# Mixed Effects Models: Concepts and Applications in Neurolmaging

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### Outline

### Concepts

- The standard fixed effects model
- What is a mixed effects model, and why do we need it?
- For simplicity, we will not address the mathematics of parameter estimation, hypothesis testing, or statistical power analysis
  - See the neuroimaging paper referenced in the next slide if you are interested

### Applications

Mixed effects models in neuroimaging research

### References

### Concepts

 A basic tutorial I recommend as an introduction to mixed effects models http://www.bodowinter.com/tutorial/bw\_LME\_tutorial2.pdf

### Applications

• Bernal-Rusiel 2013 – Statistical analysis of longitudinal neuroimage data with Linear Mixed Effects models

https://doi.org/10.1016/j.neuroimage.2012.10.065

# Concepts

# The standard fixed effects model

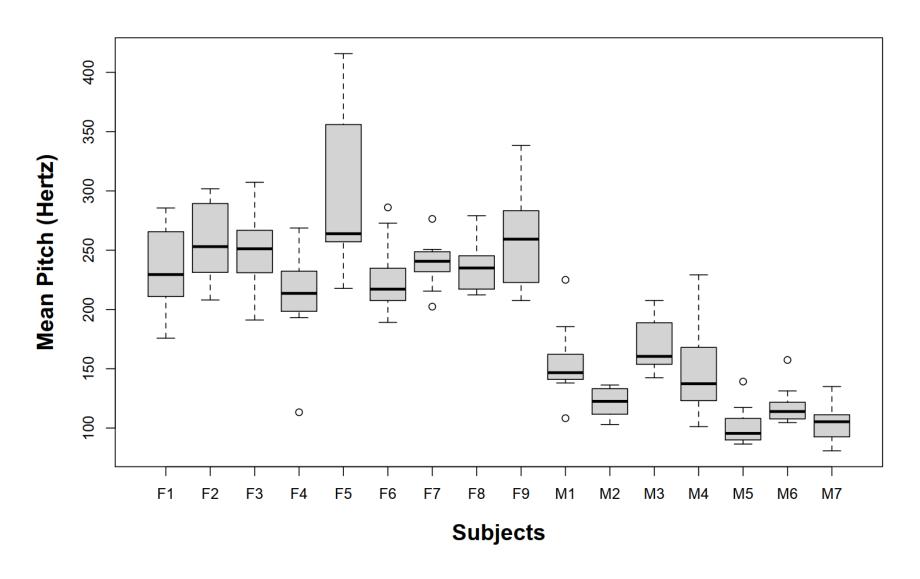
- Consider this simple experiment
  - Have a group of participants read a passage out loud, and measure their vocal pitch
  - Construct a model to predict pitch from participant demographics
- Suppose you use a linear model to predict vocal pitch from age and sex
  - $pitch \sim age + sex + \epsilon$ 
    - Age and sex are the *fixed effects* → these are your familiar independent variables
    - $\epsilon$  is the error term it represents random factors that cannot be controlled experimentally
      - Ex: hours of sleep the night before the study, nervousness during experiment, etc...
- If each study participant was measured once, then standard statistical models (ex: linear regression) could be used
  - This is because the assumption of independence between observations holds

# Motivating the mixed effects model

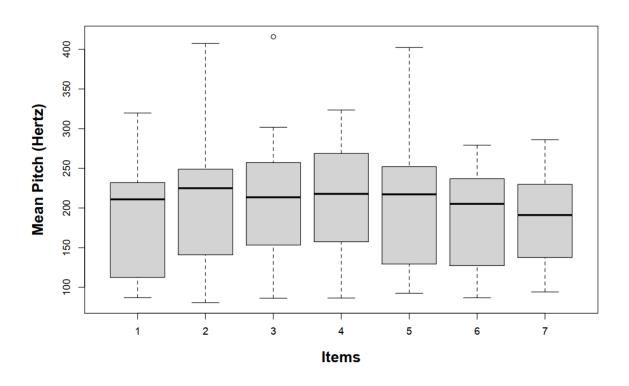
- But what if...
  - Scenario 1: each participant's pitch is measured multiple times?
    - Independence between observations no longer holds → measurements are correlated
  - Scenario 2: participants were assigned different-themed passages, as the researchers were interested in the effect of theme on pitch?
    - Passages are just a subset of possible passages, therefore modelling them as fixed effects is unnatural

- We can account for these using a mixed effects model
  - Mixed because we introduce random effects into our fixed effects model

# Motivating the mixed effects model



# Motivating the mixed effects model



# The mixed effects model

- Each participant has a different baseline pitch, which is known as a random intercept
  - Now multiple measurements from a single participant can be accounted for according to their baseline pitch
- Each passage is given a random intercept as well
  - This is done because different passages might elicit different responses, which might affect pitch
- Now suppose the effect of sex on pitch depends on the passage that is being read
  - We can model this elegantly using *random slopes*
  - *Note*: in general if you are unsure of whether to include a random effect in your model, use the likelihood ratio test to compare the likelihood of nested models

# The mixed effects model

- The new model is
  - $pitch \sim age + sex + (1|subject) + (1 + sex|passage) + \epsilon$ 
    - Age and sex are included as before
    - (1|subject) contributes the random intercept for each subject
    - (1 + sex|passage) contributes a random intercept for each passage, as well as a random slope to model the effect of sex depending on the passage

# Fundamental difference between fixed and random effects

#### Fixed effects

- Variation in the factor can be captured/exhausted
  - Ex: sex only M or F, so we can exhaust the variability in this factor

#### Random effects

- Measurements obtained are just a subset of the possible measurements we could have obtained
  - Ex: participants included in the study are just a subset of all the participants that could have been included
  - Ex: passages used are just a subset of the passages that could have been used

# Applications in Neurolmaging

# Mixed effects models for longitudinal data – key differences

- Time from baseline may be included as an effect in the model
  - This would be the case if time is thought to play an important role in the outcome of interest
    - Ex: Brain atrophy in Alzheimer's patients
  - Here time can be a fixed or random effect, depending on whether rate of atrophy is patient-dependent

 Must also account for variable timing of measurements between patients, as well as subject dropout (causing unbalanced data)

# The experiment

- Measure longitudinal hippocampal volume and entorhinal cortex thickness in a dataset consisting of
  - Alzheimer's patients
  - Subjects with mild cognitive impairment
  - Healthy controls

# Specification of fixed and random effects

#### Fixed effects

- Time from baseline
- Clinical group indicator
- Interaction between time from baseline and clinical group indicator
- Baseline age
- Sex
- APOE genotype status
- Interaction between APOE genotype status and time from baseline
- Education (in years)

#### Random effects

- Included based on the likelihood ratio test
- Random intercept and time slopes were included for all variables

### Dataset

**Table 1**Longitudinal ADNI sample characteristics.

Variable	Stable HC	Converter HC	Stable MCI	Converter MCI	AD	p-value
Number of subjects	210	17	227	166	188	
Baseline age	$75.9 \pm 5$ [60–90]	$76.7 \pm 5.1$ [63–84]	$74.8 \pm 7.7$ [55–90]	$74.7 \pm 7.1$ [55–89]	$75.2 \pm 7.5$ [55–91]	0.3464
Female %	48.1	47.1	33.48	38.6	47.3	<0.01 <sup>a</sup>
APOE-ε4 Carriers %	25.7	41.2	43.2	67.5	66	<0.0001 <sup>a</sup>
Education	$16.1 \pm 2.8$ $[6-20]$	$16.1 \pm 2.8$ [12–20]	$15.6 \pm 3.1$ [4–20]	$15.7 \pm 2.9$ [6–20]	$14.7 \pm 3.2$ [4–20]	<0.001

Baseline age (in years) and education values are in mean  $\pm$  standard deviation; Ranges are listed in square brackets; p-values indicate effects across the groups.

Key: Converter MCI, mild cognitive impairment subjects who convert to Alzheimer's disease; Converter HC, healthy controls who convert to either MCI or Alzheimer's disease.

<sup>&</sup>lt;sup>a</sup> Using Fisher's exact test; ANOVA-derived p-values were used in the other cases.

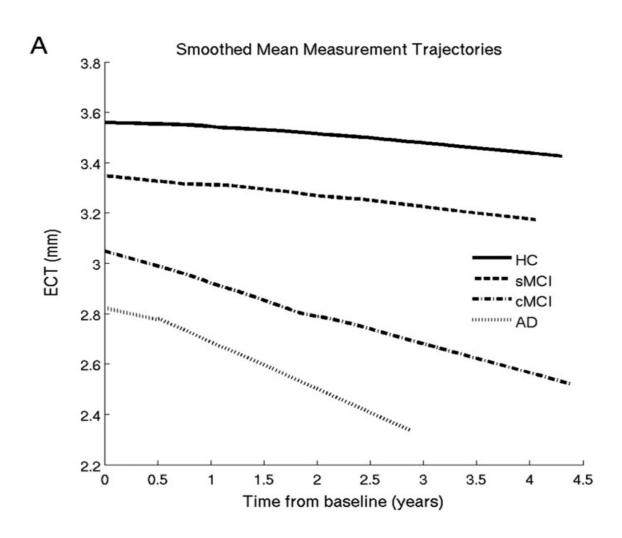
# Dataset

**Table 2** Number and timing of scans per time point by clinical group (Stable HC, N=210; Converter HC, N=17; Stable MCI, N=227; Converter MCI, N=166; AD, N=188).

Time point	Stable HC	Converter HC	Stable MCI	Converter MCI	AD	Time from baseline
Baseline	210	17	227	166	188	0
Year 0.5	197	17	194	161	166	$0.58 \pm 0.07  [0.21  0.94]$
(month 6)						
Year 1	183	17	177	153	150	$1.08 \pm 0.07  [0.68 - 1.38]$
Year 1.5	0	0	153	136	0	$1.59 \pm 0.08$ [1.26–1.92]
Year 2	129	14	108	106	96	$2.09 \pm 0.10$ [1.58-2.88]
Year 3	115	6	68	70	0	$3.09 \pm 0.09$ [2.52–3.45]
Year 4	11	0	3	10	0	$4.12 \pm 0.09$ [3.98–4.38]
Total	845	71	930	802	600	

Time from baseline (in years) is in mean  $\pm$  standard deviation; ranges are listed in square brackets.

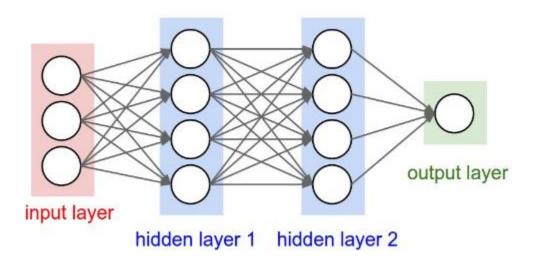
# Results



# Looking Ahead

# Neural Networks – A Possible Solution?

- Put simply, feedforward neural networks find a function between inputs and outputs
  - Ex: age, sex, clinical features to predict disease

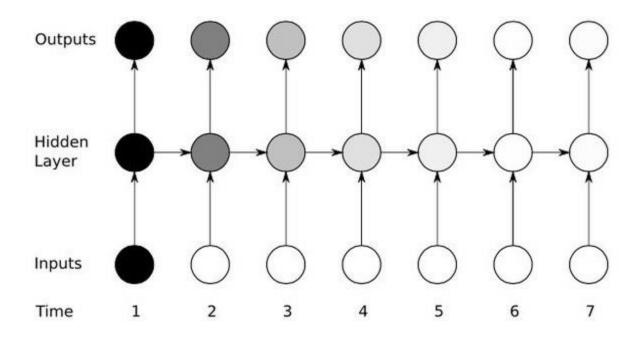


# Neural Networks — A Possible Solution?

 For longitudinal data, a similar problem arises for feedforward neural networks as with fixed effects models

- An alternative neural network architecture is the recurrent neural network (RNN)
  - This can capture the time-dependencies between measurements

# Recurrent Neural Networks



# Recurrent Neural Networks

