MODEL-BASED DRUG DEVELOPMENT

1

8 June 2011 EFSPI Statistical Leaders Meeting

Introduction

 As set out in the 1-pager, Sheiner, Senn, Lalonde and colleagues have highlighted an apparent slowness by statisticians to engage with model-based approaches to drug development

Questions

- do we agree with this negative verdict on our discipline?
- is it true, as it seems, that the kineticists have stolen a march on us?
- where is the leadership within our own profession to challenge this view?
- What, if anything, can EFSPI do here?

Models exemplified

- To motivate discussion I will highlight examples from different areas of our business
 - 1. Modelling clinical data
 - (i) Predicting relative efficacy in a new indication
 - (ii) Modelling competitor data
 - (iii) More efficient trial design
 - 2. Pre-clinical PK-PD modelling
 - 3. Biological systems modelling



1. Modelling clinical data

(i) Predicting relative efficacy in a new indication





We can calculate the probability of achieving a Target Value (PTV) for newdrug in the new indication

1. Modelling clinical data (ii) Modelling competitor data

6

Product Concept	Diabetes agent providing weight loss and/or cardiovascular benefits
Mechanism of Action	X
Strategy	Accelerated development to be in the first wave for this MOA
Competitive Landscape	Company is behind several competitors
Key Gaps in Knowledge	How to differentiate from the leading competitor?

1. Modelling clinical data (ii) Modelling competitor data

7

The leading competitor: dose-response for HbA1c%



Estimated E_{max} = 0.59 ± 0.1% with an ED₅₀ of 1.05 ± 0.8 mg for HbA1c effect

Potential outlier at 20 mg dose in diabetic naive study (-0.55% HbA1c) if removed yields

- $E_{max} = 0.67 \pm 0.04$, $ED_{50} 1.7 \pm 0.3$ mg
- variability significantly reduced, especially on ED₅₀

1. Modelling clinical data (ii) Modelling competitor data

8

The leading competitor: dose-response for % weight loss



9

Trial to investigate pain relief following two weeks treatment with *drug* in patients with knee OA



Bayesian Study Design

Use informative prior for naproxen vs. placebo

- Use elicited priors for:
 - drug vs. placebo
 - drug vs. naproxen

11

Prior for the effect of Naproxen vs. Placebo

Δ≈ N(1.6 , 0.58²)

Study_Ref	Diff	SED	Variance
AAAAAAAAA	2.0	0.33	0.11
BBBBBBBBBB	1.6	0.30	0.09
CCCCCC	2.0	0.66	0.44
DDDDDDDDD	2.1	0.83	0.68
EEEEEEEEE	1.1	0.35	0.12
FFFFFFF	1.1	0.51	0.26



□ Using this prior is equivalent to N = 54 subjects on naproxen – placebo → significant efficiency

Elicited Prior Belief for drug vs. Placebo Effect



0.8*N(-1.6, 0.58²) + 0.2*N(-0.15, (0.7²+0.58²))

13

Cumulative Predictive Posteriors for superiority to placebo when drug is Naproxen-like Cusum Predictive Posteriors



- 14
- Pre-clinical modelling is used to inform the expected dose or exposure needed to demonstrate efficacy in humans
- Defining and achieving 'efficacious exposure' (C_{eff}) is essential to create confidence that we have tested the mechanism and can walk away from a negative result in man
- □ Surprisingly, there is little agreement on how to define C_{eff}
- What it is not: the lowest drug concentration to yield a statistically significant difference from negative control (vehicle)

15

Human in vitro rat in vivo binding to target Rat in vivo down-stream pharmacology Dog in vivo efficacy

TK limit, most sensitive nonhuman species



Consistent pharmacology across species/models PK-PD well characterised Large TI C_{eff} will test the mechanism

16

- in vitro and in vivo experiments provide estimates of IC₅₀ / EC₅₀ / K_i
- These are used to construct the E_{max} curves shown
- However, these estimates may not be completely robust



- How to estimate binding affinity (K_i) for a receptor antagonist in vitro?
 - Pooling across different salt forms of drug
 - Pooling of data from different labs
 - Inclusion / exclusion of data points from assay

17



What is an appropriate estimator for Ki?

What is a 'no-regrets' dose? 10 x Ki

3. Biological systems modelling

18

- Everything starts with target selection (human biological drug target)
- We are not biologists
 - Biologists and others are building complex models to describe basic human biology
 - Hypothesised cascade / pathway linking known biological processes
 - Suggests where to intervene to achieve desired pharmacology and avoid unwanted pharmacology
 - Rely on strong assumptions and typically take data from a variety of sources
 - Statisticians should be able to scrutinize these models
- The whole field of systems biology / pharmacology needs greater statistical scrutiny

3. Biological systems modelling



3. Biological systems modelling



After two weeks of dosing, mean ADAS-cog change for monotherapy (150mg) was 3.6 points
Approved Alzheimer's drugs typically show 3-4 point improvement after 12-24 weeks Statistically significant dose-response for 150mg vs. 50mg vs. placebo (p=0.026)

Model-based drug development: Questions

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