

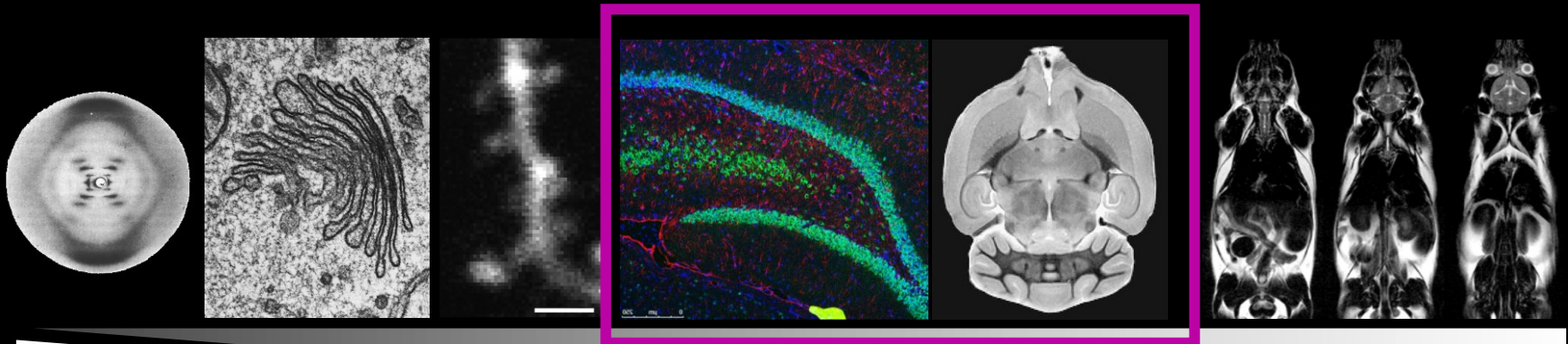


# iDISCO<sup>+</sup>: why, how and troubleshooting

Alba Vieites Prado

*Rockefeller Kavli Institute - iDISCO Tissue Clearing Workshop  
April 24-28<sup>th</sup>, 2023*

# Scales in Biology



**Spatial resolution**

**Sampling size**

How to study cm-large tissues with cellular resolution?

# Transparency & model organisms

Early development in the sea urchin  
(Cebra-Thomas)



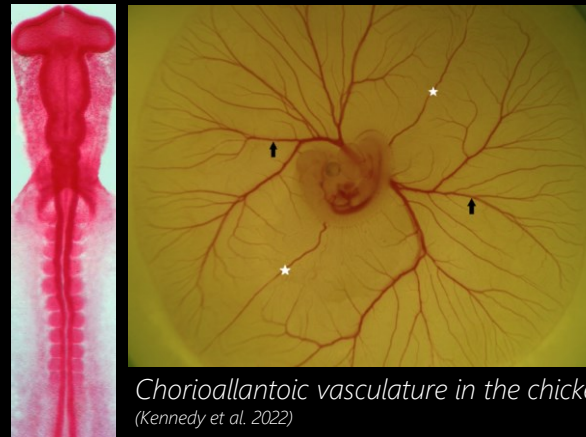
*C. elegans*  
(Mark Leaver)



Ascidians  
(Christian Gloor)



Zebrafish larvae  
(Hill, M.A. Embryology Zebrafish Development)



Chorioallantoic vasculature in the chicken  
(Kennedy et al. 2022)

Chicken embryo  
(Schoenwolf G. 2018)

# Transparency & model organisms

Sea urchin  
(Ann-Cutting)



*C. elegans*  
(Mark Leaver)



Ascidians  
(Christian Gloor)



Zebrafish



Chick

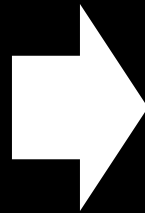
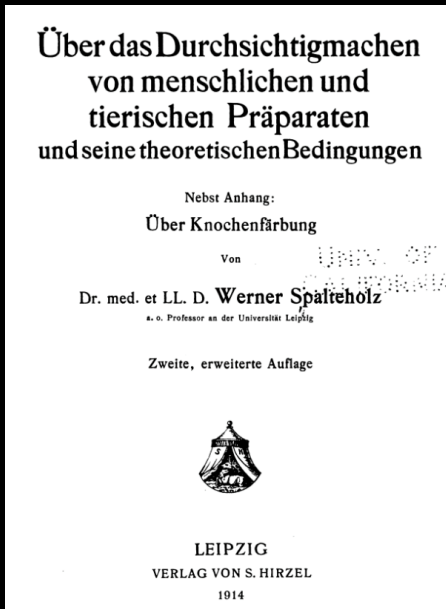


Mouse embryo

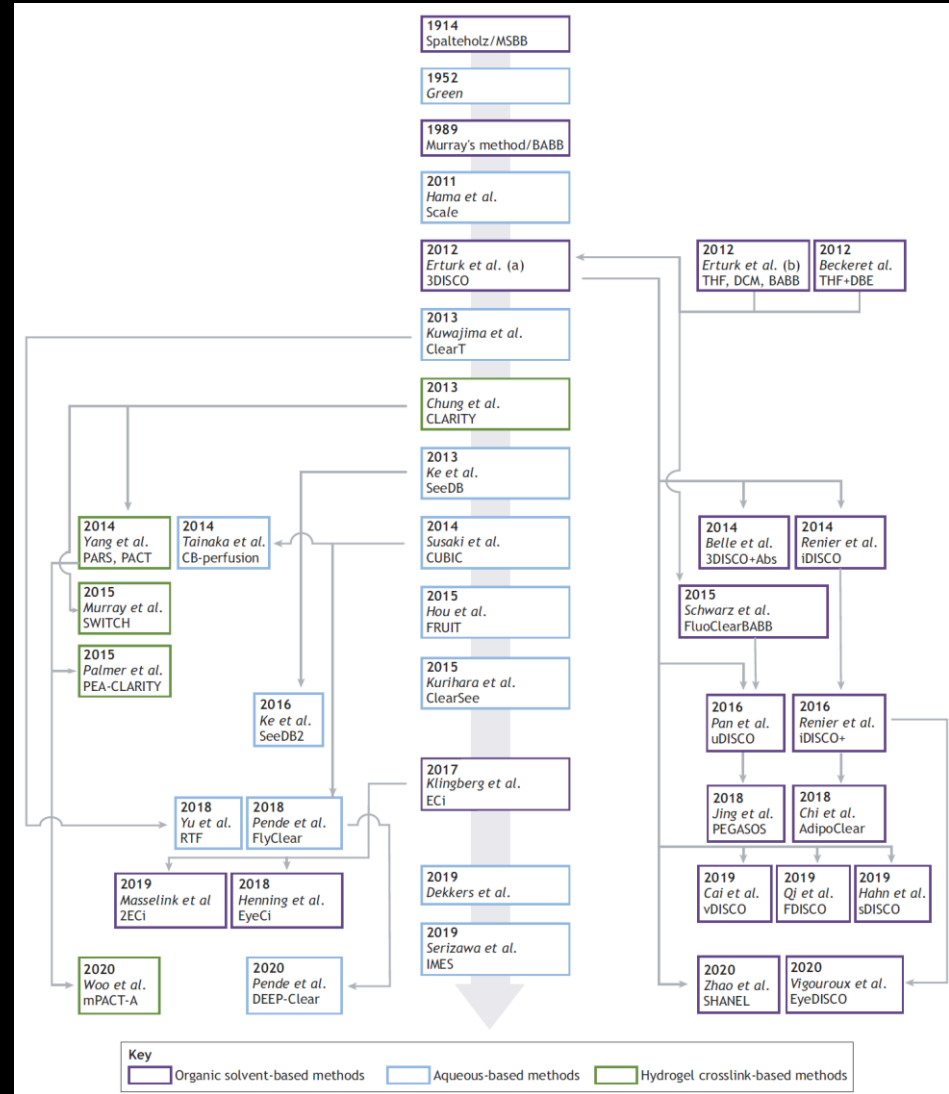
# Tissue clearing, an old concept that has just recently developed

Tissue-clearing methods used in developmental studies (2021)

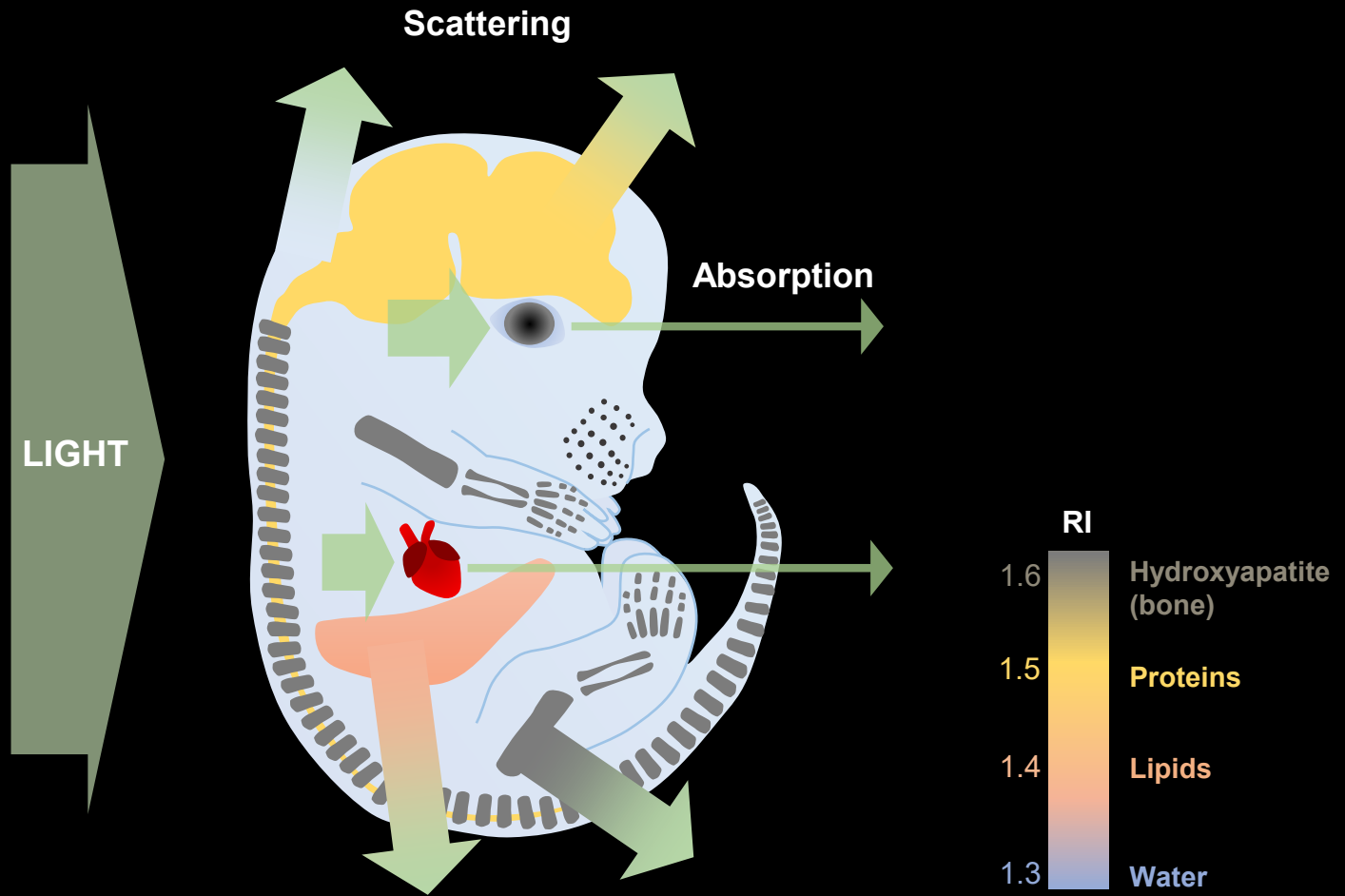
Spalteholz 1914



*"About the transparency of human and animal preparations and its theoretical conditions"*

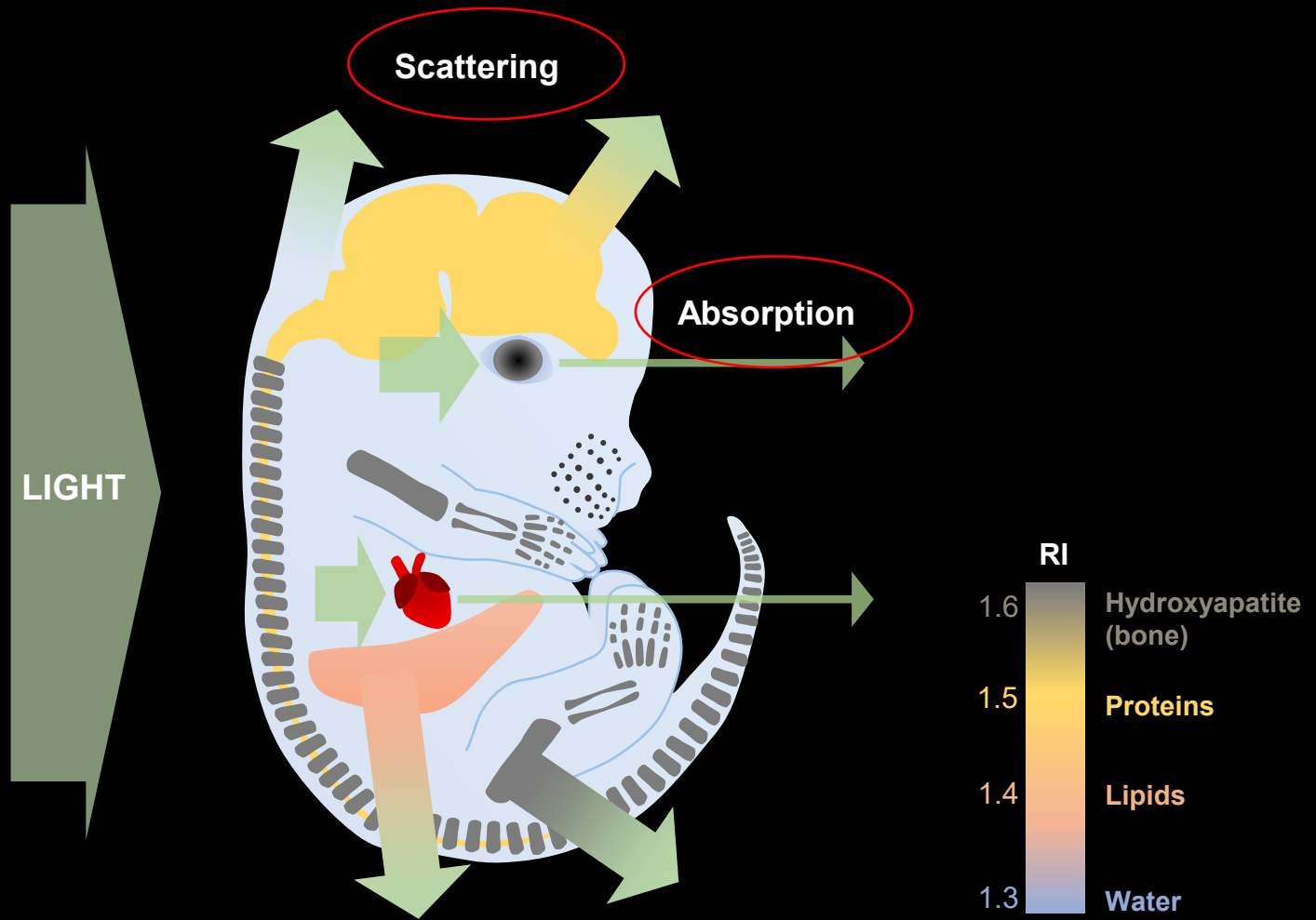


# Many methods for tissue clearing, same principle: RI homogenization



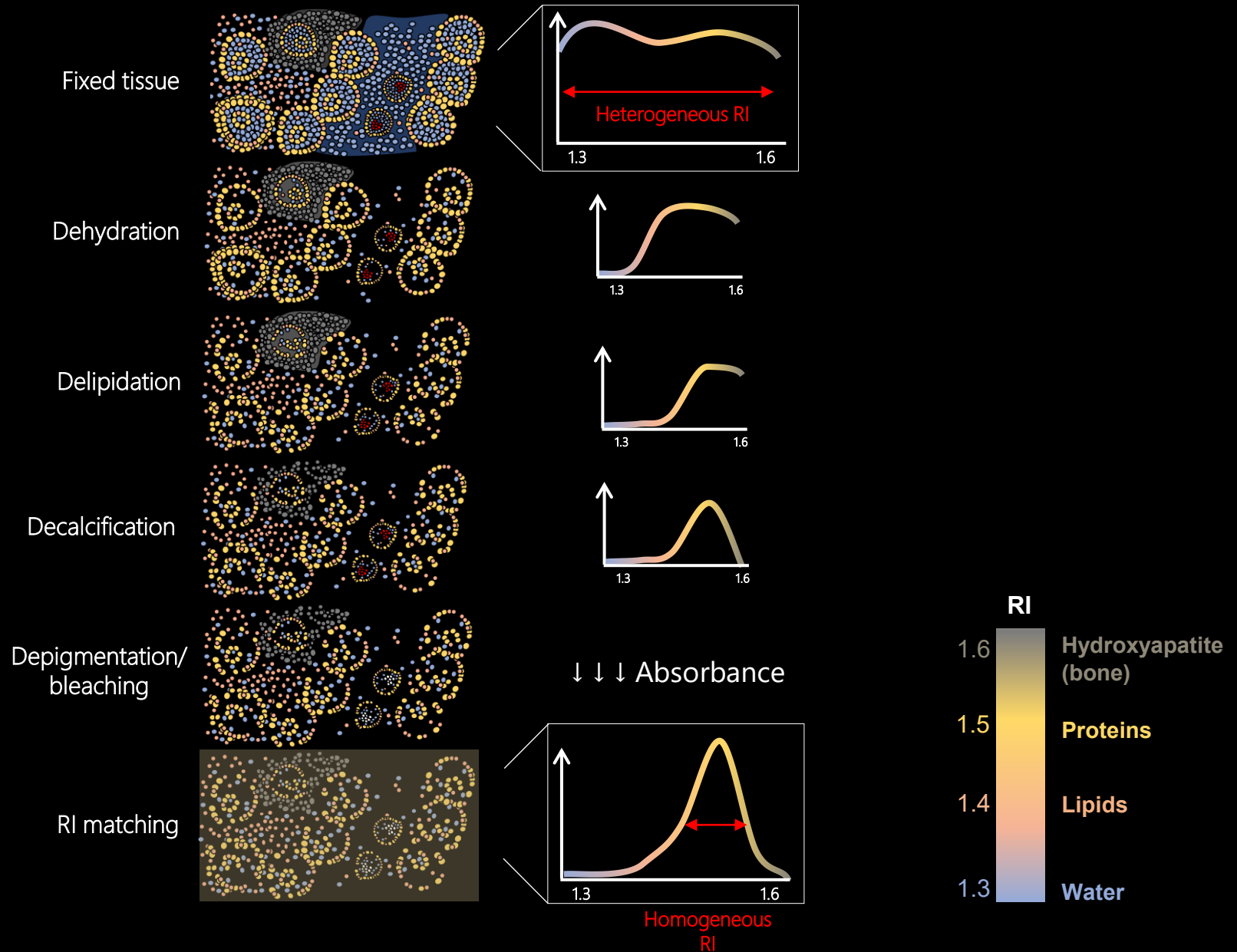
Biological samples contain a mixture of Refractive Index (RI) and pigments that perturb the light path

# Many methods for tissue clearing, same principle: RI homogenization



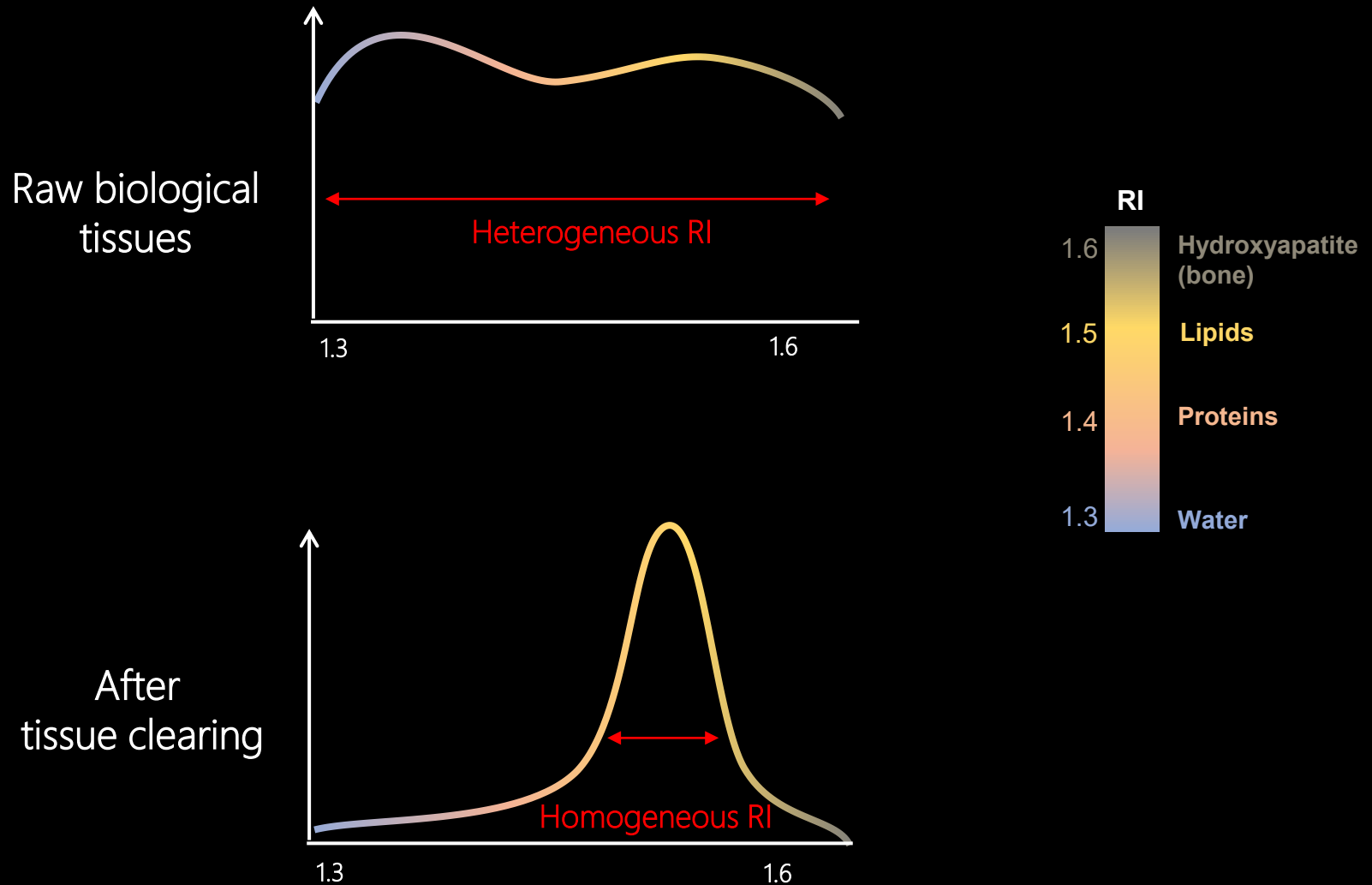
Biological samples contain a mixture of Refractive Index (RI) and pigments that perturb the light path

# Many methods for tissue clearing, same principle: RI homogenization



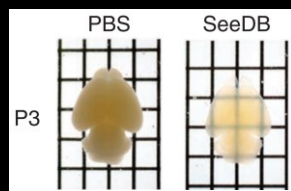


# Many methods for tissue clearing, same principle: RI homogenization



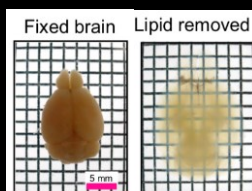
# Four families of tissue clearing protocols

## SeeDB



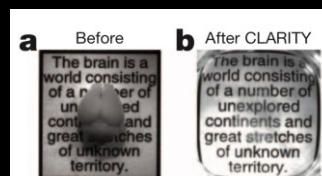
Ke, M., et al., Nat Neurosci 2013

## CUBIC



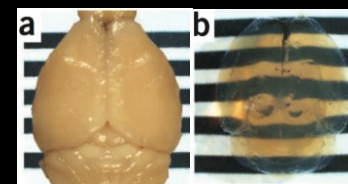
Suzaki et al, Cell 2014

## CLARITY



Chung K., et al, Nature 2013

## 3DISCO



Erturk A., et al, Nat Med 2012

### Direct RI Equilibration

#### SeeDB

Ke, M., et al.,  
Nat Neurosci 2013

FRUIT  
SeeDB2G/S  
TDE  
OptiClear  
FluoClear/RapidClear  
...

### Hyper-hydration

#### CUBIC

Suzaki, EA., et al.,  
Cell 2014

Scale U/A2  
Scale S  
CUBIC-X  
CUBIC Cancer  
CUBIC 1,2,3,4  
...

### Protein-Gel Crosslinking

#### CLARITY

Chung, K., et al.,  
Nature 2013

PARS/PACT  
Switch  
SHIELD  
ExM  
...

### Organic Solvents

#### DISCO

Ertürk, A., et al.,  
Nat. Med. 2012

FluoBABB  
**iDISCO+**  
ECi  
PEGASOS  
uDISCO  
vDISCO  
...

Protein fluorescence preservation

Scatter-free imaging

# How to choose a clearing method?

**1. Sample size**

**2. Sample type**

**3. Signal type**

**4. Imaging modality**

# How to choose a clearing method?

## 1. Sample size

Brain slice < 1mm/embryos – Refractive index matching

Mid-size samples – Classic methods (DISCO, CUBIC, Clarity)

Large samples (>2cm– uDISCO, vDISCO, perfused CUBIC)

## 2. Sample type

## 3. Signal type

## 4. Imaging modality

# How to choose a clearing method?

## 1. Sample size

Brain slice < 1mm/embryos – Refractive index matching

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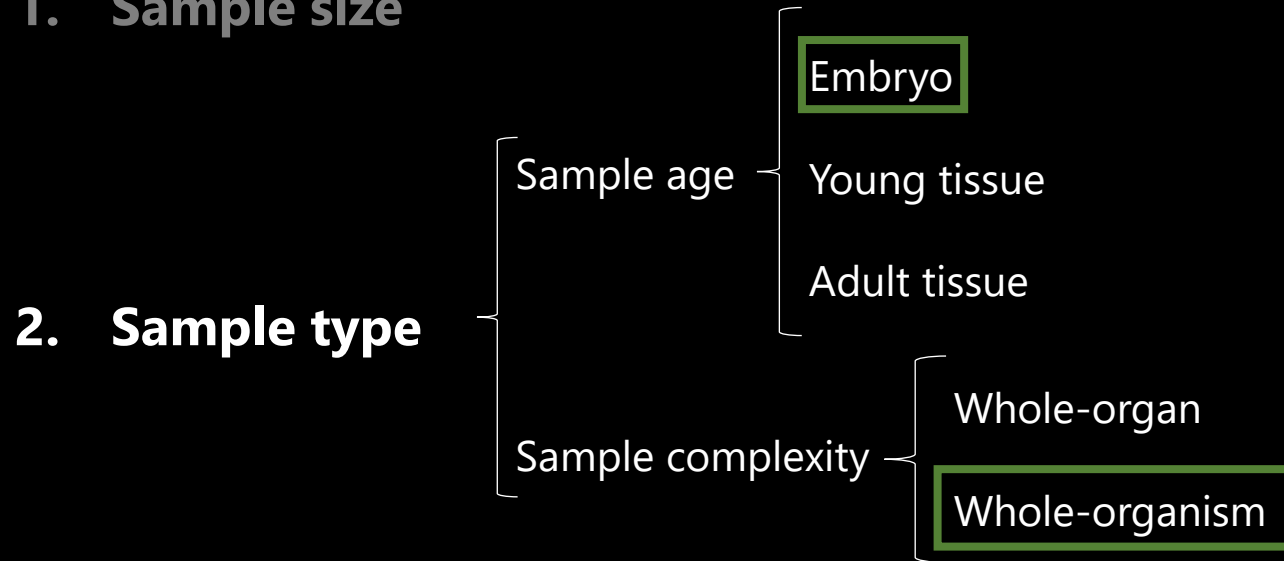
## 2. Sample type

## 3. Signal type

## 4. Imaging modality

# How to choose a clearing method?

## 1. Sample size



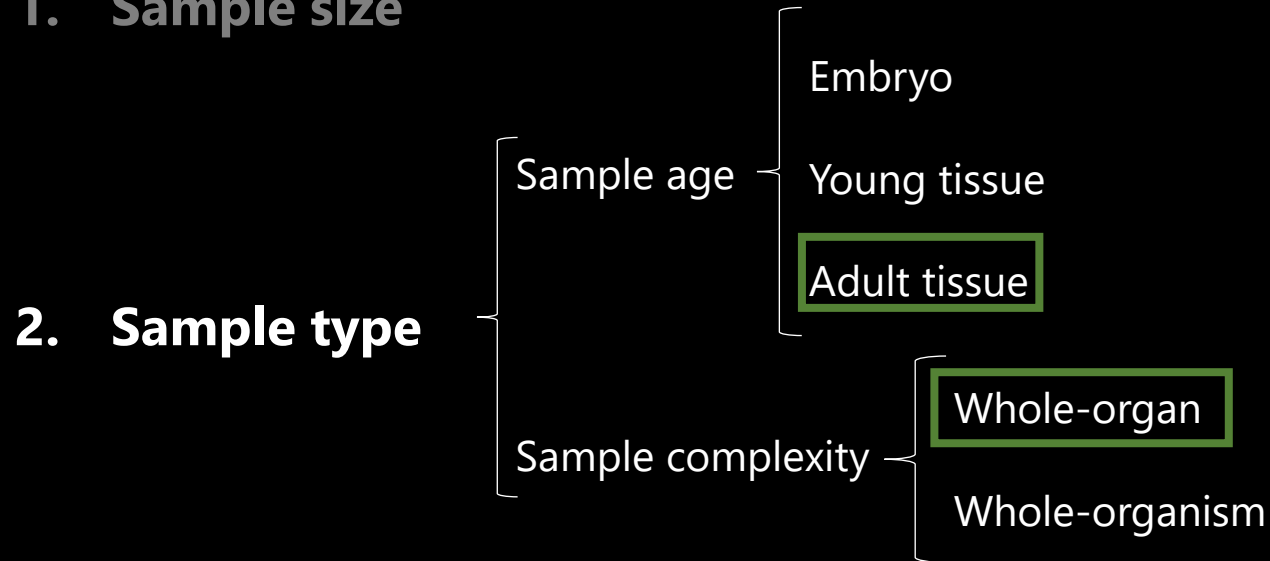
## 3. Signal type

Ex. 3DISCO, iDISCO, CUBIC, FluoClearBABB, etc

## 4. Imaging modality

# How to choose a clearing method?

## 1. Sample size



## 3. Signal type

Ex. iDISCO<sup>+</sup>, CLARITY, 2ECi

## 4. Imaging modality

# How to choose a clearing method?

## 1. Sample size

## 2. Sample type

**Tissue clearing methods are modular, adapt the chosen protocol to the features of your sample by adding:**

- **Pigment bleaching**
- **Decalcification**
- **Permeabilization**

## 3. Signal type

## 4. Imaging modality



# How to choose a clearing method?

1. **Sample size**

2. **Sample type**

3. **Signal type**

Endogenous XFP (Ex. All aqueous)

Lipophilic tracer (Ex. ClearT2, RTF, FRUIT)

Immunolabeling (Ex. iDISCO<sup>+</sup>)

4. **Imaging modality**

# How to choose a clearing method?

1. Sample size

2. Sample type

3. Signal type

Endogenous XFP (Ex. All aqueous)

Lipophilic tracer (Ex. ClearT2, RTF, FRUIT)

Immunolabeling (Ex. iDISCO<sup>+</sup>)

4. Imaging modality

# How to choose a clearing method?

1. Sample size

2. Sample type

3. Signal type

Endogenous XFP (Ex. All aqueous)

Lipophilic tracer (Ex. ClearT2, RTF, FRUIT)

Endogenous XFP+Immunolabeling (Ex. iDISCO<sup>+</sup>)

4. Imaging modality

# How to choose a clearing method?

1. Sample size

2. Sample type

3. Signal type

4. **Imaging modality**

Confocal (Ex. RI matching methods)

Light-sheet (Ex. Any method)

2 Photon microscopy (Ex. Any method?)

# How to choose a clearing method?

1. Sample size

2. Sample type

3. Signal type

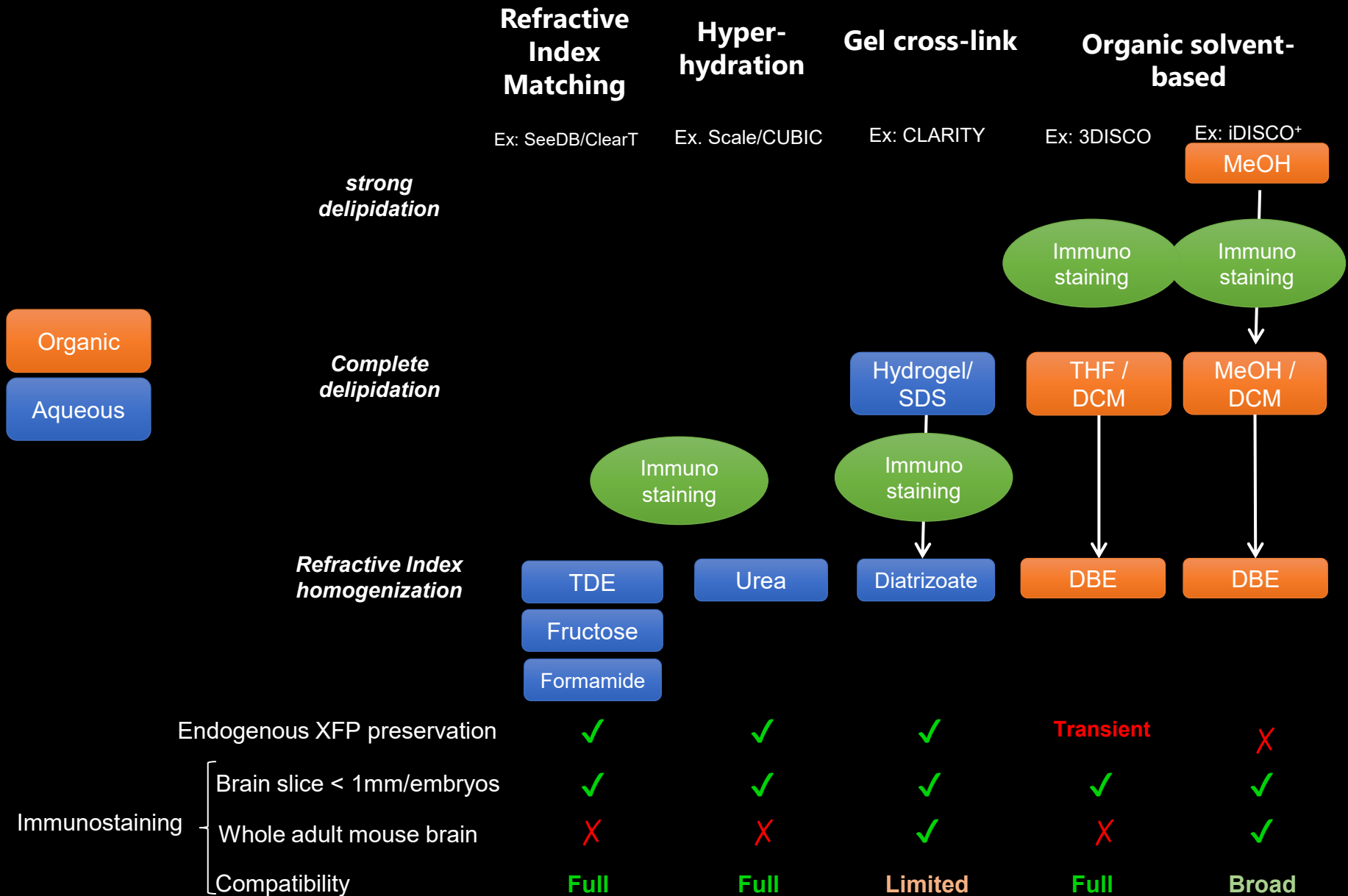
4. **Imaging modality**

Confocal (Ex. RI matching methods)

Light-sheet (Ex. Any method)

2 Photon microscopy (Ex. Any method?)

# Summary: selecting a clearing method



Endogenous XFP preservation

✓

✓

✓

**Transient**

✗

Immunostaining { Brain slice < 1mm/embryos

✓

✓

✓

✓

✓

Whole adult mouse brain

✗

✗

✓

✗

✓

Compatibility

**Full**

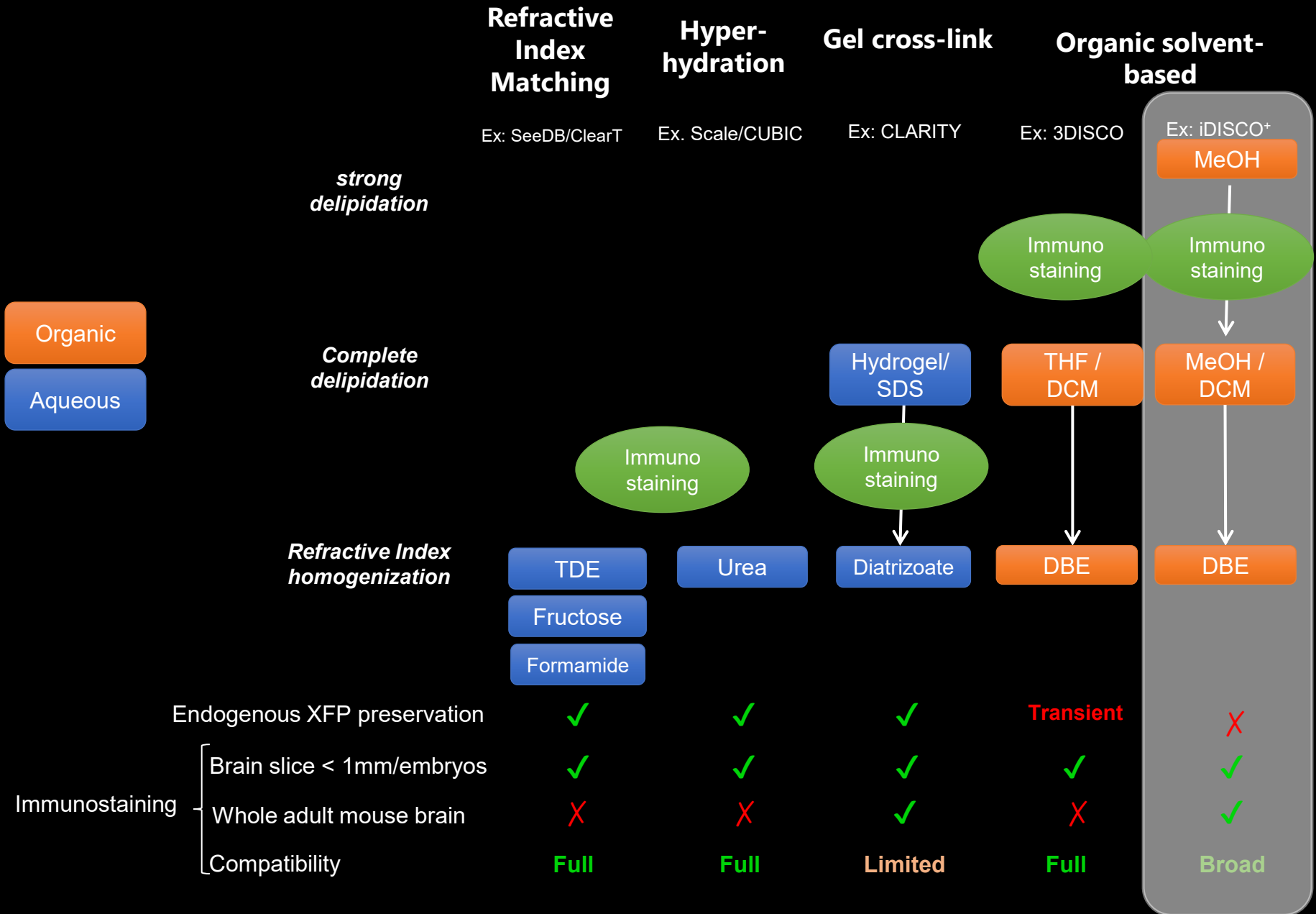
**Full**

**Limited**

**Full**

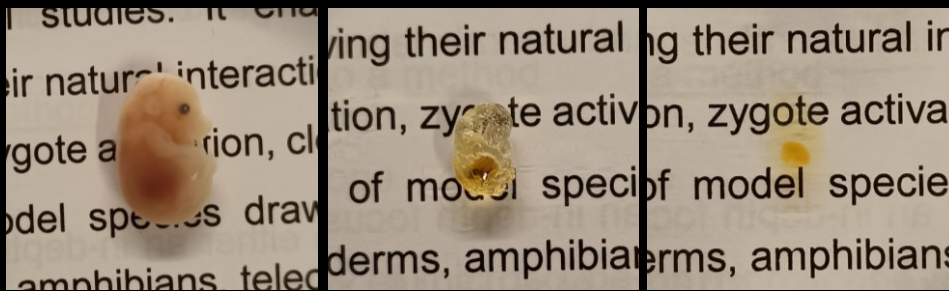
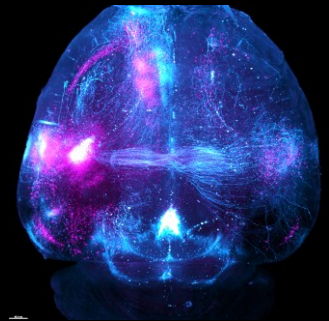
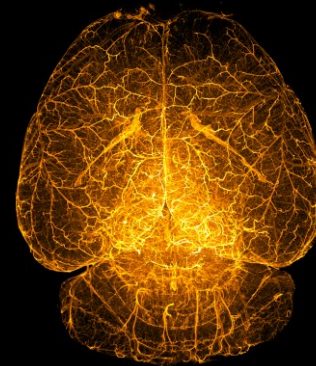
**Broad**

# Summary: selecting a clearing method



# iDISCO+

A combined method for IMMUNOLABELING and CLEARING large complex tissues

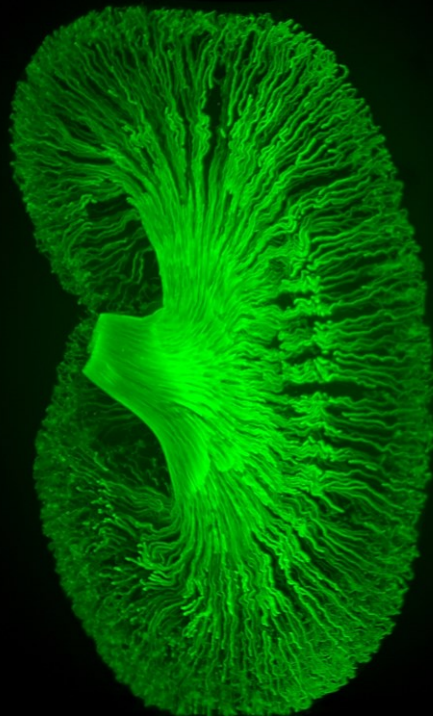
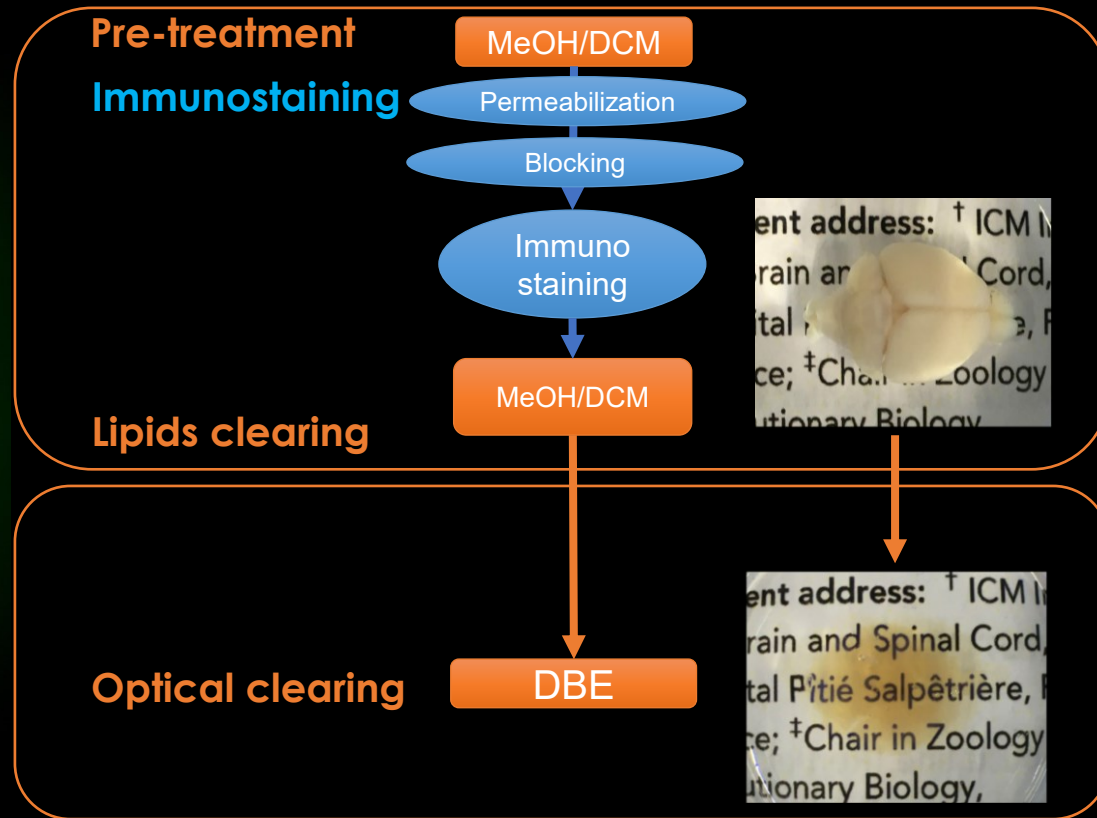




# iDISCO<sup>+</sup>

A combined method for IMMUNOLABELING and CLEARING large complex tissues

Enhanced whole-mount immunolabeling protocol + 3DISCO clearing = **iDISCO**



# iDISCO<sup>+</sup> overview

**Day « 0 » - Sample preparation**

**Days 1 to 3 – Delipidation and bleaching**

**Day 3 – Permeabilization**

**Day 4 – Blocking**

**Day 5 – Antibody incubation**

**Day 15 – Washes in PBS-TwH**

**Day 16 – Antibody incubation**

**Day 24 – Washes in PBS-TwH**

**Day 25 – Dehydration**

**Day 26 – Refractive Index matching**

# iDISCO<sup>+</sup> overview

**Day « 0 » - Sample preparation**

**Days 1 to 3 – Delipidation and bleaching**

**Day 3 – Permeabilization**

**Day 4 – Blocking**

**Day 5 – Antibody incubation**

**Day 15 – Washes in PBS-TwH**

**Day 16 – Antibody incubation**

**Day 24 – Washes in PBS-TwH**

**Day 25 – Dehydration**

**Day 26 – Refractive Index matching**

Aqueous

Organic

## Day « 0 »: Sample preparation

**Perfusion (PBS + 4% PFA, or 4%PFA alone)**

**Postfixation (PFA 4% RT)**

**Washes in PBS - 3 washes x 15min (aprox.) in PBS - RT**

# Day « 0 »: Sample preparation

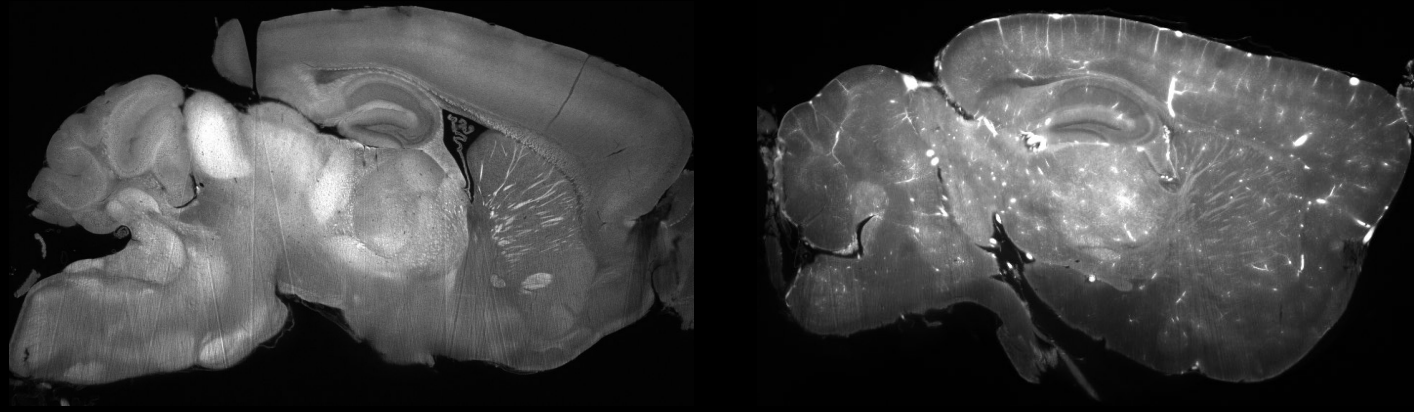
Perfusion (PBS + 4% PFA, or 4%PFA alone)

Postfixation (PFA 4% RT)

Washes in PBS - 3 washes x 15min (aprox.) in PBS - RT

## \*What can go wrong?

1. Overfixation ( ↑ tissue autofluorescence)



## Day « 0 »: Sample preparation

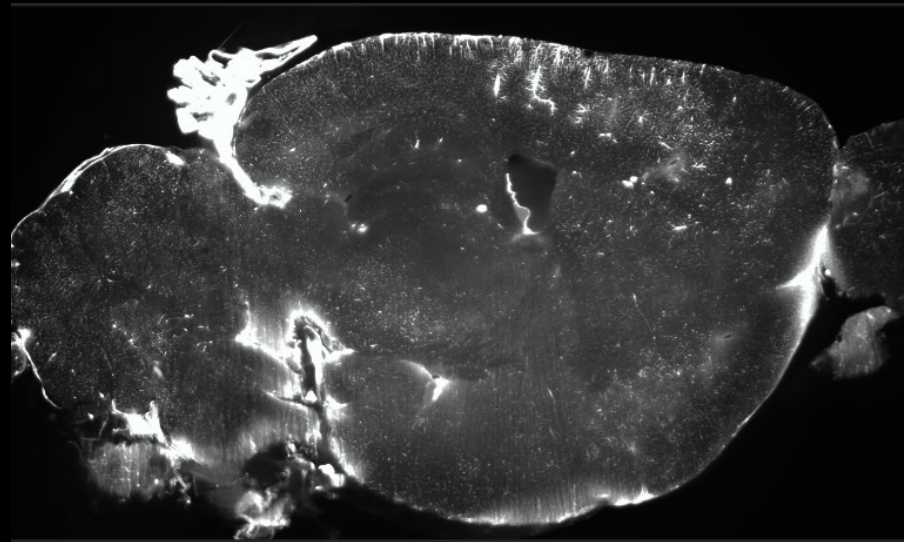
**Perfusion (PBS + 4% PFA, or 4%PFA alone)**

**Postfixation (PFA 4% RT)**

**Washes in PBS - 3 washes x 15min (aprox.) in PBS - RT**

### **\*What can go wrong?**

2. Partial perfusion (specially problematic if using Mouse 1Abs:  
↑ cross-reactivity in between blood IgGs and anti-Mouse 2°Abs)



# Day « 0 »: Sample preparation

Perfusion (PBS + 4% PFA, or 4%PFA alone)

Postfixation (PFA 4% RT)

Washes in PBS - 3 washes x 15min (aprox.) in PBS - RT

## \*What can go wrong?

3. Fungi contamination, if get this, do not continue with the protocol



**IMPORTANT:**  
All aqueous buffers of  
the protocol need to  
be supplemented with  
0,01% NaN<sub>3</sub>!

## Day 1: MeOH dehydration

MeOH 20%  $\geq$  1 hour

MeOH 40%  $\geq$  1 hour

MeOH 60%  $\geq$  1 hour

MeOH 80%  $\geq$  1,5 hours

MeOH 100%  $\geq$  1 hour

MeOH 100%  $\geq$  1,5 hours

MeOH 33%/DCM 66% over night RT

\*Ex. for adult whole-brain, to be adapted depending on the sample size

### \*What can go wrong?

1. Partial dehydration due to short steps of dehydration:

High autofluorescence in the center of the brain

Anisotropic shrinkage (short steps of dehydration)



## Day 1: MeOH dehydration

MeOH 20%  $\geq$  1 hour

MeOH 40%  $\geq$  1 hour

MeOH 60%  $\geq$  1 hour

MeOH 80%  $\geq$  1,5 hours

MeOH 100%  $\geq$  1 hour

MeOH 100%  $\geq$  1,5 hours

MeOH 33%/DCM 66% over night RT

\*Ex. for adult whole-brain, to be adapted depending on the sample size

### \*What can go wrong?

2. Brain can blow up! Skip initial MeOH dehydration in brains <P7)

## Day 2: MeOH rehydration and permeabilization

MeOH 100% x 2 long washes along the day

MeOH 80%/H<sub>2</sub>O<sub>2</sub> 20% (5 to 6% final concentration) over night 4°C

### \*What can go wrong?

1. Nothing!!

Tubes could eventually open due to O<sub>2</sub> pressure: ensure they are correctly closed.

## Day 3: MeOH washes and bleaching

MeOH 60%  $\geq$  1,5 hours

MeOH 40%  $\geq$  1 hour

MeOH 20%  $\geq$  1 hour

PBS

PBS-Tx

Permeabilization solution – 37°C over night

### \*What can go wrong?

1. Nothing really

## Day 4: Blocking

Blocking solution – 37°C over night

**\*What can go wrong?**

1. Nothing really

## Day 5: 1°Ab incubation

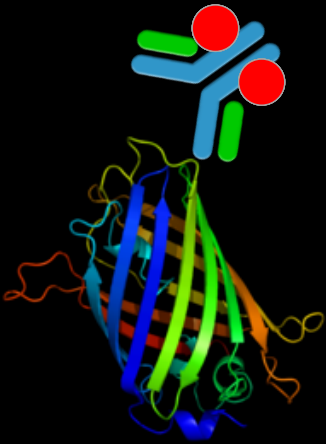
Antibody diluted in blocking solution – variable duration – 37°C

Ex. adult mouse brain 10 days, P3 brain 5 days, 1mm section 2 days, etc.

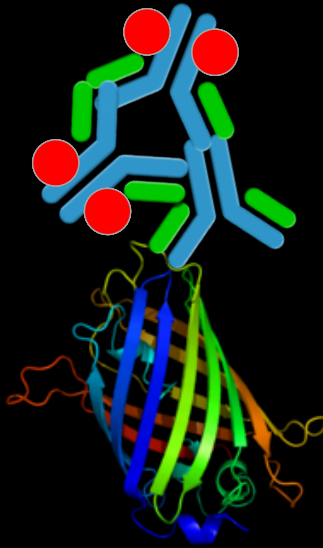
### **\*What can go wrong?**

1. Signal amplification should be optimized with the labeling strategy

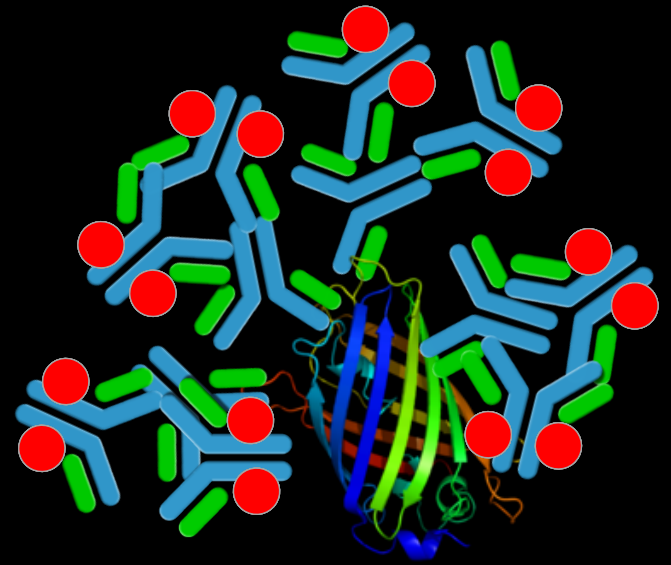
- Signal amplification



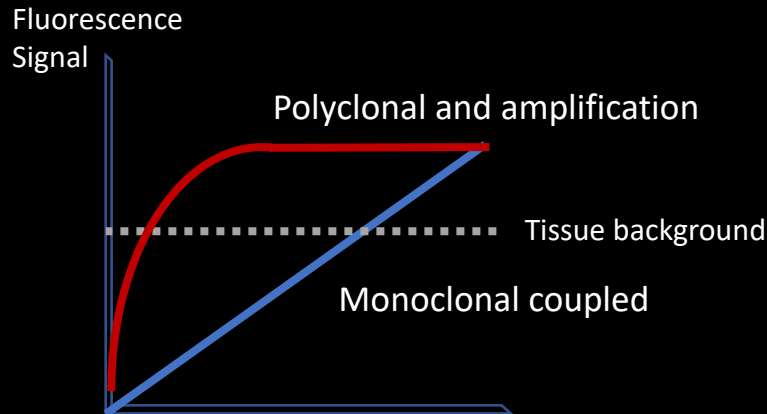
Monoclonal coupled



Monoclonal secondary Ab amplification



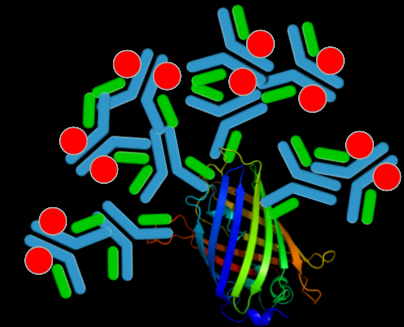
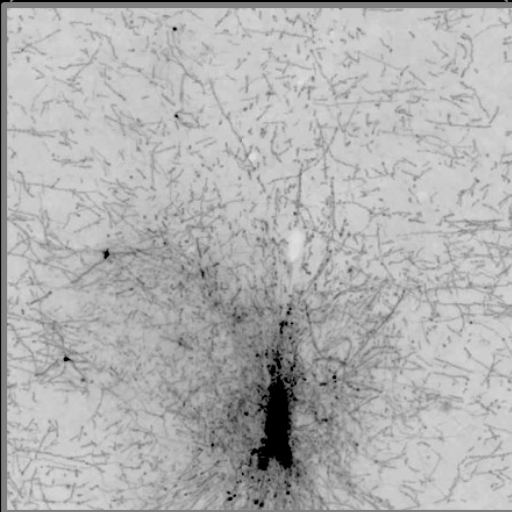
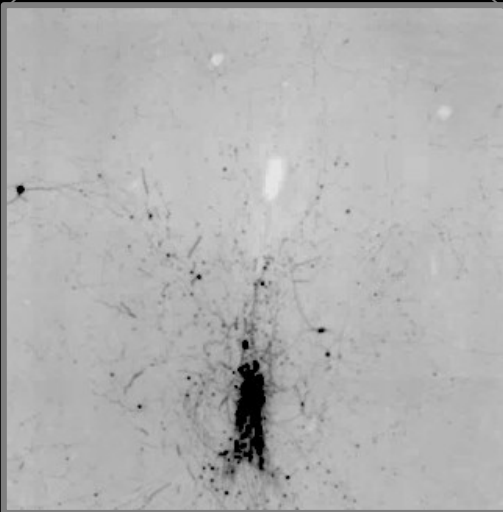
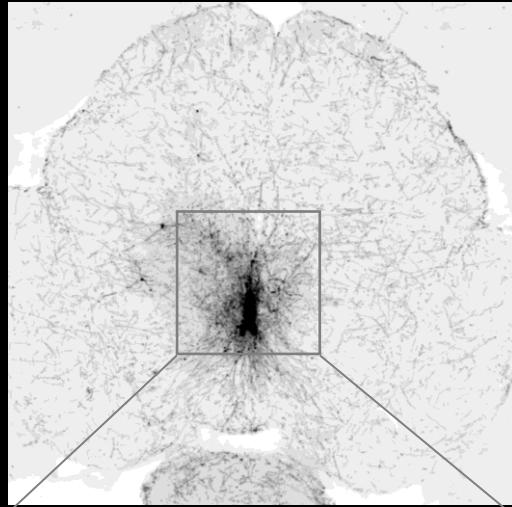
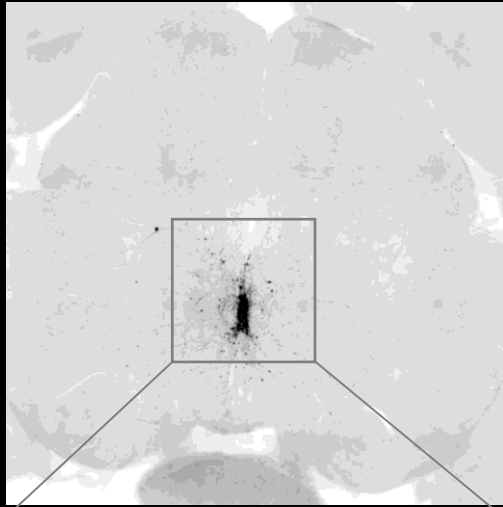
Polyclonal secondary Ab amplification



# Immunolabeling with **2-stages polyclonals** to amplify **sparse antigens**

GFP endogenous

GFP immuno

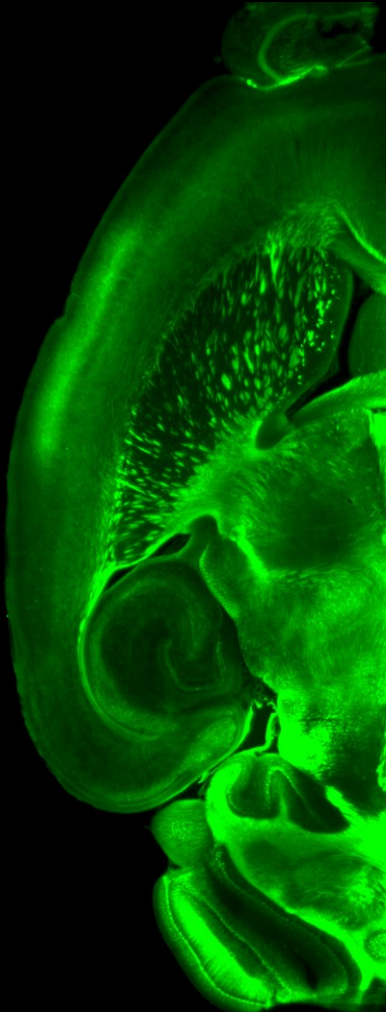


Polyclonal  
secondary Ab amplification

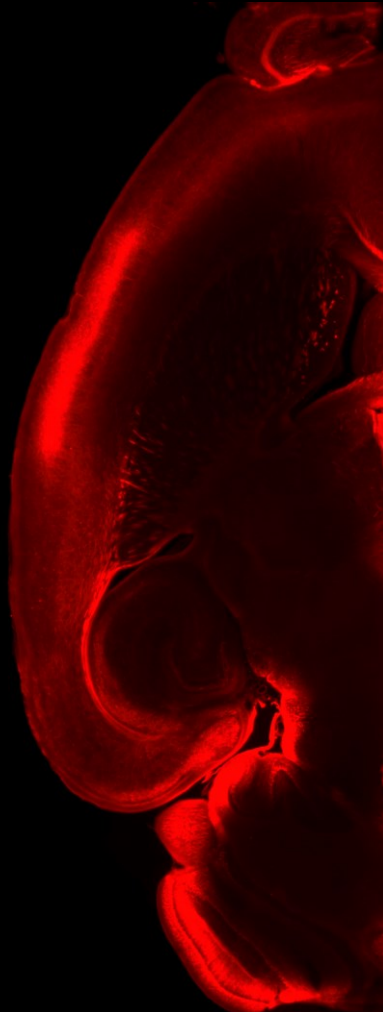
CART-cre :: AAV1-lsl-hSyn-eGFP

# Immunolabeling with **conjugated mAb** to amplify **dense antigens**

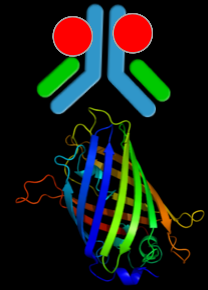
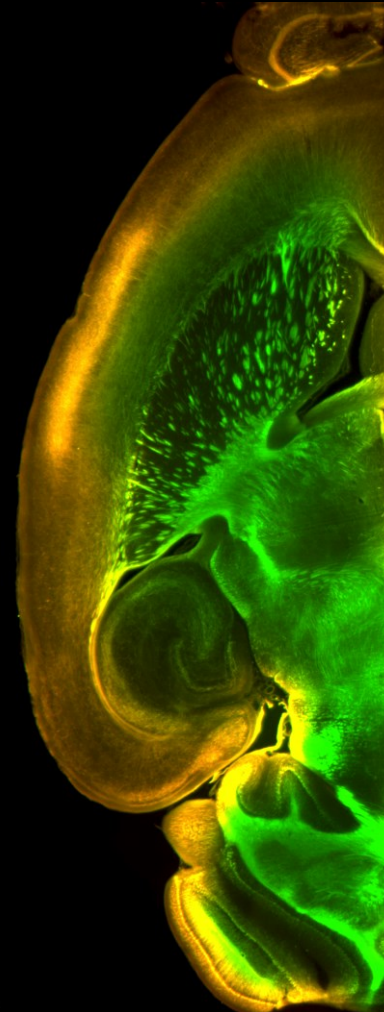
Conjugated primary



Conjugated secondary



Overlap



Direct immunodetection  
enables optimal diffusion



# Day 5: 1°Ab incubation

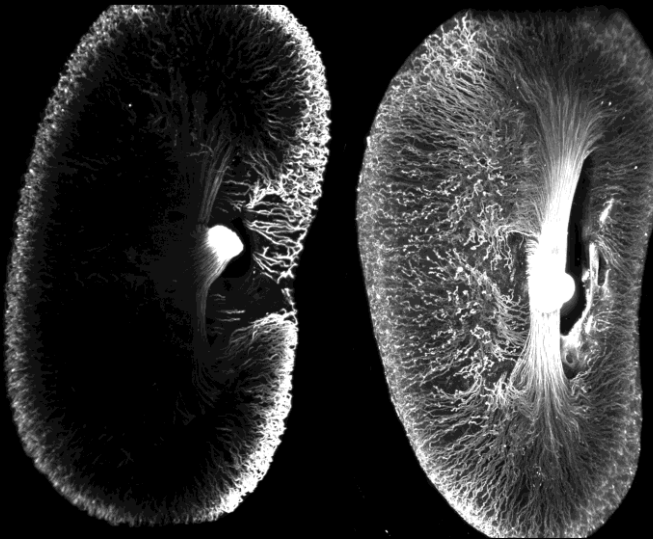
Antibody diluted in blocking solution – variable duration – 37°C

Ex. adult mouse brain 10 days, P3 brain 5 days, 1mm section 2 days, etc.

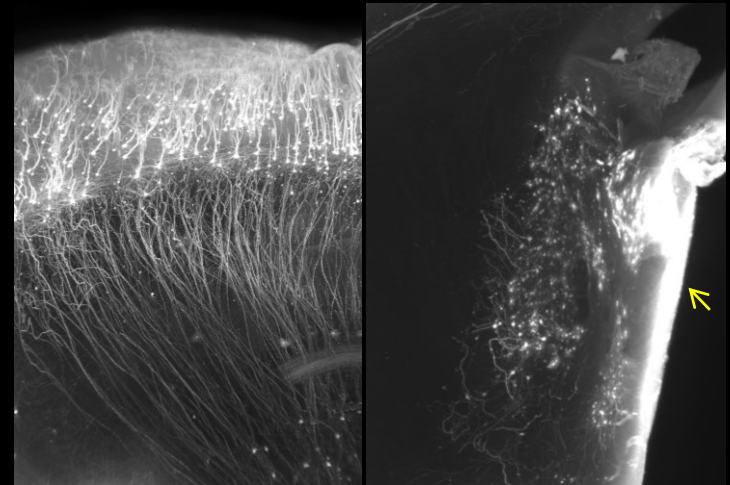
## \*What can go wrong?

### 2. Antibody concentration

“antibody depletion when concentration is too low



“ring background’ when concentration is too high



[IgG] < [antigen]

Working range

Saturating [IgG]

# Day 5: 1°Ab incubation

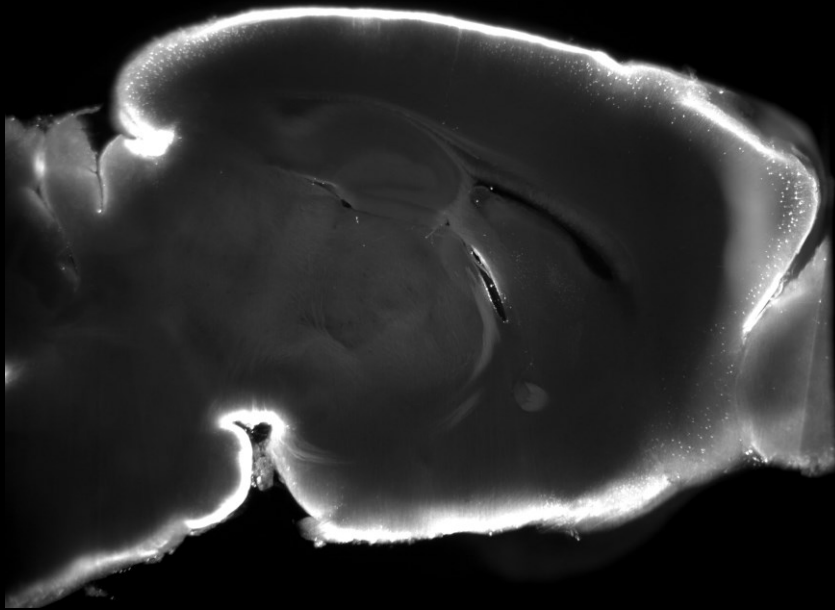
Antibody diluted in blocking solution – variable duration – 37°C

Ex. adult mouse brain 10 days, P3 brain 5 days, 1mm section 2 days, etc.

## \*What can go wrong?

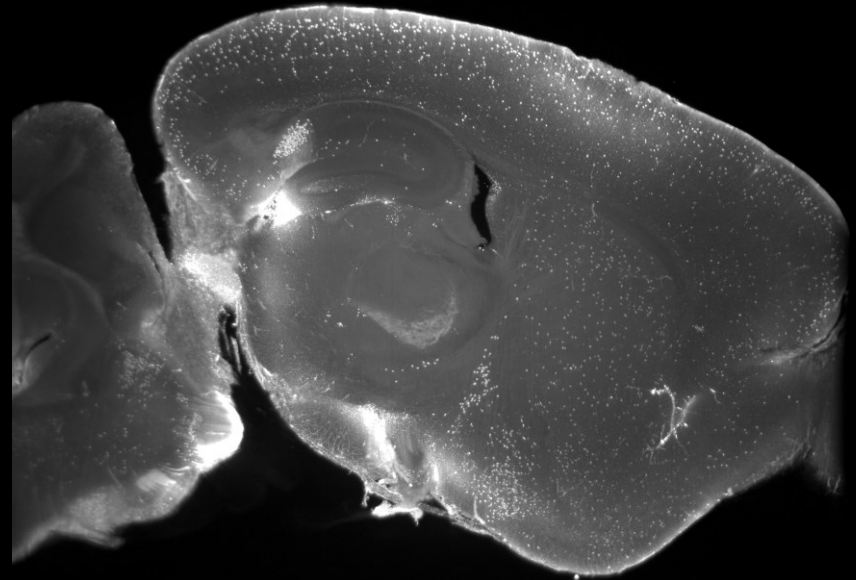
### 3. Antibody reference

Novus chicken anti-Cherry/tdTomato at 1/500<sup>th</sup>

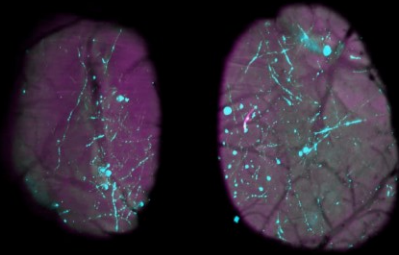


Ring surface background

Rockland rabbit anti-RFP/tdTomato at 1/500<sup>th</sup>



complete diffusion



**cFOS**  
**Autofluorescence**

# Day 15: washes

Washes in PBS-TwH at RT

**\*What can go wrong?**

Really nothing

## Day 16: 2° Ab incubation (if necessary)

Antibody diluted in blocking solution – variable duration – 37°C

Ex. adult mouse brain 8 days, P3 brain 4 days, 1mm section 1 day, etc.

### \*What can go wrong?

Much less critical than 1°Ab, just avoid this step and use 1° conjugated Abs for dense antigens

# Day 26: washes

Washes in PBS-TwH at RT

**\*What can go wrong?**

Really nothing

## Day 27: dehydration in MeOH

MeOH 20%  $\geq$  1 hour

MeOH 40%  $\geq$  1 hour

MeOH 60%  $\geq$  1 hour

MeOH 80%  $\geq$  1,5 hours

MeOH 100%  $\geq$  1 hour

MeOH 100% overnight – RT

\*Ex. for adult whole-brain, to be adapted depending on the sample size

### \*What can go wrong?

1. Short incubations will lead to incomplete dehydration and poor transparency

## Day 28: RI matching (« clearing »)

MeOH 33%/DCM 66% - 3 hours – RT

DCM 15 to 30min 2x

DBE

### \*What can go wrong?

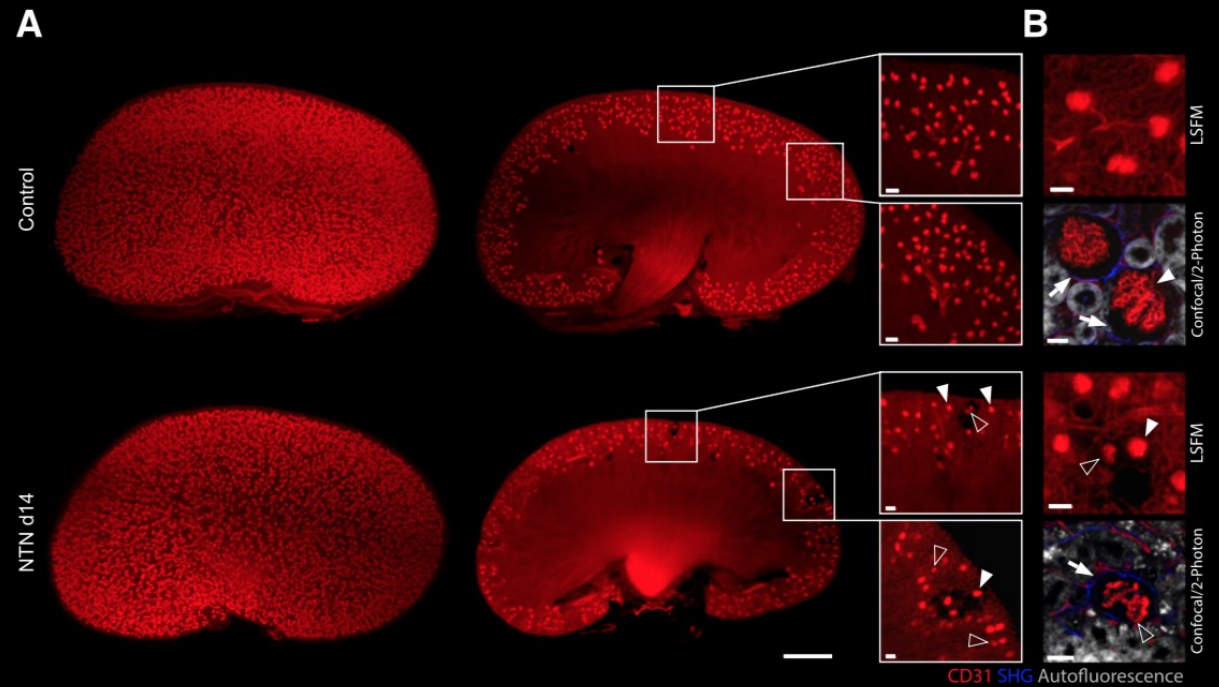
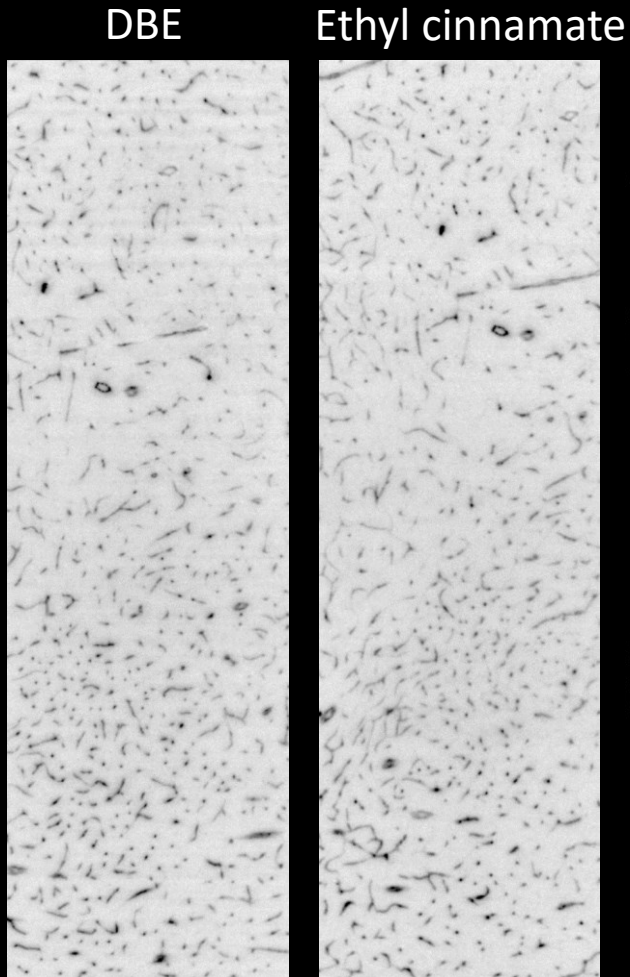
1. If there is any trace of water, the samples will look partially opaque





# Day before imaging: RI matching with Eci

Eci – overnight RT

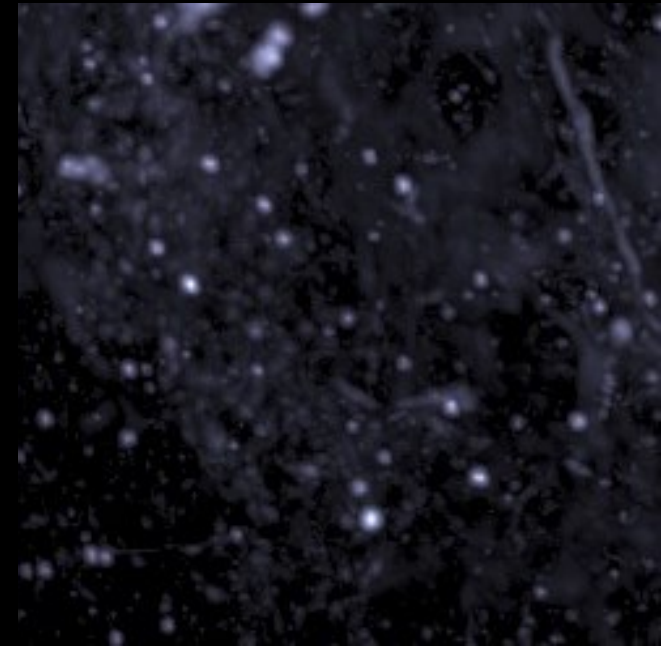
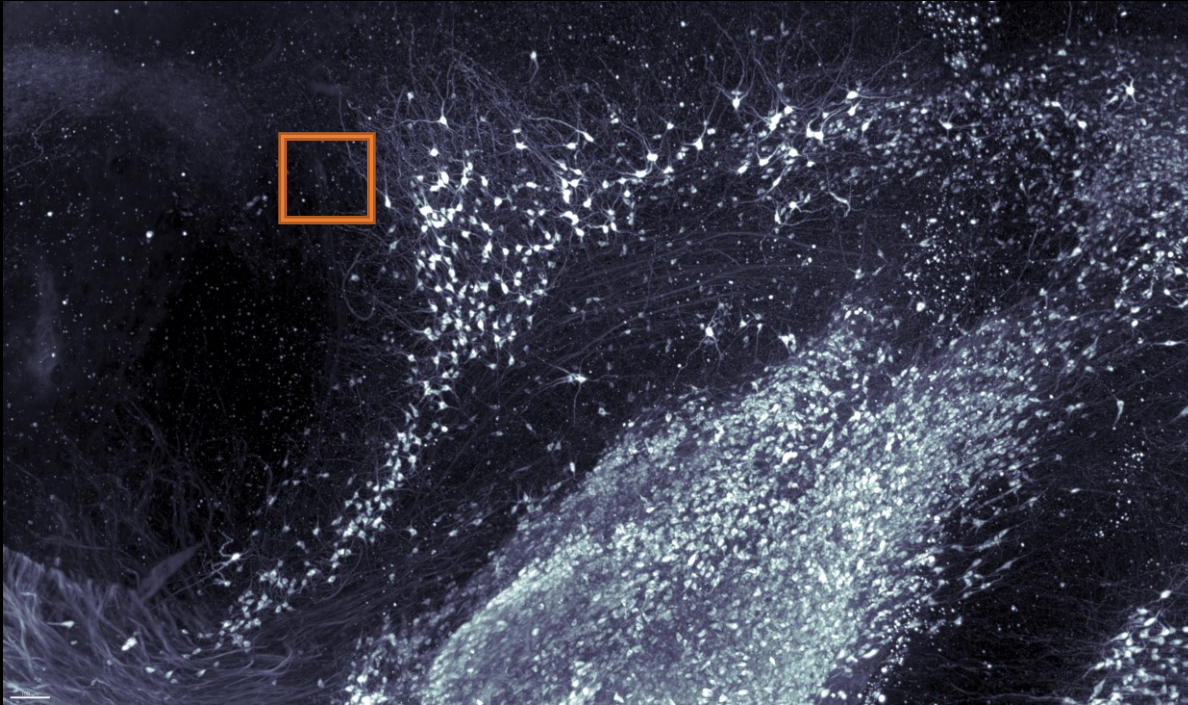


(Klingberg, J Am Soc Nephrol 2017)

4x lens, 1 laser, NA 0,148

# Multifactorial problems: Dots artefacts - primary antibody+fixation problems

Adult mouse brain, TH staining, middle optical planes max projection



2 factors :

- Keeping the sample longer in the fridge before processing increases the dots
- Dots appear when the antibody is too concentrated compared with the epitopes
- Dots are more visible when the signal is weak (one can use PBS/Methanol)

## Multifactorial problems: sample oxidation – long latencies or excess of air

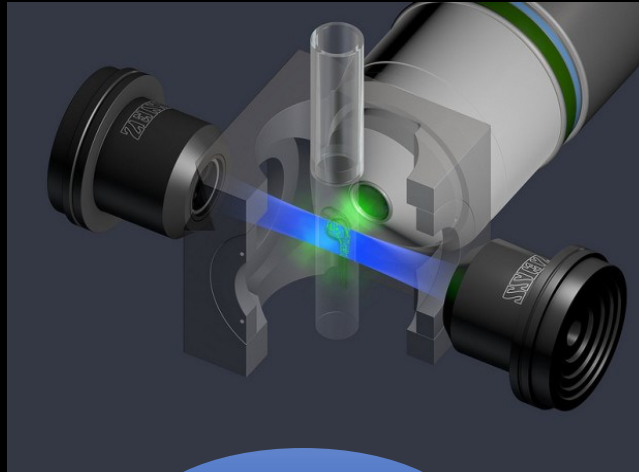


2 factors :

- Keeping the sample longer in the fridge before processing increases oxidation
- Keeping a large bubble of air in the sample's tube throughout the different steps of the protocol

## Practical tips

- Supplement all aqueous buffers with 0,01% NaN<sub>3</sub>
- Fill all tubes to the top, minimize air in the sample tube
- Find a MeOH-compatible ink labeling system for the samples
- Use a wheel shaker for all steps except bleaching
- Embed fragile samples in a 1% Agarose gel before final dehydration
- Shrink large samples with sparse labeling by doing the final dehydration in PBS/MeOH
- Choose red-shifted fluorophores



```

load averages: 0.45, 0.39, 0.37
98 processes: 89 idle, 1 on processor
CPU0 states: 0.0% user, 0.0% nice, 0.0% system, 0.0% interrupt, 100% idle
CPU1 states: 0.0% user, 0.0% nice, 0.0% system, 0.0% interrupt, 100% idle
Memory: Mem: 689/2524 act/tot, Free: 16681, Swap: 0/20321 used/tot

PID USER#PRM PRI NI CE SIZE RES STRAT WRTI TIME CPU COMMAND
26309 nicholas 2 0 1776k 4780k sleep/1 poll 0:06 0.00% mpd
16366 nicholas 2 0 1520k 4556k sleep/1 poll 1:34 0.00% mpd
23020 nicholas 2 0 4172k 2944k sleep/0 poll 0:00 0.00% mpd
2798 nicholas 2 0 3360k 1952k sleep/1 poll 0:00 0.00% scmpc
12869 root 2 0 456k 796k sleep/0 kqread 0:00 0.00% mpd
7401 uw 2 0 1540k 2540k sleep/1 select 0:00 0.00% httod
10925 root 2 0 1124k 2104k sleep/1 select 0:00 0.00% sendmail
5864 root 2 0 1184k 1168k sleep/1 poll 0:01 0.00% logmon
15102 nicholas 2 0 3304k 2260k sleep/0 select 0:02 0.00% sshd
1688 root 2 0 148k 144k idle nfsd 0:02 0.00% nfsd
26590 root 2 0 148k 144k idle nfsd 0:01 0.00% nfsd
70 nicholas 2 0 1304k 2124k sleep/0 poll 0:00 0.00% tmap
28091 root 2 0 612k 952k idle select 0:00 0.00% cron
10340 nicholas 0 0 692k 620k idle ttyn 0:00 0.00% ksh
13971 _syslogd 2 0 624k 840k sleep/0 poll 0:00 0.00% syslogd
19861 nicholas 2 0 972k 2704k sleep/1 poll 0:00 0.00% ncpc
27153 nicholas 2 0 1500k 111k sleep/0 select 0:00 0.00% emacs

```

```

-client_msg_fn_detach(struct hdr *hdr, struct client_ctx *cctx)
-client_msg_fn_detach(struct msg *msg, struct client_ctx *cctx)
{
    if (hdr->size != 0)
        if (msg->hdr.len != MSG_HEADER_SIZE)
            fatalx("bad MSG_DETACH size");
    client_write_server(cctx, MSG_EXITING, NULL, 0);
}

int
client_msg_fn_shutdown(
    struct hdr *hdr, struct client_ctx *cctx)
{
    struct msg *msg = msg_struct(client_ctx *cctx);
    if (hdr->size != 0)
        if (msg->hdr.len != MSG_HEADER_SIZE)
            fatalx("bad MSG_SHUTDOWN size");
    client_write_server(cctx, MSG_EXITING, NULL, 0);
}

```

```

20:28:31 nicholas@pelena 0 1
tmap-borders.diff
tmap-cfgour.diff
tmap-icsp-12diff.dl
tmap-icsp1.diff
tmap-icsp2.diff
tmap-nodegraph.ctd
nicholas@pelena 0 1

```

```

nicholas@pelena 0 1 *$
nicholas@pelena 0 1 *$ []
nicholas@pelena 0 1 *$

```

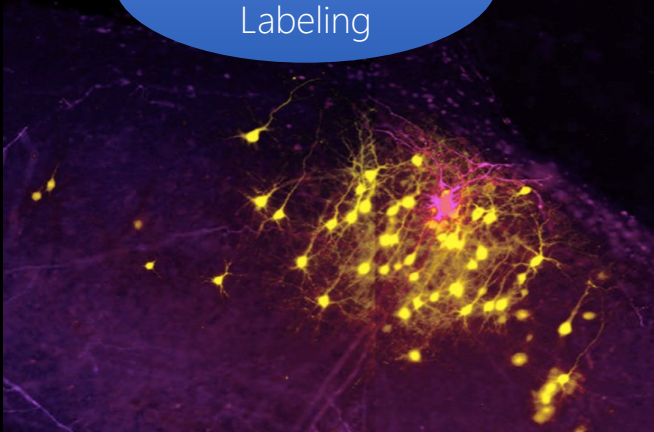
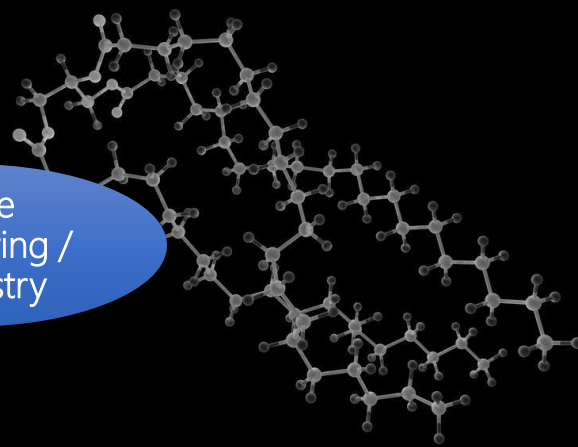
Lightsheet microscopy

Computer Sciences

Mesoscale imaging

Tissue Engineering / Chemistry

Molecular / Genetic Cell Labeling

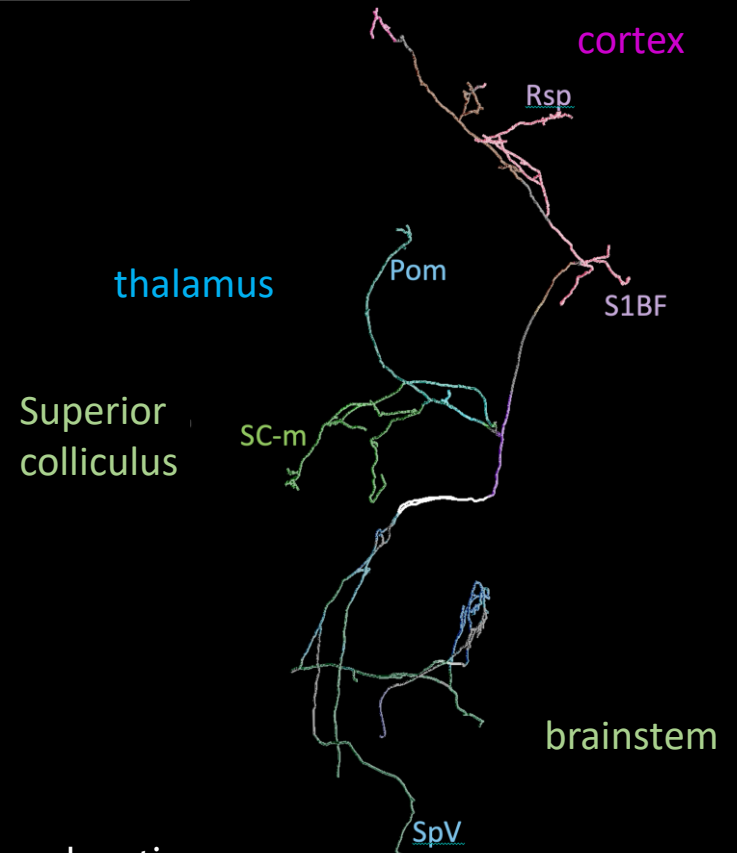
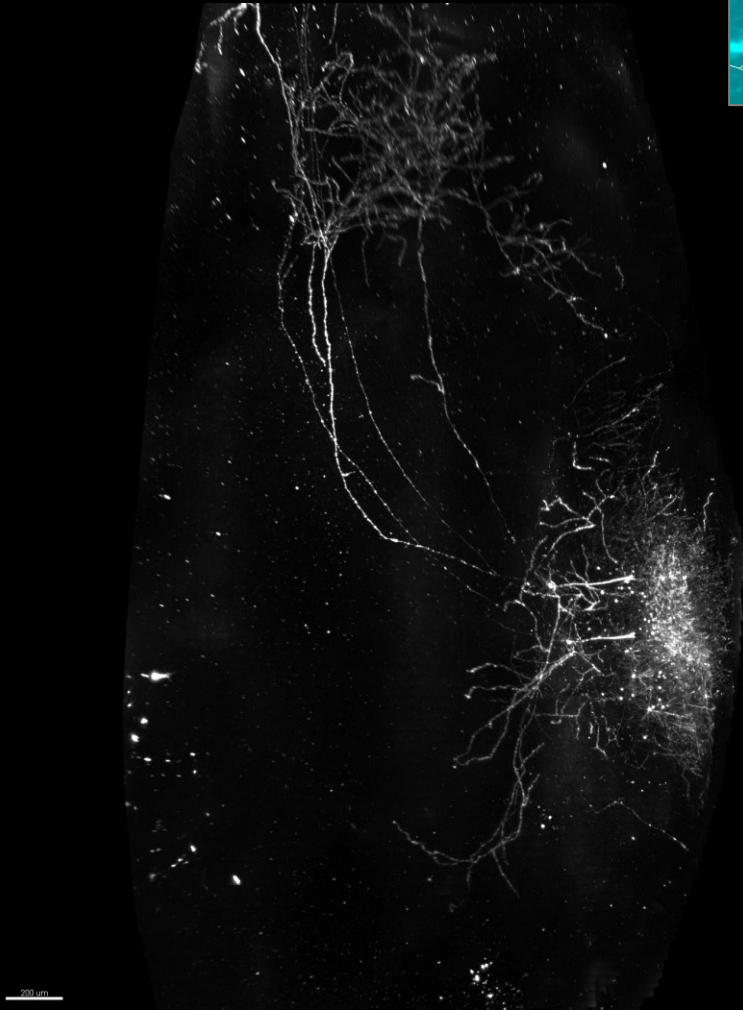
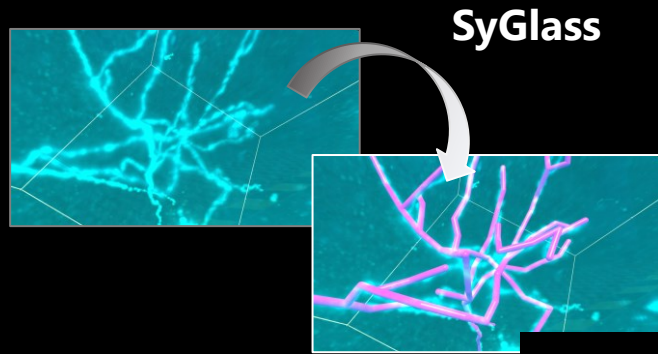


## Qualitative analysis (visualization)



Imaris

# Qualitative analysis



VR-assisted 3D image exploration

# Quantitative analysis

*ClearMap: CellMap*, module for cell detection (Renier\*, Adams\*, Kist\*, Wu\*, et al. Cell 2016)

TubeMap, module for vasculature segmentation (Kirst\*, Skriabine\*, Vieites-Prado\* et al. Cell 2020)

