



Austrian Agency for Health and Food Safety  
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# Identification of an appropriate similarity condition

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# Overview of this talk



## Setting a similarity condition

- Understanding the Active Pharmaceutical Ingredient
  - Analytical characterisation using state of the art orthogonal methods
  - Isolation and in-depth assesment of variants
- Identification of Critical Quality Attributes
- Establishment of a dedicated control strategy
  - Setting of IPC's and Specifications
  - Analytical validation
  - Process Performance Qualification runs
- Comparability protocol
  - Setting similarity conditions
- Comparability study



# When do we assess comparability?



## ☞ Changes to (adaptations of) established manufacturing processes

- Process transfer
- Scale up / down
- Adaptations – optimisations
  - Optimisation of Amino acid sequence
  - Switch to different expression host
  - Changes w/i USP – media / process / scale
  - Changes w/i DSP – media / process / cycling
  - Changes to formulation – Excipients / process

## ☞ Development of a Biosimilar

# What means comparable?



## ➤ No change in CQA's

- Initially only focused on changes of manufacturing processes
- Extended to confirmation of biosimilarity

## ➤ Comparability according to ICH Q5E

- Science driven approach
- Flexible – case by case
- Pre- and post-change product highly similar but not necessarily identical
- Statistical flexibility
  - Comparability ranges
  - Historically justified quality range
- Differences have no adverse impact on clinical safety (including immunogenicity) and efficacy (including pharmacology)

# What is the meaning of biosimilarity?



## No change of CQA's

- Highly similar quality profile, notwithstanding minor differences in clinically inactive components
- No clinically meaningful differences between the biologic product and the reference product in terms of safety, purity, and potency
- Demonstrated by extensive comparability exercise using orthogonal methods – not restricted to release and stability specification
- Analysis of degradation pathways
- Method status: Qualified for the intended purpose!
- Any differences
  - Will have to be appropriately justified with regard to their potential impact on safety and efficacy
  - Might trigger further non-clinical assessment

# Understanding of the API



## Identification of CQA's

### ↪ Efficacy related properties

- Biological functions
  - Do we understand the MOA?
  - Do we know all the interaction partners?
- Understanding of the physiological milieu
- Effects of post-translational modifications
- Micro-heterogeneity triggered alteration of biological functions
- Stability profile – stressed degradation studies

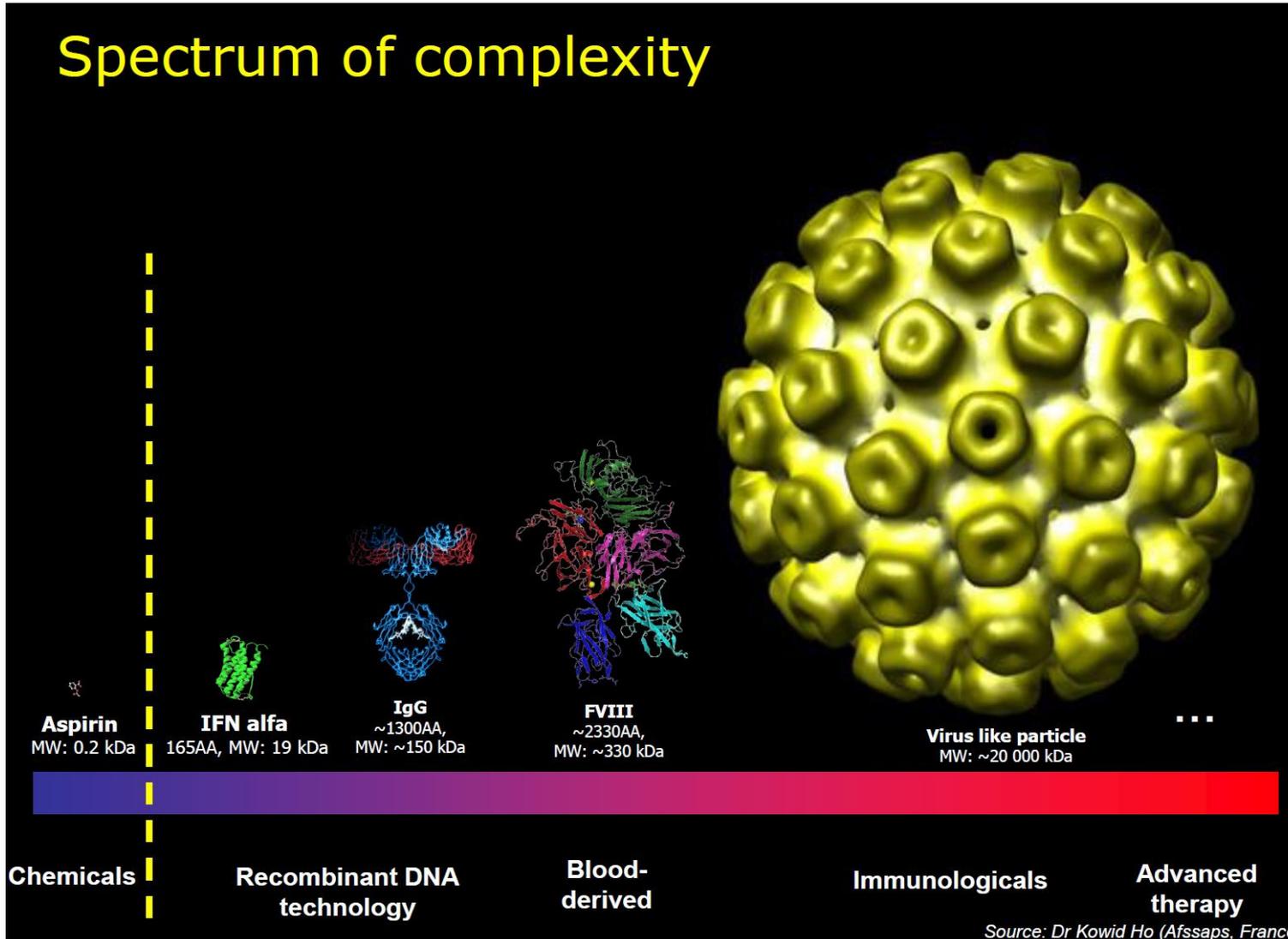
### ↪ Safety related properties

- Impurity profile
- Immunogenicity (product related and process related)

# Spectrum of Complexity

## Large molecules

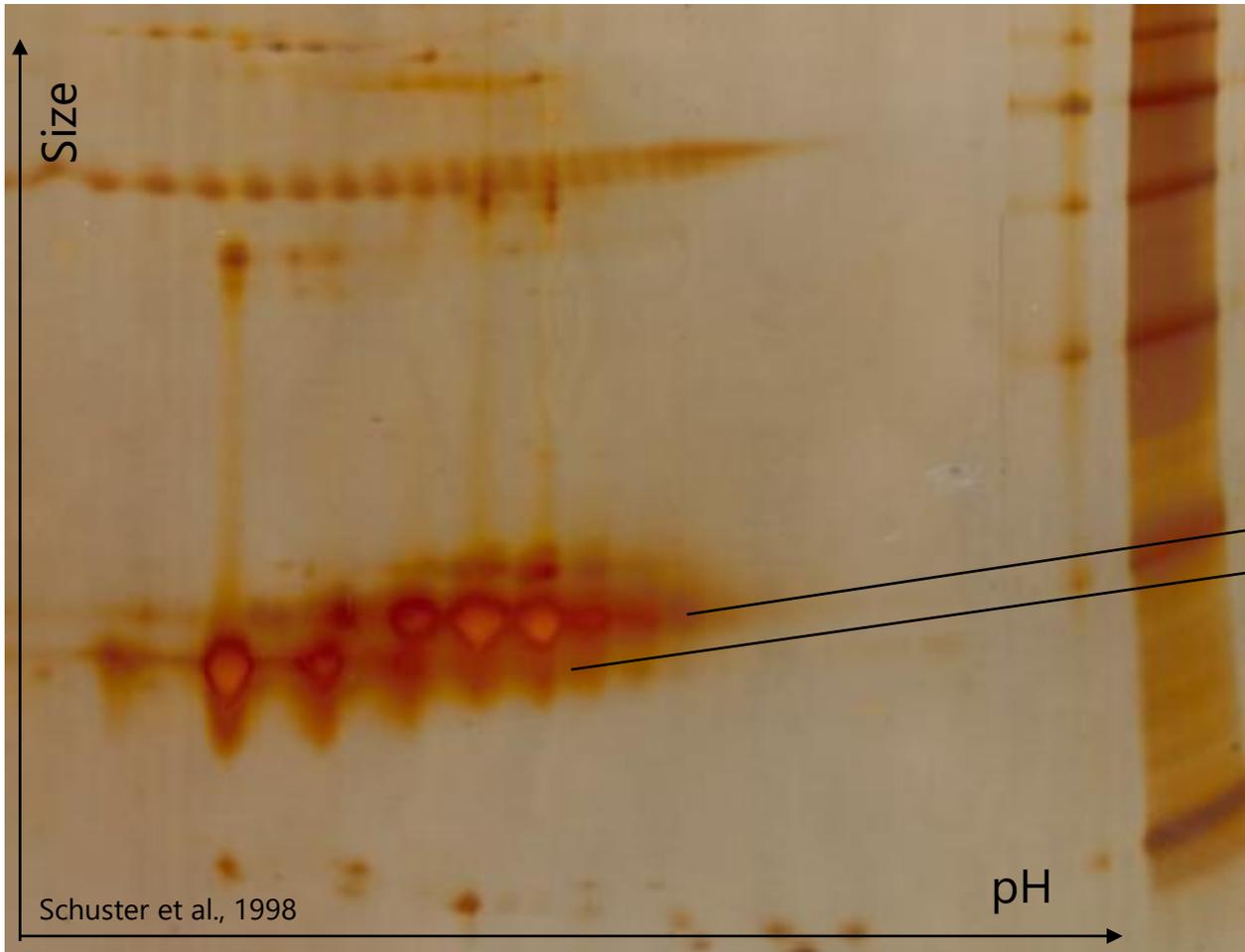
## Spectrum of complexity



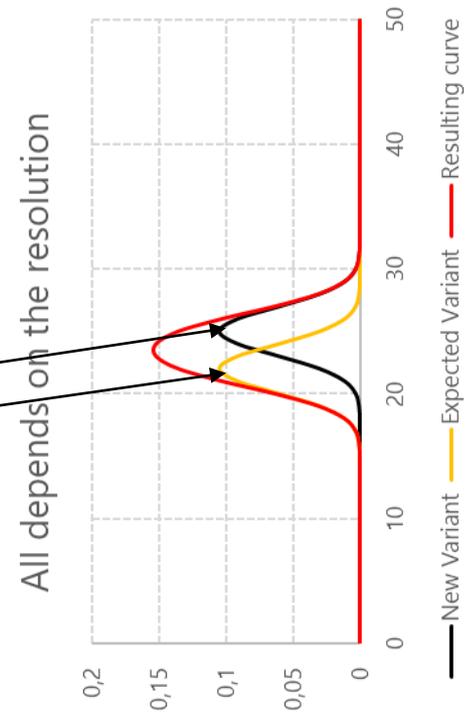
# Spectrum of Complexity

## Complex molecules

rec. human Eukaryotic Elongation Initiation Factor 5A



Schuster et al., 1998

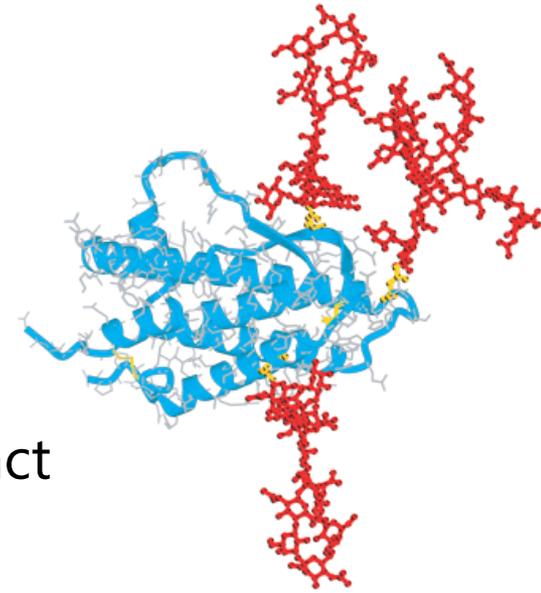


— New Variant — Expected Variant — Resulting curve

# Spectrum of Complexity

## Understanding the API

- Complex manufacturing process
    - Multiple steps
    - Black box of the expression host
  - Large size and complex molecules
  - Small process changes may have a high impact
  - Variability
    - At process level – batch to batch variability
    - Analytical variability – assay precision
    - Moving target: time dependent variability
- Instability of the API (pH, temperature, oxidation, mechanical stress,...) –
- No characterization at molecular level
    - micro-heterogeneity in structure



Courtesy of M.R. Wormald and R.A. Dwek, Oxford Glycobiology Institute, and P.M. Rudd, NIBRT

rec. human EPO

# Analytical toolbox



## Understanding the API

- Primary, secondary, tertiary and quaternary structure  
Peptide map, disulfide bridges, CD, FT-IR, crystallisation,...
- Purity: size, hydrophobicity, charge distribution  
CE, HPLC, PAGE, AUC, DLS,...
- Post-translational modifications  
N- and O-linked glycosylation, phosphorylation, proteolysis, ubiquitynation, oxidation, deamidation,...
- Content and identity
- Impurities and adventitious agents  
CE, HPLC, sterility, qPCR, endotoxins, ELISA,...
- Degradation pathways – stress studies

# Biological activity

## Understanding the API



- ↪ Set of binding and cell-based assays for characterisation and comparison of biological activity
- ↪ Assessment of all known biological functions necessary
- ↪ Ideally inclusion of all interaction partners in the physiological milieu
- ↪ Assay format
  - Bioassays, SPR, Enzymatic assays, ECL based displacement assays
- ↪ Side by side analysis required
- ↪ For mAb's
  - Fab-associated functions (e.g. neutralisation of a soluble ligand, receptor activation or blockade)
  - Fc-associated functions (e.g. ADCC, CDC, complement activation)

# Setting of similarity conditions



## Managing process changes

### Setting ranges

- Quantitative ranges where possible
- **Not be wider than variability of representative RMP batches**
- A descriptive statistical approach to establish ranges for quality attributes could be used, if appropriately justified

### Statistical approaches

- No regulatory requirement to use any specific statistical method
- Proposals for statistical evaluation need to be justified
- Raw data should always be provided to enable assessment of comparability independently from applied statistics
- Statistical package assessed on a case by case basis

# Justification of equivalence ranges



## Understanding of variability

- Micro-heterogeneity of biologics
- Analytical variability – precision and accuracy
- Batch to batch variability
- Stability – aging of biologics

## CQA's are controlled by limits

- IPC's
- Specification

## Acceptable variability

- Clinically justified – safety & efficacy
- Experienced with pre-change material or RMP
- Limitations by analytical performance

# How do we compare

## Establishment of acceptance criteria



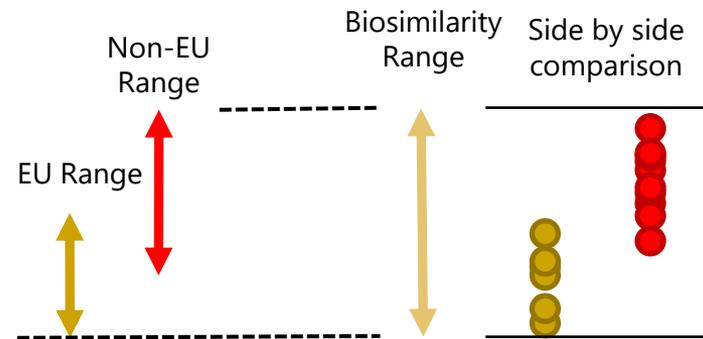
↪ Treatment of patient occurs via an individual batch ( $\neq$  mean)

↪ Pre-defined comparability range

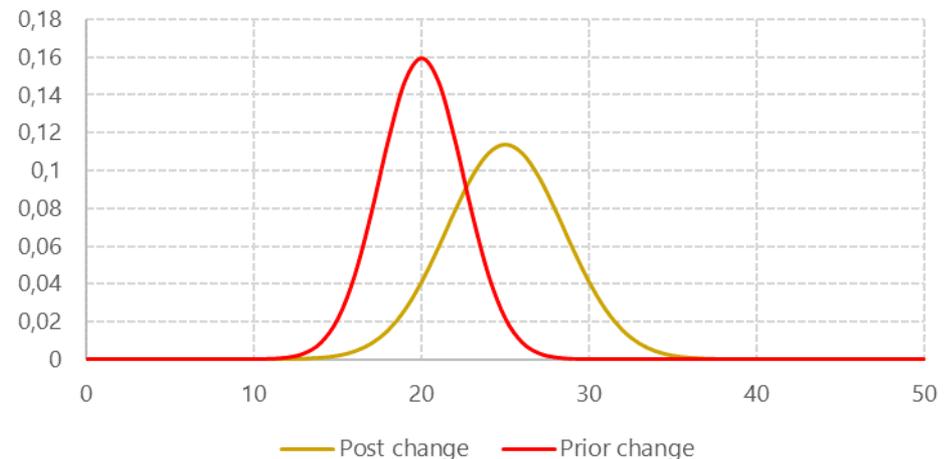
- Min-max approach
- Mean  $\pm$   $k \times$  SD

↪ Equivalence testing

- $\Delta$  of means
- $\Delta$  of variance



Comparable?



Cesar dog food: Perhaps we search out a companion that reminds us of ourselves

# Equivalence testing - comparability range



## Pros and cons

### ☞ **Equivalence testing** – pure statistical evaluation

- Complex!
- Increases objectivity
- Assumes normal distribution of analytical and process variability
- Robust towards outliers
- Focussed on mean

90% c.i. of means & w/i  $1.5 \sigma$  of the reference product

### ☞ **Comparability range** – scientifically justified

- Reflects control strategy (upper and lower acceptance limits)
- Clinically justified, thus scientific rationale
- In line with EU biosimilarity guidance - should not be wider than the range of variability of RMP unless otherwise justified

### ☞ 3 tiered approach

- Specific tier based criterion: equivalence, ranges, descriptive

# Biosimilarity



## **Selected concerns raised during the review process of biosimilar applications and in scientific advice procedures**

- ↪ Setting of biosimilarity ranges – statistical approaches  
Tolerance interval based on analysis reduced number of RMP batches resulted in too wide ranges for biosimilarity assessment
- ↪ Use of non - EEA authorised RMP requires full quality comparison of non - EEA RMP with EEA RMP
- ↪ Differences in defined quality attributes needs to be justified and might require isolation and characterisation of the isolated variants

# Limitations



## ☞ CQA's

- Number of CQA's
- Understanding of CQA's
- Inter-relations of CQA's – moieties within the API
- Clinical relevance – dose relationship mostly not established
- CQA's controlled by process & IPC's

## ☞ Sample numbers – impossible to sample all RMP lots

## ☞ Sampling might be biased

- Age of batches – shelf lives – stability indicating QA's – comparison @ EOS, normalisation?
- DP's from identical DS lots
- RMP process changes
- Batches from other markets – supportive only

# Limitations



## ↪ Statistical limitations

- Data BQL or „no new peaks“ above detection limit
- „Presence of major peaks only“
- Comparison of fingerprints

## ↪ Process-related impurities are process specific

## ↪ Statistical tools

- Data size and distribution driven
- CQA driven

# Examples

## IgG – 150 kDa and $7.2 \times 10^{16}$ possibilities



### Physicochemical characteristics

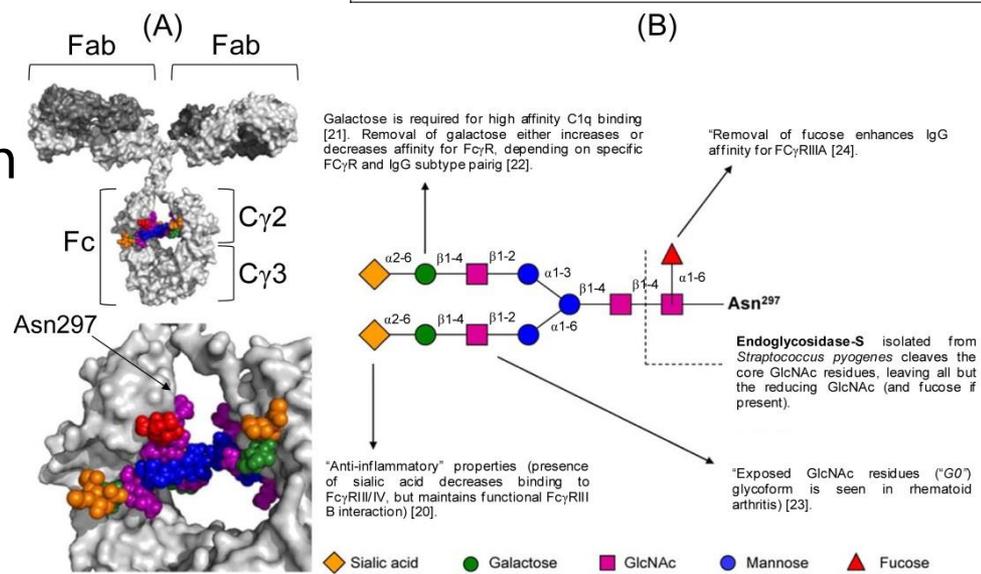
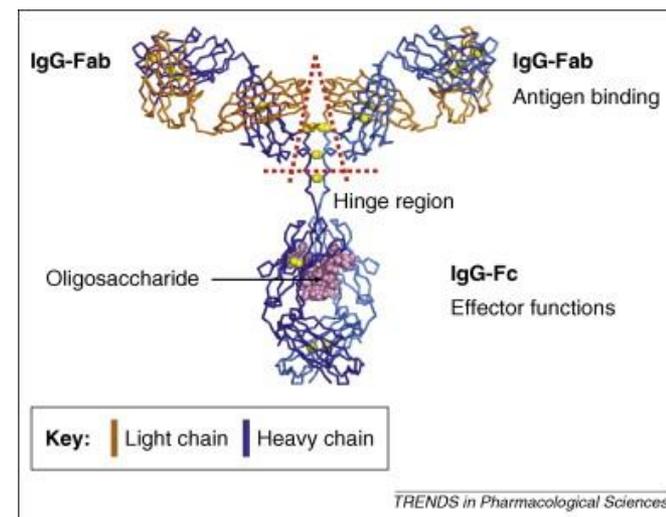
- N-glycosylation sites (2)
- Disulphide bridges (16) – shuffling/cleavage
- Deamidation, acetylation, glycation
- Methionine oxidation sites
- Pyro-Glutamic acid
- C-term lysine
- Fragmentation, aggregation

### Interaction with target antigen

Affinity, avidity, crossreactivity

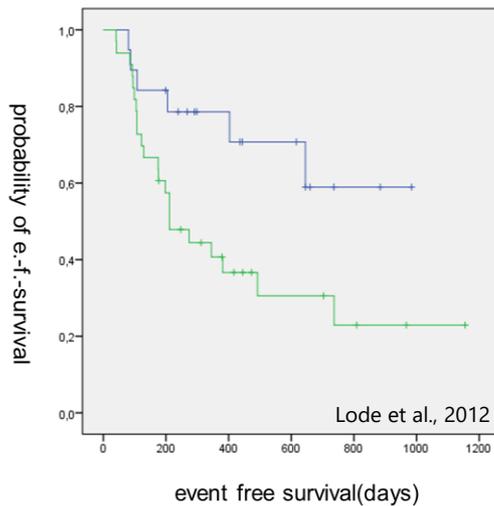
### Fc related interaction

- Effector functions
- Pharmacokinetics



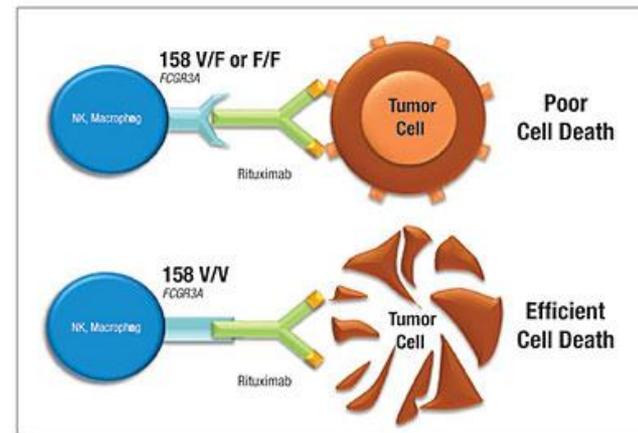
# Clinical relevance of 158 F/V polymorphism

## Dinutuximab



2A \ 3A	H/H	H/R	R/R
V/V	Blue	Blue	Green
V/F	Blue	Blue	Green
F/F	Green	Green	Green

Log-rank test: p=0.025



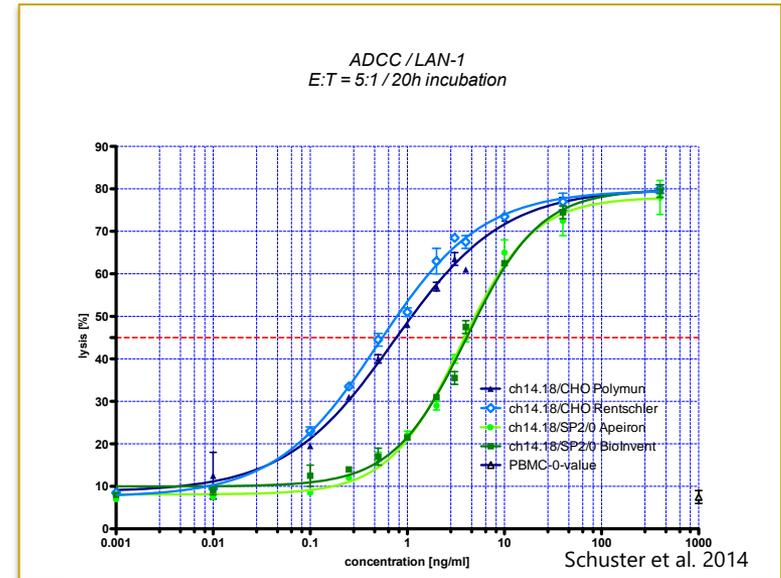
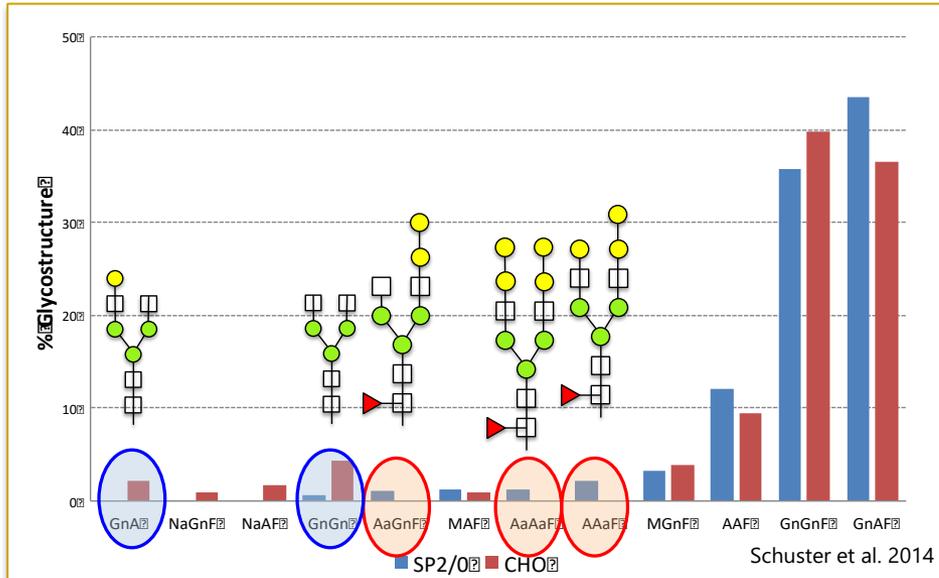
Dall'Ozzo S, et al. Cancer Research, 2004; 64:4664-4669.

- Increased affinity of mAb binding in 158 V/V genotype
- Increased ADCC for 158 V/V genotype
- Reduced overall survival for F/F genotype

## Importance of CD16 binding affinity

# Small differences – considerable effects

## N-linked oligo saccharides - Dinutuximab



- Presence of Galili epitopes on SP2/0 material
- Presence of afucosylated expression products on CHO material
- Clinical confirmation not feasible
  - No head to head comparison
  - Patient numbers - Orphan indication

# Monoclonals are complex molecules



## But well studied product-class

- ↪ The mode of action is complex and may involve contributions from multiple mechanisms
- ↪ High level of microheterogeneity
  - There will always be differences
  - Even small differences may have significant effects
  - Need to combine physicochemical results with functional assays (e.g. antigen-antibody binding assays and cell-based assays)
  - Qualification in preclinical and clinical studies
- ↪ Demonstration that differences do not impact on clinical efficacy and/or safety challenging
- ↪ But: We meanwhile know what to look at

# Biosimilar to Etanercept

Case study

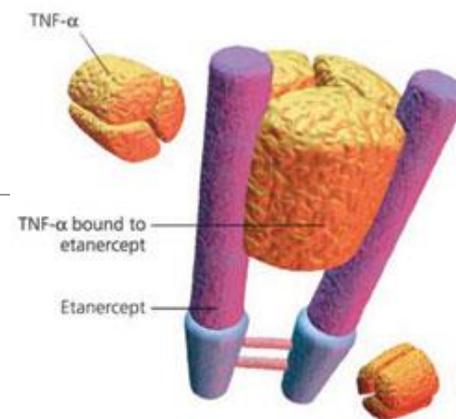
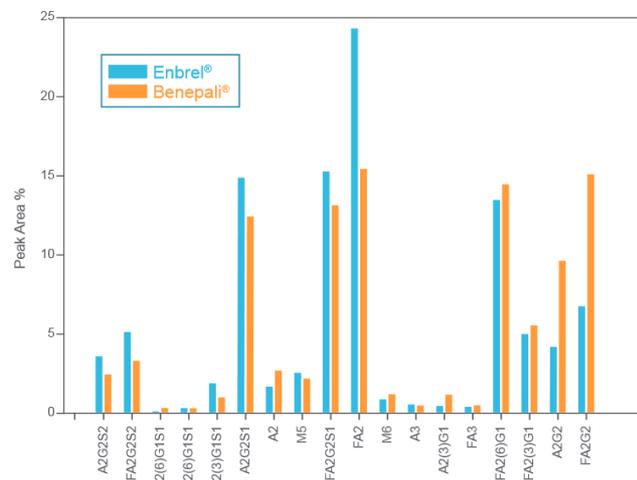
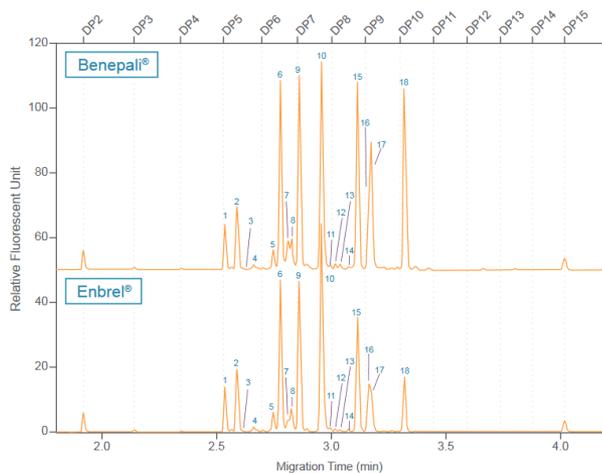


- Different N-glycosylation profile of BS
  - A-fucosylated glycan content in BS higher
  - CD16 binding and ADCC affected – Critical?



Cesar dog food: Perhaps we search out a companion that reminds us of ourselves

- TNF-alpha trap: ADCC not relevant MOA
  - Conclusion: Fucose content in this case not a CQA
  - Differences not clinically meaningful
  - No impact on the safety/efficacy



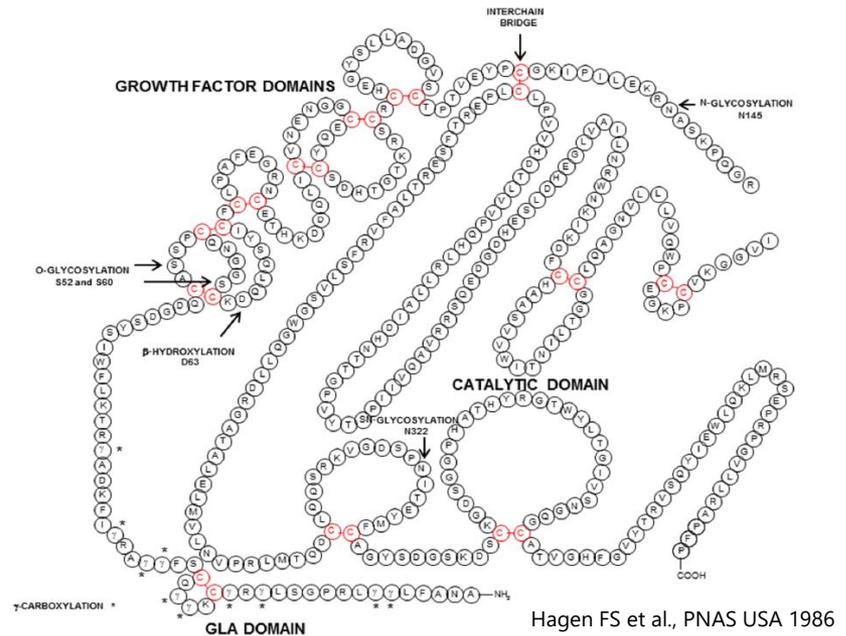
Biosimilars/News, 03.04.2020

# rhFVIIa – 406 AA and $4.5 \times 10^{15}$ possibilities



## Which QA is uncritical?

- Multiple interaction partners
- Post translational modifications
  - 2 N- and 2 O-glycosylation sites
  - 1 Phosphorylation site
  - 1  $\beta$ -hydroxylation site
  - 12 disulphide bridges
  - Activation by proteolytic cleavage
  - Light chain – 152 AA, 20 kDa
    - N-terminal gamma-glutamic acid-rich domain – 9  $\gamma$ -carboxylation sites with multiple calcium-binding sites and
    - 2 epidermal growth factor (EGF)-like domains (kringle domains)
  - Heavy chain – 254 AA, 30 kDa
    - Catalytic domain and a single calcium-binding site



# rhFVII a - Complex molecule

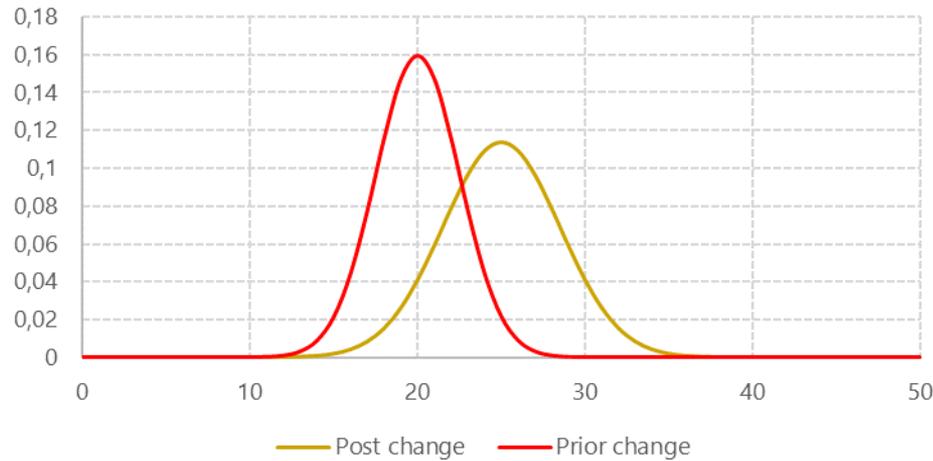


- ↪ Despite long lasting experience – less expertise
- ↪ Relevance of some post-translational modification unclear
  - Singular modification - interplay at molecular level
- ↪ So far no biosimilar approved
- ↪ What do we look at
  - In-depth characterization of the API and identification of CQA's
  - Understanding of process variability
  - Quantitative assessment of every CQA
  - Extensive assessment by functional assays
  - Detailed in-depth characterisation of isolated variants and structure-activity relationship studies
  - Clinical validation

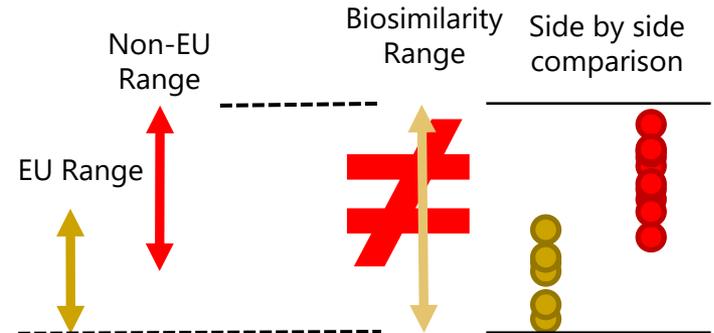
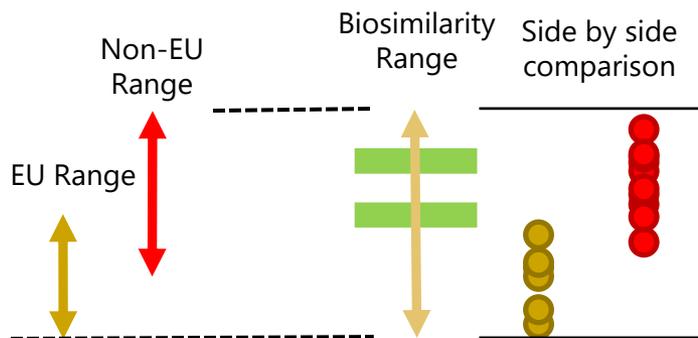
# What do we compare

## Establishment of acceptance criteria

Comparable?



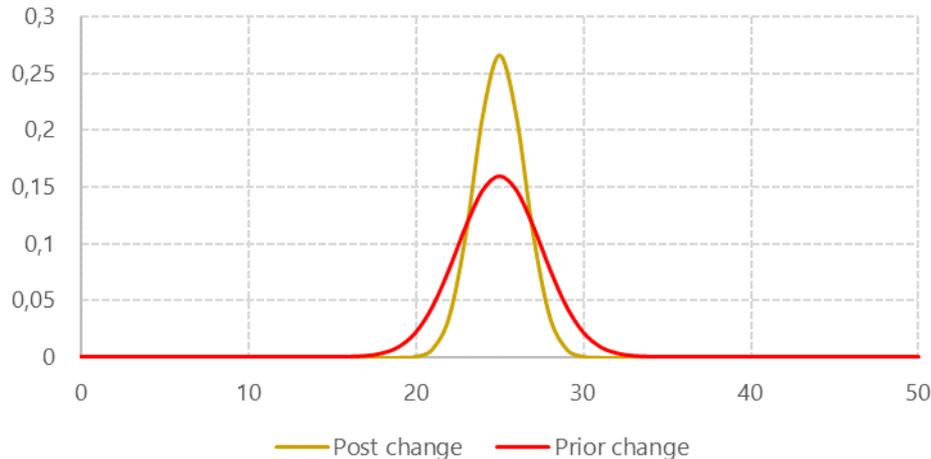
Cesar dog food: Perhaps we search out a companion that reminds us of ourselves



# Macroscopic point of view

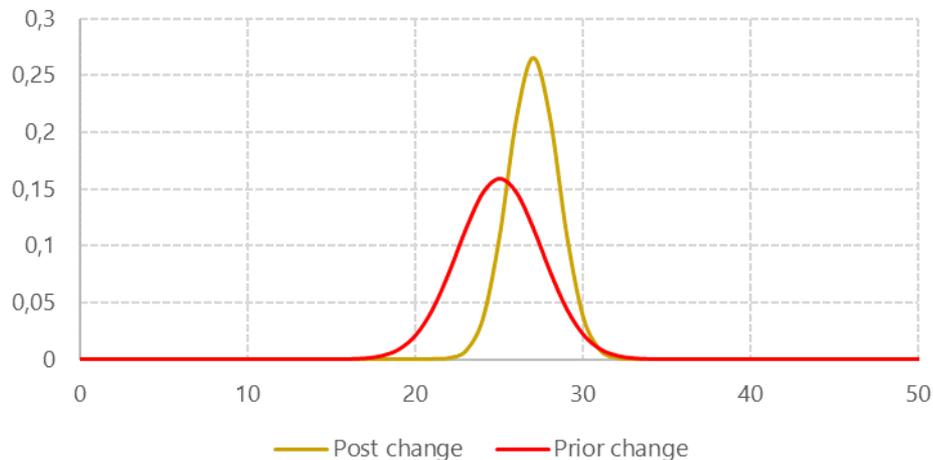
## Continuously Gaussian distributed random variables

Same mean



- Same mean, reduced variance
  - Improved analytical precision
  - Improved process variability
  - Clinically justified
  - Loss of variants unlikely

Shifted mean

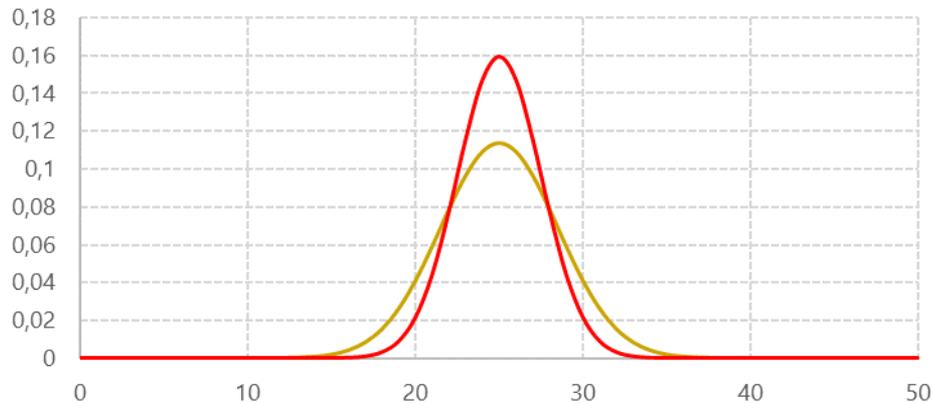


- Shifted mean, reduced variance
  - Improved analytical precision
  - Other Reference Std?
  - Reduced process variability
  - Loss of variants?
  - Increase of other variants?
  - Clinically justified?

# Macroscopic point of view

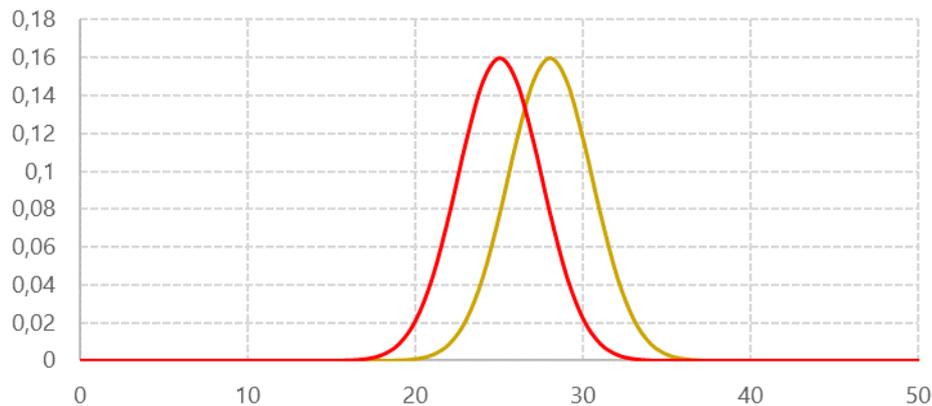
## Continuously Gaussian distributed random variables

Increased variance



- Same mean, increased variance
  - Reduced analytical precision
  - Increased process variability
  - Clinically not justified

Shifted mean



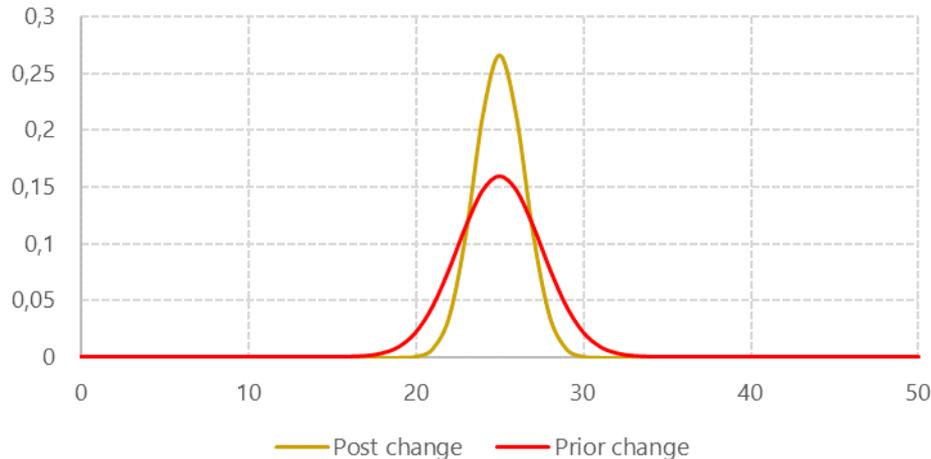
- Shifted mean, identical variance
  - Change in process
  - Clinically not justified

— Post change — Prior change

# In-depth analysis

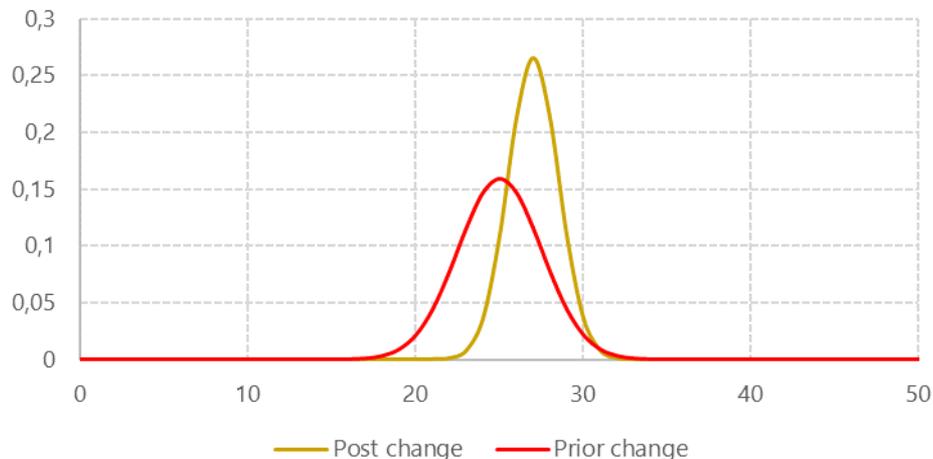
## Random distribution by analytical variability?

Same mean



- Same mean, reduced variance
  - Improved analytical precision
  - Reduced process variability
  - Clinically justified

Shifted mean

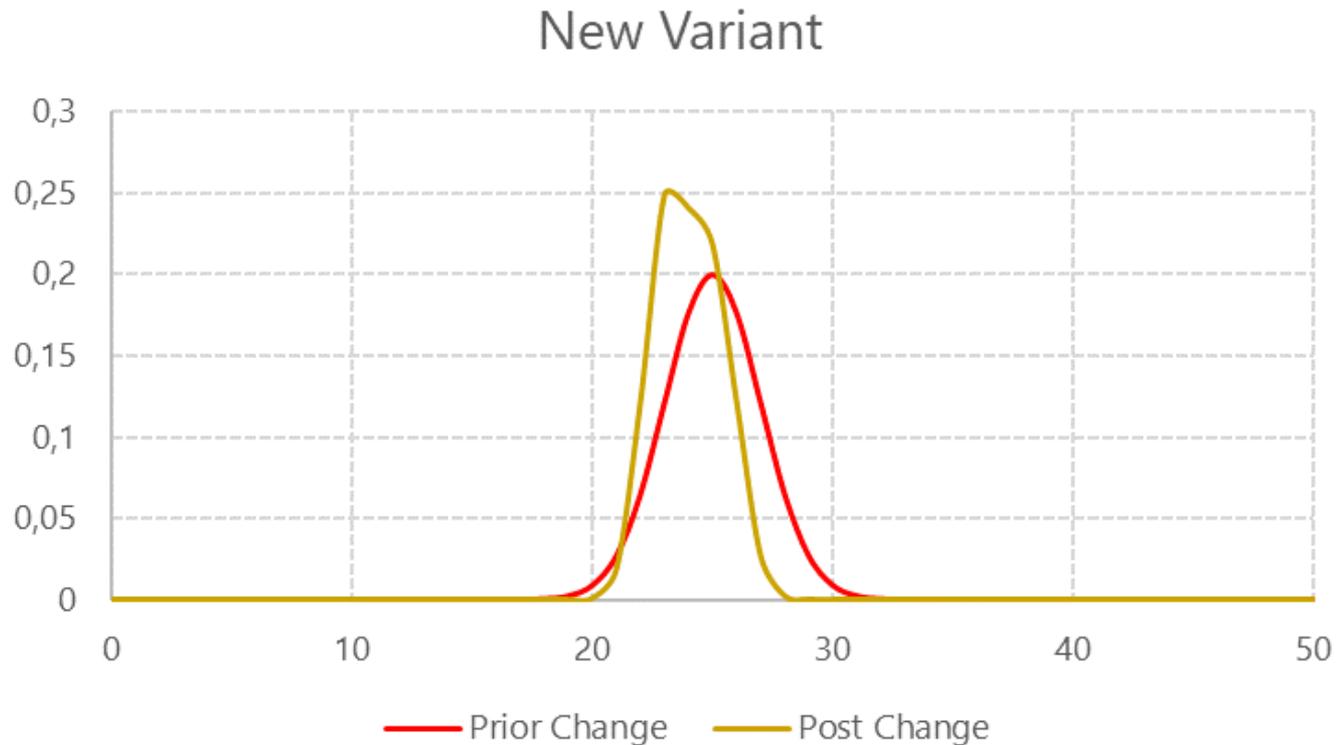


- Shifted mean, reduced variance
  - Improved analytical precision
  - Reduced process variability
  - Shifted mean contained within the original data range
  - Is it really clinically justified?

# In-depth analysis

## Different peak shape

- ↪ Shifted mean, same variance?
  - Peak shape slightly different
  - Clinically justified?

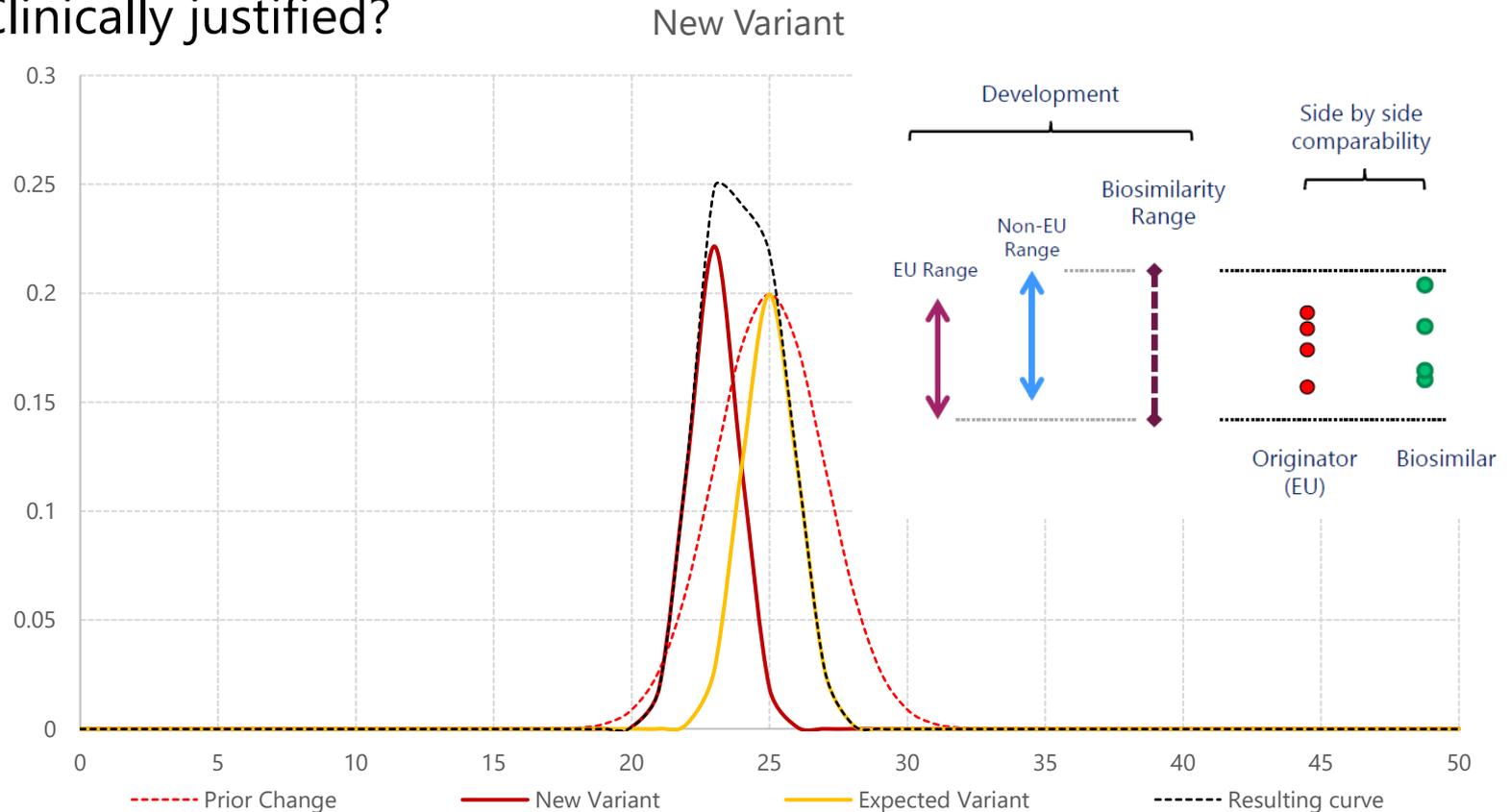


# In-depth analysis



## Discrete molecular variants!

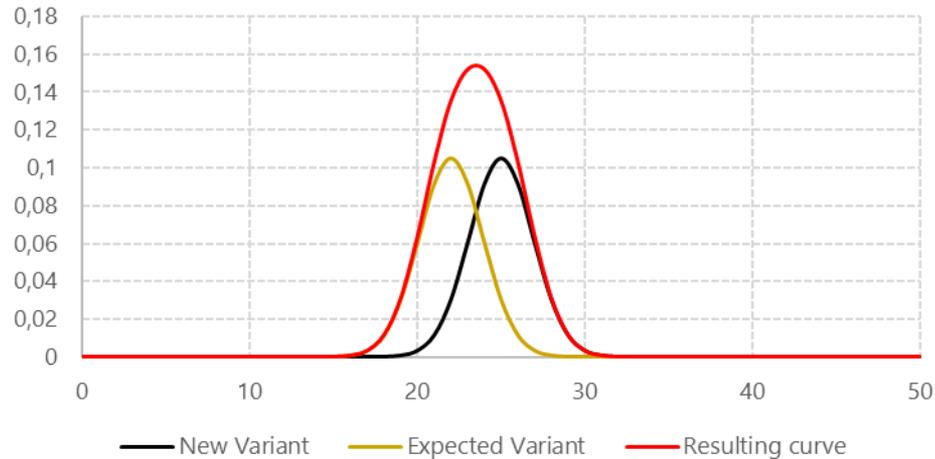
- Change in manufacturing process leads to new variant
- Impact on safety and efficacy?
- Clinically justified?



# What is behind the peaks?

## Random distribution by analytical variability?

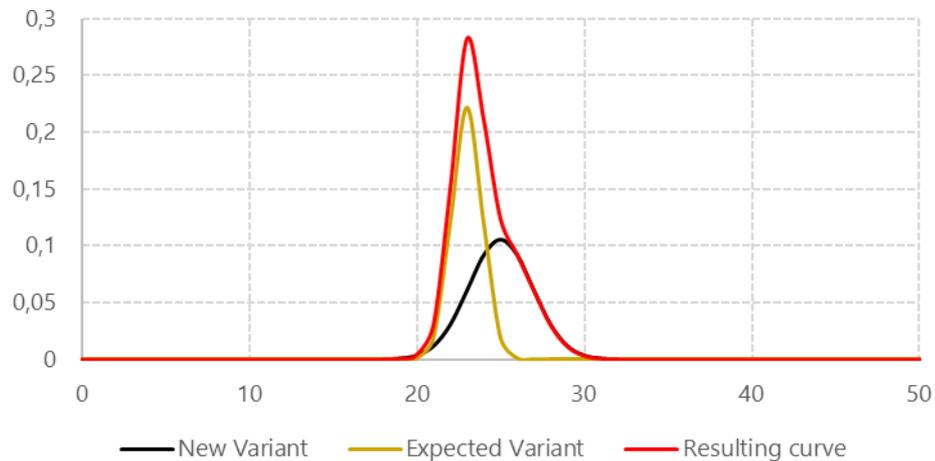
Gaussian distributed resulting peak



2 species, similar amounts

- Broad gaussian distributed peak
- Improved analytical resolution
- Orthogonal methods
- Clinically justified?

Skewed Peaks



2 species, different amounts

- Further investigations required
- Criticality?
- Improved analytical resolution
- Reduced process variability
- Clinically justified?

# It depends!



## Defining a similarity condition

- ☞ Efficacy - what is the MOA
  - Antigen traps / neutralisation of ligand-receptor interactions
  - Cellular effector functions / activation of complement cascade
  - Impact of pharmacology – recycling via FcRn
  - Biological / cellular assays
- ☞ Safety - specific process and product related impurity profile
  - Immunogenicity – aggregates, oligosaccharides
  - Side effects triggered by complement activation
  - Charge variants, de-amided, oxidized, C-terminal Lysine variants
- ☞ mAbs with Fc-triggered effector functions
  - Binding to antigen and to Fcγ, FcRn and to C1q
  - N-linked oligo-saccharides

# Establishment of a similarity condition



## Summary

- Selection of multiple API based comparability criteria
  - Scientifically justified for substance class by literature
  - Identified through in-depth product characterisation, isolation and analysis of defined variants and criticality assessment

Examples: content of variant A, amount of defined oligo-sacch. structure,...
- Analytical performance
  - Identification and establishment of the analytical portfolio
  - Assessment of analytical suitability and variability
- Process performance
  - Assessment of process variability
- Definition of acceptance criteria for head to head assessment
  - Justification by RMP / (pre-)clinical studies with representative batches

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