



# Functional Connectivity Patterns in Monolingual and Bilingual Infants

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fNIRS 2018, Tokyo

# Motivation I: Early bilingual exposure

- Experience-related effects in brain structure and function.
- Evidence for **bilingual adaptation**.

Li et al., 2014 *Cortex (Review)*

- Bilingual adaptation in **infants**.

- **Resting state FC?**

Molnar et al., 2014 *Infancy*

Ferjan-Ramirez et al., 2017 *Developmental Sci.*

Costa and Sebastián-Gallés, 2014 *Nature Rev. Neurosci. (Review)*

- Resting state FC in infants.

- Premature birth
  - Socio-economic status
  - Autism

Smyser et al., 2010 *Cereb. Cortex (fMRI)*

Gao et al., 2015a *Cereb. Cortex (fMRI)*

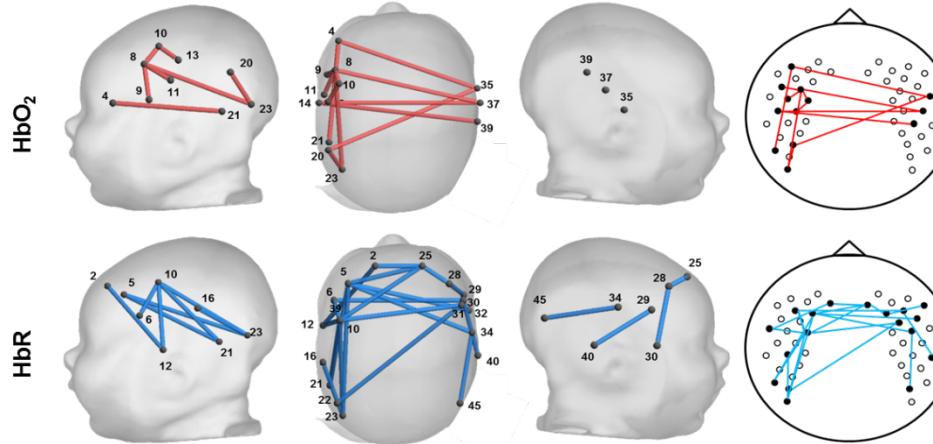
Keehn et al., 2013 *Front. Hum. Neurosci (fNIRS)*

Gao et al., 2017 *Neuroscientist (Review)*

# Motivation II: Quality assessment (QA)

- Assess the **reliability** of our previous **fNIRS** resting state FC study in **awake** 4-month old monolingual and bilingual infants.

Spanish–Basque Bilinguals > Spanish Monolinguals



- We suspect that these results might be driven by motion artifacts that were not accurately identified and removed.

# The current work: Resting state FC in asleep 4-month-old monolingual and bilingual infants

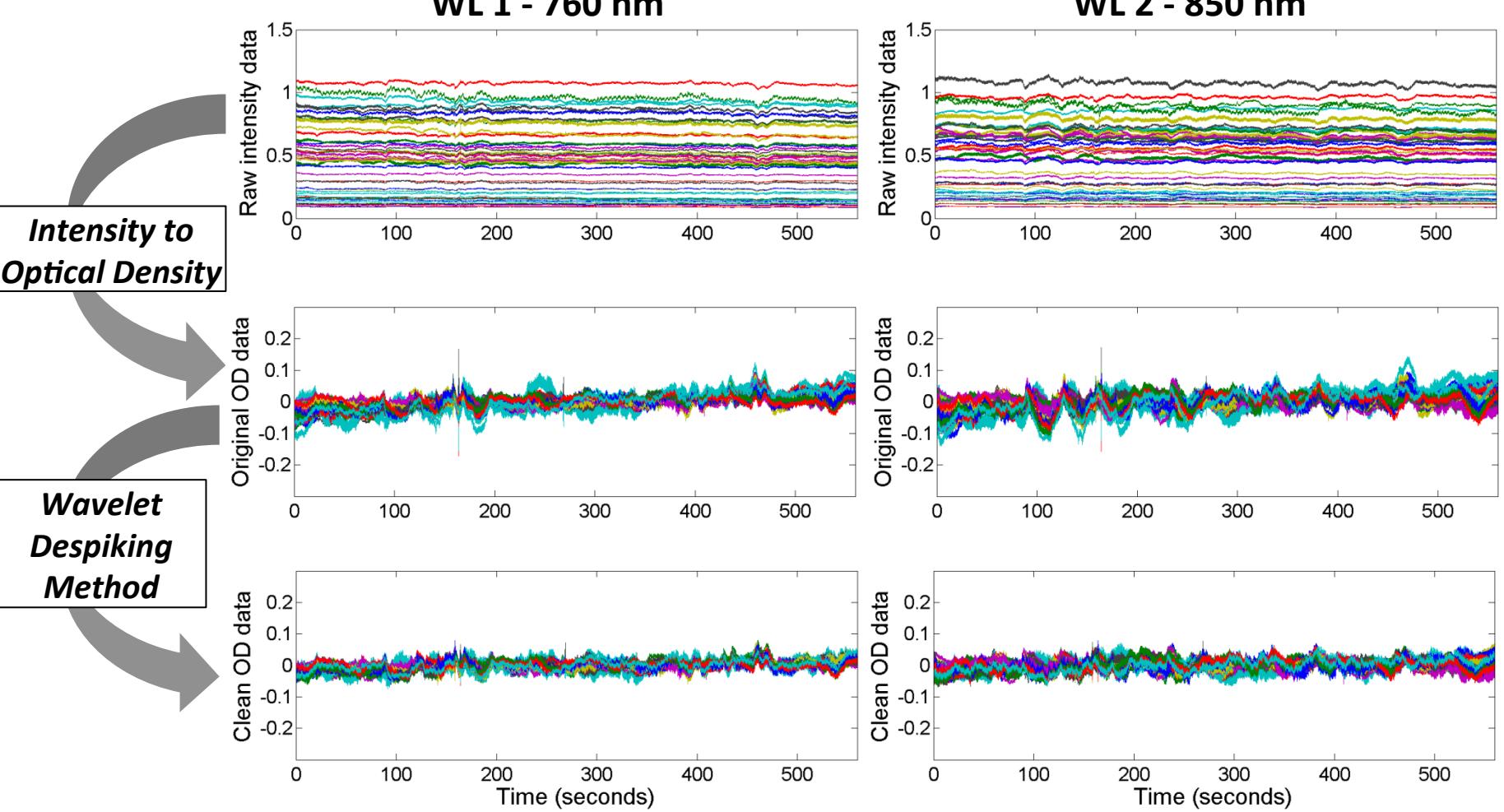
## Participants

- 36 Spanish (SP)-Basque (BQ) bilingual infants.
- 30 Spanish monolingual (>90% exposure to SP) infants
- 33 Basque monolingual (>90% exposure to BQ) infants.

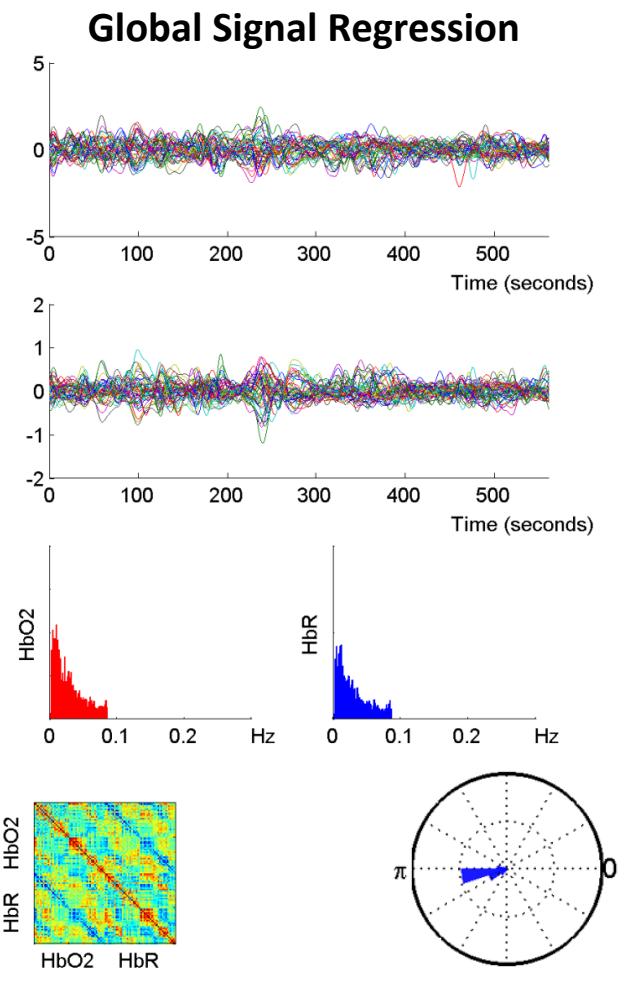
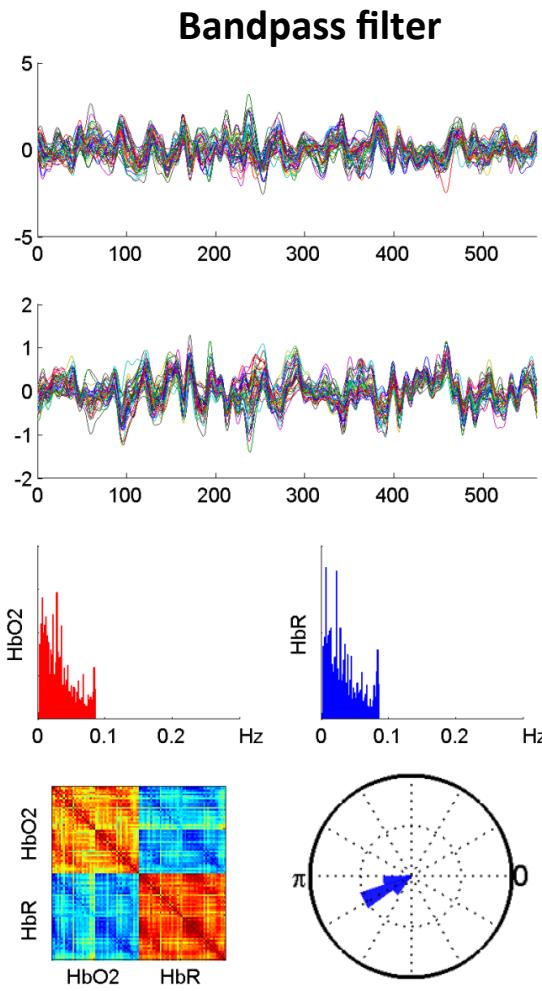
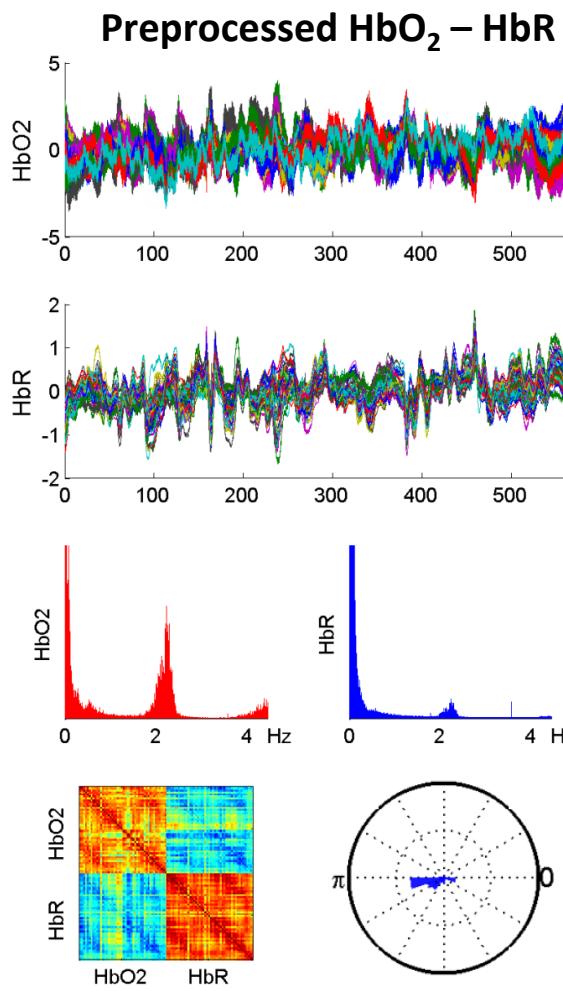
## Data acquisition

- Spontaneous hemodynamic activity during **natural sleep**.
- NIRx NIRS<sup>®</sup> system (16 sources – 24 detectors).
- **10 minutes recordings** of continuous data.

# QA: Preprocessing I



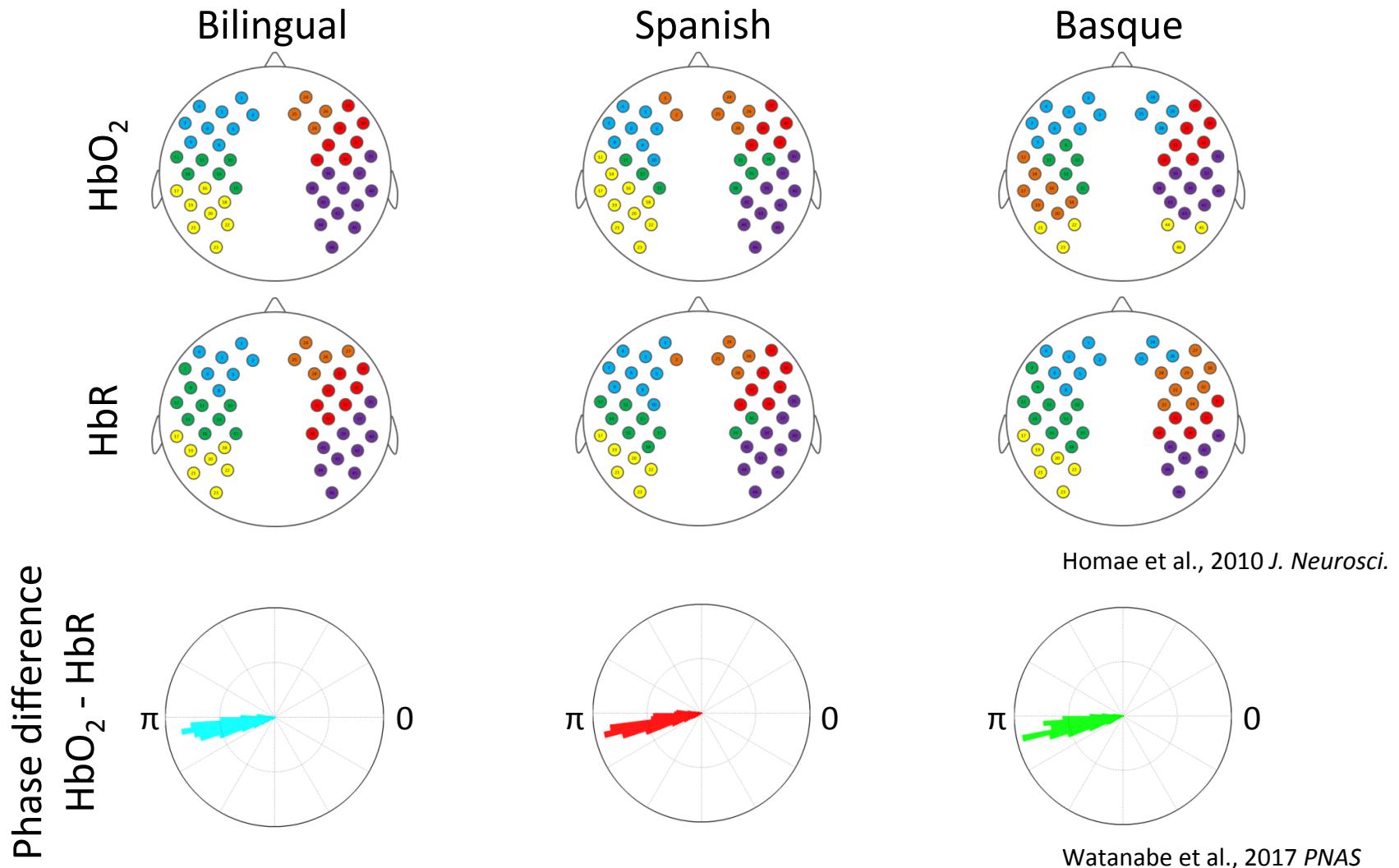
# QA: Preprocessing II



Pfeifer et al., 2018 *Front. Hum. Neurosci.*

Santosa et al., 2017 *J. Biomed. Opt.*

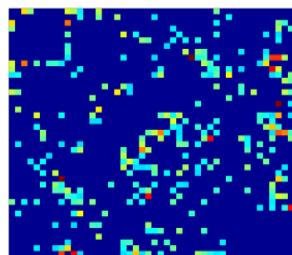
# QA: Replication of previous studies



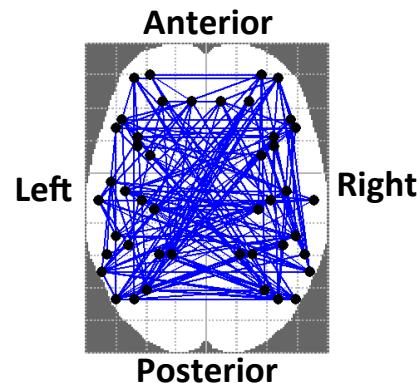
# Group effects I: Network Based Statistics

- Identify networks showing differences between groups.
  1. Compute statistics at each edge independently. Keep significant pairs.
  2. Identify networks (i.e., interconnected edges).
  3. Assess the significance of the network based on its size using non-parametric testing with participant permutation.

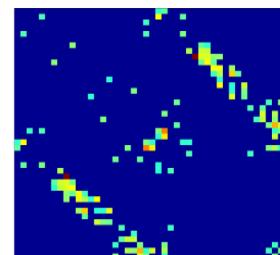
**ANOVA group (HbR)**



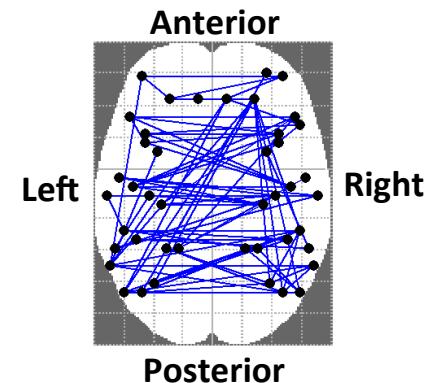
$p = 0.049$



**SP > BQ Monolinguals (HbR)**

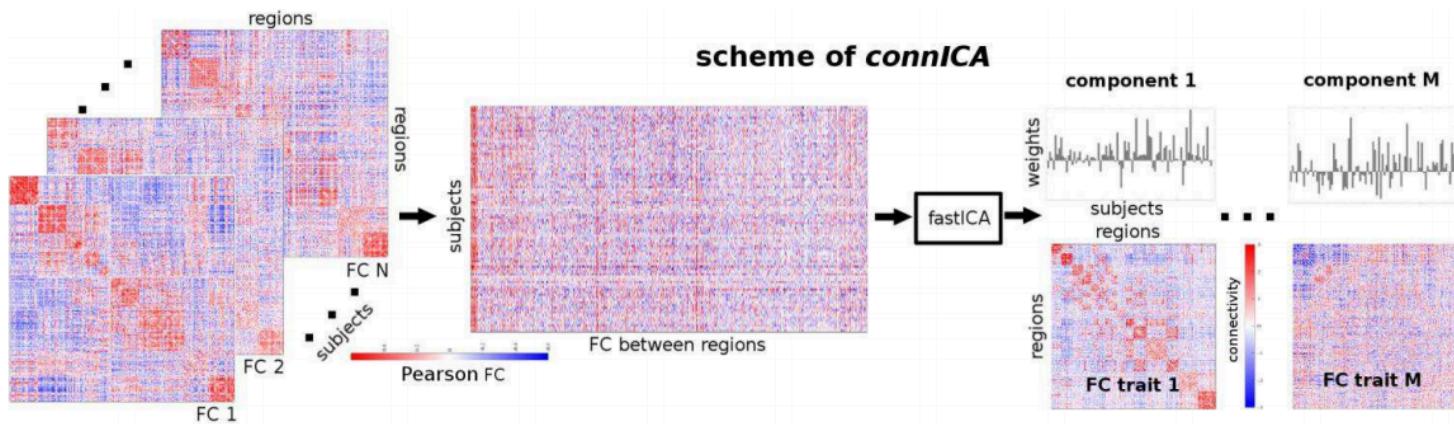


$p = 0.047$



# Group effects II: connICA

- Identify **robust independent FC patterns** in a set of individual FC matrices (i.e., connectomes).

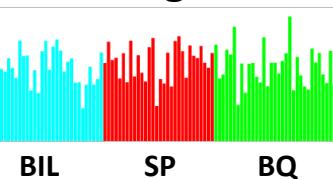


- Associated **weights** quantify the presence of the FC pattern in each participant and can be linked to individual metrics (e.g., clinical outcomes, bilingualism) to assess group differences.

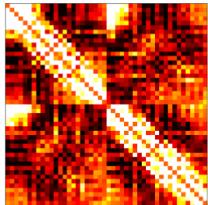
# Group effects II: connICA

HbR

Weights



FC pattern

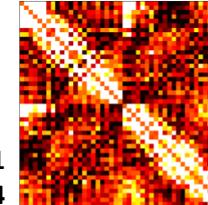


$$F(2, 96) = 0.86 \\ p = 0.4246$$

*Neighboring channels*

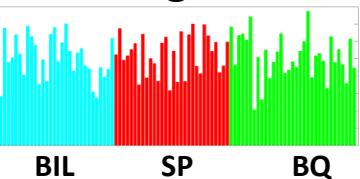
HbO<sub>2</sub>

FC pattern



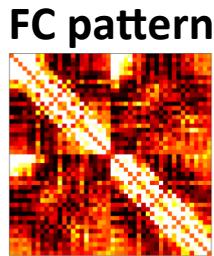
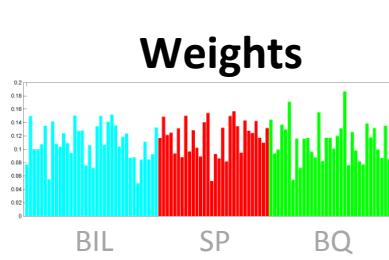
$$F(2, 96) = 0.11 \\ p = 0.894$$

Weights



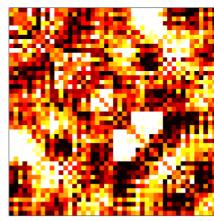
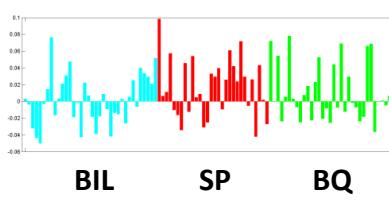
# Group effects II: connICA

HbR



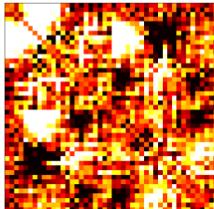
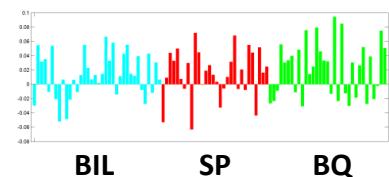
*Neighboring channels*

$$F(2, 96) = 0.86 \\ p = 0.4246$$



*Right Fronto-Temporal &  
Bilateral Fronto-Temporal channels*

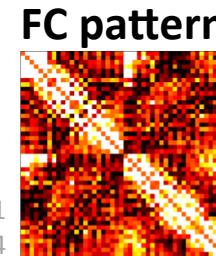
$$F(2, 96) = 1.92 \\ p = 0.152$$



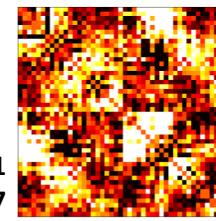
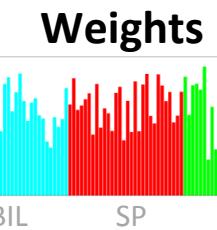
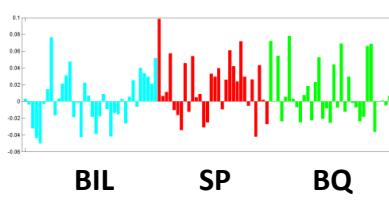
*Left Fronto-Temporal &  
Bilateral Fronto-Temporal channels*

$$F(2, 96) = 0.59 \\ p = 0.5546$$

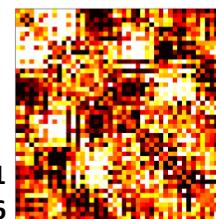
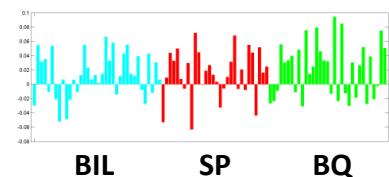
HbO<sub>2</sub>



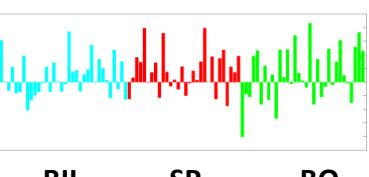
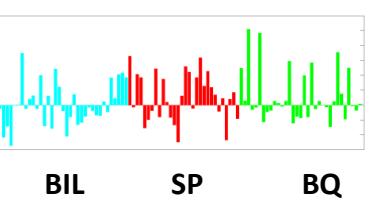
$$F(2, 96) = 0.11 \\ p = 0.894$$



$$F(2, 96) = 0.91 \\ p = 0.4047$$

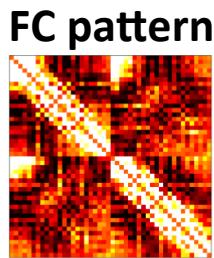
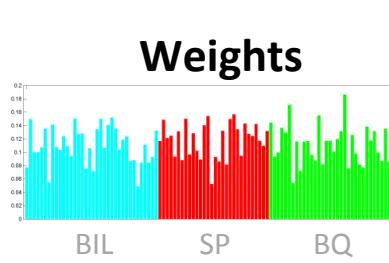


$$F(2, 96) = 0.51 \\ p = 0.6026$$

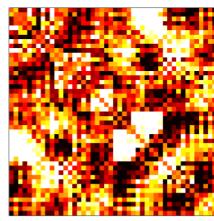


# Group effects II: connICA

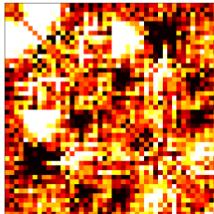
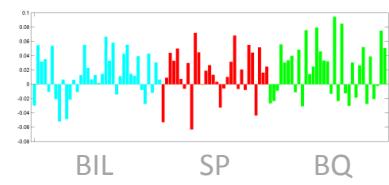
HbR



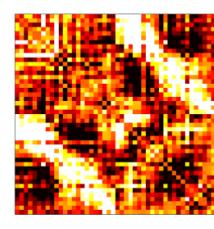
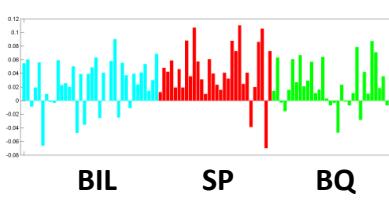
$F(2, 96) = 0.86$   
 $p = 0.4246$



$F(2, 96) = 1.92$   
 $p = 0.152$

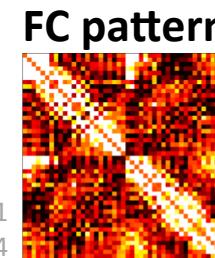


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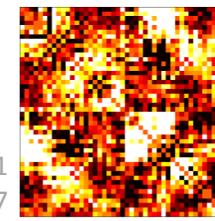


$F(2, 96) = 3.26$   
 $p = 0.0426$   
 $SP > BQ$   
 $p = 0.0497$

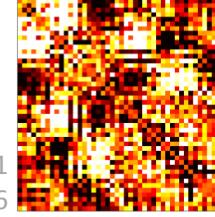
HbO<sub>2</sub>



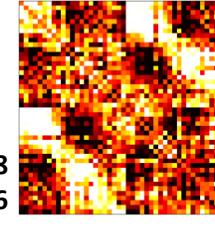
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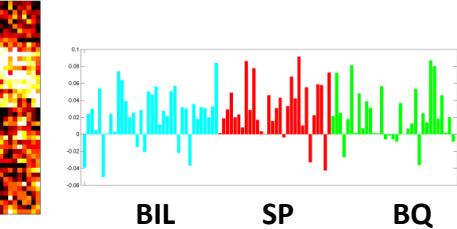
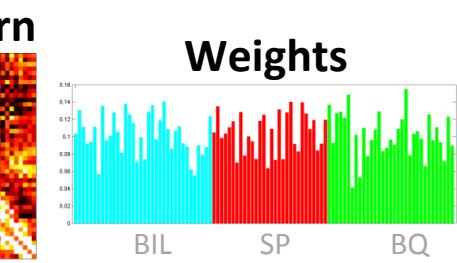
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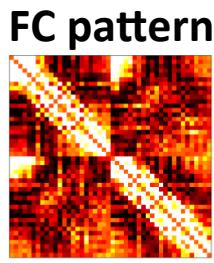
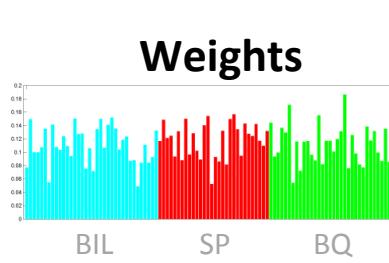


$F(2, 96) = 0.8$   
 $p = 0.4516$

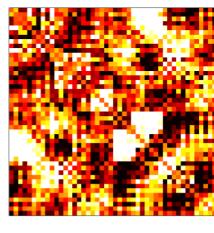
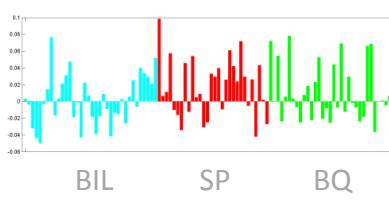


# Group effects II: connICA

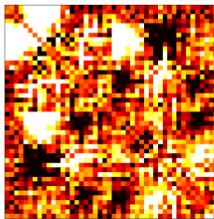
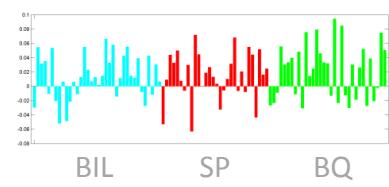
HbR



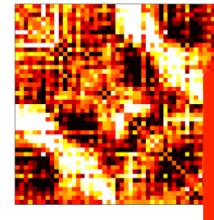
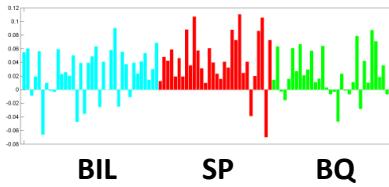
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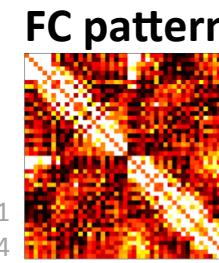


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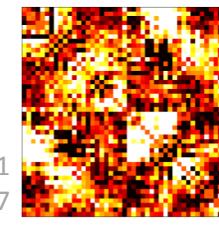


$F(2, 96) = 3.26$   
 $p = 0.0426$   
SP > BQ  
 $p = 0.0497$

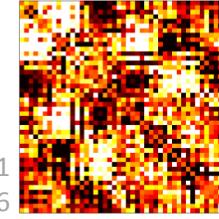
HbO<sub>2</sub>



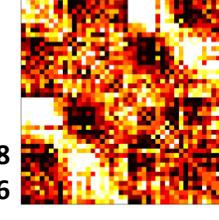
$F(2, 96) = 0.11$   
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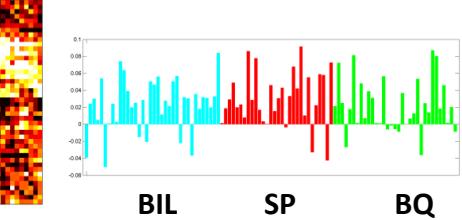
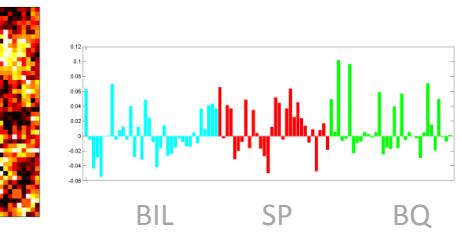
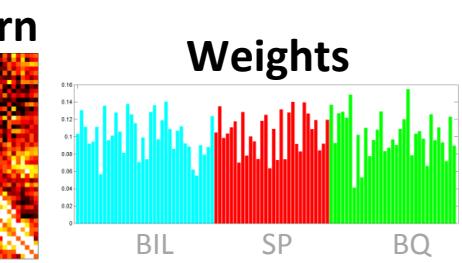
$F(2, 96) = 0.91$   
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$F(2, 96) = 0.51$   
 $p = 0.6026$



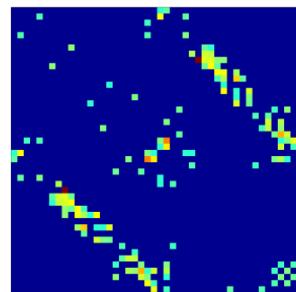
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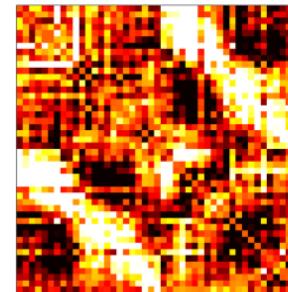
# Discussion I: Early bilingual exposure

- **Marginal differences** between groups, but **consistent** results across methods.

**SP monolinguals > BQ monolinguals**



NBS (HbR)



connICA (HbR)

- Feasibility of **NBS** and **connICA** methods to assess group differences in fNIRS resting state FC studies.

# Discussion II: Quality assessment

- Relevance of the current work:
  - Large sample size in three experimental groups.
  - Recording duration and data quality.
  - Replication of previous studies.

**<https://github.com/borjablanco/fNIRS-2018>**

- Impact of **data quality** and **data preprocessing** in infant fNIRS resting state FC studies.
- Importance of unified data processing criteria that ensure **consistency** across studies and **replication**.



# Thank you

**<https://github.com/borjablanco/fNIRS-2018>**

*Monday October 8<sup>th</sup>. Poster session III – 66  
Effect of prewhitening in resting-state near infrared spectroscopy data.  
B. Blanco, M. Molnar and Cesar Caballero-Gaudes*

# *Supplementary information*

# Early bilingual exposure

- Experience-related effects in brain structure and function (e.g., socioeconomic status, formal education, music training).
- Bilingual language acquisition.
  - **Language processing (bilingual ≠ monolingual).**
  - **Language acquisition (bilingual ≠ monolingual).**

Bialystok et al., 2017 *Psychol. Bull*

Costa and Sebastián-Gallés, 2014 *Nature Rev. Neurosci.*

- Bilingual adaptation in children and adults:

- **Structural imaging**

Mohades et al., 2015 *PLoS One*;  
Li et al., 2014 *Cortex (Review)*

- **Functional task-based imaging**

Arredondo et al., 2016 *Developmental Sci.*;  
Abutalebi et al., 2012 *Cereb. cortex*

- **Resting state FC**

Berken et al., 2016 *J. Neurosci.*;  
Grady et al., 2015 *Neuropsychologia*

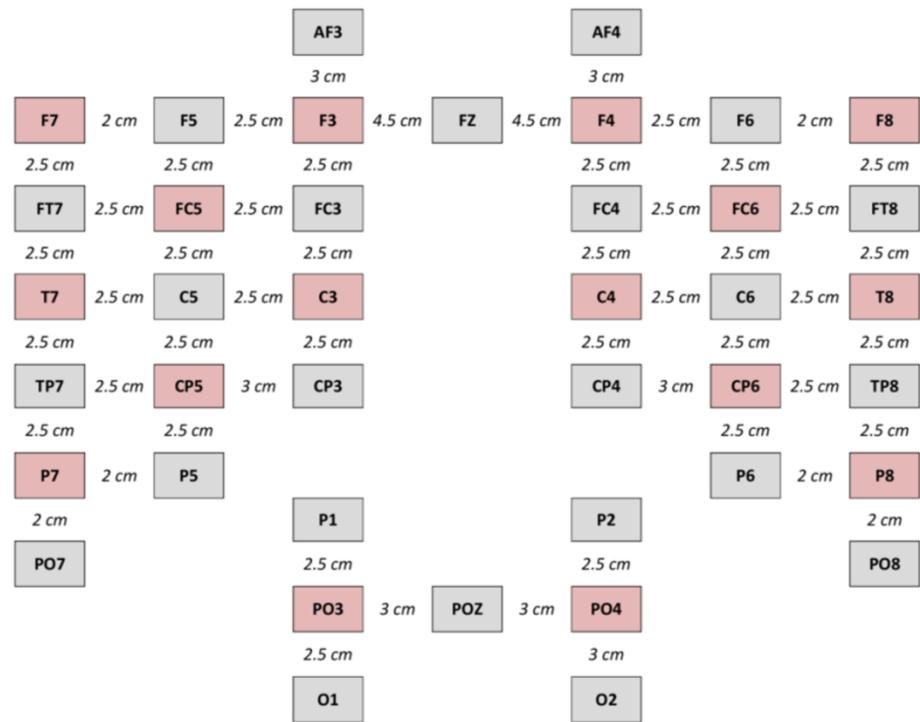
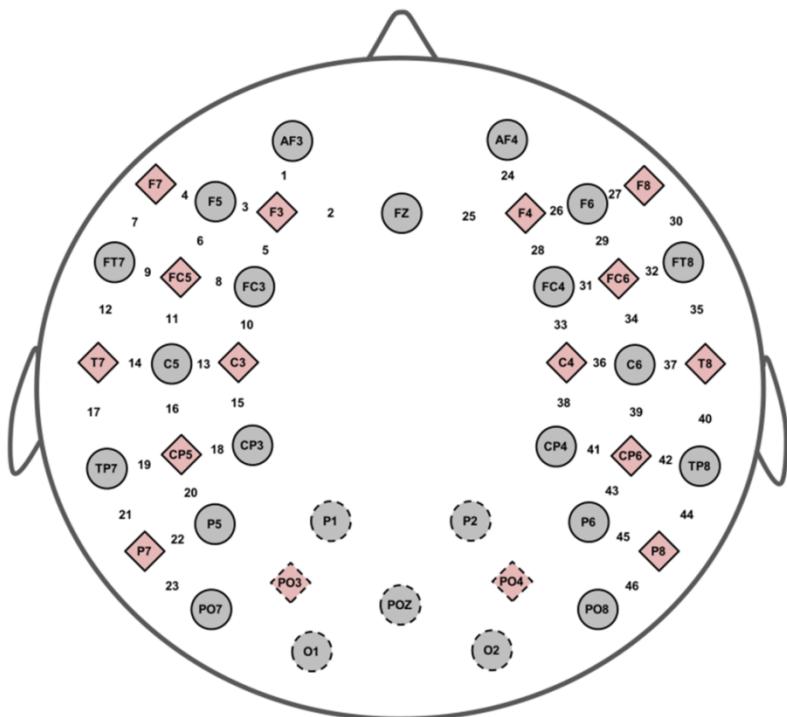
# Early bilingual exposure

- Bilingual adaptation in infants:
  - **Behavioral studies**  
Kovacs & Mehler, 2009a *PNAS*  
Sebastian-Galles et al., 2012 *Psychol. Sci.*  
Molnar et al., 2014 *Infancy*
  - **Functional task-based imaging**  
Ferjan-Ramirez et al., 2017 *Developmental Sci.*  
Nacar-Garcia et al., 2018 *Sci Rep.*
  - **Resting state FC?**
- Resting state FC in infants:
  - **Premature birth**  
Smyser et al., 2010 *Cereb. Cortex (fMRI)*  
Kwon et al., 2015 *Neuroimage (fMRI)*
  - **Socio-economic status**  
Gao et al., 2015a *Cereb. Cortex (fMRI)*
  - **Autism**  
Keehn et al., 2013 *Front. Hum. Neurosci (fNIRS)*

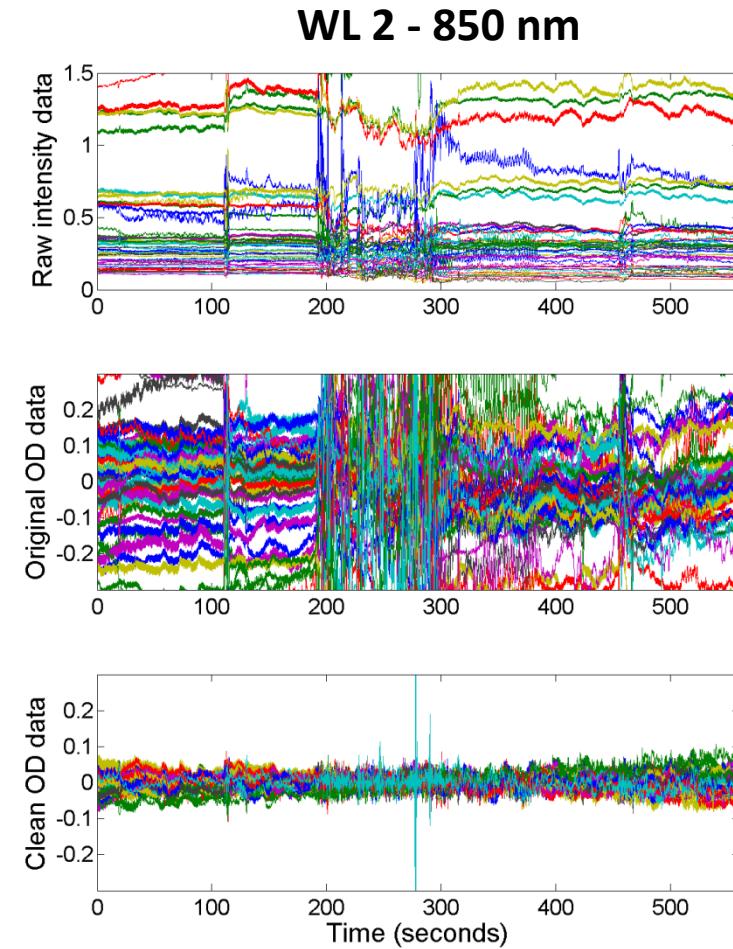
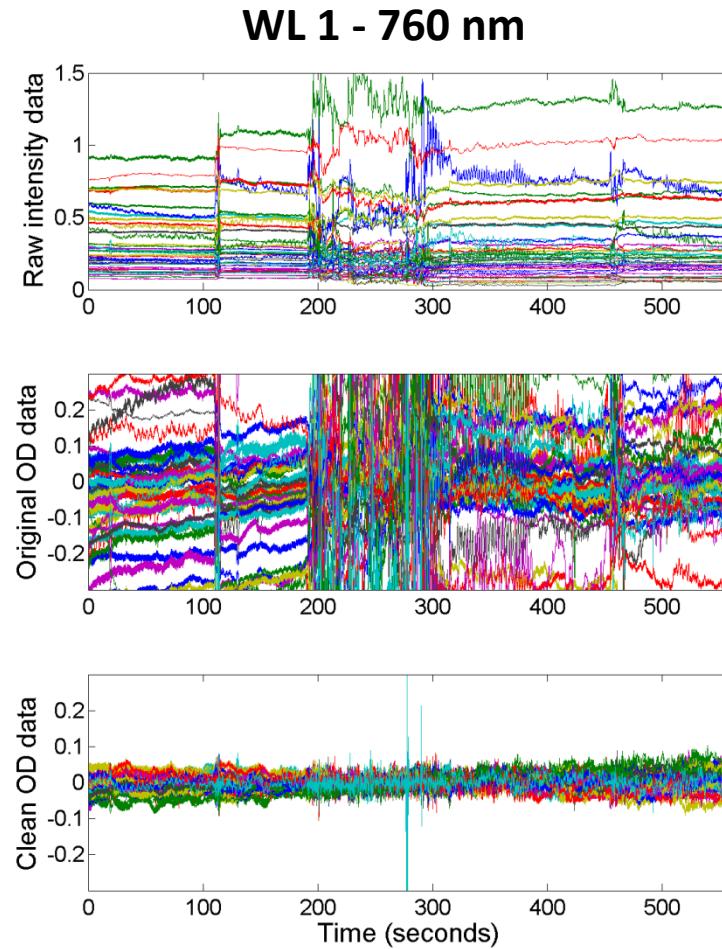
# *Participants*

- **Total:** 123 healthy full-term infants.
- Not able to fall sleep (n = 16).
- Regular exposure to English (n = 1).
- Datasets shorter than 10 minutes (n = 2)
- Low data quality (n = 5).
  
- **Final sample:** 99 healthy full term infants
- Bilinguals (n = 36, 21 girls).
- Spanish (n = 30, 13 girls).
- Basque (n = 33, 17 girls).

# Optical probe

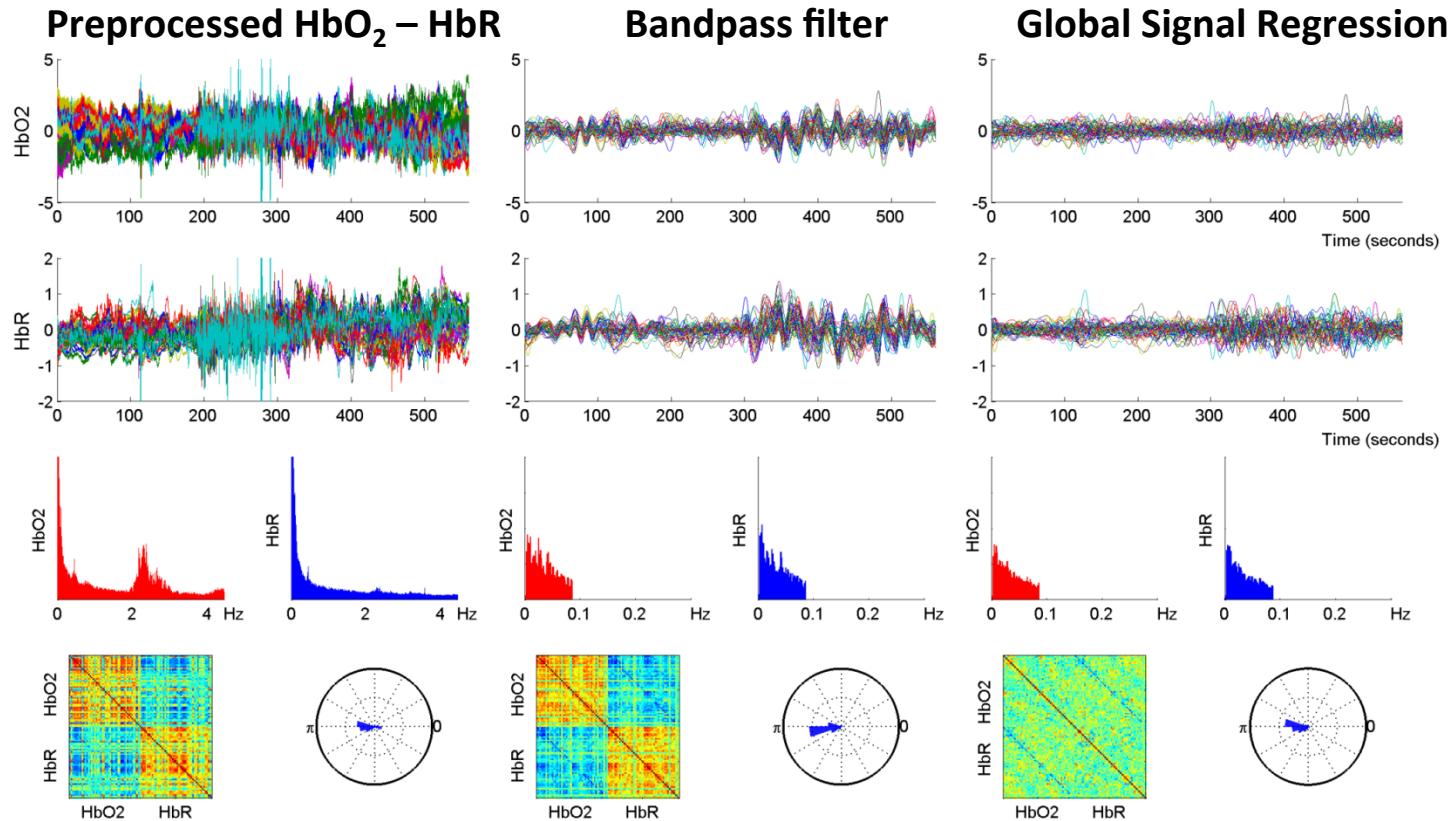


# Example of a rejected participant



<https://github.com/borjablanco/fNIRS-2018>

# Example of a rejected participant



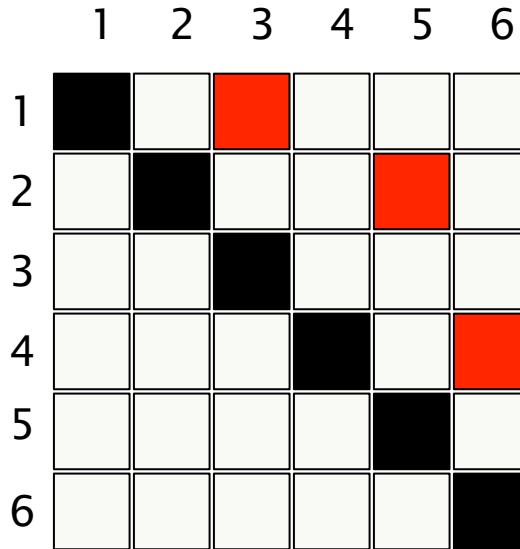
<https://github.com/borjablanco/fNIRS-2018>

# **Network Based Statistics**

1. Compute statistics for every pair of channels obtaining a matrix of F or T-values (i.e., equivalent to a F-test or a two-sample *t*-test at each channel pair).
2. Retain statistics of significant connections higher than a pre-specified threshold.
3. Identify clusters of connections (*networks*) among the connections exceeding the threshold (i.e., interconnected edges), and compute the size of the network.

**Two different measures of size:** Extent (number of connections)  
Intensity (strength of the connections)
4. Compute significance of the network based on permutation testing where participants are randomly interchanged between the groups. Repeat the previous steps for each permutation.
5. The size of the networks found in each permutation is stored to construct the null distribution of the *p*-values.
6. NBS offers higher sensitivity than standard pairwise testing approaches maintaining strict control of false positives at the network level (FWER).

# *Network Based Statistics*



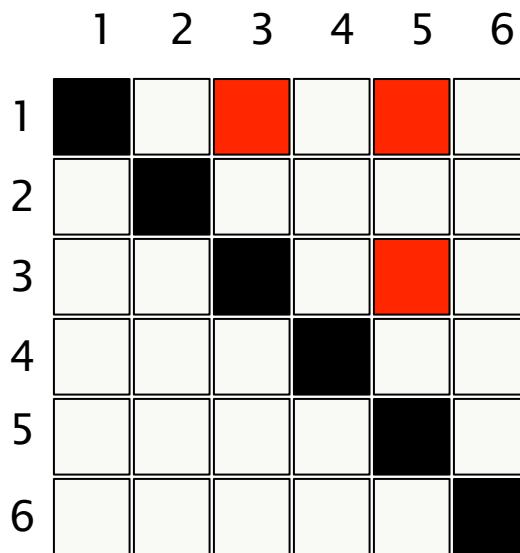
Connections higher than the prespecified threshold but not interconnected edges:

CH1 - CH3

NBS

CH2 - CH5

CH4 - CH6



Connections higher than the prespecified threshold and interconnected edges:

CH1 - CH3

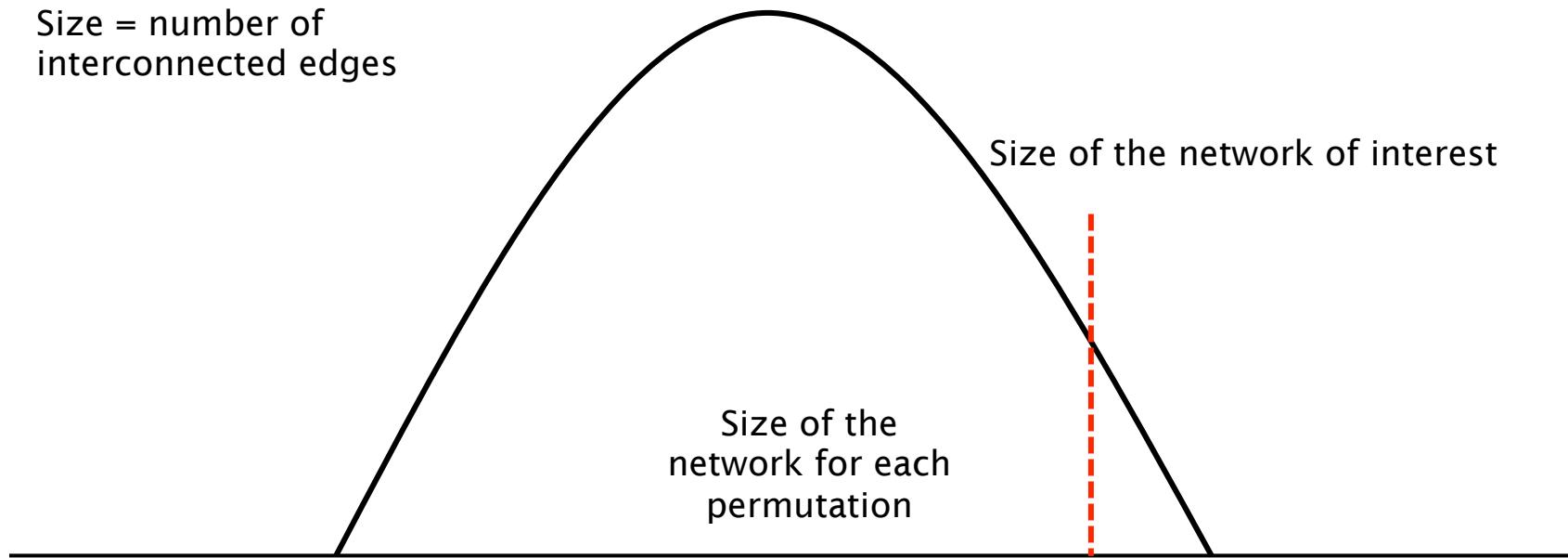
NBS

CH1 - CH5

CH3 - CH5

# ***Network Based Statistics***

Size = number of interconnected edges

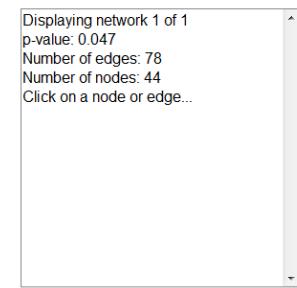
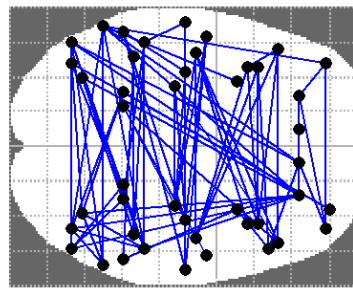
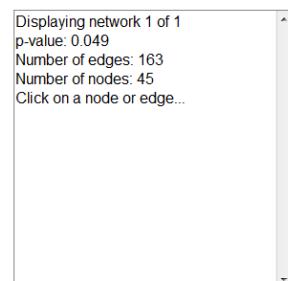
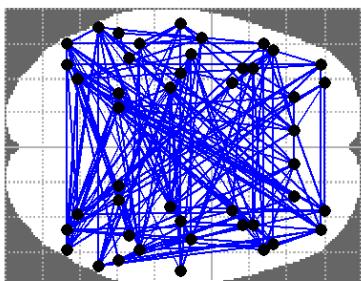
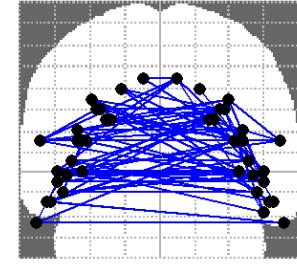
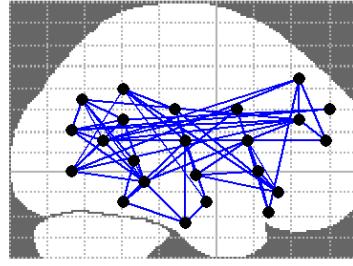
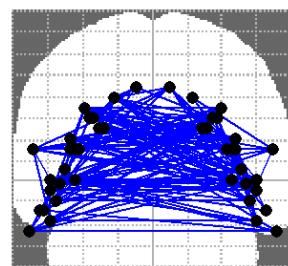
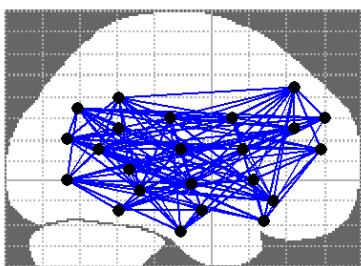


Size of the  
network for each  
permutation

Size of the network of interest

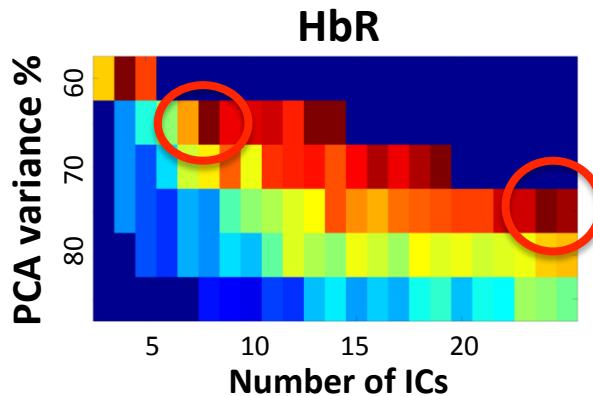
$$P \text{ value} = \frac{\text{Number of networks larger than the network of interest}}{\text{Number of permutations}}$$

# *NBS: results*



# Parameters: connICA

- Two sets of parameters for explained variance (PCA) and number of ICs. **PCA 65% - number of ICs 8: global components**  
**PCA 75% - number of ICs 24: focal components**

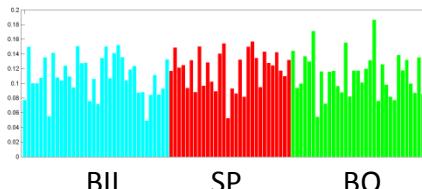


- Assess only FC patterns that appear across  $\text{HbO}_2$  and  $\text{HbR}$ .
- Use the associated weights to test differences between groups.
- ANOVA & multiple comparison of group means (Tukey's HSD).

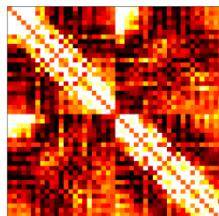
# connICA: PCA 65% - number of IC 6

HbR

Weights



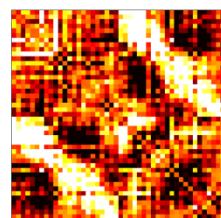
FC pattern



*RSNs component*

$$F(2, 96) = 0.86 \\ p = 0.4246$$

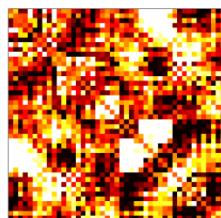
$$F(2, 96) = 0.11 \\ p = 0.894$$



*Bilateral component*

$$F(2, 96) = 3.26 \\ p = 0.0426 \\ SP > BQ \\ p = 0.0497$$

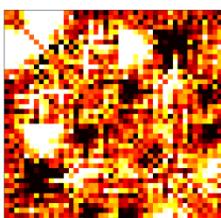
$$F(2, 96) = 0.8 \\ p = 0.4516$$



*Right FT - Bilateral FT*

$$F(2, 96) = 1.92 \\ p = 0.152$$

$$F(2, 96) = 0.91 \\ p = 0.4047$$



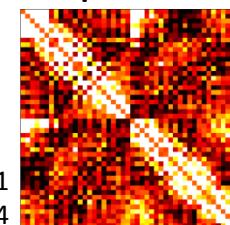
*Left FT - Bilateral FT*

$$F(2, 96) = 0.59 \\ p = 0.5546$$

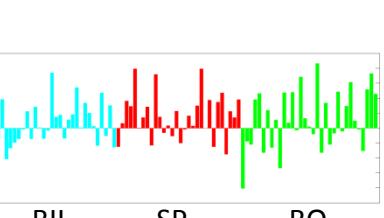
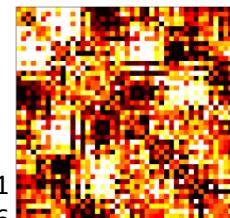
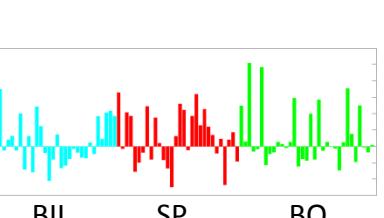
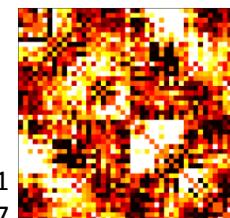
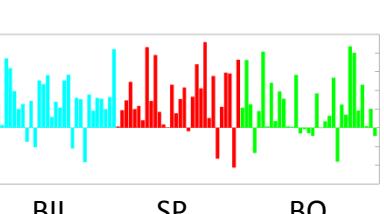
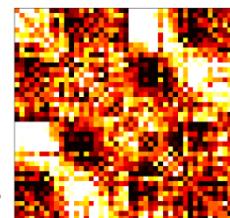
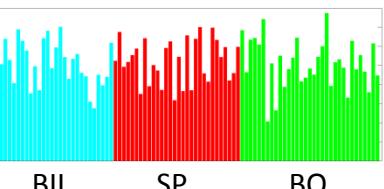
$$F(2, 96) = 0.51 \\ p = 0.6026$$

HbO<sub>2</sub>

Weights

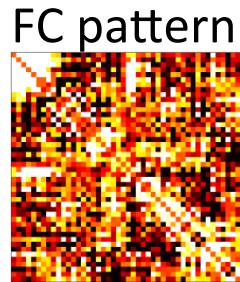
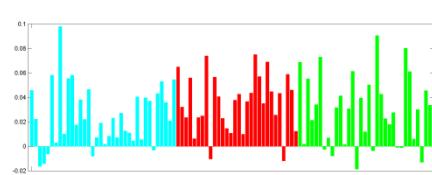


FC pattern



# connICA: PCA 75% - number of IC 24

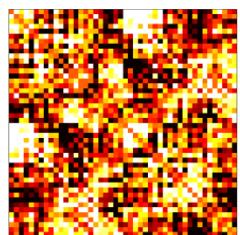
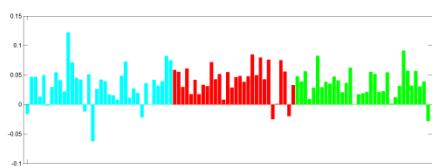
HbR



*Fronto-temporal right*

$$F(2, 96) = 0.95 \\ p = 0.3894$$

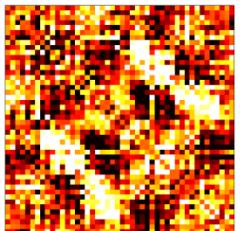
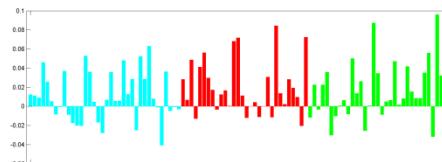
$$F(2, 96) = 3.51 \\ p = 0.0339 \\ \text{BIL} > \text{BQ} \\ p = 0.0424$$



*Bilateral Fronto-temporal*

$$F(2, 96) = 2.1 \\ p = 0.1278$$

$$F(2, 96) = 0.29 \\ p = 0.7505$$

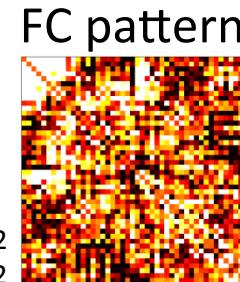


*Bilateral component\**

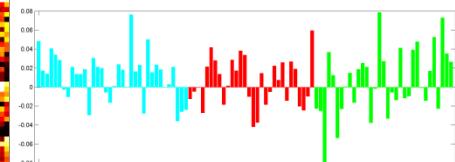
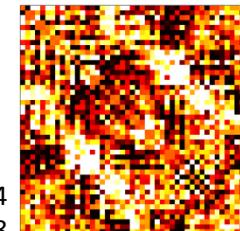
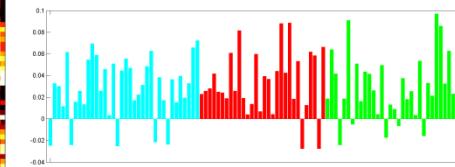
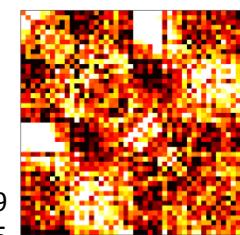
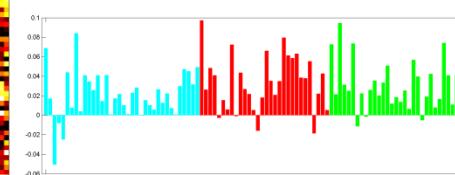
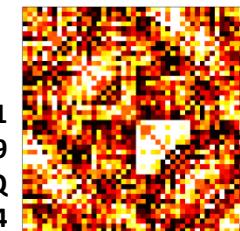
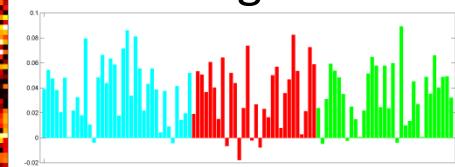
$$F(2, 96) = 1.54 \\ p = 0.2195$$

$$F(2, 96) = 1.34 \\ p = 0.2678$$

HbO<sub>2</sub>



Weights



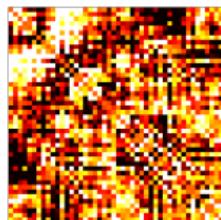
# connICA: PCA 75% - number of IC 24

HbR

Weights



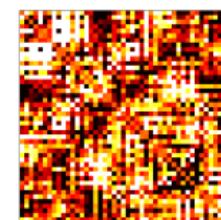
FC pattern



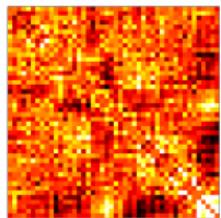
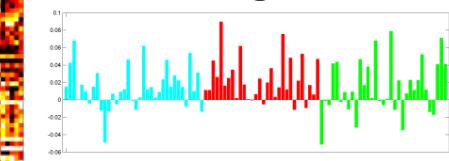
*Temporal left*

HbO<sub>2</sub>

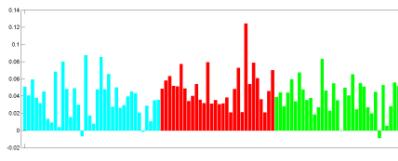
FC pattern



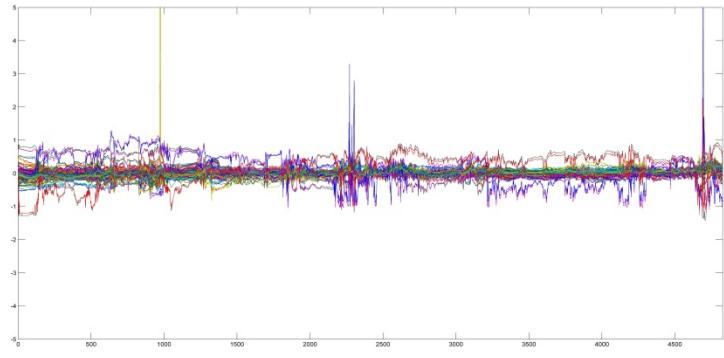
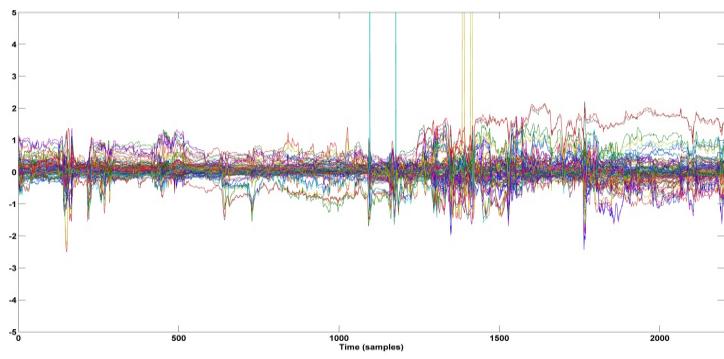
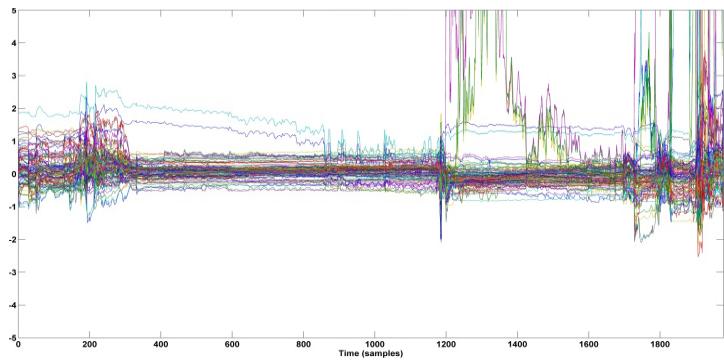
Weights



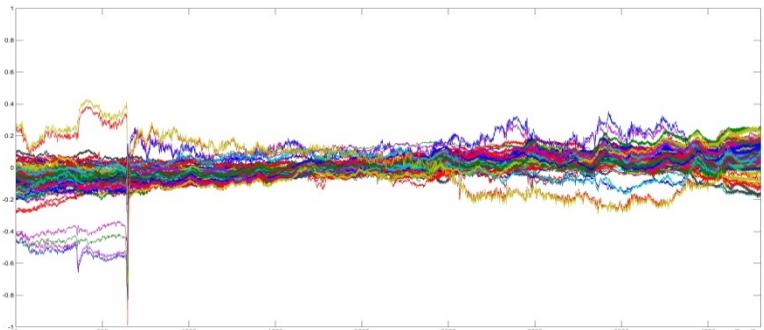
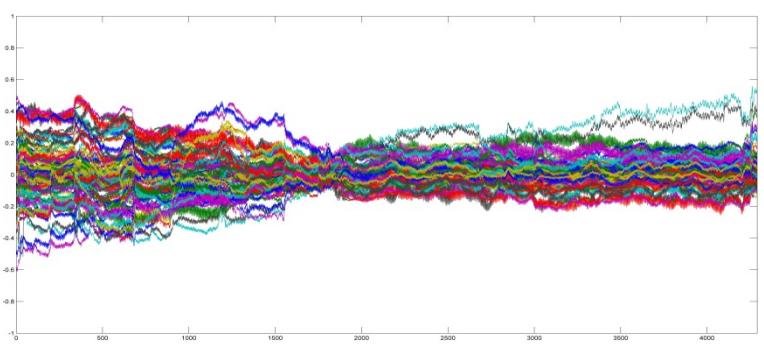
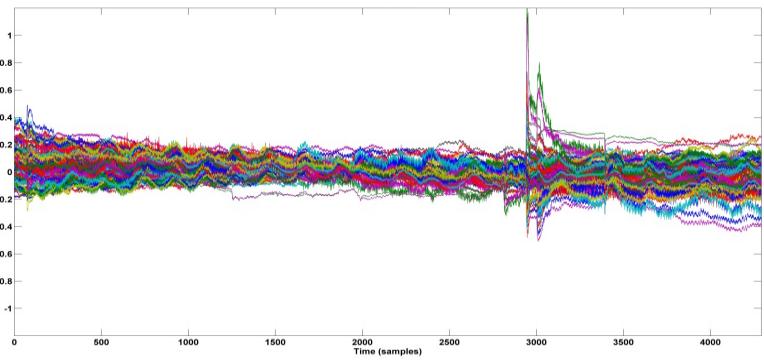
*Parietal left*



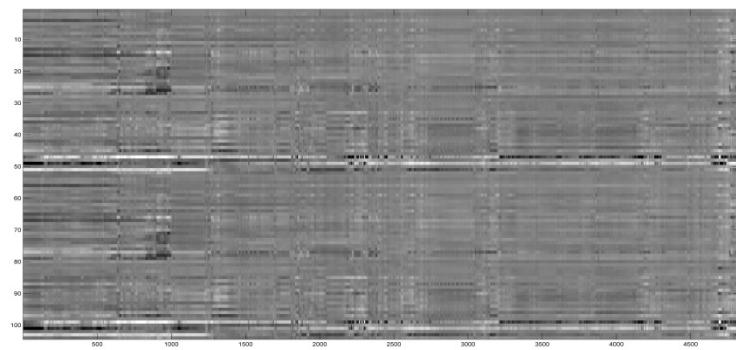
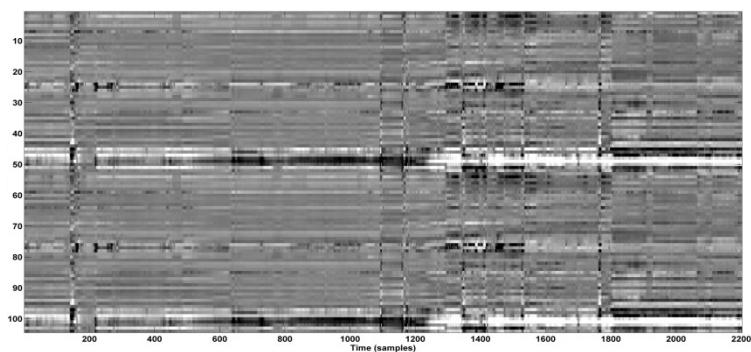
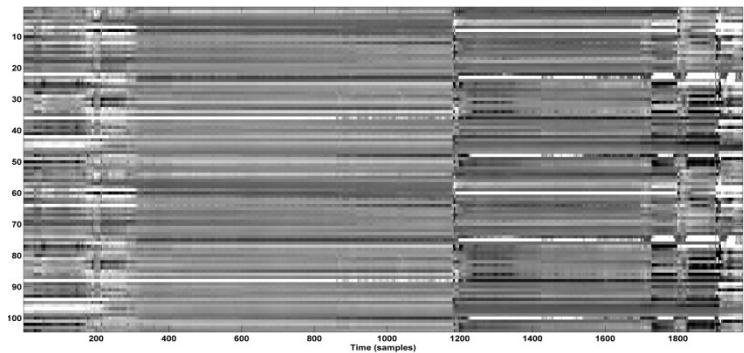
# *Awake infants*



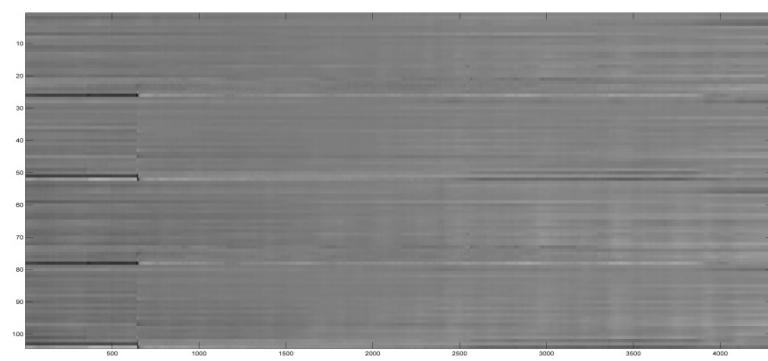
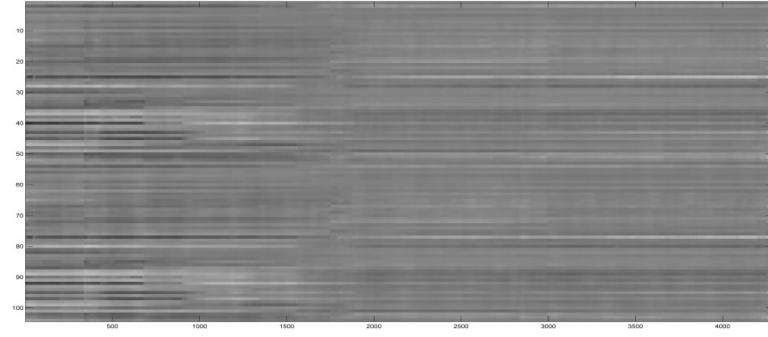
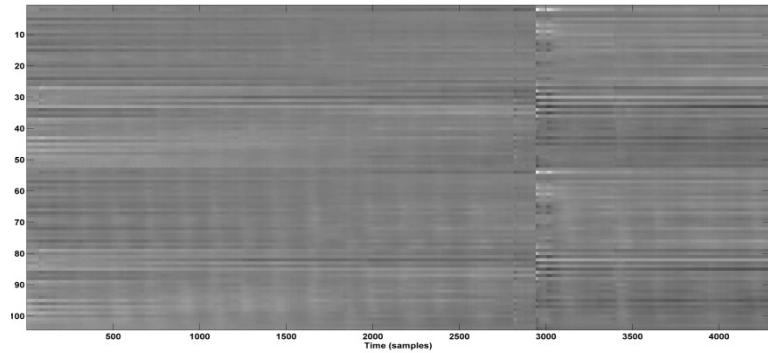
# *Asleep infants*



# *Awake infants*

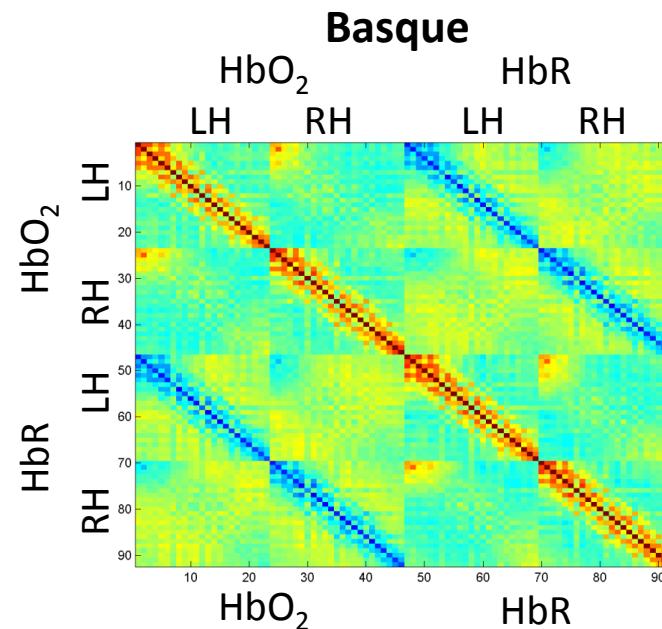
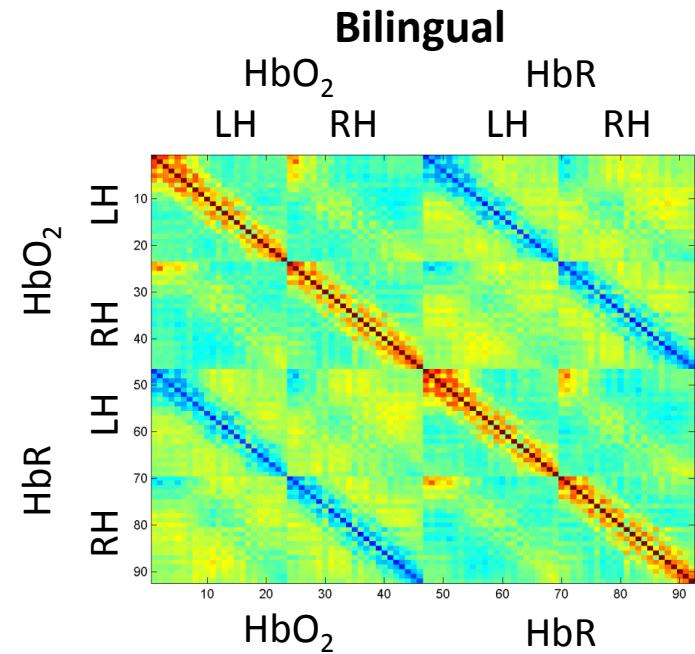
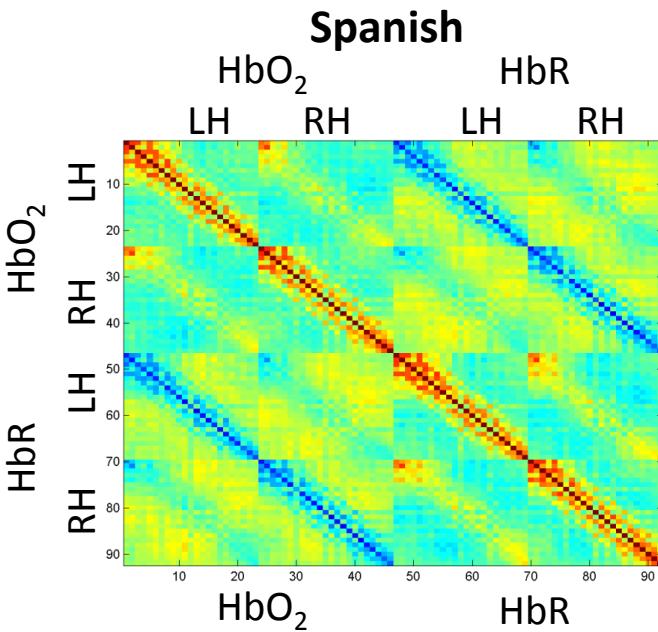


# *Asleep infants*



Average functional connectivity matrices in each group for:

- HbO<sub>2</sub>
- HbR
- Between HbO<sub>2</sub> and HbR



# 3 clusters

