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I. Background

Emotion specificity of physiological activation has been long debated in psychological research (Kreibig, 2010; Levenson, 1992)

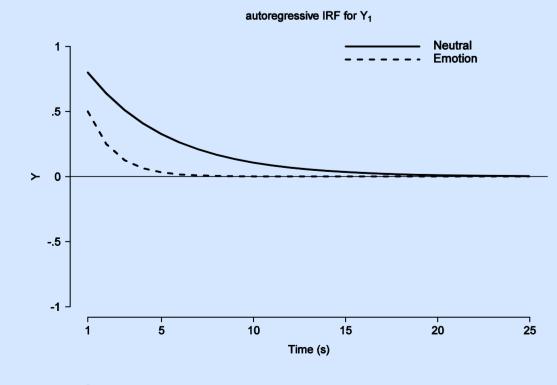
Five theoretical models can be distinguished:

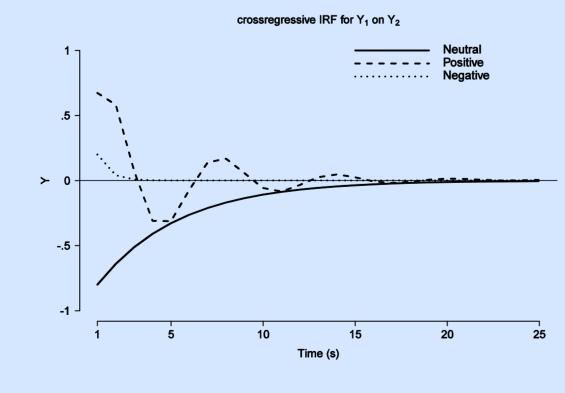
- 1. Null model = non-emotional (neutral) and emotional physiology are indistinguishable
- 2. Emotion model = physiological activation is different for non-emotional and emotional episodes
- 3. Valence model = physiological activation is different for positive and negative emotional episodes
- 4. Avoidance-approach model = physiological activation is different for approach emotions and avoidance emotions
- 5. Specificity model = physiological activation is specific for each emotion

II. Research Questions

Dimensions of specificity:

- ✓ A first dimension describes the **degree of emotional distinction**, as explained above in the five theoretical models
- ✓ A second dimension concerns the aspects of the data which are affected by the specificity
 - Most often, one looks merely at mean physiological activation
 - We extend this dimension to specificity in the mean, variance and dynamics of process (the latter is interesting in the context of emotional inertia; Kuppens et al., 2010)
 - Dynamics can be visualized with impulse response functions:





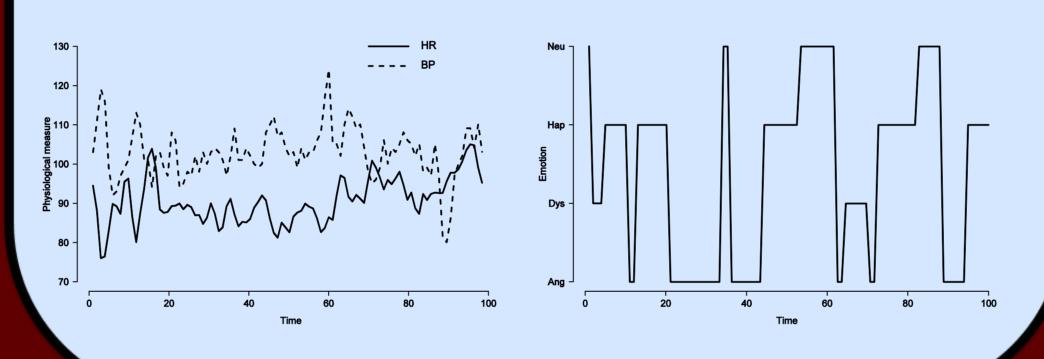
Research questions:

- ✓ **Dimension 1:** Which theoretical model on specificity in physiology is dominant? Is this different for depressed and healthy subjects?
- ✓ **Dimension 2:** On what aspects of the data can we interpret specificity? Mean, variance or dynamics?

III. Data Collection

Oregon adolescent interaction data

- ✓ Conducted at Oregon Research Institute by Nicholas Allen & Lisa Sheeber
- √ 33 depressed + 24 healthy adolescents
- ✓ Real-life social interaction task with parents
- ✓ Second-to-second measurements
- Cardiovascular: heart rate (HR) and blood pressure (BP)
- Behavioral: LIFE coding system was applied to code verbal & non-verbal behavior with emotions {Neutral, Happy, Dysphoric, Angry}
- ✓ Example data:



IV. Analysis

General approach requires advanced analysis using time series analysis and model selection

State space modeling:

- ✓ Estimation of latent states of the observed processes based on Bayesian Kalman filter
- ✓ Analysis of dynamic properties of states
- Extended with regime switching component, which allows one to estimate different sets of parameters for different time episodes (e.g., emotion episodes)

Deviance information criterion (DIC):

- ✓ Bayesian model selection criterion to select the optimal model among various estimated models
- ✓ Comparable to AIC and BIC: model fit + model complexity penalty

V. Question 1

"Which theoretical model on specificity in physiology is dominant? Is this different for depressed and healthy subjects?"

- ✓ For each individual, five models were estimated and compared
- ✓ Each model was characterized with a different configuration of regimes
- ✓ The best model was selected for each individual, based on the value of DIC
- ✓ The **null model is most popular**, rarely more complex models chosen
- ✓ Finding is very similar for healthy controls and depressed adolescents

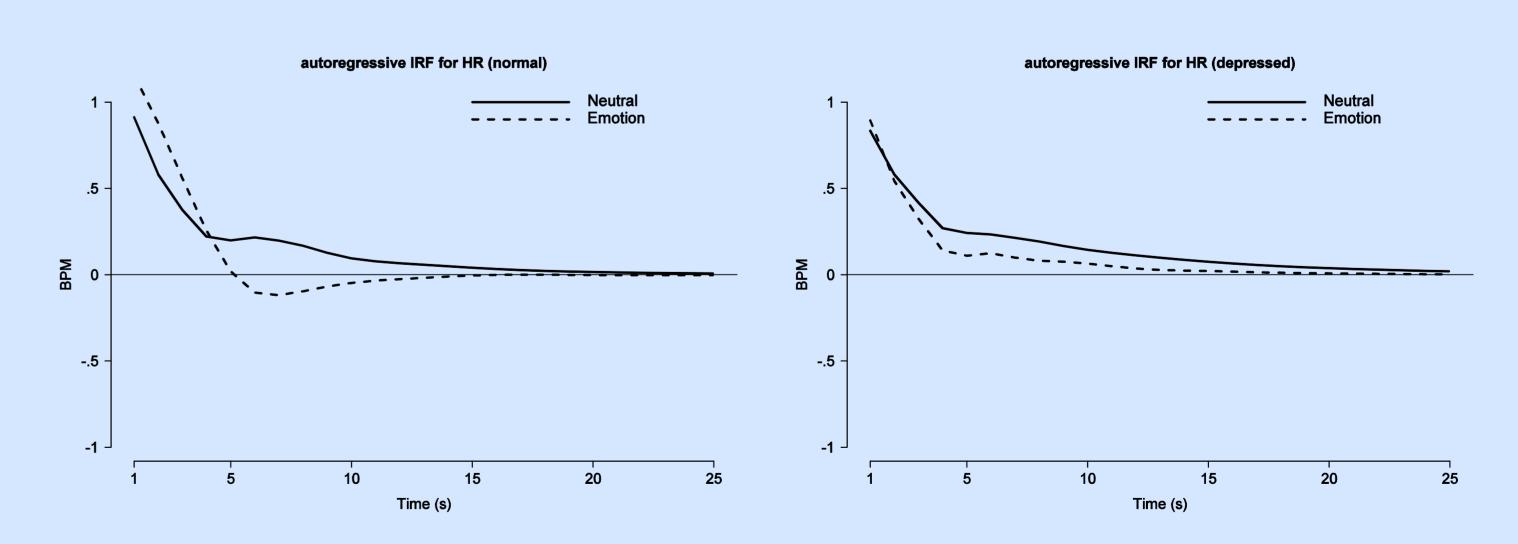
	Neutral	Нарру	Angry	Dysphoric	% controls	% depressed
M1: Null model					.96	.85
M2: Emotion model					.04	.09
M3: Valence model					0	0
M4: Approach/avoidance model					0	.06
M5: Specificity model					0	0

VI. Question 2

"On what aspects of the data can we interpret specificity?

Mean, variance or dynamics?"

✓ Study of cases is difficult, as there are few cases for which data implies specificity and no clear general patterns of specificity could be identified ✓ Illustrative comparison of HR impulse response functions for a healthy control and a depressed adolescent for the selected emotion model:



✓ Future challenge

- Narrow down by comparing models that estimate specificity for only means, variances or dynamics
- Identifying explanatory variables that are related to presence or absence of specificity