### A short teaser

#### MODELING with STOCHASTIC DIFFERENTIAL EQUATIONS and MIXED EFFECTS

Workshop on Brain Dynamics and Statistics: Simulation versus Data, (26/02/2017 - 03/03/2017, Banff, Alberta)

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#### Introduction

Differential equations are a frequently used tool for modeling the generating mechanisms behind longitudinal data. Often, the dynamics are complex and require multidimensional nonlinear models with several unknown parameters. Moreover, especially in biological systems, certain parameter values may vary across subjects and subjectspecific models are asked for. Subject-wise parameter estimation can however, be highly inefficient and unreliable if not enough subject-specific data is available (not enough repeated measurements). Thus, inference methods will lack sufficient power. The methodology of random effects allows data pooling and thereby facilitates more reliable estimation, while maintaining subject-specificity in modeling. The inclusion of stochasticity into the model itself, i.e., using stochastic differential equations (SDEs), adds further to the robustness of parameter inference, since the added system noise accounts for the (inherent) model uncertainty. We propose an approach for maximum-likelihood estimation in SDE models that include mixed effects (i.e., random and fixed effects). A striking (computational) benefit of the suggested approach is that the likelihood is explicitly available provided the parameters enter the drift term linearly. The model may however still be nonlinear in the state. This specific model class comprises a wide range of well-known statistical models (see below of some examples) and thus opens up for various applications, in particular in biology

Covariates may be included as well to adjust for different experimental conditions or subject-specific information, such as age or gender.

### Model

N independent r-dim. processes  $X^i = (X^i_t)_{0 \leq t \leq T^i}, i = 1, \ldots, N,$  governed by the SDE

#### Applications

#### Modeling the neural excitability

The Fitzhugh-Nagumo model is a two-dimensional approximation of the four-dimensional Hodghin-Hudely equations and often applied to model the regenerative fining mechanism in an excitable neuron. Neural fining is complex interplay of numerous cell processes and to account for various unexplained noise sources, a stochastic FINM model can be considered (Ansen et al., 2012). Y represents the membrane potential of a neuron and Z the recovery.

		N = 20		N = 50		N = 100	
true value		rel, bias	RMSE	rel, bias	RMSE	rel, bias	RMS
e	0.10	0.002	0.035	0.003	0.022	0.003	0.01
5	0.50	-0.004	0.057	0.001	0.033	0.001	0.02
$1/\epsilon$	10.00	-0.001	0.352	-0.003	0.216	-0.003	0.15
sle	5.00	-0.006	0.240	-0.002	0.135	-0.002	0.10
2	1.50	-0.000	0.050	-0.000	0.031	-0.001	0.02
η	1.20	-0.001	0.051	-0.001	0.031	0.000	0.02
	2.25	-0.065	0.731	-0.048	0.469	-0.017	0.34
diag({\black})	1.00	-0.053	0.312	-0.025	0.197	-0.013	0.14
	0.04	-0.042	0.016	-0.035	0.010	-0.012	0.00
	0.04	-0.064	0.015	-0.007	0.010	-0.007	0.00

 $dY_t = \frac{1}{\varepsilon} (Y_t - Y_t^3 - Z_t + s) dt + \sigma_1 dW_{1,t},$  $dZ_t = (\gamma Y_t - Z_t + \eta) dt + \sigma_2 dW_{2,t}.$ 

After the re-parametrization  $\mu=(1/\epsilon,s/\epsilon,\gamma,\eta),$  the model for neuron i can be written as

$$d\begin{pmatrix} Y_t^i \\ Z_t^i \end{pmatrix} = \begin{bmatrix} \begin{pmatrix} 0 \\ Z_t^i \end{pmatrix} + \begin{pmatrix} Y_t^{-i} - (Y_t^{-i})^3 - Z_t^i & 1 & 0 & 0 \\ 0 & 0 & Y_t^i & -1 \end{pmatrix} \end{bmatrix} (\mu + \phi^i) dt + \Sigma dW_t^i$$

with random effects  $\phi^i \sim \mathcal{N}(0, \Omega)$  and unknown  $\Omega$ .



Trace plots of four realizations of the stochastic FHN model with random effects. The corresponding (rounded) realized parameter values of  $\mu + \phi'$  are (8.86,4.29,1.50,1.39), (10.16,7.49,1.73,1.40), (9.96,5.49),1.10,107), (9.26,5.17,1.84,1.01).

### Take two popular modeling frameworks...



Showing: Number of publications by year (data accessed in early July 2016).

## $\dots$ and combine them

Data: Observations of N independent r-dim. processes  $(X_t^i)_{0 \le t \le T^i}$ .

Data model:

(old) Stochastic differential equation (with fixed effects)

 $dX^i_t = F(t, X^i_t, \mu) dt + \Sigma(t, X^i_t) dW^i_t \ \rightsquigarrow \ \text{ estimate } \mu$ 

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Actually, we can also include some covariate information  $(D_t^i)$ ...:

$$dX_t^i = F(t, X_t^i, D_t^i, \mu, \phi^i) dt + \Sigma(t, X_t^i) dW_t^i$$

## Two key reasons for increased popularity

### (1) Both frameworks very useful - their combination all the more





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SDEs capture model insufficiencies, random effects enable pooling while still being subject-specific.

(2) Both frameworks challenging - their combination all the more Goal: obtain the MLE  $\hat{\theta}_N = \arg \max_{\theta} L(\mathbf{X}; \theta)$   $\theta = (\beta, \vartheta)$ 

**Likelihood:**  $L(\mathbf{X}; \theta) = \prod_{i=1}^{N} L(X^i; \theta) = \prod_{i=1}^{N} \int L(X^i; \beta, \phi^i) p(\phi^i; \vartheta) d\phi^i$ 

intractability  $\cdot$  intractability = (intractability)<sup>2</sup>

# Good news for numerous applications

- In certain cases, the likelihood is **explicit**.
- Dynamics can be **non-linear** in the state.
- We can even include **covariate** information (gender, age, exp. condition, ...).
- Many well-known models fall into this certain class.
- The MLE is consistent and asymptotically normal (T fix, N large).
- We can do hypothesis testing.

# Possible applications to neuroscience

# (a) EEG data

- $\rightarrow\,$  avoid averaging over trials and/or subjects
- $\rightarrow\,$  keep "pooling", but model inter-trial / inter-subject variability with random effects

Data N subjects, J trials each,  $X_t^i = (X_t^{i,1}, \dots, X_t^{i,J})' \in \mathbb{R}^{r \cdot J}$ ,

$$dX_t^i = F(t, X_t^i, D_t^i, \mu, \phi^i) dt + \Sigma(X_t^i) dW_t^i$$

## (b) Stochastic FHN model with mixed effects

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Hope to see you at my poster!

