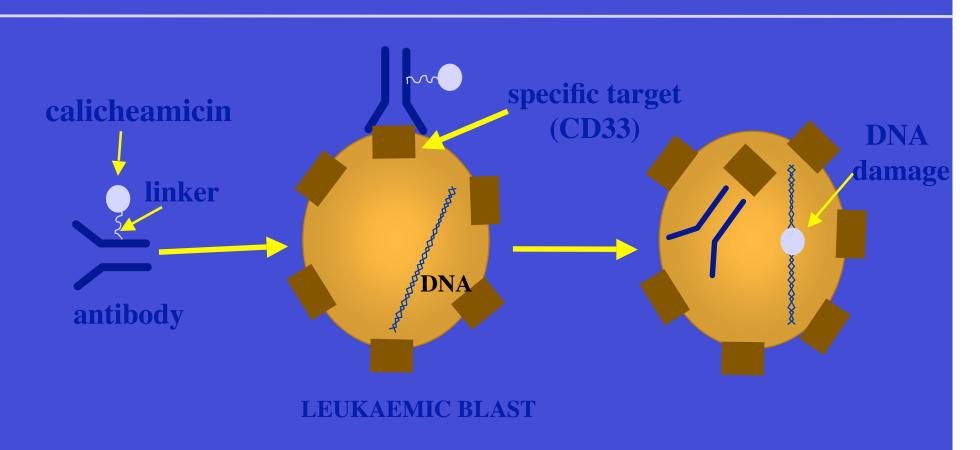


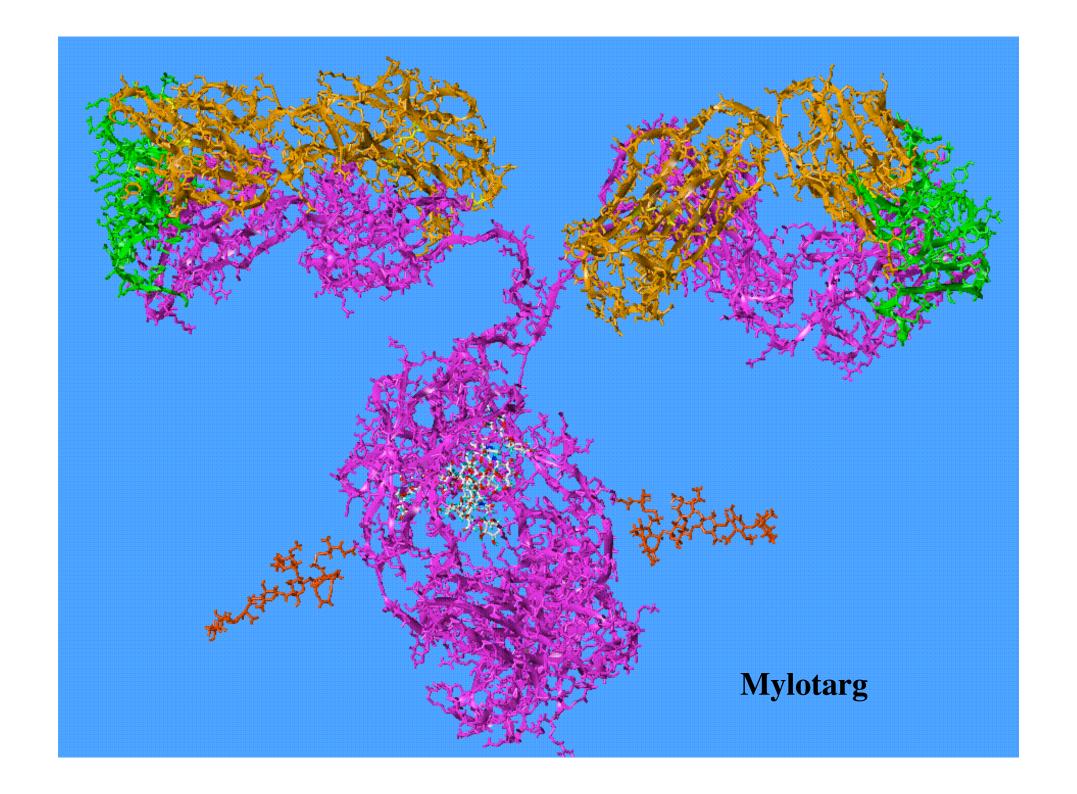


Antibody Fragments Evaluated for Delivery of 90 Y



Mylotarg[™] - a pioneering treatment for acute myeloid leukaemia





5 melvial NDC 0008-4510-01 MYLOTARG™ (gemtuzumab ozogamicin for Injection) 5 mg/vial For IV infusion only DO NOT ADMINISTER AS IV PUSH OR BOLUS Rx only ▼ Wyeth Laboratories Division of Wyeth-Ayerst Pharmaceuticals inc.

