

A DTI-based tractography study of effects on brain structure associated with prenatal alcohol exposure in newborns

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OHBM, Hamburg, 2014

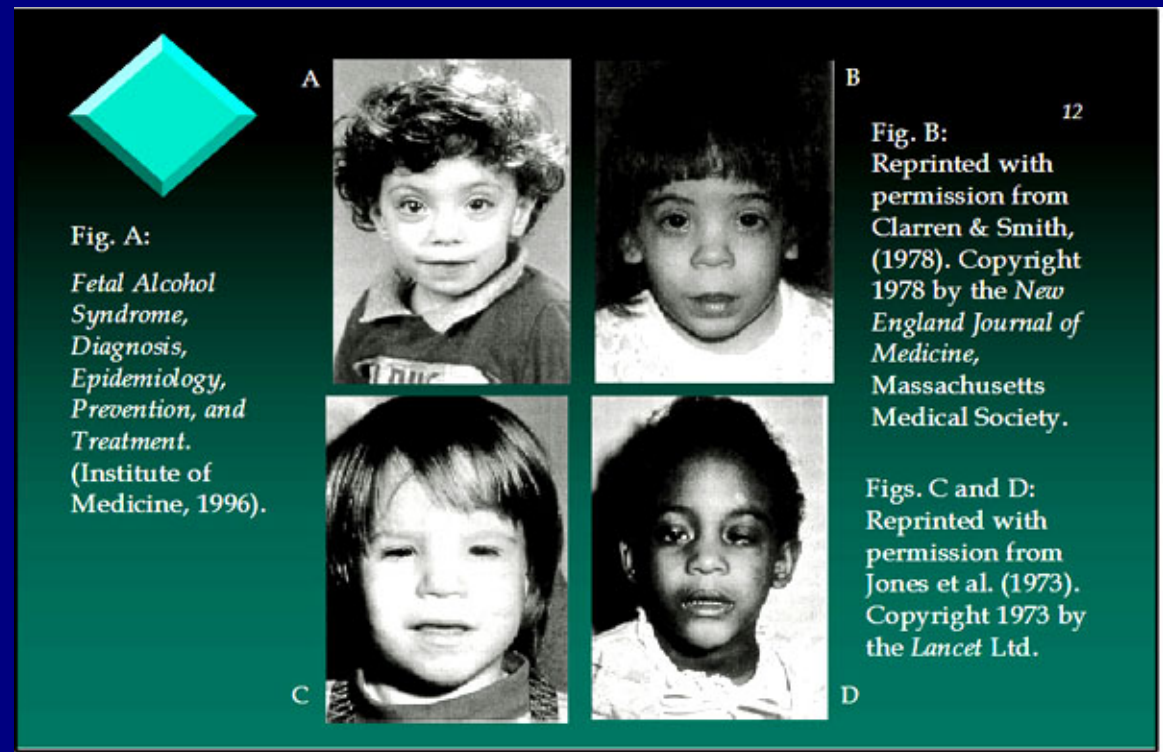
Outline

- Prenatal alcohol exposure (PAE) in brief
- Methods: diffusion tensor imaging (DTI) and tractography
- Setup for this DTI – PAE newborn study
- Newborn infant study results

→ see also Taylor et al. poster #3241, W/Th

Prenatal alcohol exposure (PAE)

- Alcohol is a teratogen, disrupting healthy embryonic and fetal development.
 - leads to various **Fetal Alcohol Spectrum Disorders (FASD)**
- FASD occurs in children whose pregnant mothers binge drank
 - e.g., ≥ 4 drinks/occasion and/or ≥ 14 drinks/wk
- Results in *poor*:
 - academic performance
 - language/math skills
 - impulse control
 - abstract reasoning
 - memory, attention and facial and skeletal dysmorphology



PAE and FASD assessment

Traditional/clinical assessments:

- the degree of facial and skeletal dysmorphology
 - For example, changes in lip, philtrum and nasal structures



Different racial/ethnic groups typically show varied changes

- cognitive deficits
- eye-blink conditioning

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Different racial/ethnic groups typically show varied changes

- cognitive deficits
 - eye-blink conditioning
- semi-quantitative, convolve many factors (upbringing, education, etc.), and do not locate brain changes

Goals of this study

To:

- 1) Use **neuroimaging** to compare structural brain development in newborns with PAE to that of HC newborns.
- 2) **Quantitatively** examine WM properties across the brain
- 3) Relate changes in **(localized) WM properties** with PAE, controlling for several confounding effects
→ examine several, and see which is/are (most) significant

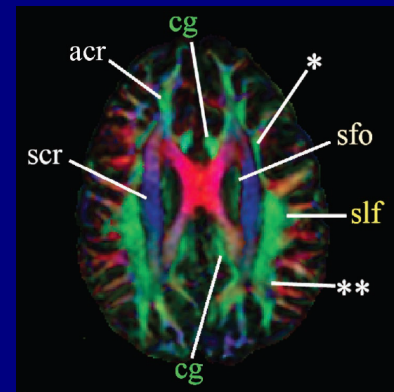
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Tools: **diffusion tensor imaging (DTI) + tractography**

- A) **delineate** similar WM ROIs across all subjects
- B) **quantify** structural properties (FA, MD, T1, ...)
- C) **statistical modeling for comparisons**
 - at whole brain, network and ROI levels



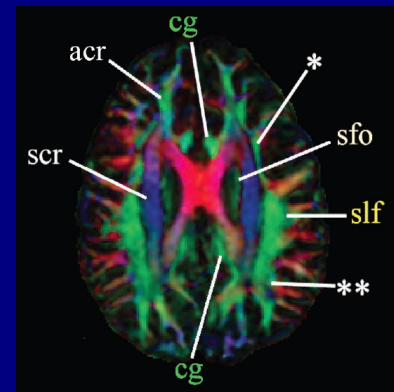
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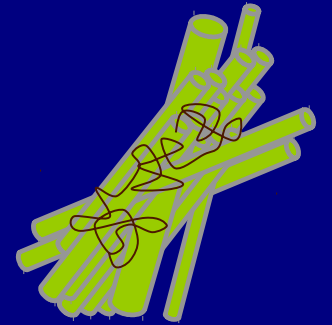


This is the first study to use DTI and tractography to examine PAE effects in the neonatal/infant period.

Local structure via diffusion MRI

(In brief)

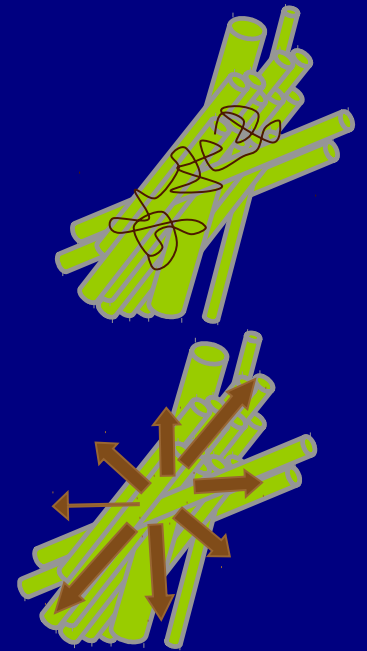
1) Random motion of molecules affected by local structures



Local structure via diffusion MRI

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- 1) Random motion of molecules affected by local structures
- 2) Statistical motion measured using diffusion weighted MRI



Local structure via diffusion MRI

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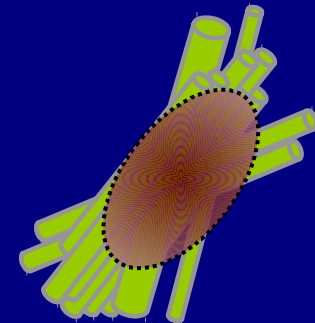
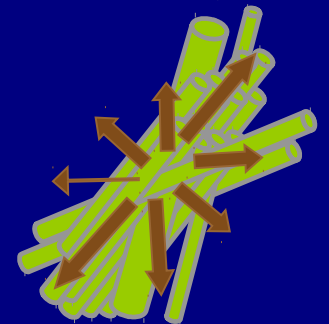
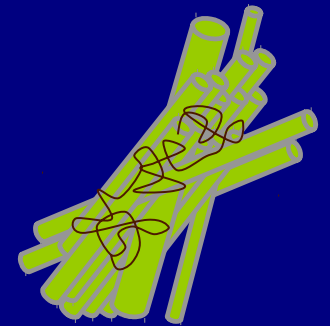
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2) Statistical motion measured using diffusion weighted MRI

3) Bulk features of local structure approximated with various reconstruction models, mainly grouped by number of major structure directions/voxel:

+ one direction:

DTI (Diffusion Tensor Imaging) \rightarrow ellipsoid



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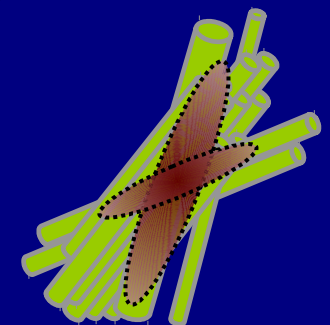
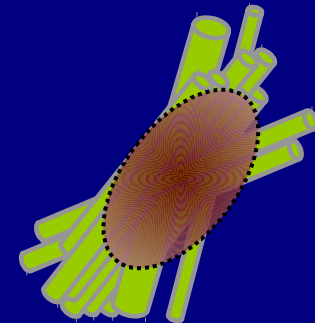
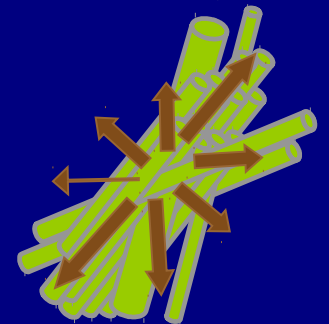
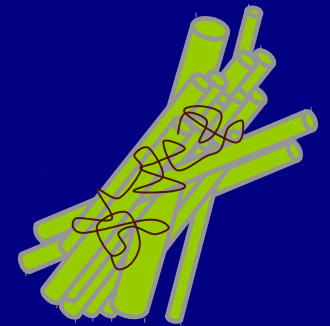
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HARDI (High Angular Resolution Diffusion Imaging)

Qball, DSI, ODFs, ball-and-stick, multi-tensor, CSD, ...



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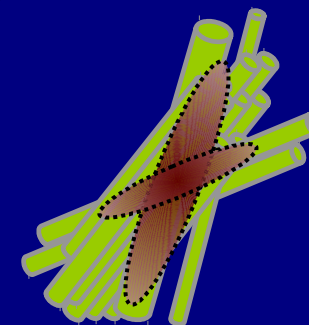
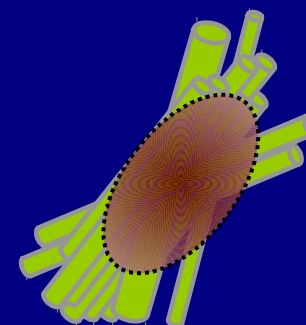
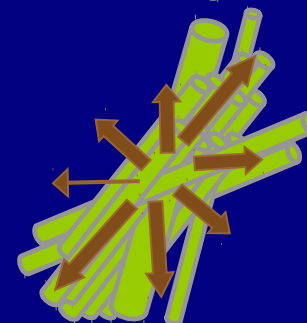
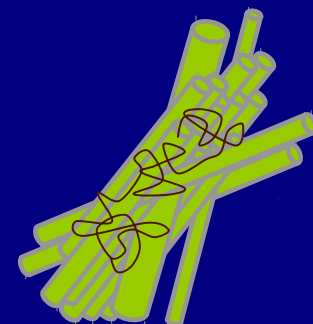
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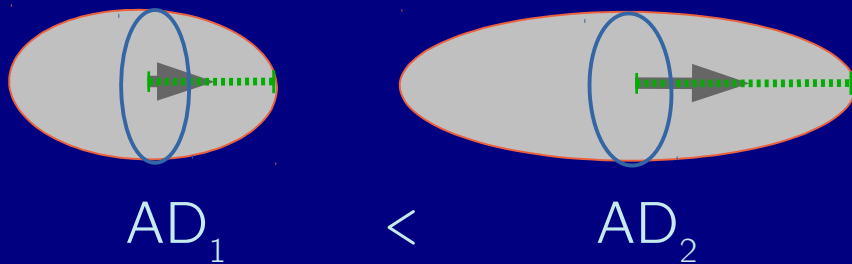
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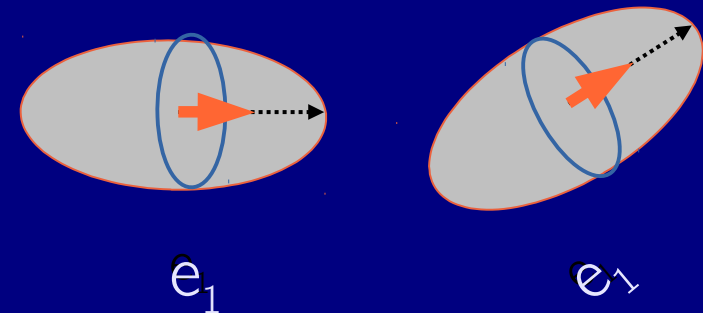
“Big 5” DTI ellipsoid parameters

Main quantities of diffusion (motion) surface

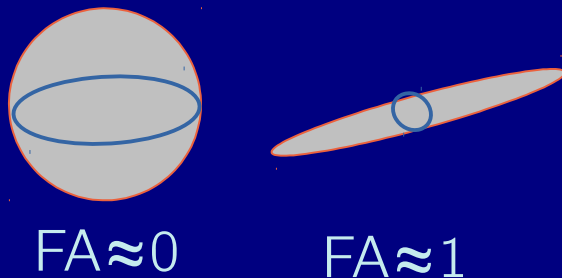
first eigenvalue, λ_1
 $= \lambda_1$, parallel/axial diffusivity, AD



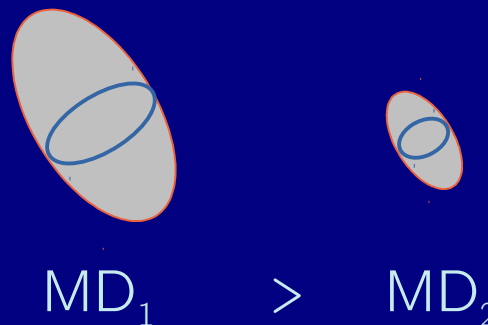
first eigenvector, e_1



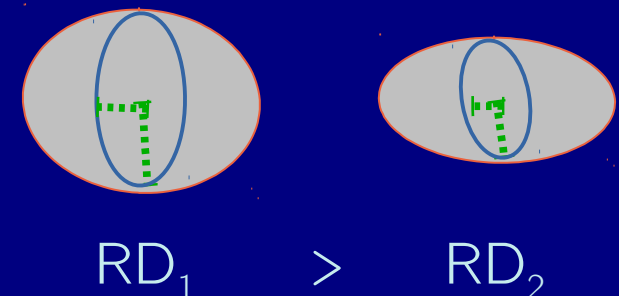
Fractional anisotropy, FA



Mean diffusivity, MD



Radial diffusivity, RD



Interpreting DTI parameters

General literature:

FA: measure of fiber bundle coherence and myelination

- in adults, $FA > 0.2$ is proxy for WM
- in infants, $FA > 0.1$ is proxy for WM¹

MD, AD, RD: local density of structure

e_1 : orientation of major bundles

¹e.g., Dubois et al., 2006

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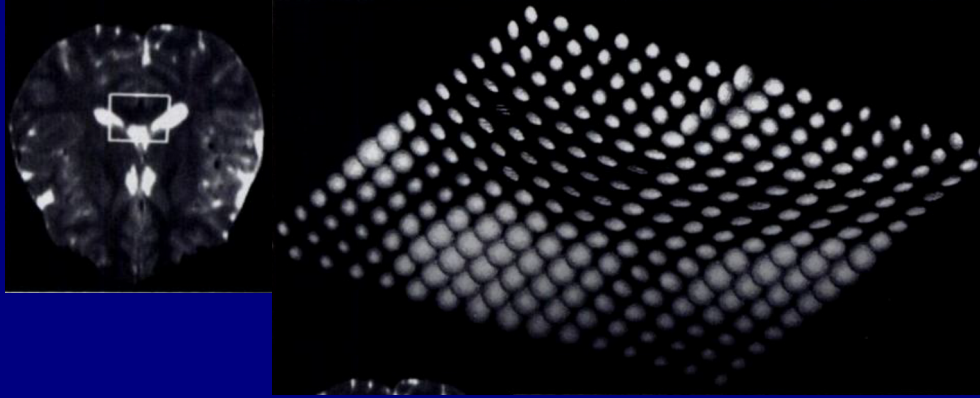
Cautionary notes:

- Degeneracies of structural interpretations
- Changes in myelination may have small effects on FA
- WM bundle diameter \ll voxel size
 - don't know location/multiplicity of underlying structures
- More to diffusion than just structure-- i.e., fluid properties
- Noise, distortions, etc. in measures

¹e.g., Dubois et al., 2006

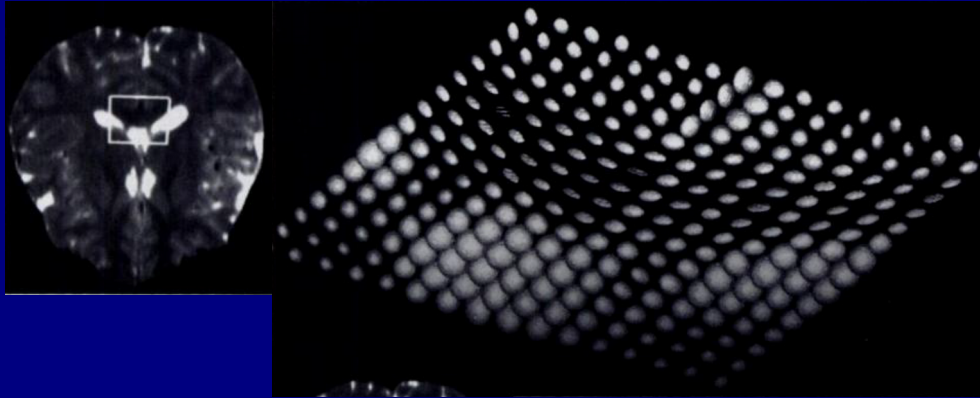
Local DTs \rightarrow extended tracts

Field of local diffusion parameters



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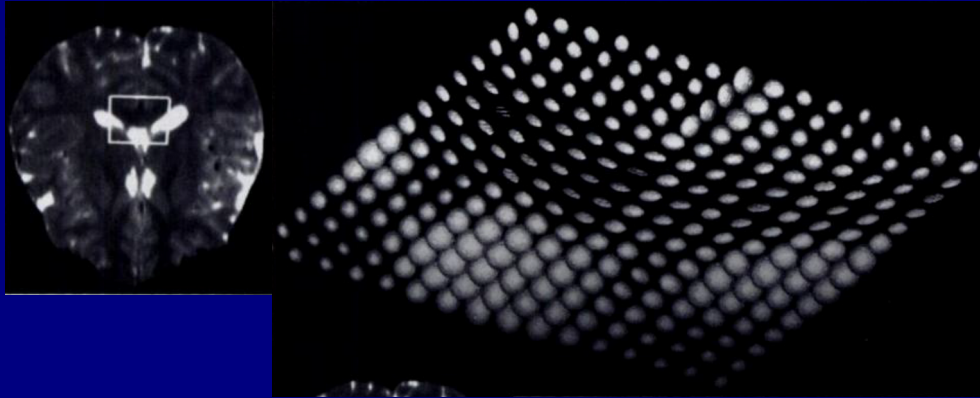


\rightarrow individual ellipsoids

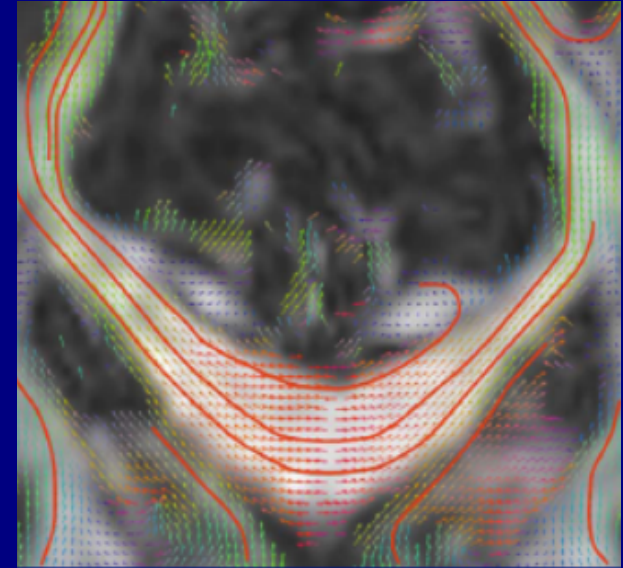


Local DTs \rightarrow extended tracts

Field of local diffusion parameters



Connect to form extended tracts



\rightarrow individual ellipsoids

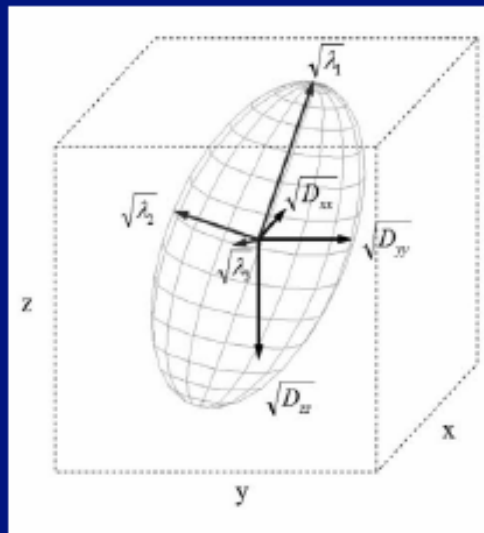


\rightarrow linked structures

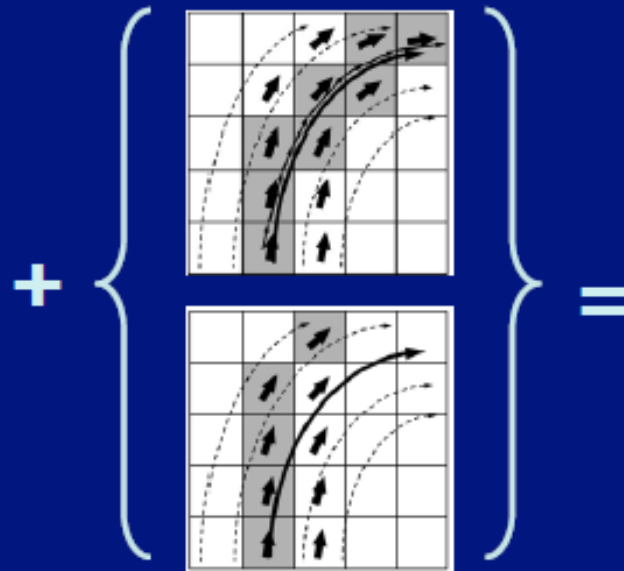


Tractography

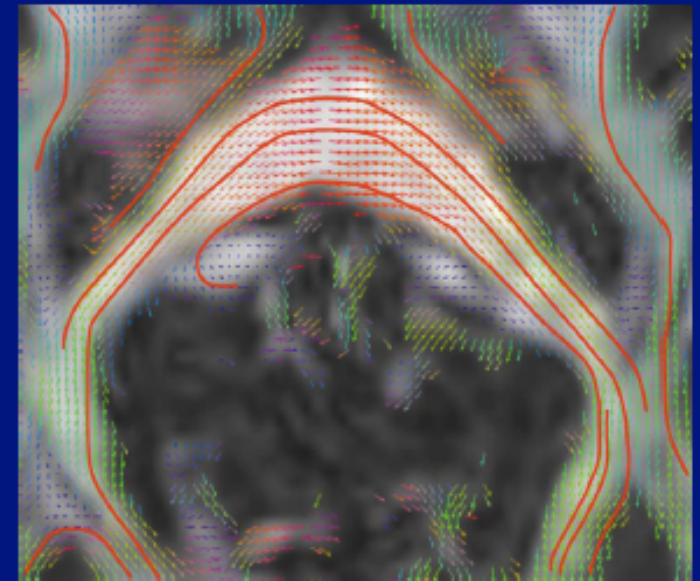
Estimate WM structure (fiber tract locations)



ellipsoid measures
(~*smoothing of
real structures*)



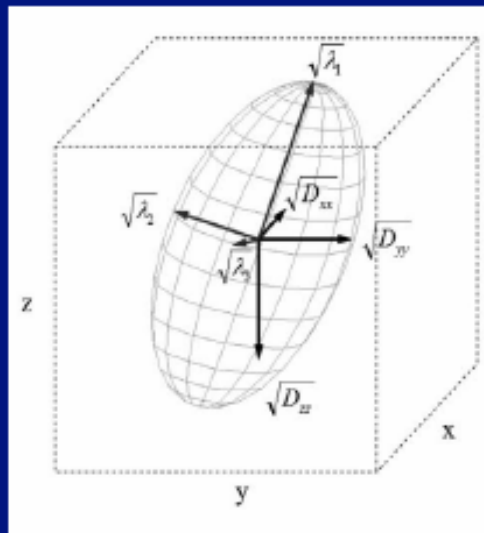
some kind of algorithm
for connecting



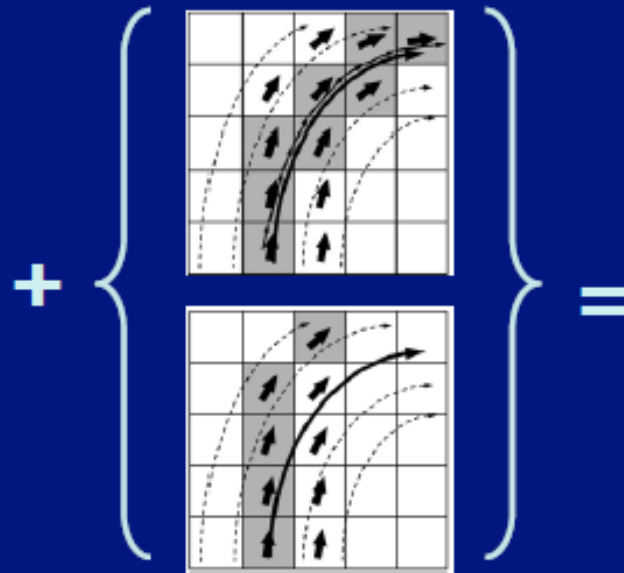
estimate spatial
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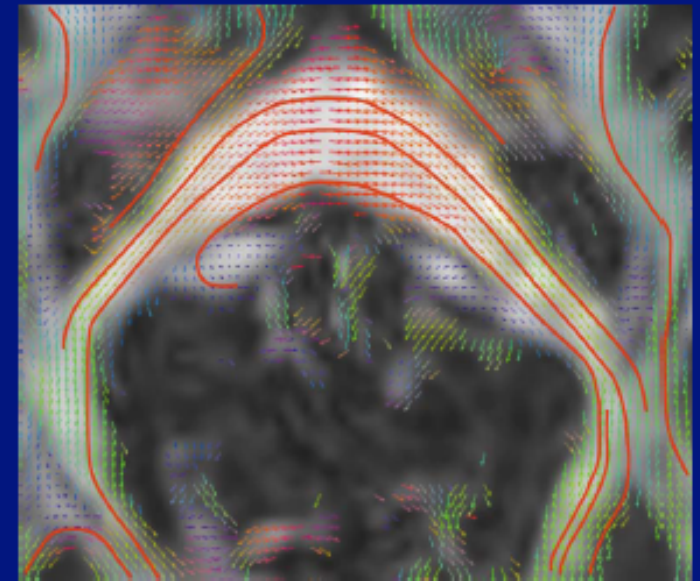
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Tracking WM fibers

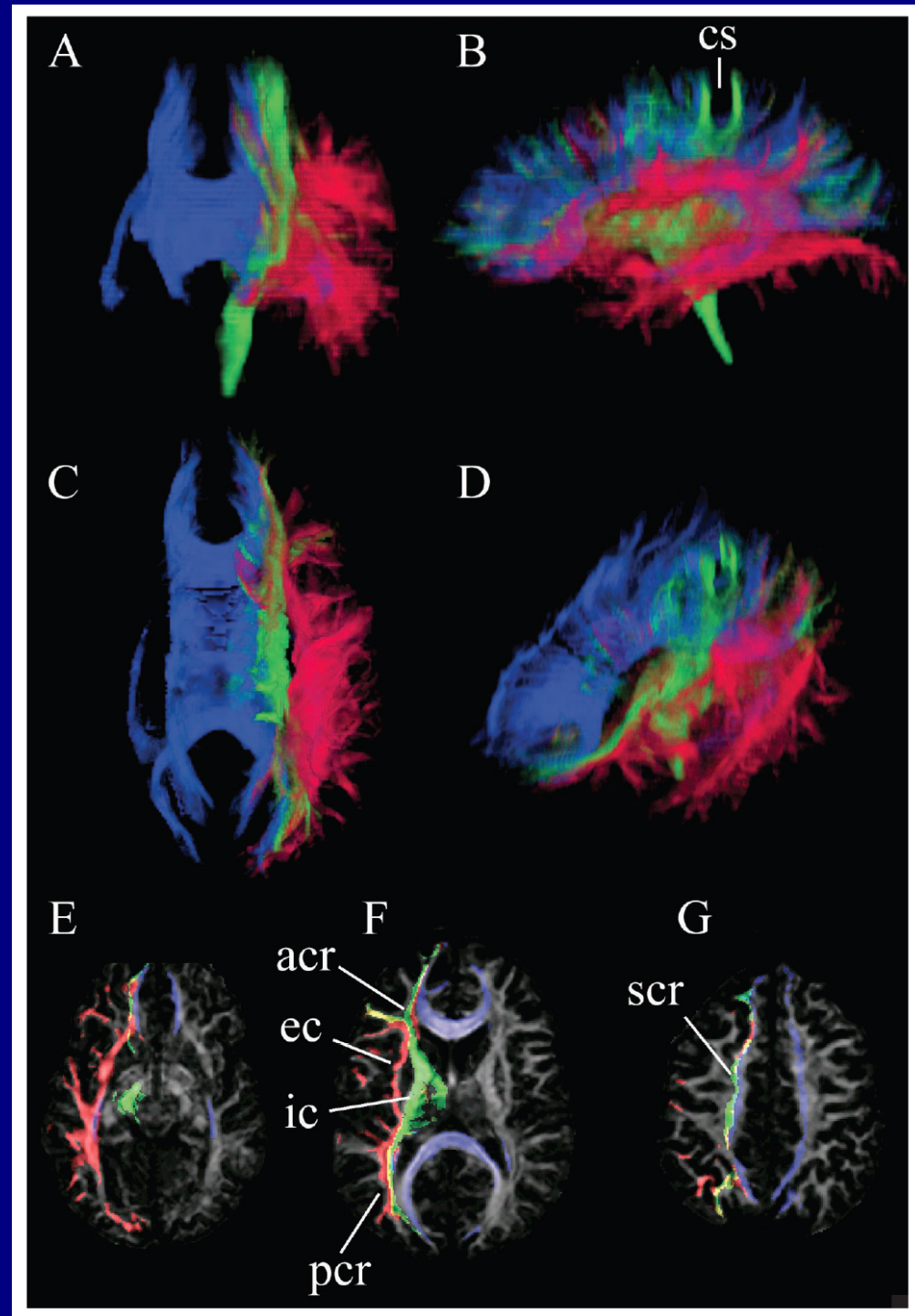
Tracking can be a useful alternative to maps/atlas for finding characteristic subsets, families or networks of the same WM bundles within each subject, for example¹:

Transcallosal
Projection
Association

Here, we use the FATCAT² tracking tools available in AFNI.

→ see Saad et al. poster
#3543 W/Th

¹Wakana, et al., 2004; ²Taylor & Saad, 2013



The subjects

- Nonsedated newborn subjects (<47 days after birth), same community
 - 11 PAE (6 female, 5 male)
postconception age range 36-44 wk (median 42 wk)
 - 9 HC (3 female, 6 male)
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¹May et al., 2007

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 - communities around wine regions have some of highest rates of alcoholism/PAE in the world¹
- Characteristics from followback interviews² and checkups include:
 - maternal detailed drinking, cigarette use, age at scan.
 - infant postconception age, sex, intracranial volume.

¹May et al., 2007; ²Jacobson et al., 2002

The scanning

- 3T Siemens Allegra + 170.9 mm circ. polarized birdcage RF coil
- Diffusion weighted imaging (DWI)
 - 2 mm isotropic voxels, whole brain coverage
 - Twice-refocused SE-EPI sequence
 - Two DWI sets with opposite phase (AP/PA) encoding
+ each: 4 $b=0$ and 30 $b=1000$ s mm^{-2} images
 - Processing included:
motion correction using FSL, susceptibility-distortion
correction^{1,2}, outlier rejection.
- Anatomical images
 - 1 mm isotropic voxels, whole brain coverage
 - multiecho FLASH sequence³
 - T1 and PD maps generated with Freesurfer-mri_ms_fitparms

¹Andersson et al., 2003; ²Rohde et al., 2004; ³van der Kouwe et al., 2008

The measures

fractional anisotropy, FA

mean diffusivity, MD

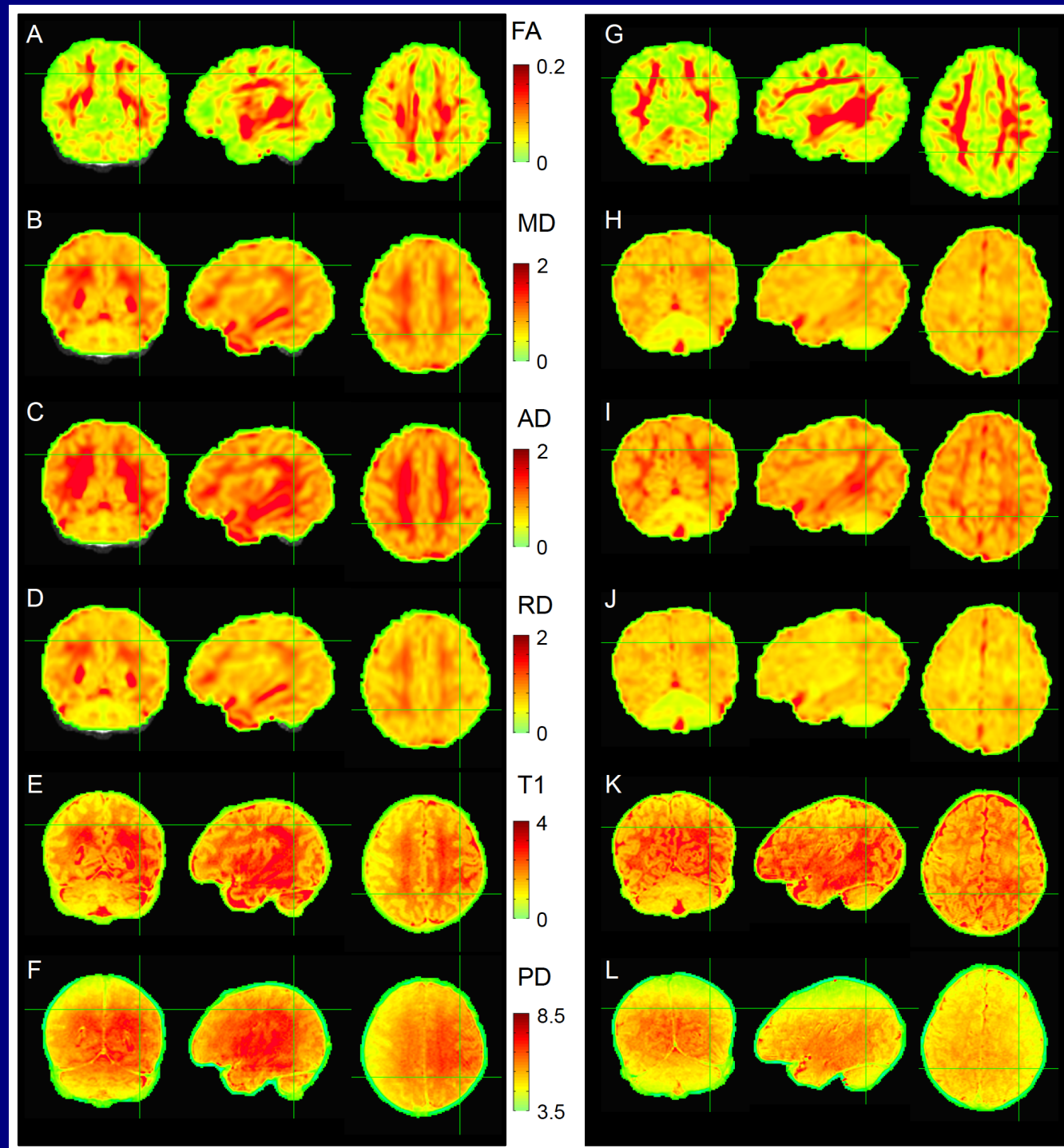
axial diffusivity, AD
(i.e., L1)

radial diffusivity, RD

T1 relaxation time, T1

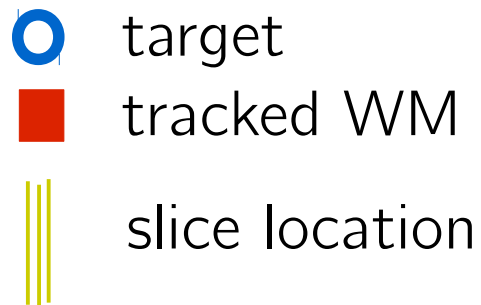
proton density, PD

...

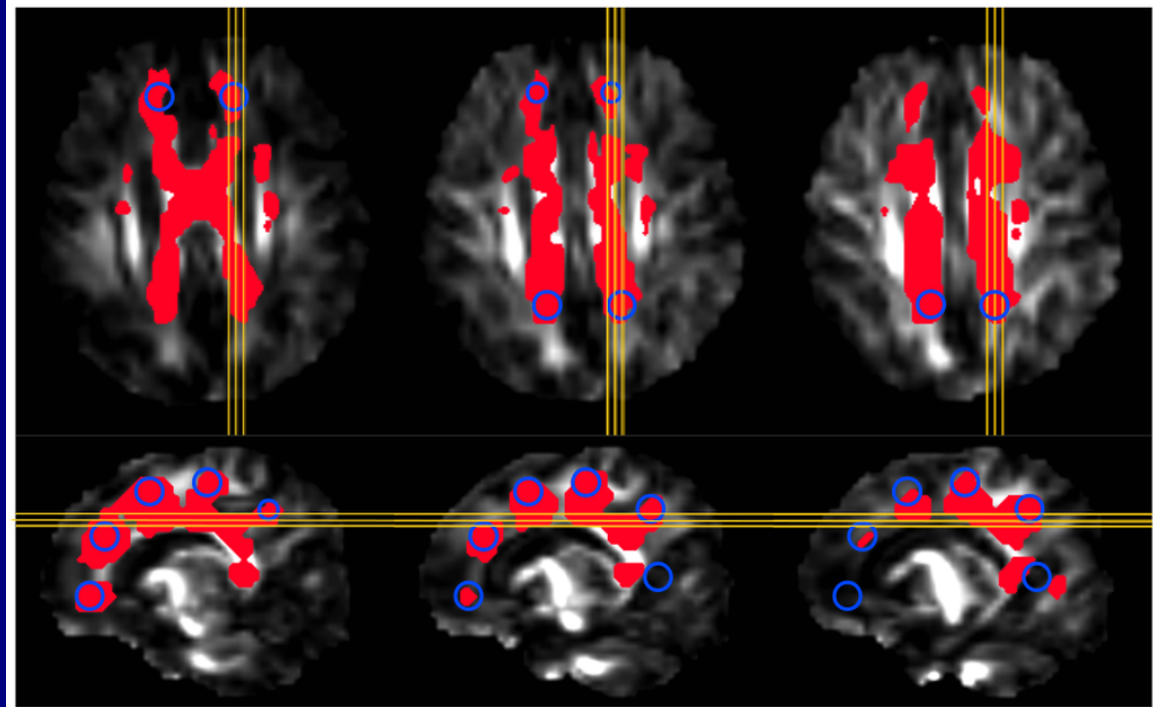


The measures

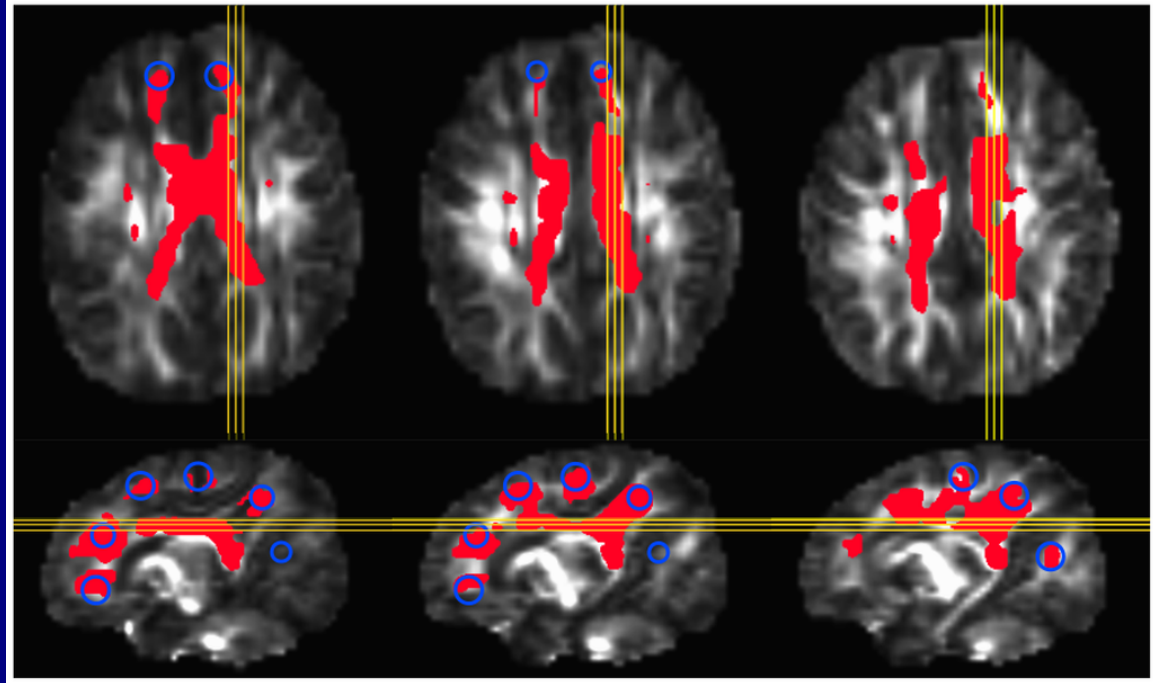
... and normalized volumes
(= the number of voxels in a
WM ROI connecting a target pair
divided by subject intracranial
volume)



HC



PAE



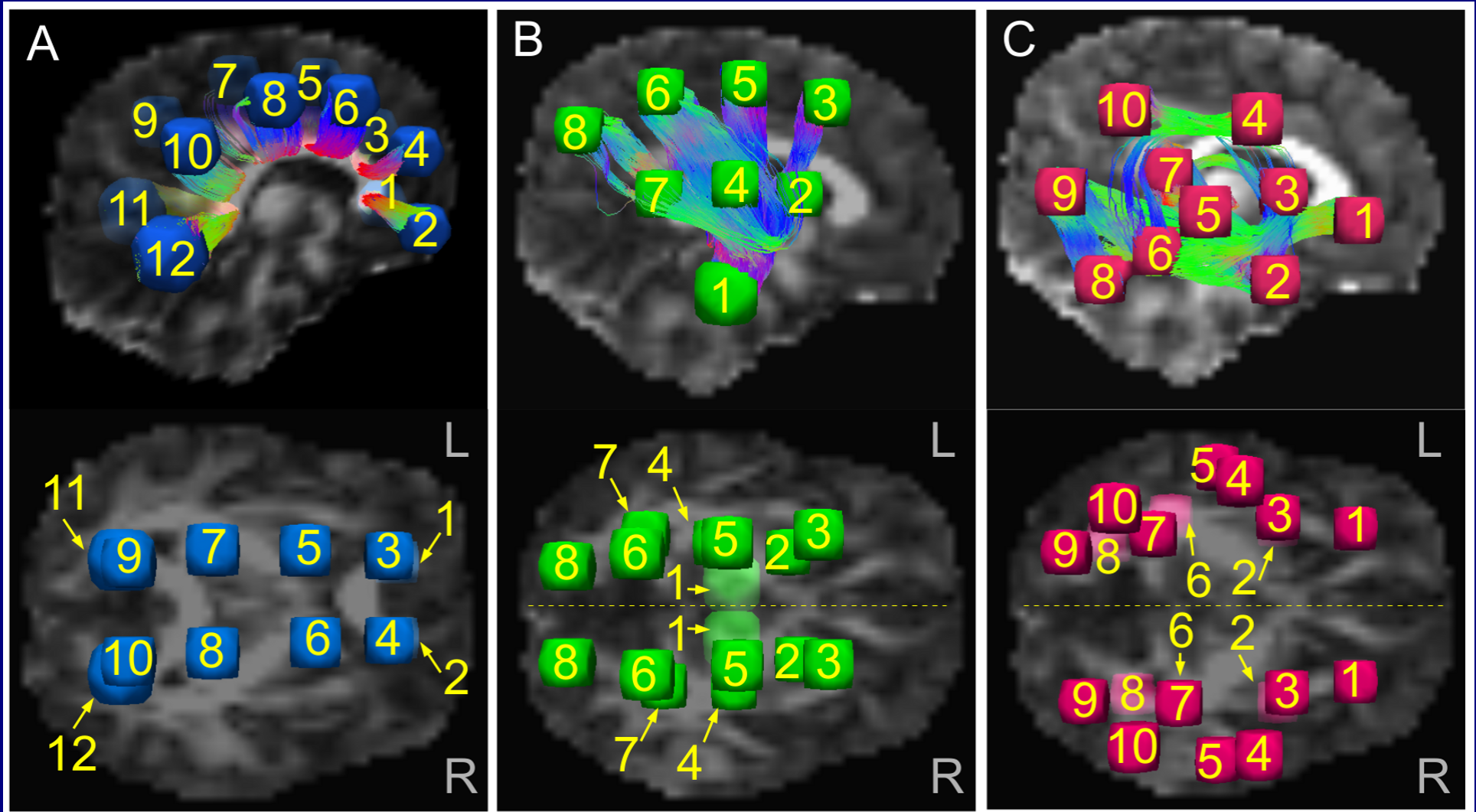
Setting up DTI-tractography

Location of targets for tractography: 5 WM networks.

CC and Cor. Rad.
(CCCR)

Projection
(L/R-PROJ)

Association
(L/R-ASSOC)



Analysis Steps

1) Place network targets

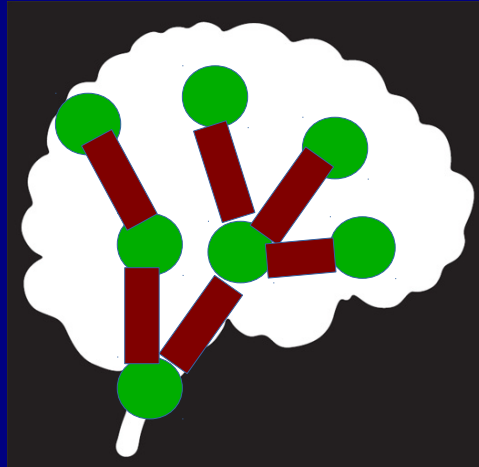


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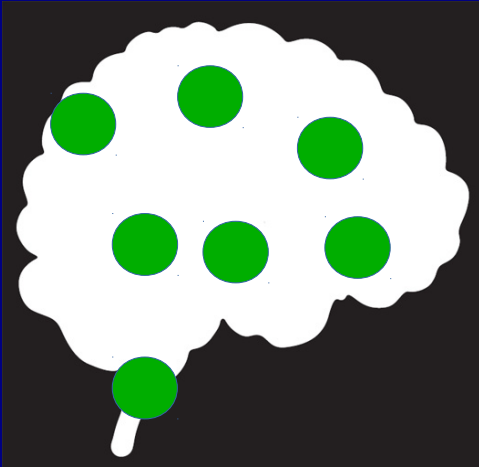


2) Probabilistic tracking

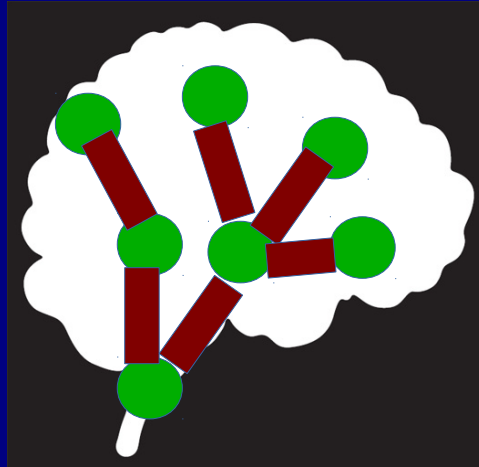


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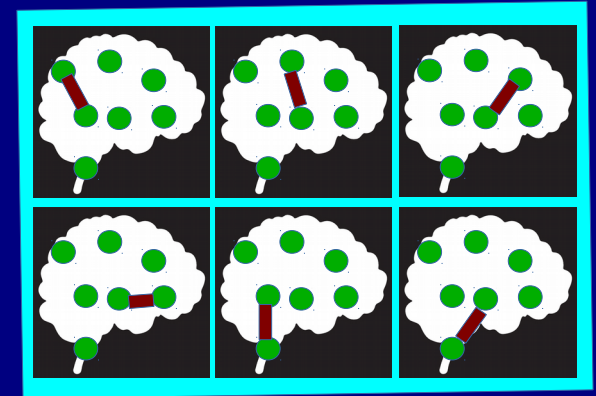
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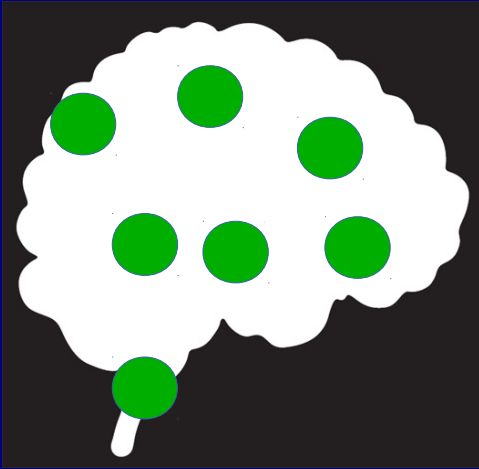


3) set of WM ROIs \rightarrow set of repeated measures

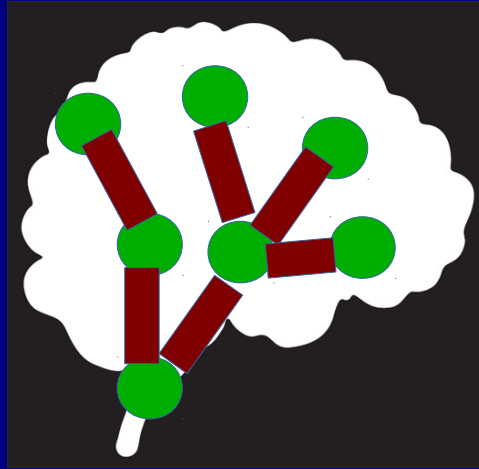


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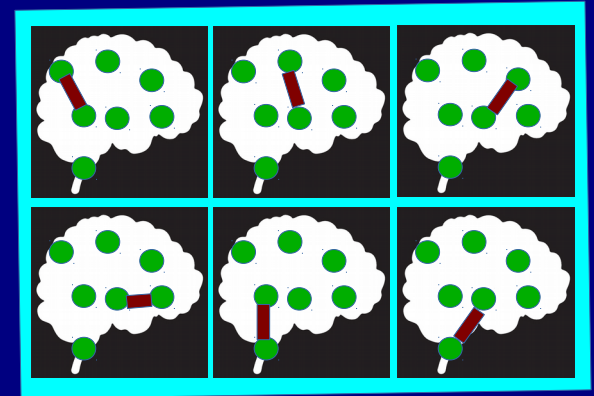
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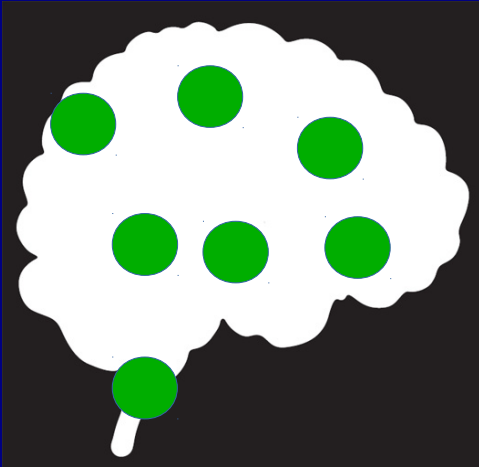
4) Multivariate model

- FA_1, FA_2, FA_3, \dots
- alc
- infant age
- infant sex
- maternal age
- maternal cig/day

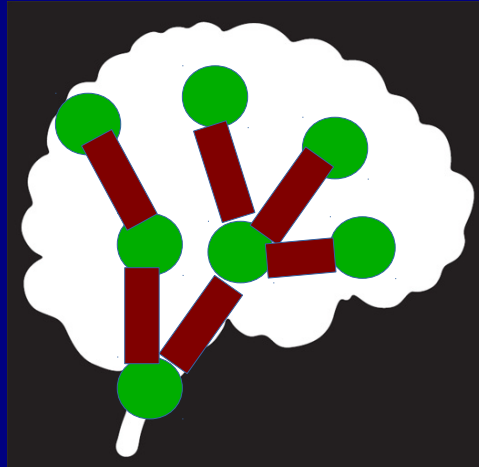
→ see Chen et al. poster #3606 W/Th

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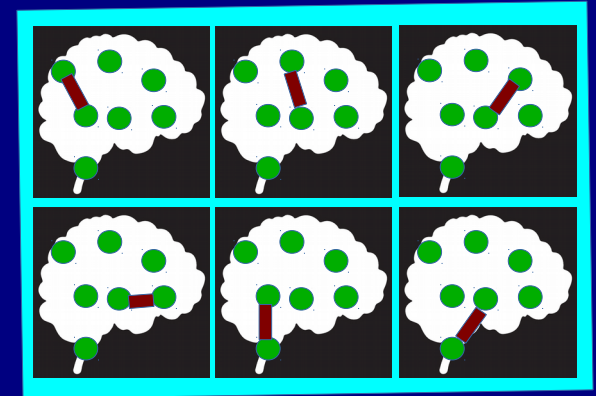
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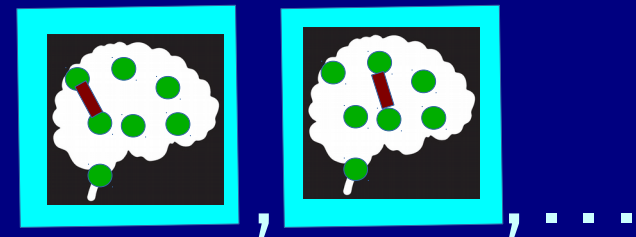


4) Multivariate model

- FA_1, FA_2, FA_3, \dots
- alc
- infant age
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- maternal age
- maternal cig/day

5) Follow-up GLM for each WM ROI

- FA
- alc
- infant age
- infant sex
- maternal age
- maternal cig/day



→ see Chen et al. poster #3606 [W/Th](#)

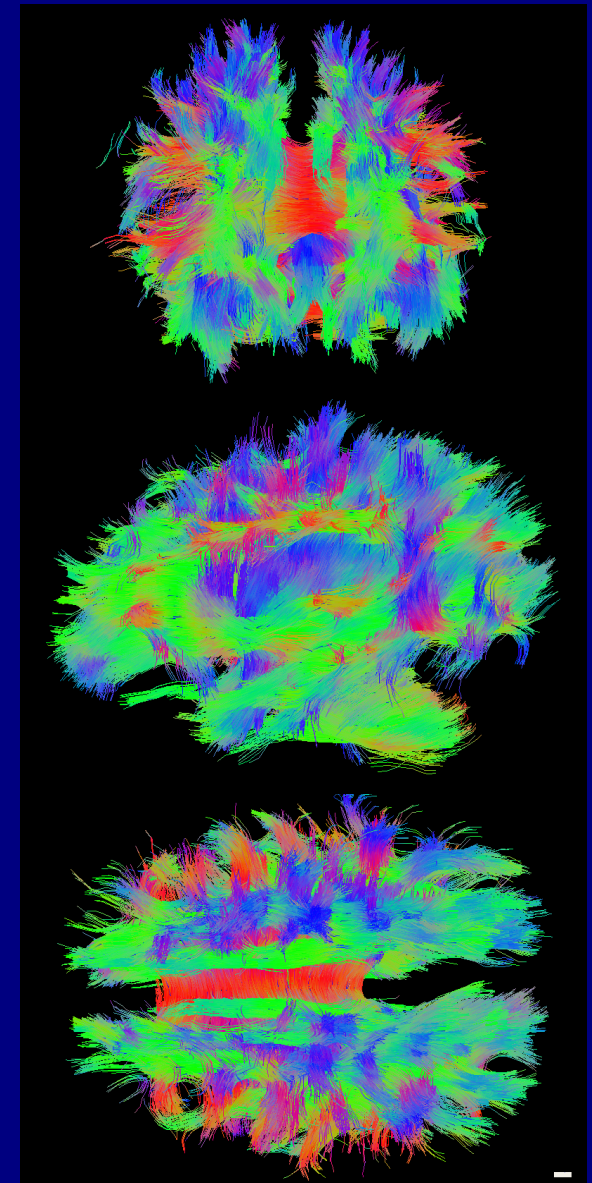
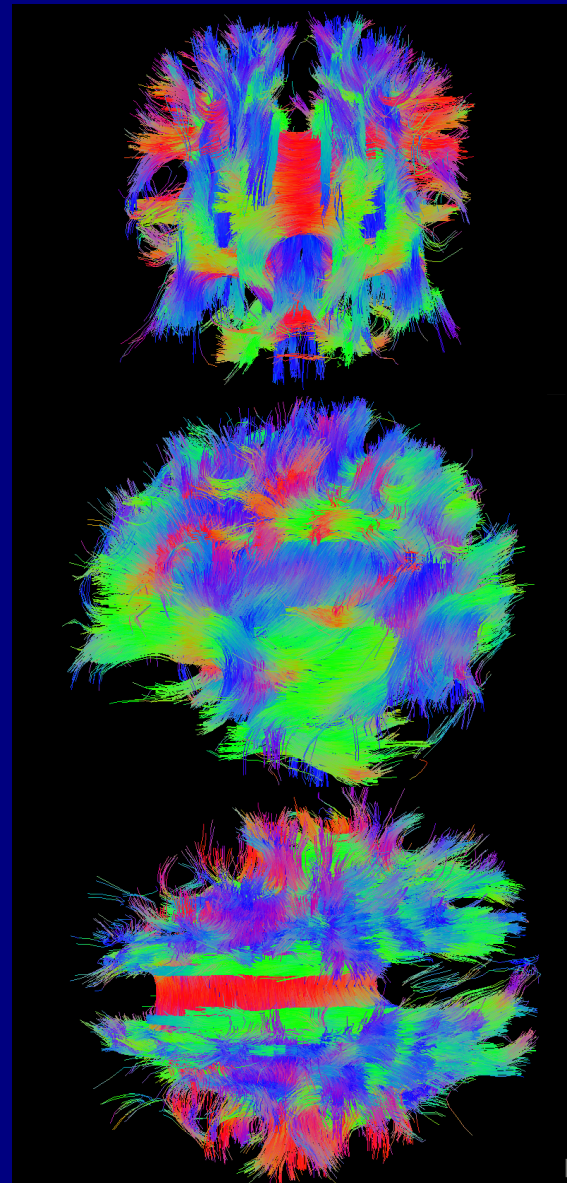
I) Results: whole brain

HC

PAE

A) Preliminary whole brain tracking:

- no major obvious differences
- no missing regions



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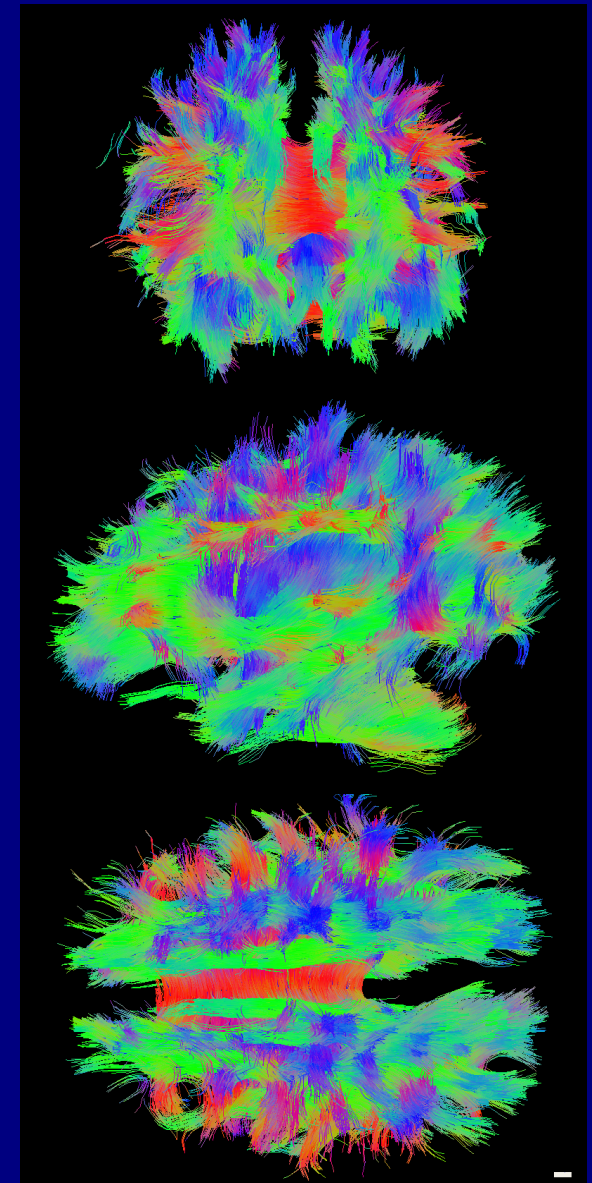
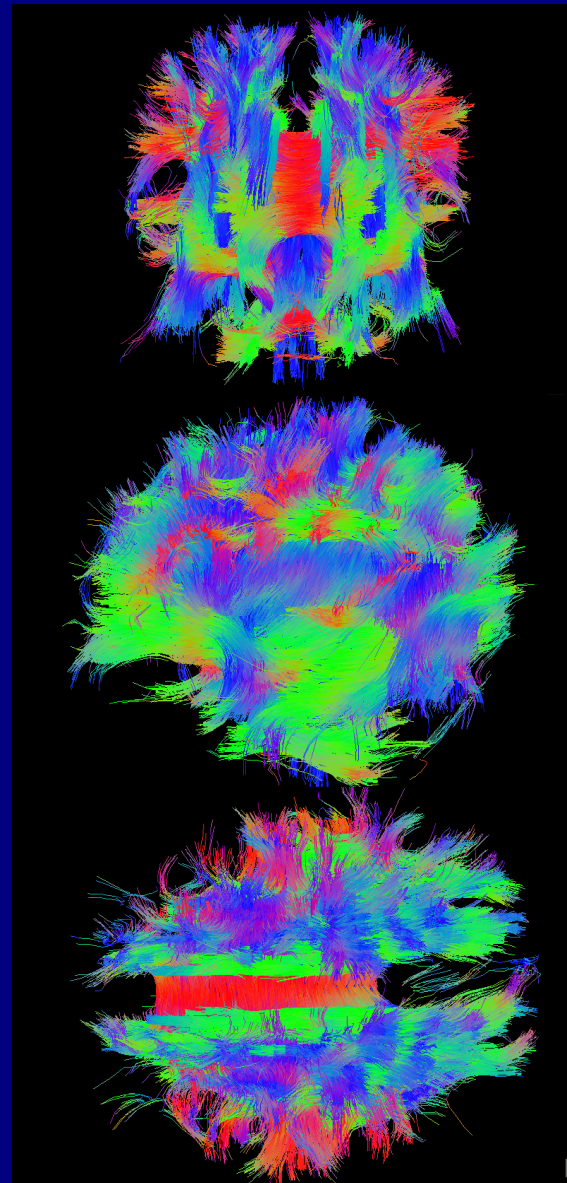
- no major obvious differences
- no missing regions

B) Brain volume and WM volume strongly associated with age.

C) Brain volume associated (negatively) with alcohol exposure

HC

PAE



II) Results: network level

The questions:

- 1) which WM networks are affected by PAE?
- 2) which parameters show effects most strongly?

Answer using:

- (for each network) a multivariate GLM for
 - set of DTI parameters
 - alcohol (frequency: binge/wk)
 - infant age (wks since conception)
 - infant sex (M/F)
 - maternal age (yrs)
 - maternal cigarette smoking (cig/day).

II) Results: network level

The questions:

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Parameters showing at least trends ($p < 0.1$) →

← Networks

| Network | FA | | | | MD | | | | AD | | | | PD | | | |
|---------|---------|---------------|-----------------|-------|---------|---------------|-----------------|----------|---------|---------------|-----------------|----------|------|---------------|-----------------|-------|
| | var. | β_{med} | $F(df_N, df_D)$ | p | var. | β_{med} | $F(df_N, df_D)$ | p | var. | β_{med} | $F(df_N, df_D)$ | p | var. | β_{med} | $F(df_N, df_D)$ | p |
| CCCR | | | | | alc | -0.70 | 8.6 (1, 14) | 0.011* | alc | -0.72 | 14.0 (1, 14) | 0.002** | cig | 0.47 | 3.5 (1, 14) | 0.083 |
| L-PROJ | cig | 0.12 | 4.2 (11, 4) | 0.091 | mat_age | 0.56 | 5.5 (1, 14) | 0.034* | mat_age | 0.53 | 6.3 (1, 14) | 0.025* | cig | 0.52 | 4.0 (1, 14) | 0.066 |
| | | | | | alc | -0.41 | 3.9 (10, 140) | 0.000*** | alc | -0.52 | 4.1 (10, 140) | 0.000*** | | | | |
| R-PROJ | age | 0.33 | 8.6 (13, 2) | 0.109 | mat_age | 0.37 | 4.4 (1, 14) | 0.056 | mat_age | 0.44 | 6.5 (1, 14) | 0.023* | cig | 0.48 | 3.4 (1, 14) | 0.085 |
| | | | | | alc | -0.41 | 1.9 (12, 168) | 0.035* | alc | -0.45 | 2.7 (12, 168) | 0.002** | | | | |
| | | | | | age | -0.41 | 5.8 (1, 14) | 0.031* | age | -0.39 | 5.3 (1, 14) | 0.038* | | | | |
| L-ASSOC | sex | | | | sex | -0.20 | 4.3 (1, 14) | 0.056 | sex | -0.39 | 5.9 (1, 14) | 0.029* | cig | 0.49 | 3.6 (1, 14) | 0.080 |
| | mat_age | -0.16 | 9.2 (13, 2) | 0.103 | alc | -0.65 | 6.0 (7, 8) | 0.011* | alc | -0.66 | 8.1 (1, 14) | 0.013* | | | | |
| R-ASSOC | alc | 0.23 | 1.8 (7, 98) | 0.090 | mat_age | 0.44 | 3.8 (1, 14) | 0.071 | age | -0.16 | 2.5 (6, 84) | 0.030* | | | | |
| | | | | | alc | -0.62 | 10.2 (1, 14) | 0.007** | mat_age | 0.43 | 4.7 (1, 14) | 0.048* | cig | 0.5 | 3.5 (1, 14) | 0.082 |
| | | | | | | | | | cig | -0.29 | 3.9 (1, 14) | 0.068 | | | | |

* $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

II) Results: network level

The questions:

- 1) which WM networks are affected by PAE?
- 2) which parameters show effects most strongly?

Parameters showing at least trends ($p < 0.1$) →

← Networks

| | FA | | | | MD | | | | AD | | | | PD | | | |
|----------------|---------|---------------|-----------------|-------|---------|---------------|-----------------|-----------------|---------|---------------|-----------------|-----------------|------|---------------|-----------------|-------|
| Network | var. | β_{med} | $F(df_N, df_D)$ | p | var. | β_{med} | $F(df_N, df_D)$ | p | var. | β_{med} | $F(df_N, df_D)$ | p | var. | β_{med} | $F(df_N, df_D)$ | p |
| CCCR | | | | | alc | -0.70 | 8.6 (1, 14) | 0.011* | alc | -0.72 | 14.0 (1, 14) | 0.002** | cig | 0.47 | 3.5 (1, 14) | 0.083 |
| L-PROJ | cig | 0.12 | 4.2 (11, 4) | 0.091 | mat_age | 0.56 | 5.5 (1, 14) | 0.034* | mat_age | 0.53 | 6.3 (1, 14) | 0.025* | | | | |
| | | | | | alc | -0.41 | 3.9 (10, 140) | 0.000*** | alc | -0.52 | 4.1 (10, 140) | 0.000*** | cig | 0.52 | 4.0 (1, 14) | 0.066 |
| | | | | | mat_age | 0.37 | 4.4 (1, 14) | 0.056 | mat_age | 0.44 | 6.5 (1, 14) | 0.023* | | | | |
| R-PROJ | age | 0.33 | 8.6 (13, 2) | 0.109 | alc | -0.41 | 1.9 (12, 168) | 0.035* | alc | -0.45 | 2.7 (12, 168) | 0.002** | cig | 0.48 | 3.4 (1, 14) | 0.085 |
| | mat_age | -0.16 | 9.2 (13, 2) | 0.103 | age | -0.41 | 5.8 (1, 14) | 0.031* | age | -0.39 | 5.3 (1, 14) | 0.038* | | | | |
| | | | | | sex | -0.20 | 4.3 (1, 14) | 0.056 | sex | -0.39 | 5.9 (1, 14) | 0.029* | | | | |
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| | | | | | mat_age | 0.44 | 3.8 (1, 14) | 0.071 | age | -0.16 | 2.5 (6, 84) | 0.030* | | | | |
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* $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

→ Statistically significant alcohol exposure associations in ~every WM network

II) Results: network level

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Parameters showing at least trends ($p < 0.1$) →

← Networks

| Network | var. | FA β_{med} | $F(df_N, df_D)$ | p | var. | MD β_{med} | $F(df_N, df_D)$ | p | var. | AD β_{med} | $F(df_N, df_D)$ | p | var. | PD β_{med} | $F(df_N, df_D)$ | p |
|---------|---------|--------------------------------|-----------------|-------|---------|---------------------|-----------------|----------|---------|---------------------|-----------------|----------|------|---------------------|-----------------|-------|
| CCCR | | | | | alc | -0.70 | 8.6 (1, 14) | 0.011* | alc | -0.72 | 14.0 (1, 14) | 0.002** | cig | 0.47 | 3.5 (1, 14) | 0.083 |
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| | | | | | alc | -0.41 | 3.9 (10, 140) | 0.000*** | alc | -0.52 | 4.1 (10, 140) | 0.000*** | cig | 0.52 | 4.0 (1, 14) | 0.066 |
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| | | | | | | | | | cig | -0.29 | 3.9 (1, 14) | 0.068 | | | | |

* $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

→ Increased alcohol exposure:
decreased AD
(and decreased MD)






III) Results: ROI level

The question:

1) where are most significant AD-alcohol relations in each network?

Answer using:

- (for each ROI) a GLM for
 - single DTI parameter
 - alcohol (frequency: binge/wk)
 - infant age (wks since conception)
 - infant sex (M/F)
 - maternal age (yrs)
 - maternal cigarette smoking (cig/day).

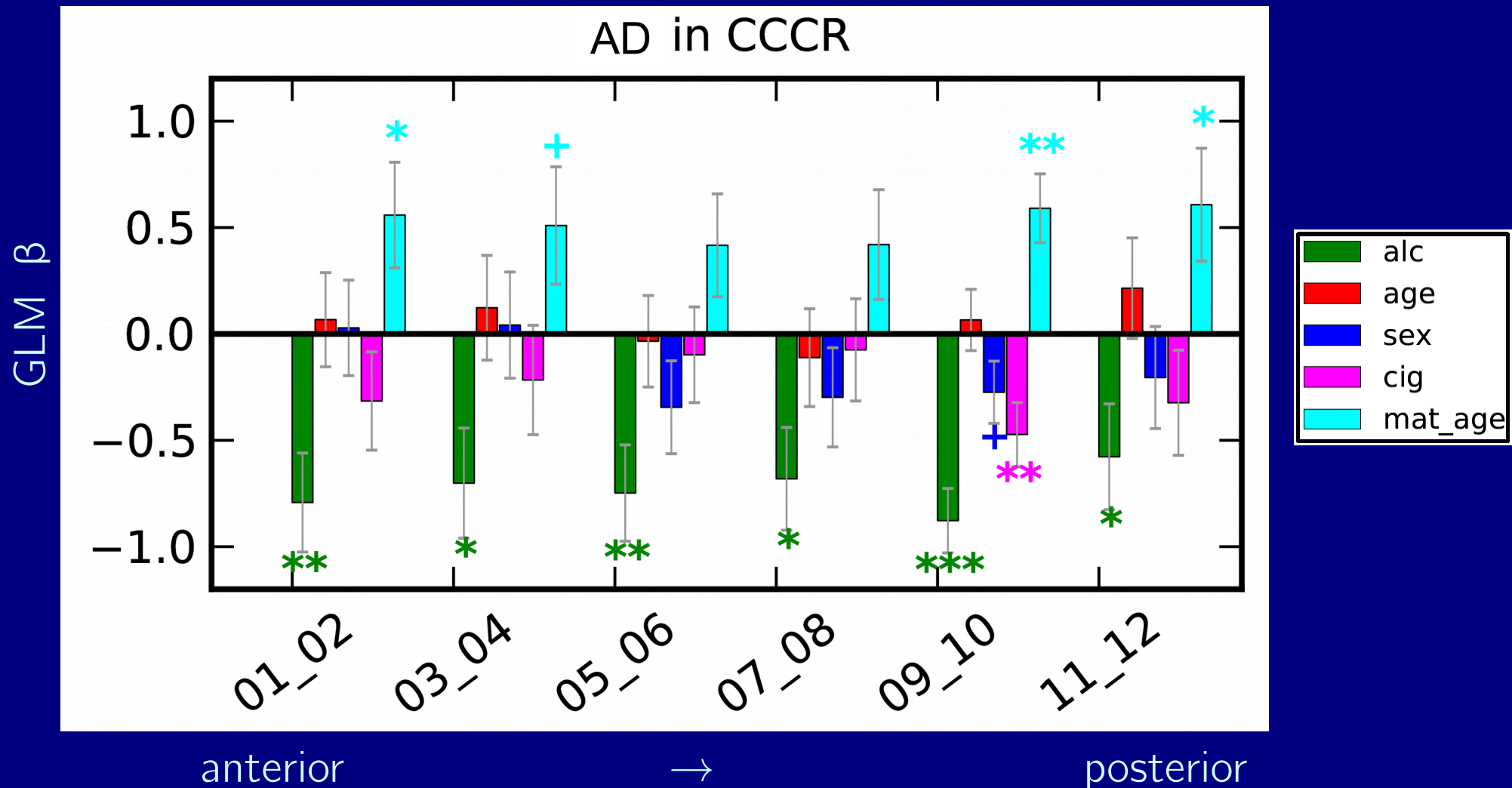
| | |
|---------------------------------------------------------------------------------------|---------|
|  | alc |
|  | age |
|  | sex |
|  | cig |
|  | mat_age |

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The question:

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Transcallosal (CC and corona radiata)

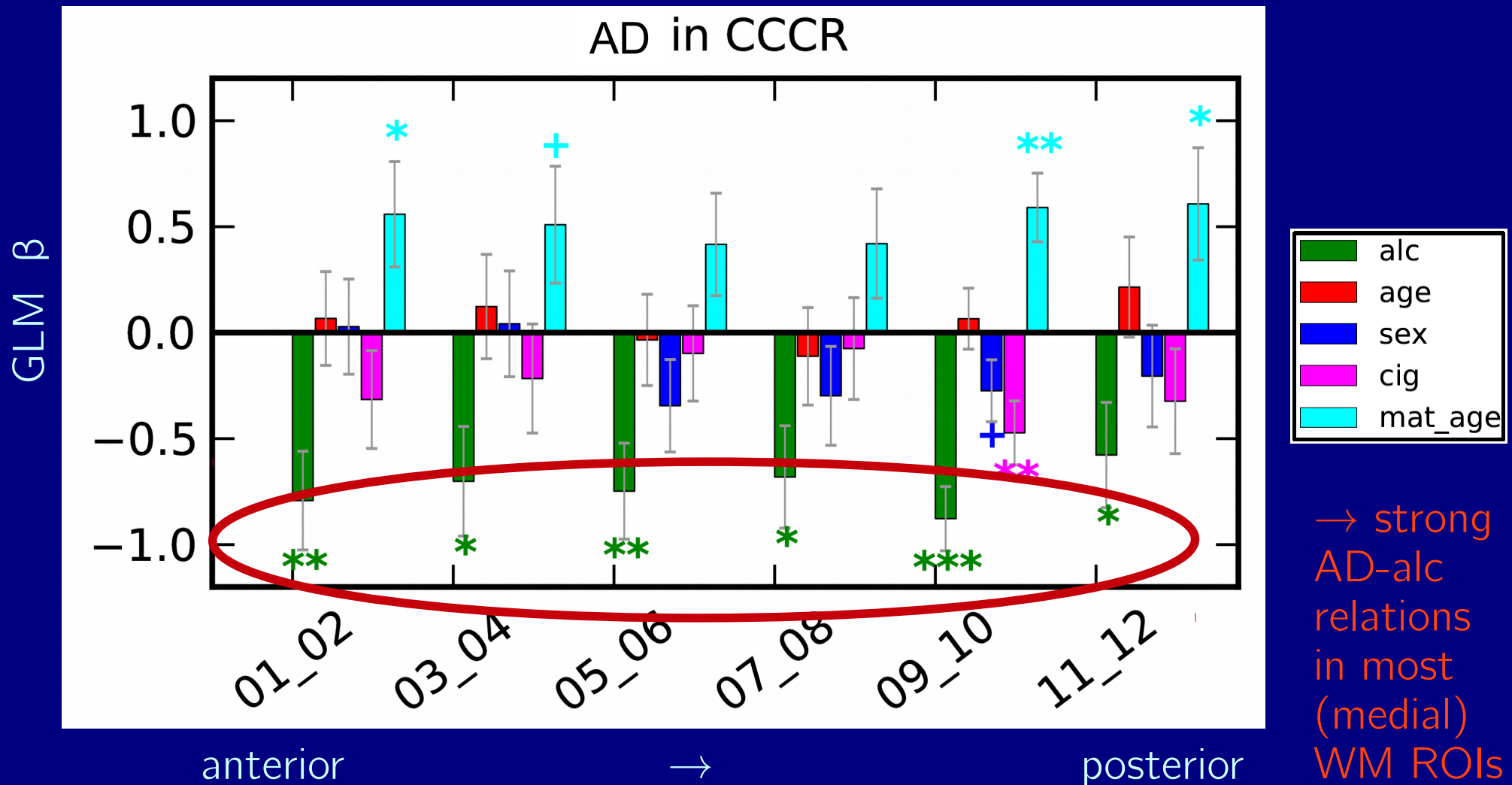


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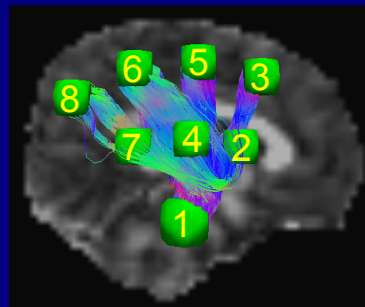
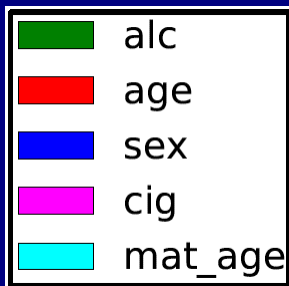
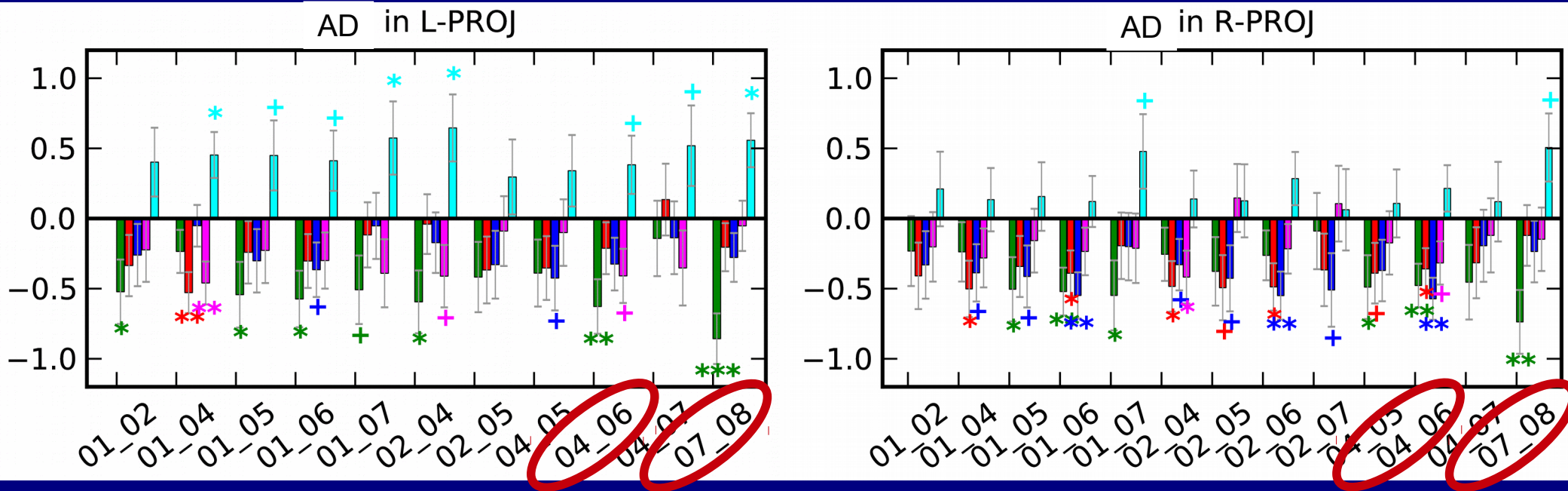


III) Results: ROI level

The question:

1) where are most significant AD-alcohol relations in each network?

L and R Projection



→ strong (symmetric)
AD-alc relations in
posterior WM ROIs

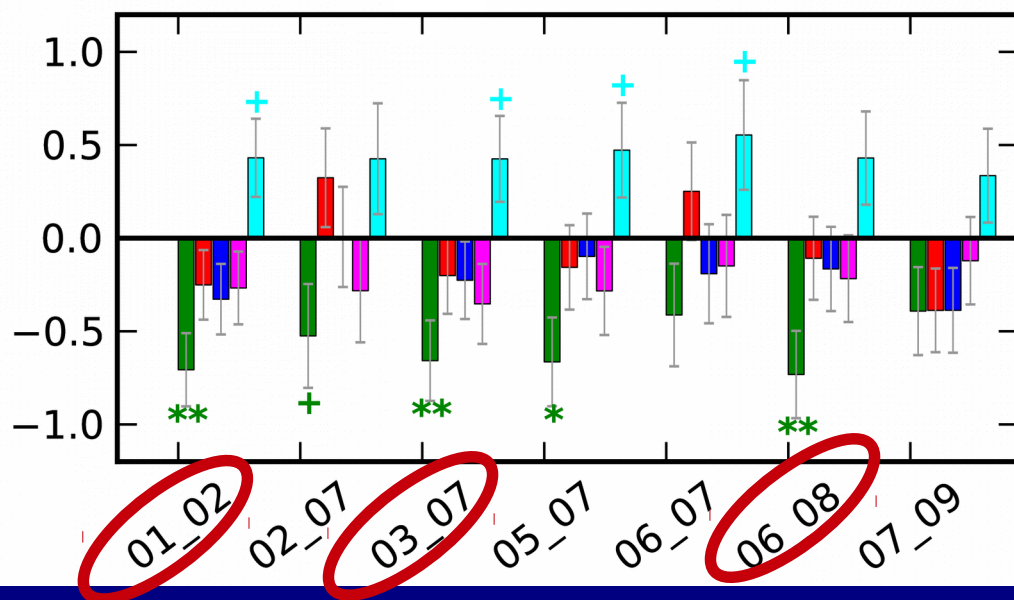
III) Results: ROI level

The question:

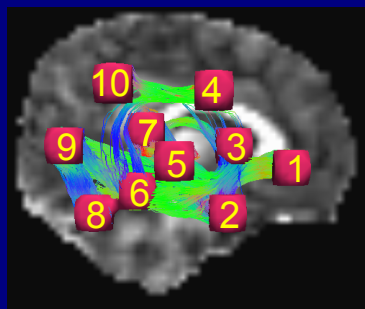
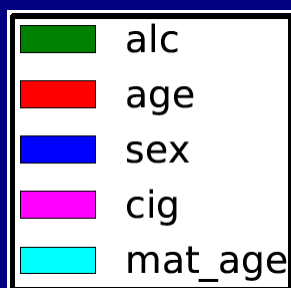
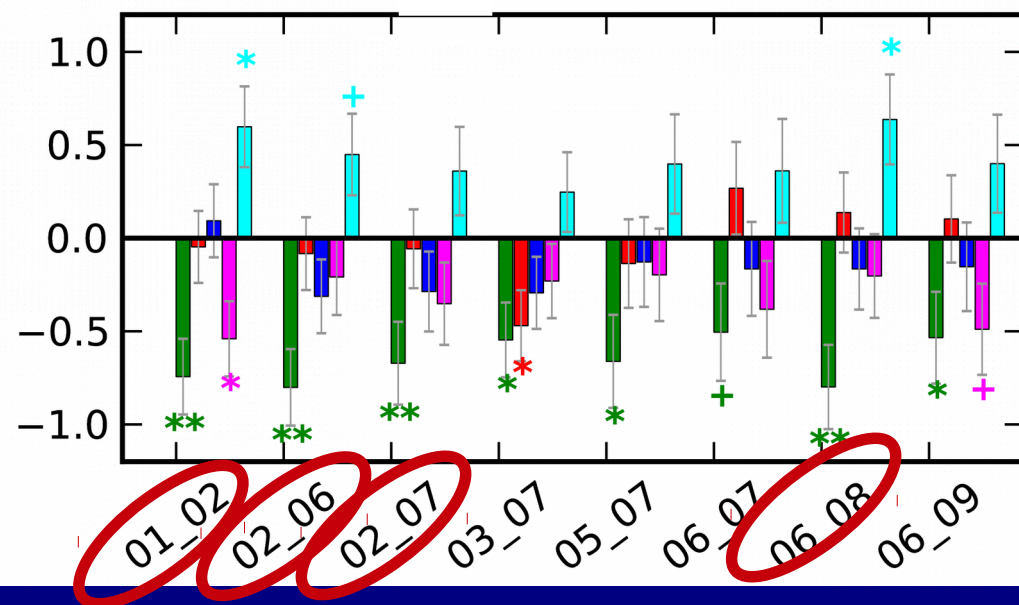
1) where are most significant AD-alcohol relations in each network?

L and R Association

AD in L-ASSOC



AD in R-ASSOC



→ strong AD-alc relations in (symmetric) ~medial or inferior WM ROIs

Conclusions

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 - likely due to locations of early WM maturation/myelination
- decreased AD has been associated with WM inflammation and damage in animal models; disruption of linear neurofilaments; reduction of fast transport; and axolemmal atrophy^{1,2}.

Conclusions

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 - decreased AD has been associated with WM inflammation and damage in animal models; disruption of linear neurofilaments; reduction of fast transport; and axolemmal atrophy^{1,2}.
- Further work will continue with more newborn subjects and investigating, e.g., lateralization; as well as with older (pediatric) subjects to investigate developmental WM trajectories.

¹*Wu et al., 2007; Beaulieu et al. 2002*

Many thanks to all study coauthors:

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Pia Wintermark

NIMH/NIH, USA

Gang Chen

Wayne State University, USA

Sandra W. Jacobson

Joseph L. Jacobson

Extra info.

| | HC (<i>n</i> = 9) | | PAE (<i>n</i> = 11) | | <i>p</i> -value |
|----------------------------------------------|--------------------|-------|----------------------|-------|-----------------|
| | Mean/% | SD | Mean/% | SD | |
| <i>Infant characteristics</i> | | | | | |
| Sex (% female) | 33.3 | | 54.5 | | 0.075 |
| Gestational age at birth (wk) | 38.7 | 1.9 | 38.6 | 2.0 | 0.939 |
| Postpartum age at scan (wk) | 3.0 | 1.6 | 2.6 | 1.5 | 0.589 |
| Postconception age at scan (wk) | 41.6 | 2.1 | 41.2 | 2.3 | 0.658 |
| Birthweight (g) | 2753.3 | 478.4 | 2754.1 | 413.5 | 0.997 |
| <i>Maternal characteristics</i> | | | | | |
| Maternal age at delivery (yr) | 24.0 | 4.7 | 29.8 | 4.6 | 0.012 * |
| Parity | 1.2 | 1.2 | 2.3 | 1.5 | 0.105 |
| Education [†] (yr) | 10.6 | 0.9 | 9.7 | 1.0 | 0.807 |
| Marital status (% married) | 33.3 | | 18.2 | | 0.176 |
| Smoking (cig/day) | 4.3 | 2.5 | 6.5 | 4.8 | 0.218 |
| Marijuana (days/wk) | 0.0 | 0.1 | 0.2 | 0.5 | 0.509 |
| <i>Extent of alcohol consumption</i> | | | | | |
| <i>At conception</i> | | | | | |
| oz AA/day [‡] | 0.02 | 0.05 | 2.25 | 1.89 | 0.002 ** |
| oz AA/occasion [‡] | 0.13 | 0.38 | 4.70 | 3.51 | 0.001 ** |
| frequency (days/wk) | 0.02 | 0.05 | 0.42 | 0.30 | 0.001 *** |
| <i>Across pregnancy</i> | | | | | |
| oz AA/day [‡] | 0.01 | 0.03 | 1.68 | 1.52 | 0.004 ** |
| oz AA/occasion [‡] | 0.13 | 0.40 | 4.66 | 1.35 | 0.000 *** |
| frequency (days/wk) | 0.01 | 0.03 | 0.33 | 0.26 | 0.001 ** |
| <i>Infant global parameters</i> | | | | | |
| Number DWIs ^a | 21.7 | 4.4 | 22.4 | 3.6 | 0.698 |
| Total intracranial volume (cm ³) | 469.4 | 60.8 | 434.6 | 43.5 | 0.152 |
| WM volume (cm ³) | 200.7 | 23.0 | 182.9 | 25.6 | 0.123 |
| WM fraction | 0.43 | 0.04 | 0.42 | 0.03 | 0.535 |

HC = healthy control; PAE = prenatal alcohol exposure; SD = standard deviation; AA = absolute alcohol; * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

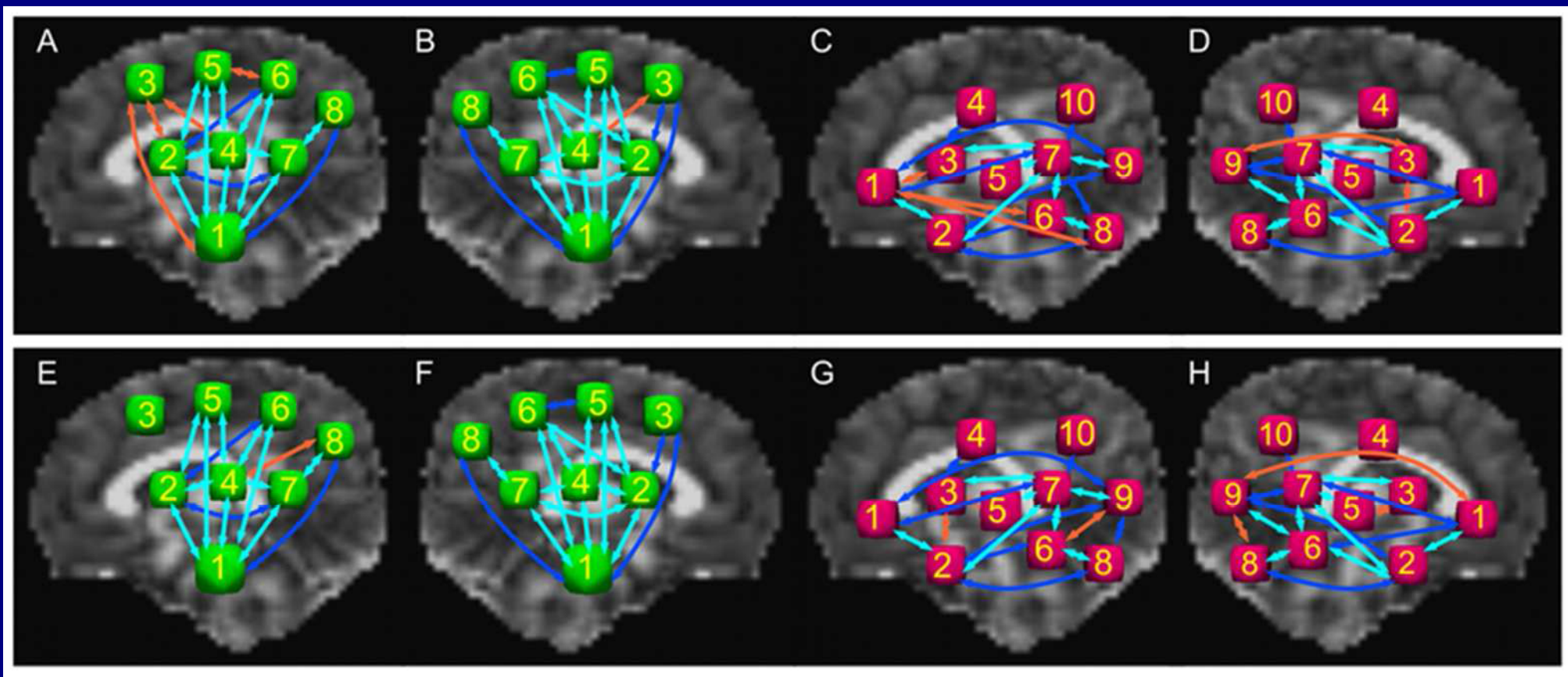
[†]Education missing for one HC mother.

[‡]Measures of oz AA for one PAE mother were Winsorized (values > 3 SD above mean were recoded).

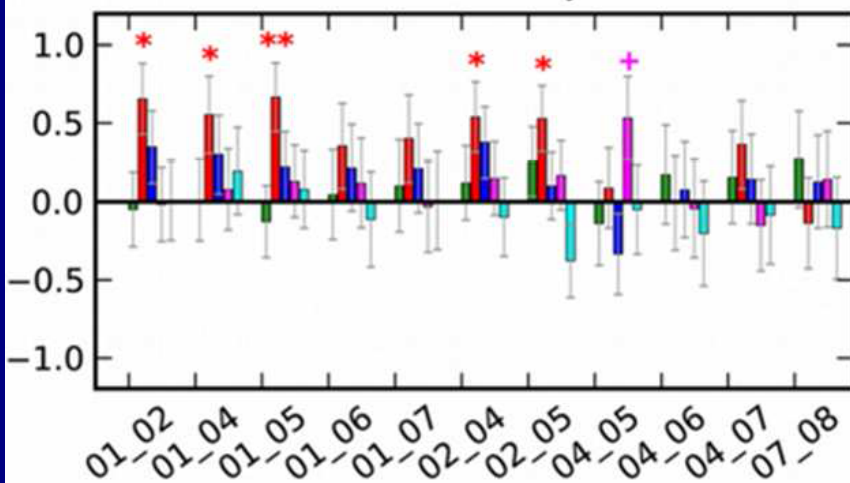
^aNumber of diffusion weighted images (DWIs) after deleting any dropout/motion-corrupted volumes.

| Variable | Predictors | | | | | | | | | |
|--------------|------------|---------|----------|---------|----------|---------|----------|---------|----------|---------|
| | alc | | age | | sex | | cig | | mat_age | |
| | <i>r</i> | β | <i>r</i> | β | <i>r</i> | β | <i>r</i> | β | <i>r</i> | β |
| Brain volume | -0.26 | -0.36* | 0.76*** | 0.77*** | -0.39 | -0.21 | -0.08 | -0.14 | -0.40 | -0.13 |
| WM volume | -0.30 | -0.21 | 0.65** | 0.54* | -0.35 | -0.24 | -0.07 | -0.03 | -0.51* | -0.23 |
| WM fraction | -0.14 | 0.12 | 0.03 | -0.15 | -0.04 | -0.09 | -0.04 | -0.11 | -0.30 | -0.52 |

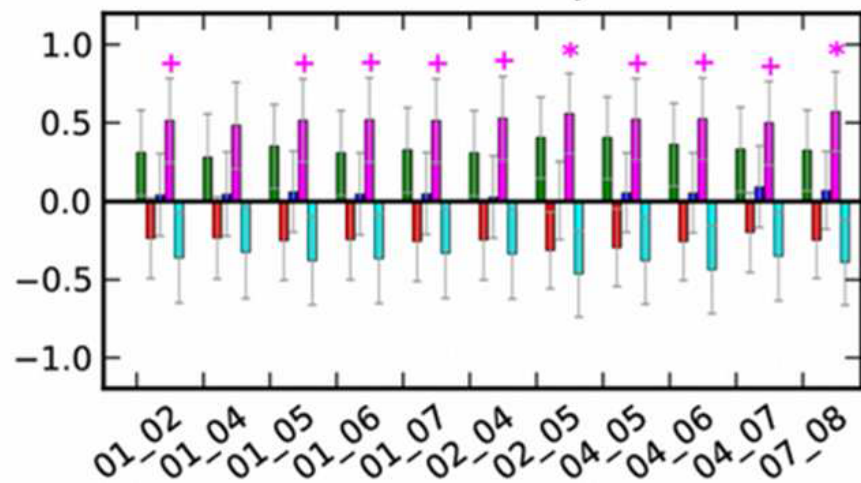
For each predictor, *r* represents the simple correlation between each predictor and the outcome; β is the standardized regression coefficient after adjustment for all the other predictors; * $p<0.05$; ** $p<0.01$; *** $p<0.001$. alc = frequency of maternal drinking; age = infant age since conception; sex: male=0, female=1; cig = maternal smoking (cig/day); mat_age = maternal age at delivery.



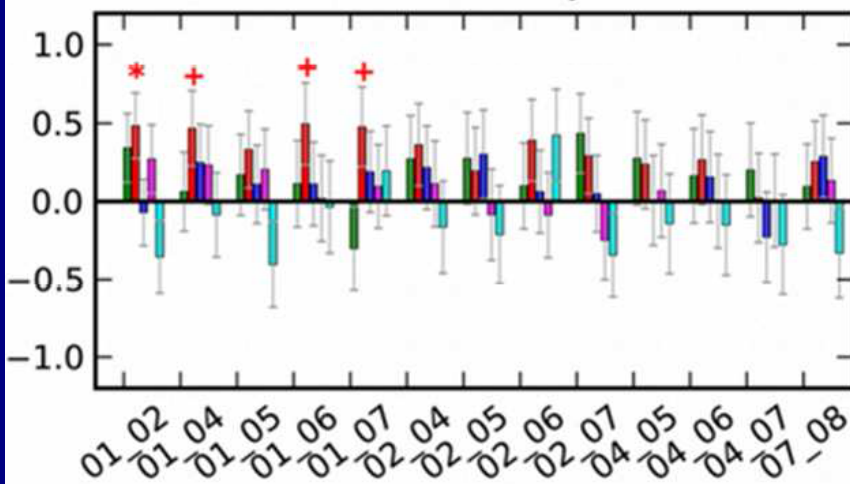
FA in L-PROJ



PD in L-PROJ



FA in R-PROJ



PD in R-PROJ

