



Bayesian Model Comparison for Dose Response Studies

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Disclaimer



- Expository work on recent developments on BMCPMod by Frank Fleischer, Sebastian Bossert, Qiqi Deng, Christina Loley, Jana Gierse (2022)
 Pharmaceutical Statistics.
- MCPMod is well established statistical method. BMCPMod is a Bayesian extension to MCPMod where we assume prior distribution on the mean response.
- No GSK clinical trail data were used. Simulated data has been used for this project.



Contents



- Drug Development
- Introduction to Dose Response
 - Pairwise comparison and ANOVA
 - Model-Based comparison
 - Dose Response Models
- 3 Model Comparison Procedure and Modelling method
- Bayesian Model Comparison Procedure and Modelling method
- Simulation Study



Drug Development



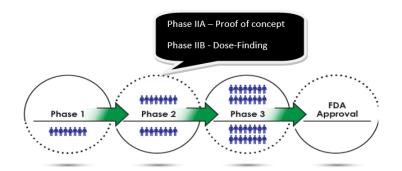


Figure: Drug Development



Drug Development



Why Phase II trials are important?

- Historically initially marketed drugs have later been recognized as excessive doses.
- It is believed by many that high attrition rate in Phase III is largely driven by inadequate dose selection.
- Selection of dose(s) to advance into the Phase III is one of the most challenging decisions during drug development.
- Dose response studies have a huge impact on dose selection process for the success of Phase III trial.
- It reduces the resources/cost/time.





Key Definitions



- Dose is the amount of drug administered to achieve a desired benefit (e.g., 500mg paracetamol).
- **Response** is an observation or effect shown by body to the drug.
- MED (Minimum Effective Dose) is the smallest dose above which a desired efficacy outcome is obtained.
- MTD (Maximum Tolerated Dose) is the dose beyond which the drug cannot be tolerated (due to high safety signals). Doses below the MTD are safe.
- Therapeutic window: If MED<MTD then the interval (MED, MTD) is called the Therapeutic window. Doses can be prescribed in this window and optimal dose(s), if any, should lie in this window.



Objectives



- To understand the underlying dose response relationship
- To identify MED, intermediate dose(s) and MTD.
- To find an optimal dose for Phase III that will be
 - high enough to be effective yet
 - low enough to have acceptable safety



ANOVA



ANOVA Test

- Step1: Testing the difference between means of different dose groups.
 - Hypotheses:
 - $H_0: \mu_1 = \mu_2 = \dots = \mu_4$
 - ullet $H_a:\mu_l
 eq\mu_m$, where μ_l and μ_m are two sample means out of all samples.
- Step2: Post hoc Tests
 - ullet If H_0 is rejected then proceed with pairwise comparison of mean response and compare it with adjusted critical value.



ANOVA



Limitations of ANOVA approach

- Doses are considered as qualitative factor in traditional ANOVA approach.
- This leads us selecting dose more or less than the actual dose required.
- It has a very vague idea about the shape of the dose response relationship.
- In Anova the means are joined by straight line. We can observe an un-smooth fit.
- For a better understanding of dose response relationship modelling approach is required in Dose Finding.



Model-Based comparison



- The modelling approach to dose finding is based on an assumed functional relationship between the response and the dose, treated as a quantitative variable.
- The model fitted is used to test for the presence of dose response and to estimate an adequate dose(s) to achieve a desired response



Dose Response Models



- Model: $f(d,\theta) = \theta_0 + \theta_1 f^0(d,\theta^0)$
- Emax Model: $E_0 + E_{\max} \frac{d}{(ED_{50} + d)}$
- Linear Model: $E_0 + \delta d$
- Exponential Model: $E_0 + E_1 \exp(\frac{d}{\delta})$
- Logistic Model:

$$E_0 + E_{max} \frac{1}{(1 + exp[\frac{ED_{50} - d}{\delta}])}$$

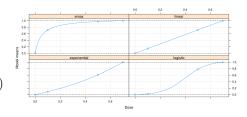


Figure: Dose Response Models

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- Multiple Comparison Procedure Modelling (MCP-Mod) Technique approach was proposed by Bretz, Pinheiro, and Branson (2005) and was approved by FDA.
- MCP-Mod combines both into single method
 - MCP Stage: Select best model(s) with appropriate contrasts controlling FWER from a class of candidate models.
 - Mod Stage: Model using best model and find target dose(s) of interest.







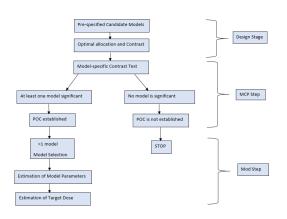


Figure: MCPMod - Overview

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July 1, 2022 13/32





- Response Y_{ij} is observed for a set of parallel groups of patients.
- k active doses groups: d_1,\ldots,d_k plus placebo dose group: $d_0\to k+1$ groups.

$$Y_{ij} = \mu_{d_i} + \epsilon_{ij}, \quad \epsilon_{ij} \stackrel{\text{i.i.d}}{\sim} N(0, \sigma^2) \quad i = 0, ..., k, j = 1, ... n_i$$

- ullet μ_{d_i} : is the mean dose response at different dose levels d_0,\dots,d_k
- ullet n_i cohort size of each dose group d_i
- ullet ϵ_{ij} : Error term; assumed to be independent and identically distributed.
- Mean dose response is represented as $u_{d_i} = f(d_i, \theta)$ for some dose-response model f(.).
- ullet θ : parameter vector of the unknown dose-response model f.

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- The main aim is to find the best-fitting model(s) from the set of candidate models.
- Null hypotheses

$$H_0^m:c_m'\boldsymbol{\mu}=0$$

for all $m \in \{1, 2, ..., M\}$ for some contrast c_m such that $\sum c_{mi} = 0$

Alternative Hypotheses

$$H_1^m: \boldsymbol{c}_m' \boldsymbol{\mu} \neq 0$$

for at least one $m \in \{1, 2, \dots, M\}$

ullet Single contrast test for testing the m^{th} model:

$$T_m = \frac{c'_m \overline{Y}}{\sqrt{S^2 \sum_{i=1}^k c_{mi}^2 / n_i}}, \quad m = 1, \dots, M$$







- ullet If H_1 is true for any model m then T_m follows non-central t-distribution with non-centrality parameter.
- A significance test for the above hypotheses testing:

$$T_{\mathsf{max}} = \max_{m \in \{1, 2, \dots, M\}} T_m.$$

- If $T_{max} \ge q$, a significant dose response signal is established.
- q is the adjusted critical value.
- If POC is not established, that is, no significant model => stop.
- If at least one model is significant, we choose the "best" model out the set of candidate models (this the model selection approach, the model averaging approach can also be followed).



Optimal Contrast



- Optimal contrast maximizes the chance of rejecting the associated null hypothesis, that is maximizes the non-centrality.
- Under balanced sample size allocation $n_i = n$, $i = 0, 1 \dots, k$

$$c_m = \frac{\boldsymbol{\mu} - \bar{\mu} \mathbf{1}}{||\boldsymbol{\mu} - \bar{\mu} \mathbf{1}||}$$
$$= \frac{\boldsymbol{\mu}^0 - \bar{\mu}^0 \mathbf{1}}{||\boldsymbol{\mu}^0 - \bar{\mu}^0 \mathbf{1}||}$$

ullet μ^0 is vector of standardized mean dose response.





Bayesian-MCPMod approach

$$Y_{ij} = \mu(d_i) + \epsilon_{ij}, \quad \epsilon_{ij} \stackrel{\text{i.i.d}}{\sim} N(0, \sigma^2) \quad i = 0, ..., k, j = 1, ... n_i$$

Prior is considered as normal distribution

$$oldsymbol{\mu} \sim \mathcal{N}_{K+1}\left(oldsymbol{ heta}_0, rac{\sigma^2}{oldsymbol{n}_0} I
ight)$$

ullet Posterior μ is

$$\mu_i \mid \mathcal{Y} \sim \mathcal{N}\left(\frac{\frac{\theta_{0,i}}{\sigma^2/n_{0,i}} + \frac{\bar{V}_i}{\sigma^2/n_i}}{\frac{n_{0,i}}{\sigma^2} + \frac{n_i}{\sigma^2}}, \frac{1}{\frac{n_{0,i}}{\sigma^2} + \frac{n_i}{\sigma^2}}\right) = \mathcal{N}\left(\frac{n_{0,i}\theta_{0,i} + n_i\bar{y}_i}{n_{0,i} + n_i}, \frac{\sigma^2}{n_{0,i} + n_i}\right)$$





- Placebo group: $\mu_{d_0} \sim w_1 N(\theta_{0,0,1},\sigma^2/n_{0,0,1}) + \dots + w_4 N(\theta_{0,0,4},\sigma^2/n_{0,0,4})$
- Active dose groups: one component conjugate normal prior

$$\boldsymbol{\mu} \sim w_1 \cdot \mathcal{N} \left(\begin{pmatrix} \theta_{0,0,1} \\ \theta_{0,1} \\ \vdots \\ \theta_{0,K} \end{pmatrix}, \begin{pmatrix} \frac{\sigma^2}{n_{0,0,1}} & \frac{\sigma^2}{n_{0,1}} & \cdots & \frac{\sigma^2}{n_{0,K}} \end{pmatrix} I \right) + \ldots +$$

$$w_L \cdot \mathcal{N} \left(\begin{pmatrix} \theta_{0,0,4} \\ \theta_{0,1} \\ \vdots \\ \theta_{0,K} \end{pmatrix}, \begin{pmatrix} \frac{\sigma^2}{n_{0,0,4}} & \frac{\sigma^2}{n_{0,1}} & \cdots & \frac{\sigma^2}{n_{0,K}} \end{pmatrix} I \right)$$

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$$\mu \mid \mathcal{Y} \sim w_{1}^{*} \mathcal{N} \begin{pmatrix} \frac{n_{0,0,1}\theta_{0,0,1} + n_{0}\bar{Y}_{0}}{n_{0,0,1} + n_{0}} \\ \frac{n_{0,1}\theta_{0,1} + n_{1}\bar{Y}_{1}}{n_{0,1} + n_{1}} \\ \vdots \\ \frac{n_{0,K}\theta_{0,K} + n_{K}\bar{Y}_{K}}{n_{0,K} + n_{K}} \end{pmatrix}, \begin{pmatrix} \frac{\sigma^{2}}{n_{0,0,1} + n_{0}} \cdots \frac{\sigma^{2}}{n_{0,K} + n_{K}} \end{pmatrix} I \\ + \dots + \begin{pmatrix} \begin{pmatrix} \frac{n_{0,0,4}\theta_{0,0,4} + n_{0}\bar{Y}_{0}}{n_{0,0,1} + n_{0}} \\ \frac{n_{0,1}\theta_{0,1} + n_{1}\bar{Y}_{1}}{n_{0,1} + n_{1}} \\ \vdots \\ \frac{n_{0,K}\theta_{0,K} + n_{K}\bar{Y}_{K}}{n_{0,K} + n_{K}\bar{Y}_{K}} \end{pmatrix}, \begin{pmatrix} \frac{\sigma^{2}}{n_{0,0,4} + n_{0}} \cdots \frac{\sigma^{2}}{n_{0,K} + n_{K}} \end{pmatrix} I \\ \vdots \\ \frac{n_{0,K}\theta_{0,K} + n_{K}\bar{Y}_{K}}{n_{0,K} + n_{K}\bar{Y}_{K}} \end{pmatrix}$$

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$$\mathbf{c}_{m}^{T}\boldsymbol{\mu} \mid \mathcal{Y} \sim \underbrace{w_{1}^{*}\mathcal{N}\left(\mathbf{c}_{m}^{T}\boldsymbol{\theta}_{\mathcal{Y}}^{1}, \sum_{i=0}^{K} c_{m,i}^{2}\tau_{1}[i,i]\right)}_{\sim Z_{1}} + \cdots + \underbrace{w_{4}^{*}\mathcal{N}\left(\mathbf{c}_{m}^{T}\boldsymbol{\theta}_{\mathcal{Y}}^{4}, \sum_{i=0}^{K} c_{m,i}^{2}\tau_{4}[i,i]\right)}_{\sim Z_{4}}$$

$$P\left(\mathbf{c}_{m}^{T}\boldsymbol{\theta} > 0 \mid \mathcal{Y}\right) = \sum_{l=1}^{4} w_{l}^{*} \cdot P\left(Z_{l} > 0\right),$$

$$= \sum_{l=1}^{4} w_{l}^{*}\boldsymbol{\Phi}\left(\frac{\mathbf{c}_{m}^{T}\boldsymbol{\theta}_{y}^{l}}{\sqrt{\sum_{i=0}^{K} c_{m,i}^{2}\tau_{l}^{2}[i,i]}}\right) = p_{m}^{y}.$$

Significant test is established if $p^* = \max_{m=1,...,M} p_m^y$ is greater than an adjusted critical value $1 - \alpha^*$.

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Example:

- k = 4 active dose groups and a placebo group: $d_0 = 0, d_1 = \frac{1}{20}, d_2 = \frac{3}{20}, d_3 = \frac{10}{20}, d_4 = 1$
- Standard deviation $\sigma = 1.1$
- ullet M=6 candidate models
 - Three different Emax models
 - One Exponential model
 - One Linear model
 - One Logistic model
- $\alpha = 0.05$
- Overall sample size n=300





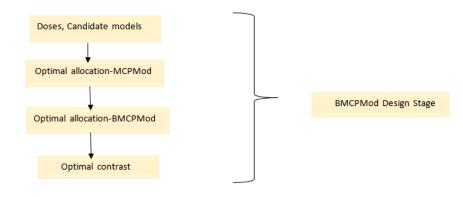


Figure: BMCPMod Design Stage





Prior Information:

Placebo group has three different conjugate priors

prior	weights	$Mean(heta_{0,0,i})$	$ESS(n_{0,0,i})$	ESS_{elir}
informative	0.5	0	30	9.5
weakly informative 1	0.25	0.25	2	
weakly informative 2	0.25	-0.25	2	

Table: Placebo prior information

- We consider vaguely informative prior for active groups.
- active dose groups: prior mean: $\theta_{0,i} = 0$, ESS: $n_{0,i} = 1$





Expected local-information-ratio effective sample size

- $ESS_{elir} = E_{\mu}(\frac{i(p(\mu))}{i_F(\mu)})$
- $i_F(\mu)=-\sum_i n_i E(\frac{\partial^2}{\partial \mu^2} log f(y_{i1}|\mu))$ is the Fisher information of the sampling model
- $i(p(\mu)) = -\frac{d^2 \log p(\mu)}{d\mu^2}$ is the prior distribution of $\mu.$
- If the prior is a k components mixture distribution, $p(\mu) = \sum_{j=1}^K w_j p_j(\mu)$
- \bullet ESS_{elir} is calculated using function ess of RBesT package in R.



Optimal allocation



• Experimental optimal design is represented as

$$\xi = \{(d_1 \cdots d_k), (w_1 \cdots w_k)\}\$$

- Fishers Information matrix is $M(\xi,\theta) = \sum_{i=1}^k w_i \frac{\partial f(d,\theta)}{\partial \theta} \frac{\partial f(d,\theta)}{\partial \theta} \frac{\partial f(d,\theta)}{\partial \theta}$
- D-optimal Design using General Equivalence Theorem (GET)
 - ullet Design maximizes $log\{\det[oldsymbol{M}(\xi)]\}$
 - $d(x,\xi) = f(x)^T [M(\xi)]^{-1} f(x) \le p$
- Optimal allocation is calculated using function optDesign of DoseFinding package in R.





• Optimal allocation for MCPMod:

Dose	0	$\frac{1}{30}$	$\frac{3}{30}$	$\frac{10}{30}$	1
N=300	80	33	44	48	95

• Optimal allocation for BMCPMod:

Dose	0	$\frac{1}{30}$	$\frac{3}{30}$	$\frac{10}{30}$	1
N=300	72	34	46	50	98



Optimal Contrast



Optimal contrast for unbalanced sample size:

$$oldsymbol{c}_m \propto oldsymbol{S}^{-1} \left(oldsymbol{\mu}_m^0 - rac{oldsymbol{\mu}_m^0{\,}'oldsymbol{S}^{-1} oldsymbol{1}}{1'oldsymbol{S}^{-1} oldsymbol{1}}
ight)$$

- \bullet μ_m^0 is the vector of standardized mean dose response.
- $S = diag(n_0, n_1, \dots, n_k)$ is the matrix of inverse of the cohort sizes.
- We normalize the optimal contrast so that $||c_m||=1$.





Optimal contrast for BMCPMod allocation:

	Emax1	Emax2	Emax3	exponential	logistic	linear
0	-0.685	-0.837	-0.613	-0.379	-0.499	-0.425
0.03	-0.182	-0.006	-0.204	-0.172	-0.229	-0.186
0.1	-0.021	0.142	-0.094	-0.206	-0.262	-0.201
0.33	0.215	0.221	0.174	-0.120	0.231	-0.050
1	0.672	0.480	0.737	0.877	0.759	0.861

Table: BMCPMod Optimal Contrast





ullet The posterior probabilities for M=6 models

Models	posterior probabilities
Emax1	0.3107429
Emax2	0.4103961
Emax3	0.2813772
Exponential	0.2122869
Linear	0.2528265
Logistic	0.2225135

Table: Posterior Probabilities

- Adjusted critical value is 0.14823
- 0.4103961 > 0.14823 POC is established.



References



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Thank You