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# **Antibody Humanization**

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## Murine monoclonal Abs





We can engineer the antibody so that it "looks" human.



### Antibodies are modular





# Chimeric antibody

Human Constant domain

Murine variable domains



# Chimeric antibody

Human Constant domain

Murine variable domains

Still antigenic!



### Chimeric vs humanised



#### Chimeric vs humanised



How much can we humanise without loosing affinity?

## Humanization strategies

CDR grafting: Use only the murine CDRS

SDR grafting: CDR + structurally important residues (Canonical Structures)

Resurfacing: Change all surface residues (apart to CDR) to human

Superhumanization: Check for each segment in the Ab and add mutation to maximize local similarity to the human

### Humanization strategies

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# Humanization strategies

CDR grafting: Use only the murine CDRS

usually the antibody loses affinity

SDR grafting: CDR + structurally important residues (Canonical Structures)

often binding is poor

# Antibody Paratope

The binding probability of each residue depends on all the environment.

Even distant residues might change the binding mode

# Framework residues affect antigen binding (Mouse)



# Framework residues affect antigen binding (Mouse)



### Humanization process















# **Basic Protocol**

- 1) Identify Murine germlines
- 2) Identify the murine CDRs (IMGT, Kabat, etc.)
- 3) Identify Human acceptor
  - i- germlines
  - ii- complete antibody
- 4) Graft the CDR
- 5) Identify all backmutations
- 6) Select Backmutations to include
- 7) Test production, affinity, immunogenicity etc.
- 8) If results are not good, go to 6

### **Backmutations**

How to priorities backmutations:

- Are important for the structure (Vernier zone, L/H interface)
- Modify the charge
- Prevent clashes and cavities
- Do not increase the risk of immunogenicity
- Improve expression (literature)
- No additional PTMs or digestion sites

### Conclusion

Antibody humanization is needed for any Ab drug

It is still a "trial and error" procedure

non CDR residues can greatly affect the binding

Structural analysis is extremely important