

Analysis of in vivo responses by mixed-effect models reference

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Objectives. In *in vivo* experimentation, the large intra-group variability between animals is a major obstacle that prevents to **detect significant**. therapeutic effects of treatment factors. Our objective is to assess a new statistical method able to better estimate and characterize the additive effects of the combination of an oncolytic virus (TG6002) and the prodrug flucytosine (5-FC) associated in an anti-cancer treatment. The experimental data are kinetics of tumor growth collected during in vivo assays carried out on mice.

$$y_{ij} = x(t_{ij}, \theta_{\mathbf{i}}) + \epsilon_{\mathbf{ij}}, \quad \forall \mathbf{i} = \mathbf{1}, \dots, \mathbf{r}, \forall \mathbf{j} = \mathbf{1}$$
$$x(t_{ij}, \theta_{\mathbf{i}}) = k_{1,i}(t_{ij} + t_0) + x_{0,i} + k_{2,i}(t_{ij} + \theta_{\mathbf{i}}) = \lambda + \mathbf{c}_{\mathbf{i}}\beta + \theta_{\mathbf{i}}, \ \theta_{\mathbf{i}} \sim \mathcal{N}(\mathbf{0}, \mathbf{\Omega}), \quad \forall \mathbf{i} = \mathbf{1},$$
$$\theta_{\mathbf{i}} = (x_{0,i} \quad k_{1,i} \quad k_{2,i} \quad \tau_i)^{\mathbf{T}}$$
$$\epsilon_{ij} = \sigma \ e_{ij}, \ e_{ij} \stackrel{i.i.d.}{\sim} \mathcal{N}(0,1), \quad \forall i = 1, \dots, \tau_i$$
$$c_i = (c_{c,i} \quad c_{d,i} \quad c_{0,i})$$

$x_{0,i}$	Initial diameter	C_{C}	TG6002 0
$k_{1,i}$	First growth rate	c_d	5-FC pres
$k_{2,i}$	Second growth rate	c_O	Initial dia
$ au_i^{-,\cdot}$			

Our model structure fits correctly all the 104 observed growth responses. The confidence on the estimation results is such that we can detect two tratement effects. Indeed, we show a 3% reduction of the therapeutic response time due to 5-FC and a division by five of the growth delay with TG6002.

Results confirm the relevance of mixed-effect kinetic models to increase the power of statistical tests applied to in vivo studies and efficacy of the combination TG6002 with 5-FC.

Reference

T. Bastogne, A. Samson, P. Vallois, S. Wantz-Mézières, S. Pinel, D. Bechet, and M. Barberi-Heyob. Phenomenological modeling of tumor diameter growth based on a mixed effects model. Journal of Theoretical Biology, 262:544-552, 2010.



Analysis of in vivo Responses by Mixed-Effect Models L. BATISTA^{1,3,4}, B. BASTIEN², T. BASTOGNE^{1,3,4}, J. FOLOPPE², P. ERBS² ¹CRAN, ²TRANSGENE, ³INRIA BIGS, ⁴CYBERnano

Method

- $\mathbf{L}, \ldots, \mathbf{n_i},$
- $t_0 \tau_i$
- $\ldots, \mathbf{r}, \ldots$
- concentration factor esence factor ameter variation

The experimental set up is decomposed into 4 main steps:

- 1. a full factorial design of experiments is proposed. TG6002 was tested at four concentrations in combination or not with 5-FC. Eight groups of 13 mice are randomly affected to each experimental condition.
- $r, \forall j=1,\ldots,n_i,$ 2. we propose a mixed-effect model to describe the kinetic growth of the mean tumor diameter.
 - 3. the model parameters are determined with a **maximum likelihood estimato**r based on an expectation-maximization algorithm.
 - 4. effects of the two examined components are assessed with a **Wald test**.

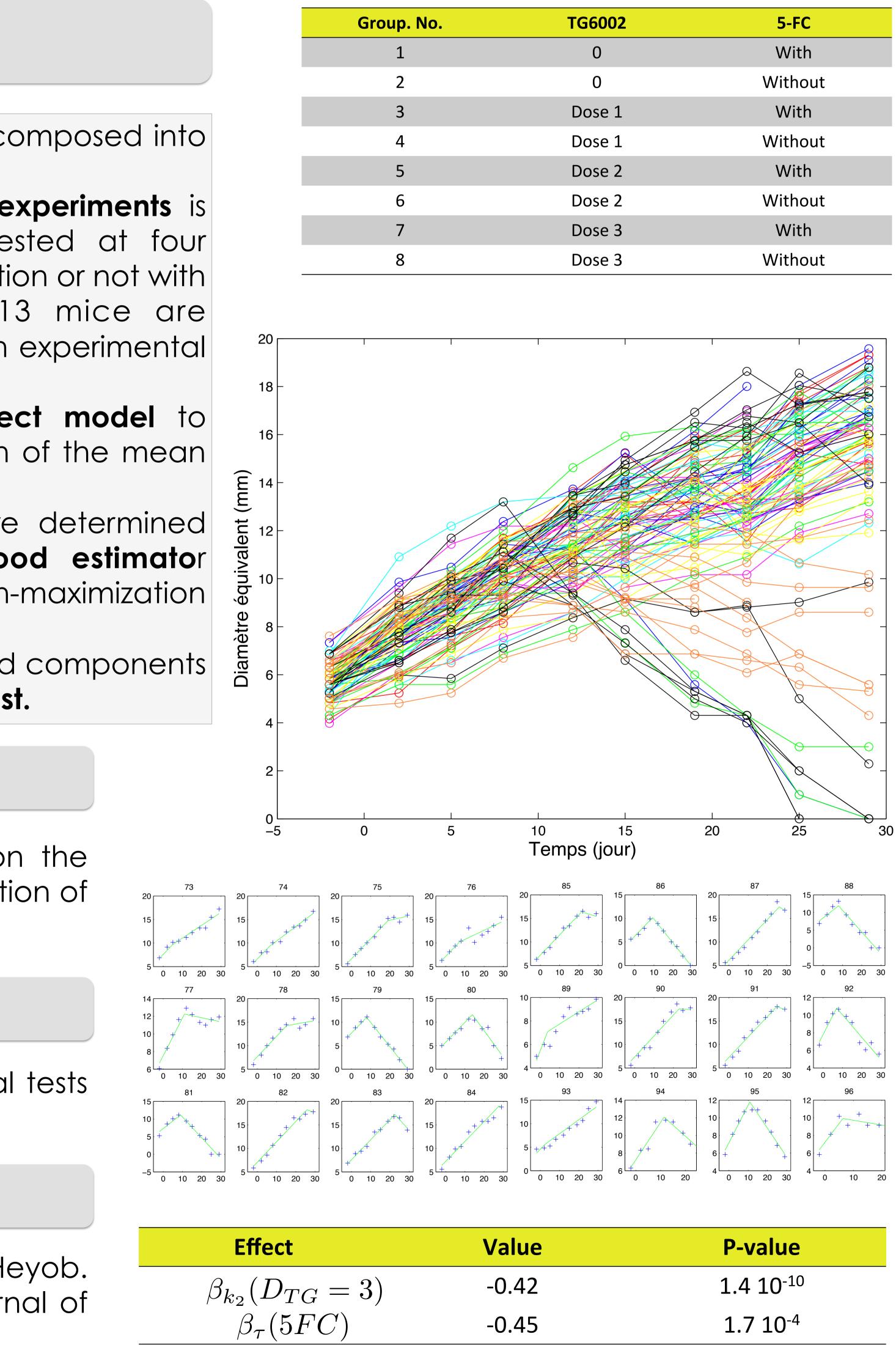
Results

Conclusion

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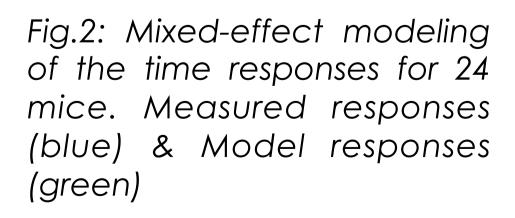




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Fig.1: Longitudinal Data: growth kinetics of the tumor equivalent diameter (mm) for 104 mice distributed between 8 groups of treatment: group i (magenta), group 2 (red), group 3 (cyan), group 4 (blue), group 5 (yellow), group 6 (green), group 7 (orange), group 8 (black)



Tab.2: Estimated effects of the therapeutic factors on the post-treatment growth rate (k_2) and response delay (τ).Only the 2 most significant effects are indicated here.