Setting specifications...

... or when statisticians can play the Good, the Bad, and the Ugly...

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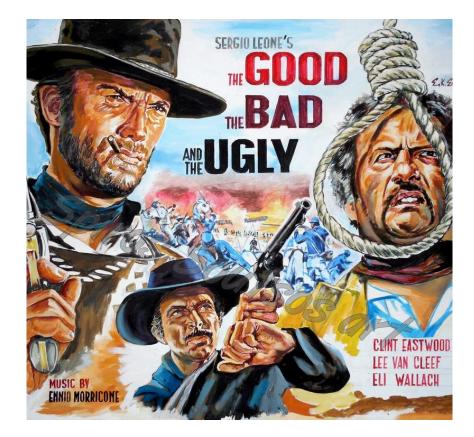


What's good, what's bad?

- Setting specifications = define a threshold which determine what is good and what is bad...
- As statisticians, we can derive intervals, probabilities, but to determine what is good and what is bad is over our scope

 Very high risks in case of bad choice... High pressure from scientists, process owners and authorities

 Lack of worldwide agreement on principles and practices



Setting specifications limits, in theory

Product safety & efficacy

Non-clinical / clinical data to define clinically relevant specifications directly related to product safety and efficacy

Specs

Manufacturing

Data covering the full range of variability expected for the product Include process, analytical variability, stability data, ect.

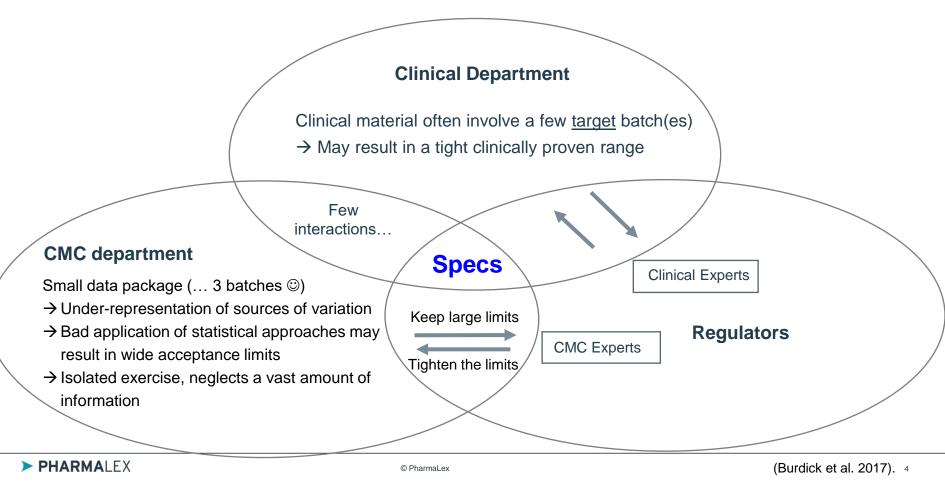
Appropriate methods to estimate variabilities and derive limits (DoE, DS, ect.)

Regulatory (QbD framework)

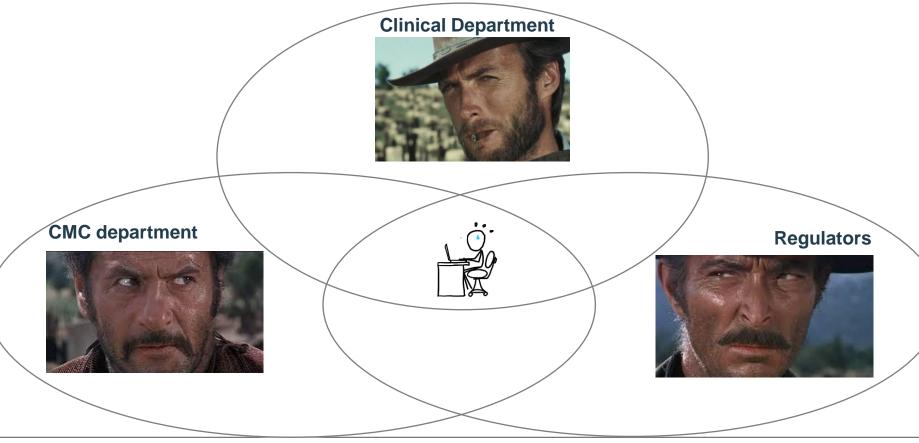
Development of specs is a process driven by data obtained throughout the product life cycle following a control strategy

→ Balance the risks

Setting specifications limits, in practice



Setting specifications limits, a Mexican standoff...



REAL STORY, BUT ALL RESULTS AND FIGURES BASED ON SIMULATED

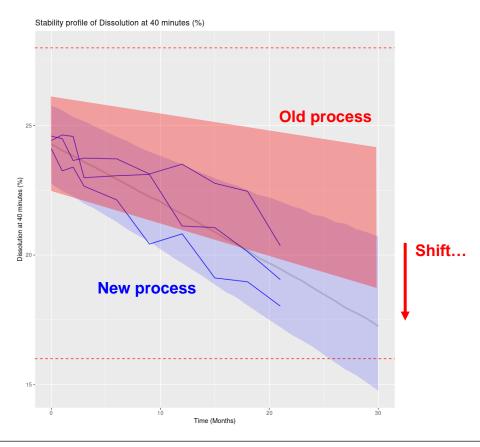
DATA FOR CONFIDENTIALITY PURPOSES

Illustration of a (hot) request about specifications limits

 After a change in the manufacturing process, a shift in dissolution was observed in stability

New batches degrade quicklier than those before the change...

Is there a risk of Out-Of-Specification?



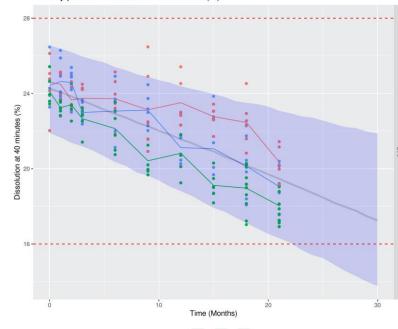
Prediction using a hierarchical model (Bayesian implementation)

$$y_{ij} = A + \alpha_i + B \times T_{ij} + \beta_i \times T_{ij} + \varepsilon_{ij}$$

 y_{ij} = Value for i^{th} batch at j^{th} time point

- A = overall Intercept
- α_i = random effect of the *i*th batch: ~ N(0, σ_{α}^2)
- B = overall Slope
- $\begin{array}{ll} \beta_i & = {\rm random \ effect \ of \ the \ slope \ of \ the \ } i^{th} \ {\rm batch:} \\ & \sim {\rm N}(0, \ \sigma_{\beta}{}^2) \end{array}$
- $T_{ij} = j^{th}$ stability time point for i^{th} batch
- ε_{ij} = Residual Variability ~ N(0, σ_{ε}^2)

With uncorrelated α_i and β_i



Batch

🔶 A 🔶 B 🔶 C

Stability profile of Dissolution at 40 minutes (%)

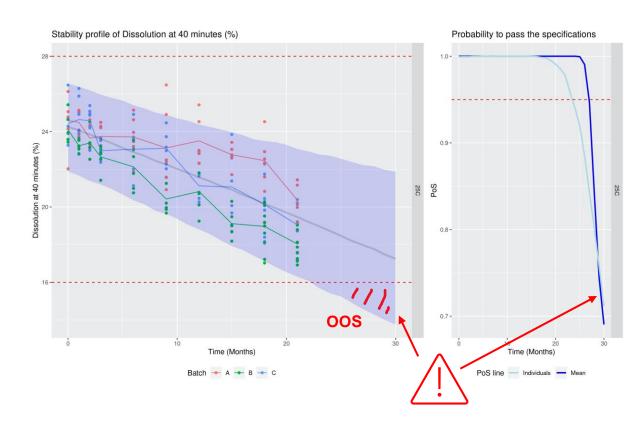
High chances of OOS in one year...

$$y_{ij} = A + \alpha_i + B \times T_{ij} + \beta_i \times T_{ij} + \varepsilon_{ij}$$

 $y_{ij} = \text{Value for } i^{th} \text{ batch at } j^{th} \text{ time point}$

- A = overall Intercept
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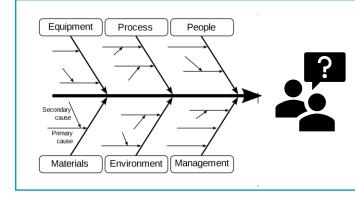
With uncorrelated α_i and β_i



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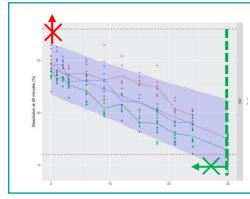
How to avoid OOS with the new process?



No root cause identified despite extensive investigations...

Need to update specifications limits





Not possible to release batches higher because of technical limitations

No lower historical batches available

Not possible to shorten the SL

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A Bioequivalence study was initiated to establish product safety

- A clinical Bioequivalence study was initiated
- The study included ~ 70 subjects
- The study was performed using the batch with the steepest degradation path
- The study duration was ~ 4 months



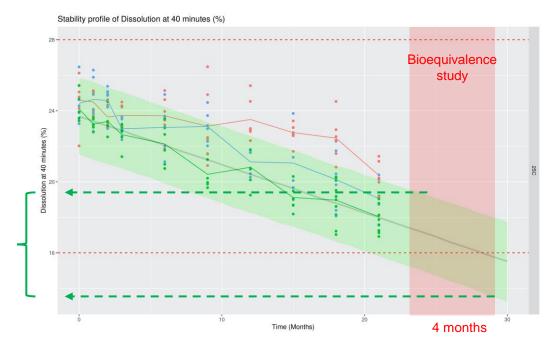
THE INVENTION OF CLINICAL TRIALS

Linking patient and stability data, the strategy



Patients will be exposed to the batch **A** in the clinical trial when that batch will be aged between 24 and 28 months

- → Back predict the dose range patients have been exposed during the whole study
- → If there is no concern in both safety and efficacy in the BA study: we have a good rationale to redefine the specs ☺

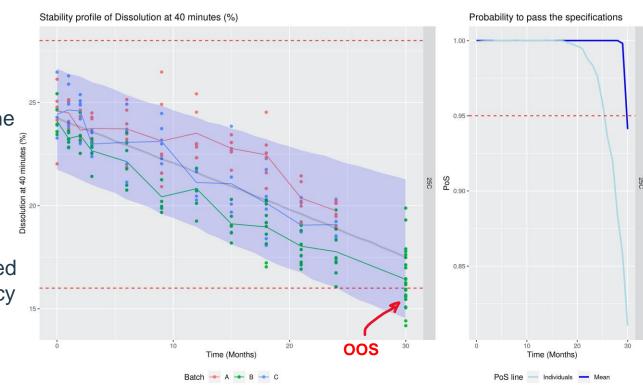


And indeed at 30 months...

- Predictions were proven accurate
- OOS were observed for the Batch A at 30 months

BUT

The BA study was achieved without any safety / efficacy concern ^(C)

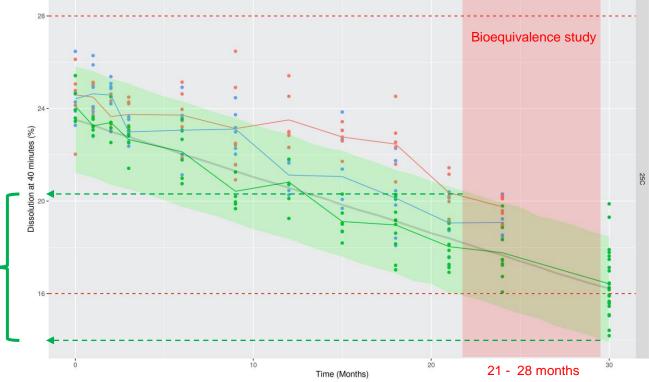


First step: get a clinically relevant dissolution range

Stability profile of Dissolution at 40 minutes (%)

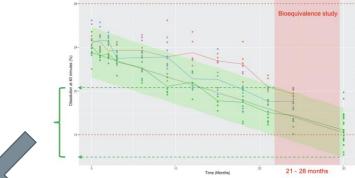
The dose range patients have been exposed during the whole BA study is back-predicted

Dissolutions values ranging from 14 to 20% are clinically safe and efficient ©



Second step: based on the predicted dissolution values, compute the Q-value following on USP-711

IVR time	LSL ('Stated range' in the USP-711 acceptance table 2)
40 minutes	?



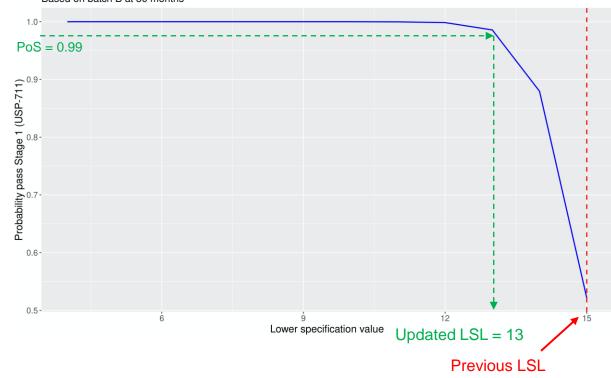
Acceptance Table 2

Level	Number Tested	Criteria
L	6	No individual value lies outside each of the stated ranges and no individual value is less than the stated amount at the final test time.
L2	6	The average value of the 12 units $(L_1 + L_2)$ lies within each of the stated ranges and is not less than the stated amount at the final test time; none is more than 10% of labeled content outside each of the stated ranges; and none is more than 10% of labeled content below the stated amount at the final test time.

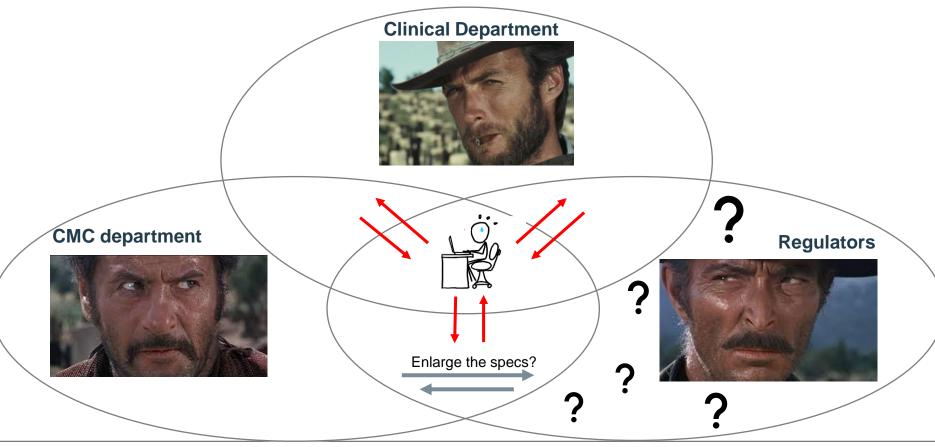
Second step: Back-computation of the Q-value

The new LSL (i.e Q-value) is back-predicted using simulations

- The new LSL is designed to be
 - clinically-relevant
 - process-wise achievable
 - USP-711 compliant



And last but not least, convince regulators...



Take home messages

- Authorities are moving from compliance-driven specifications limits to a risk-based life cycle approach
- But still there is no a globally harmonized guidance available on the topic... But this gives you a great flexibility!
- Flexibility to use all available data to balance the producer and consumer risks
- Flexibility to set provisional limits and to update the limits based on data from post-marketing

Regulatory (QbD framework)

Development of specs is a process driven by data obtained throughout the product life cycle following a control strategy



Many thanks for your attention! Any questions?



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