Introduction to EEG/MEG and the FieldTrip toolbox

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What is FieldTrip

a MATLAB toolbox for the analysis of MEG, EEG and animal electrophysiology data

can import data from many different file formats

contains algorithms for spectral analysis, source reconstruction, statistics, connectivity, …

Talk outline

Recording signals from the brain

Temporal and spatial features of the signals

Analyzing those signals with FieldTrip

Background on the FieldTrip toolbox

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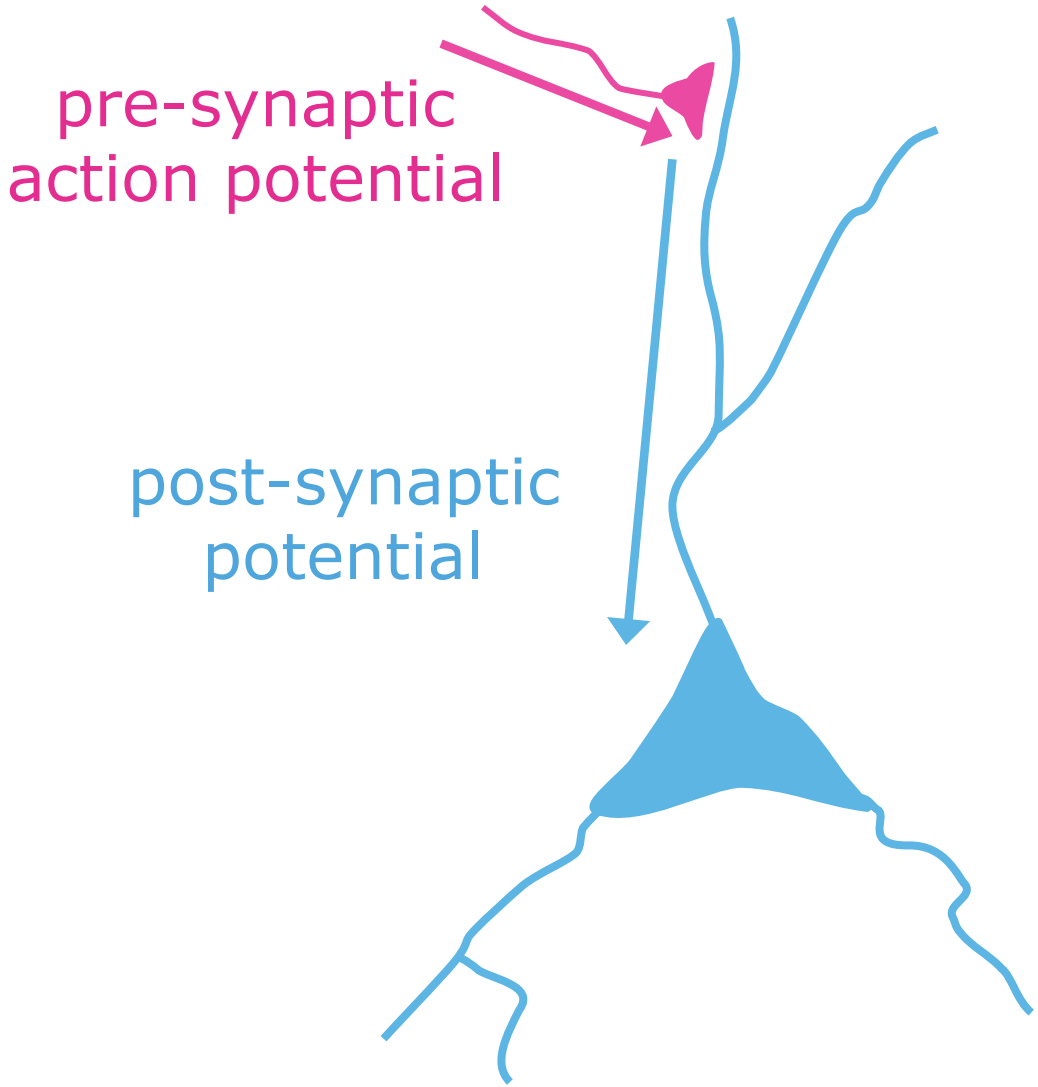
What produces the electric current and magnetic field



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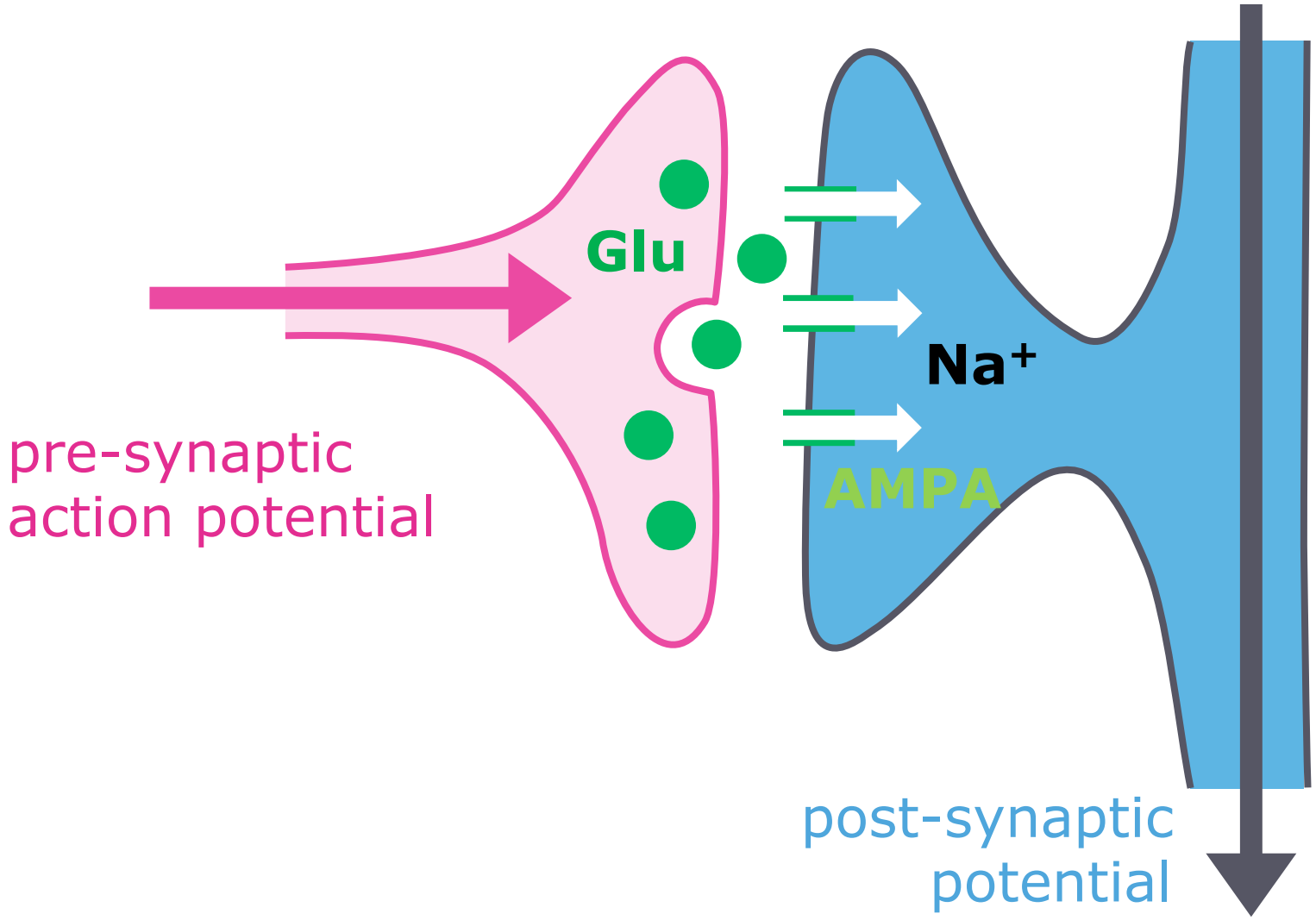

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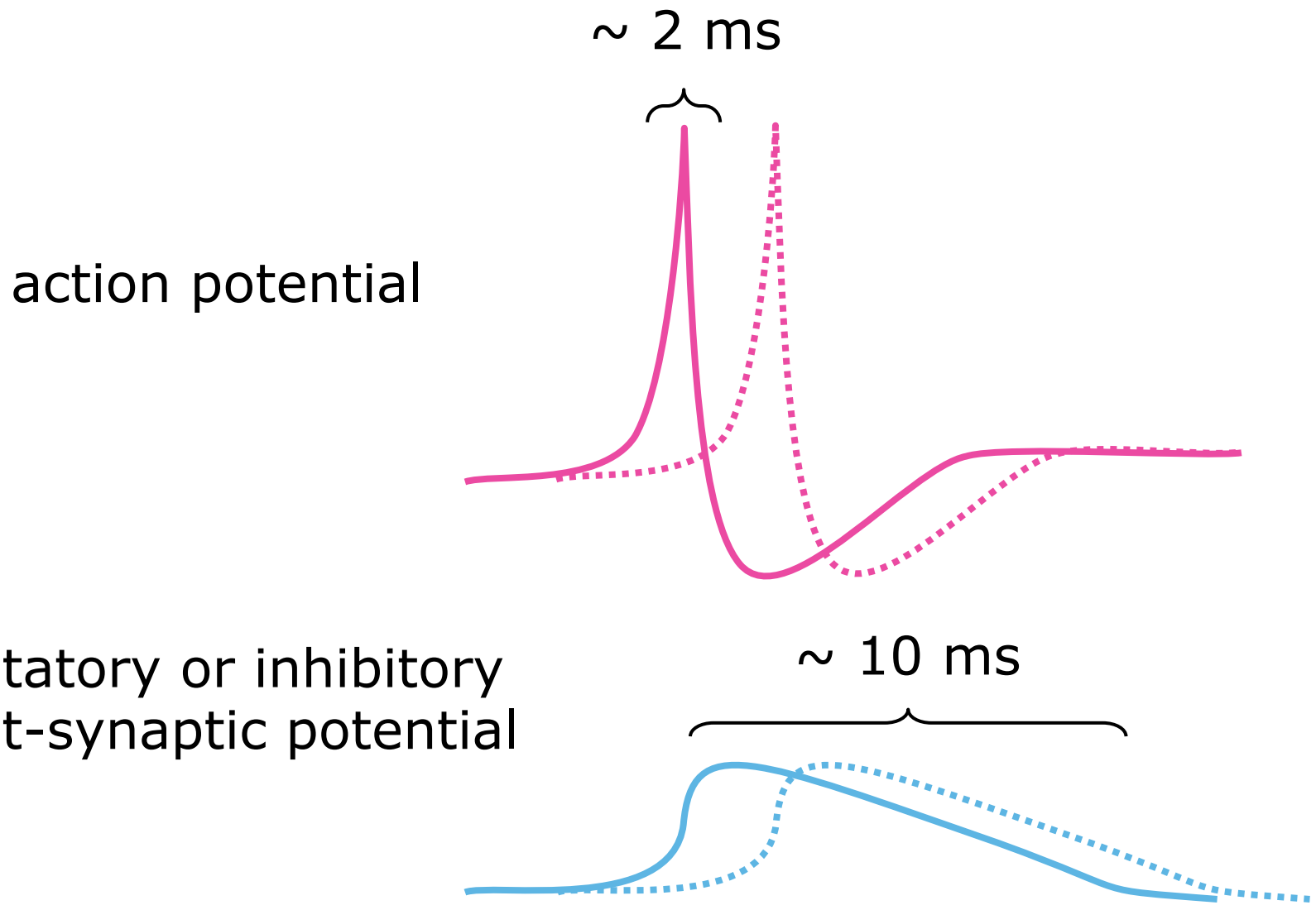

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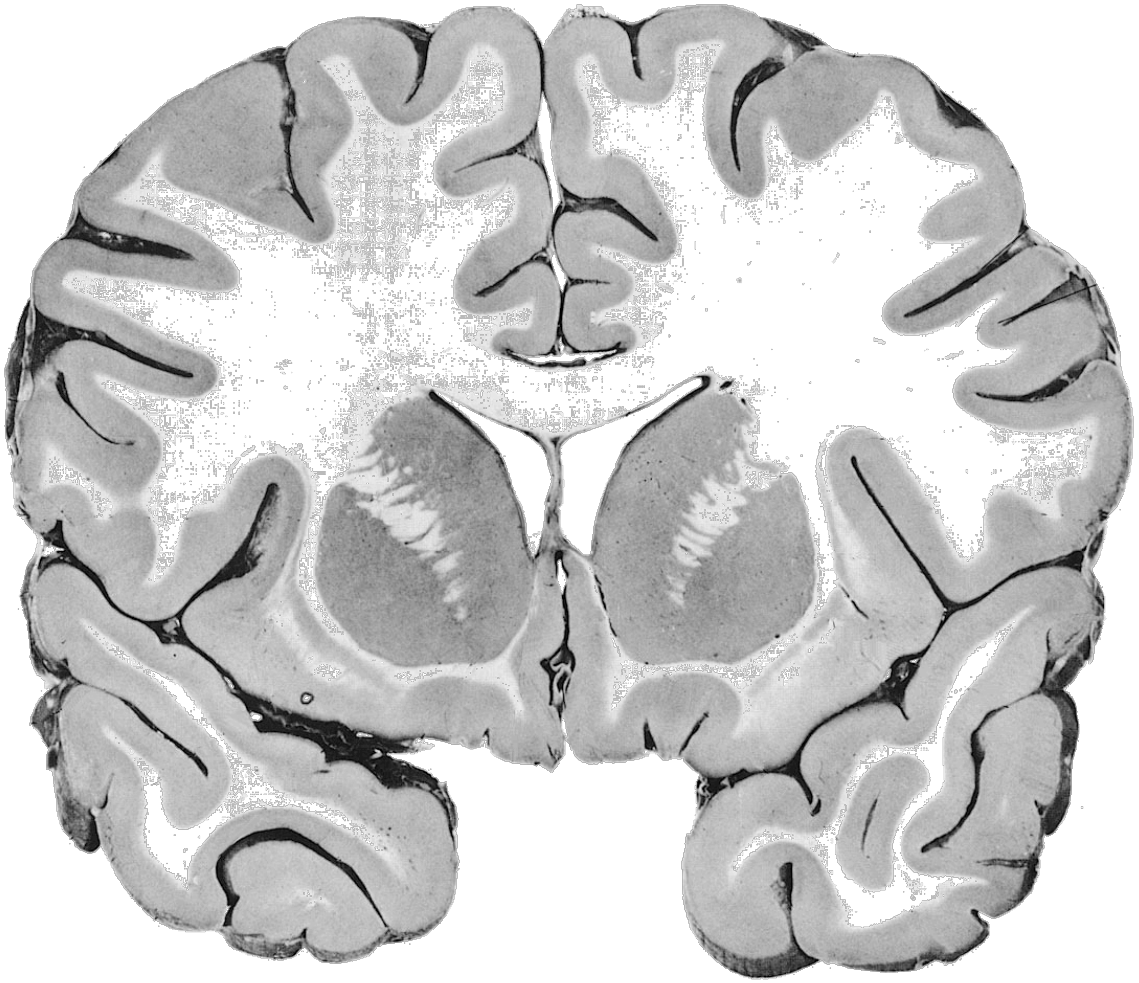

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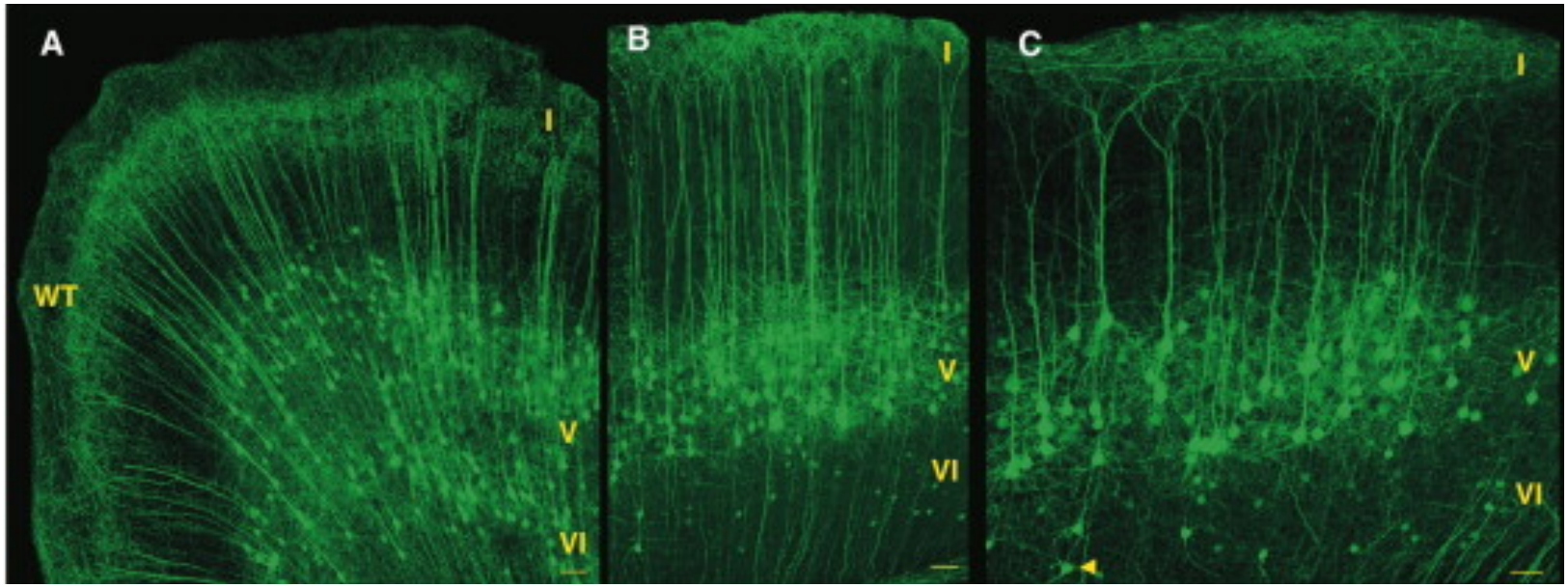

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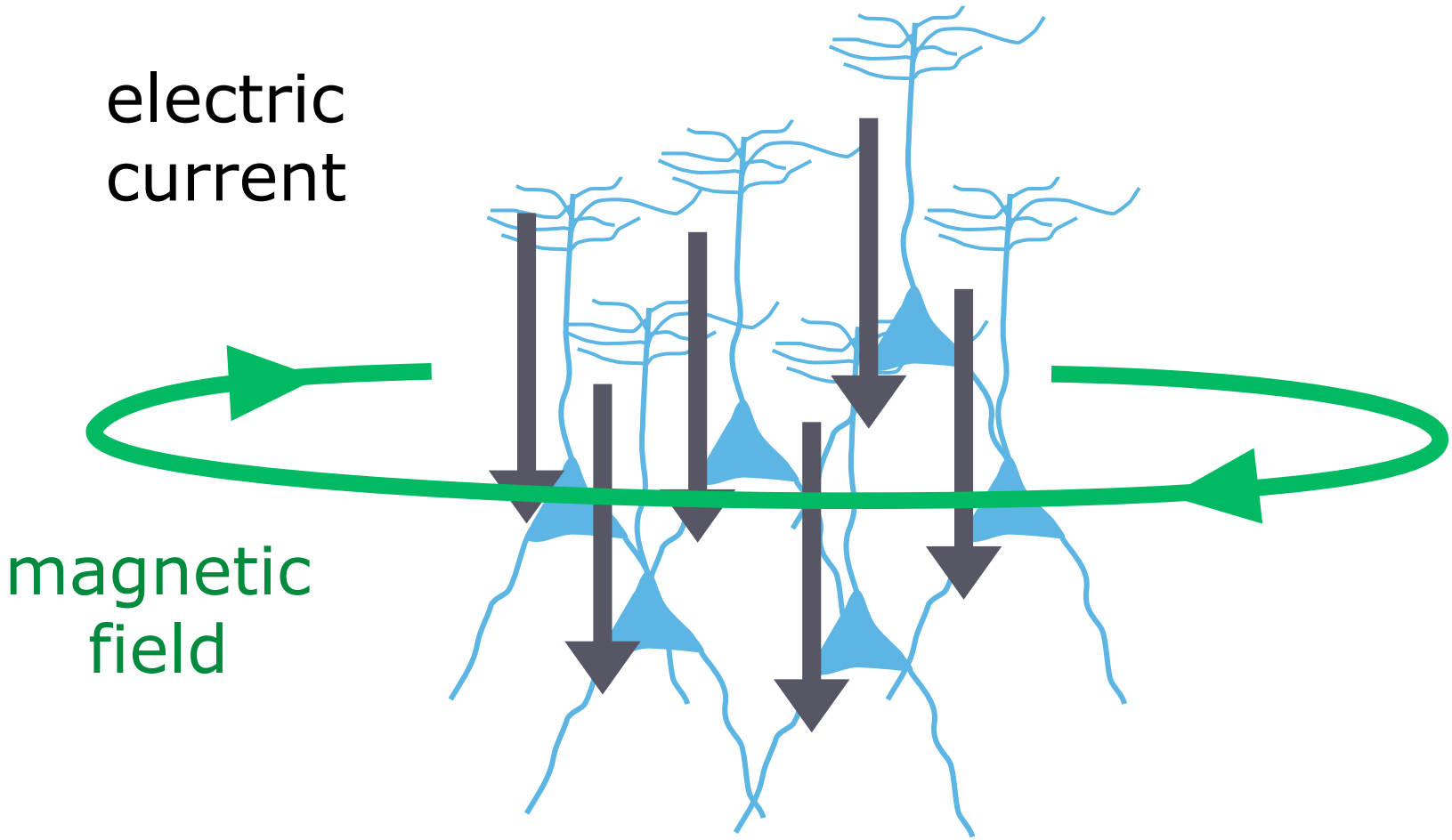
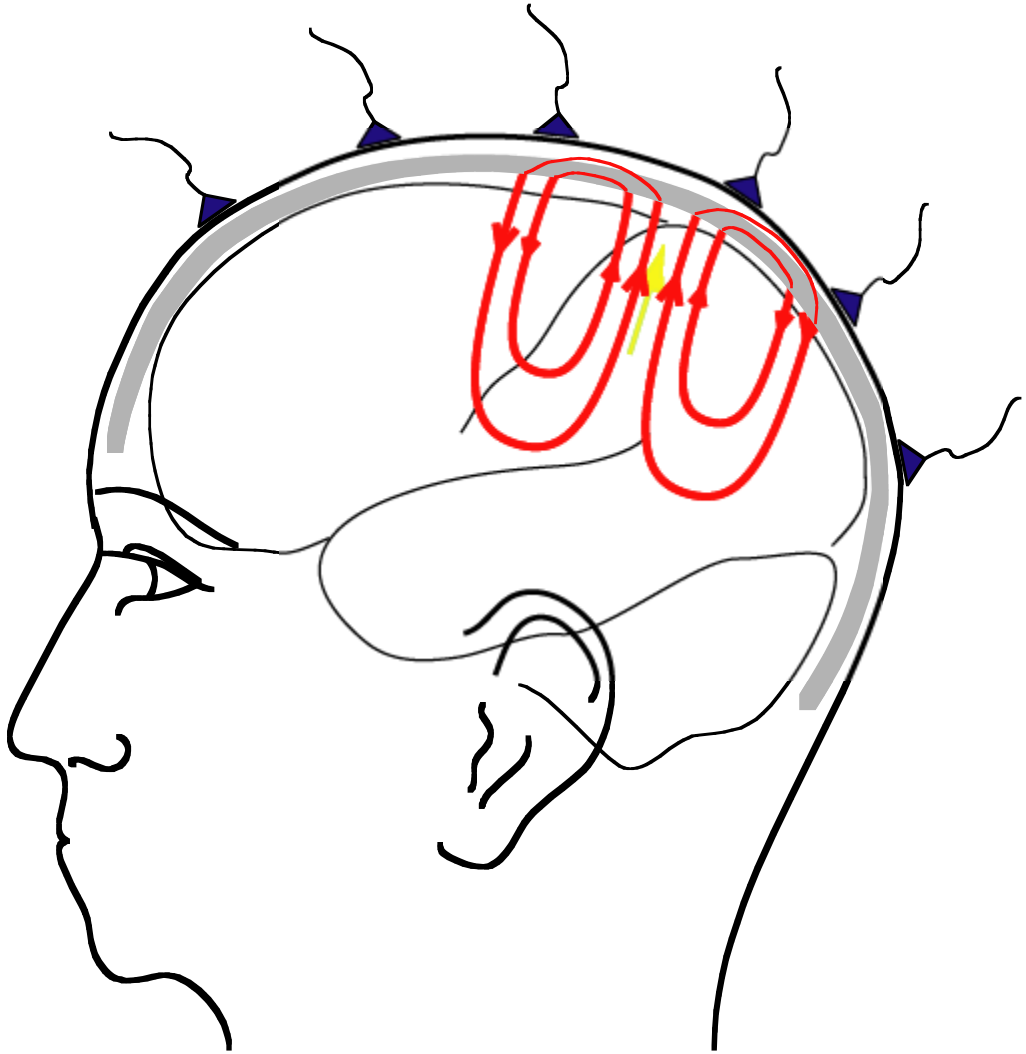
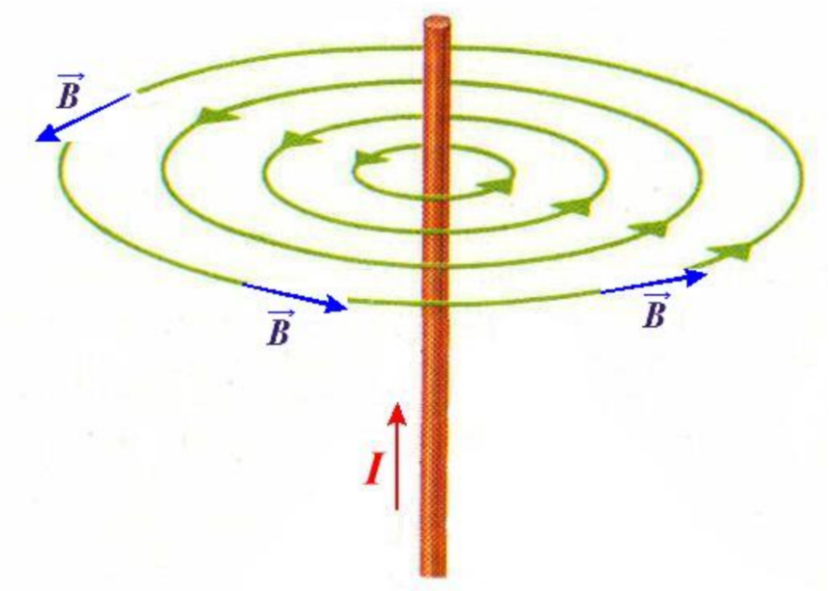
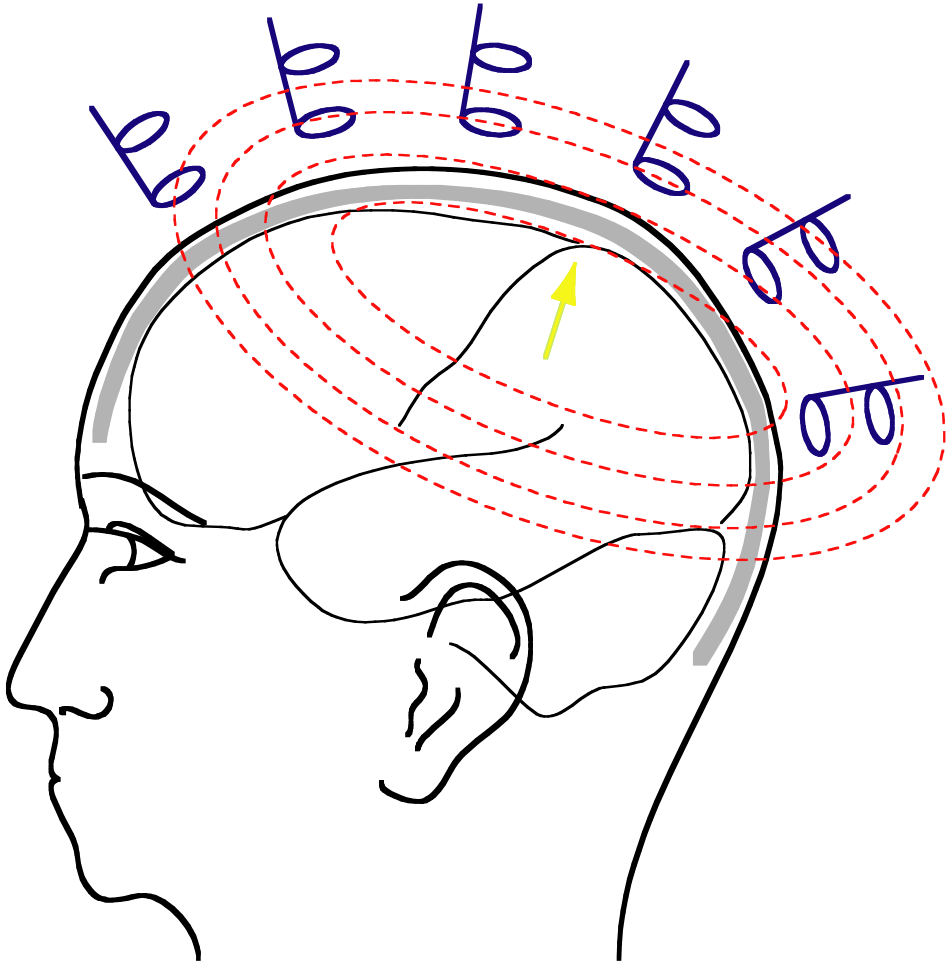


EEG volume conduction



Electric current → magnetic field

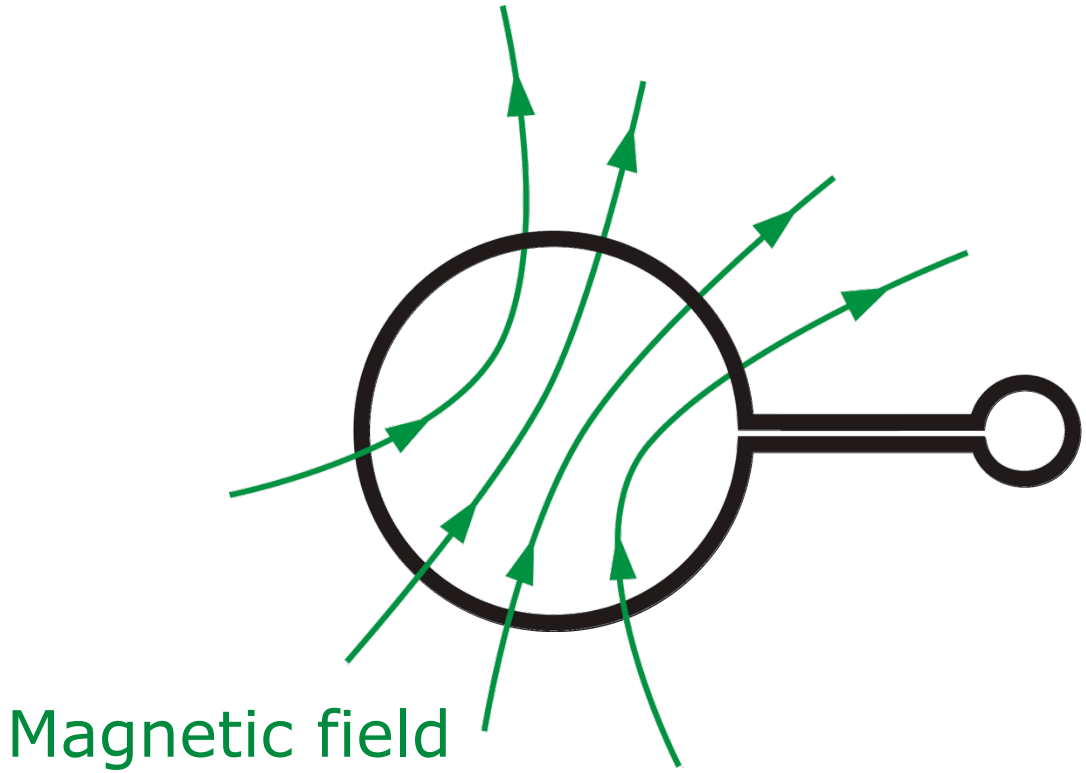




Technical challenges of MEG

- Sensitive magnetic detectors
  - superconductive sensors
- Deal with environmental noise
  - shielding
  - sensor design

MEG sensors - magnetometer



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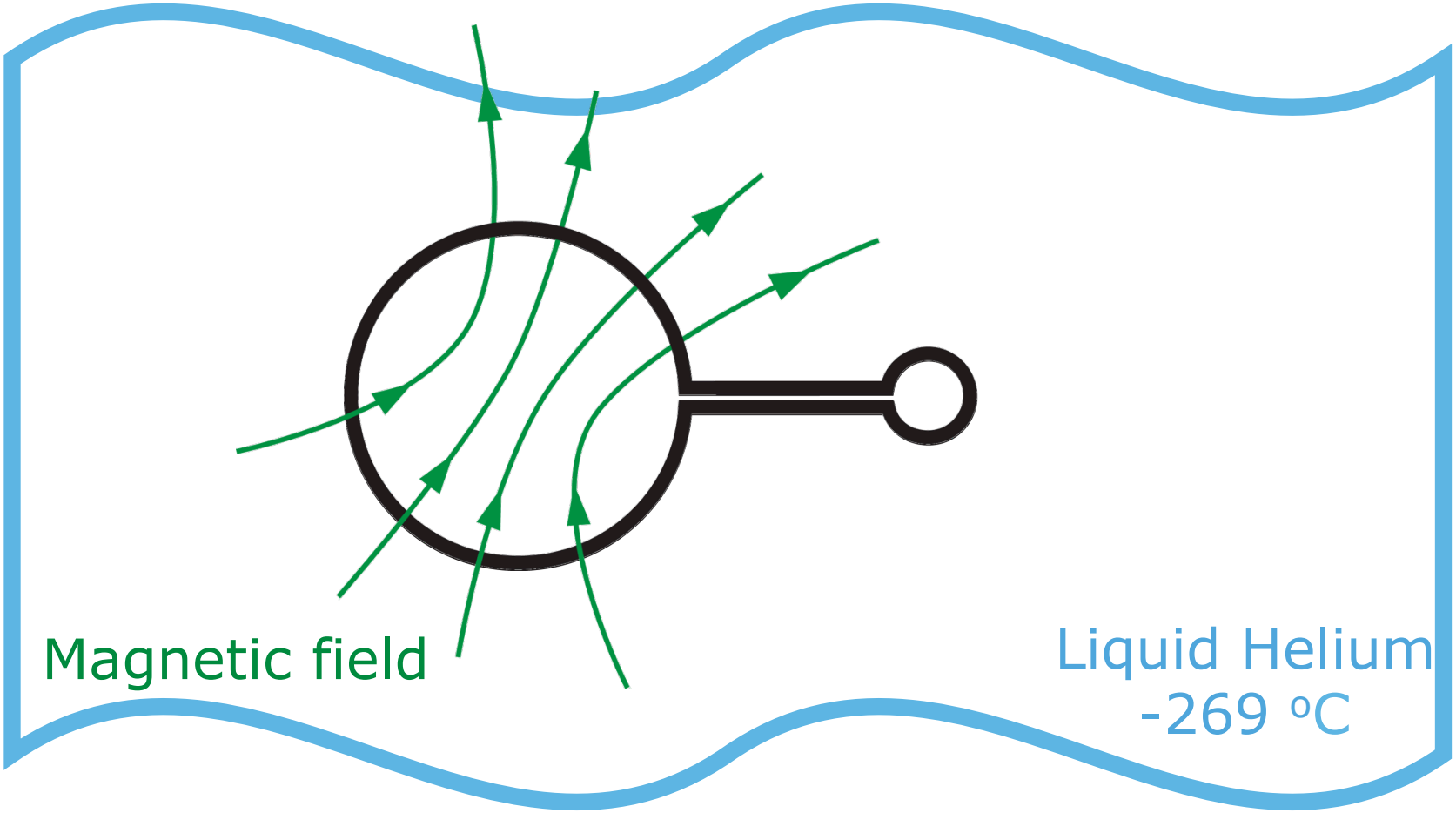

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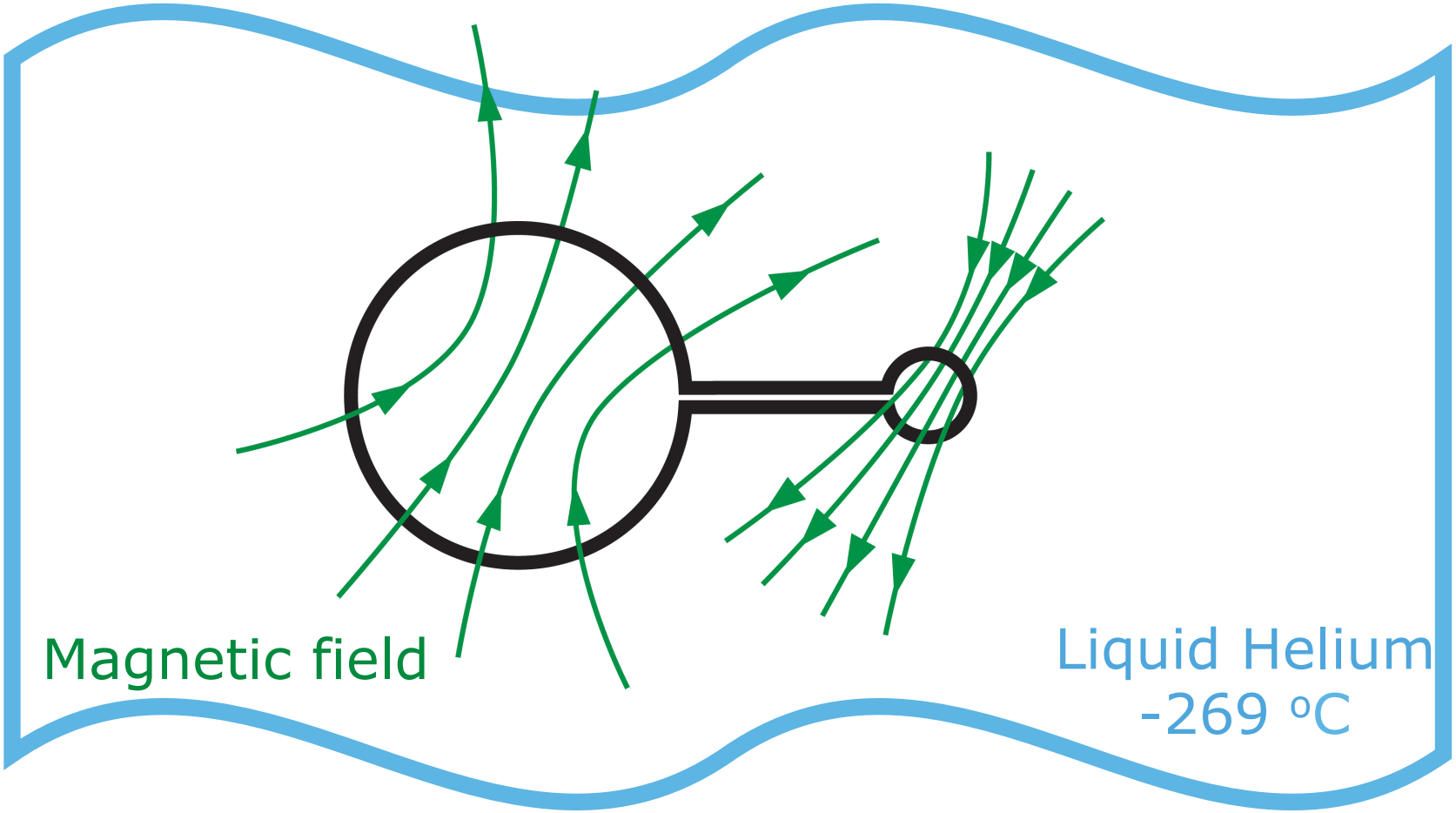

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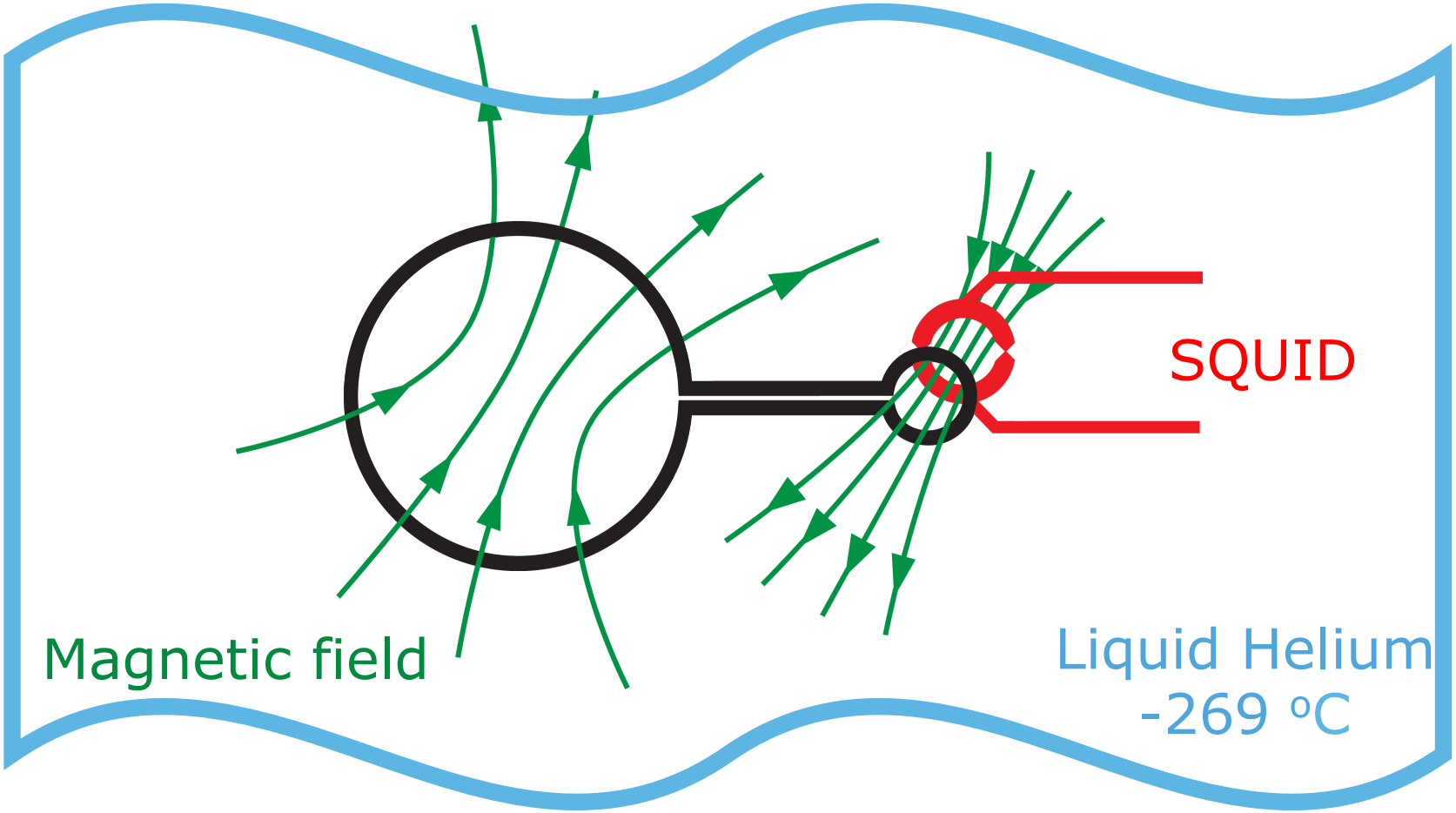
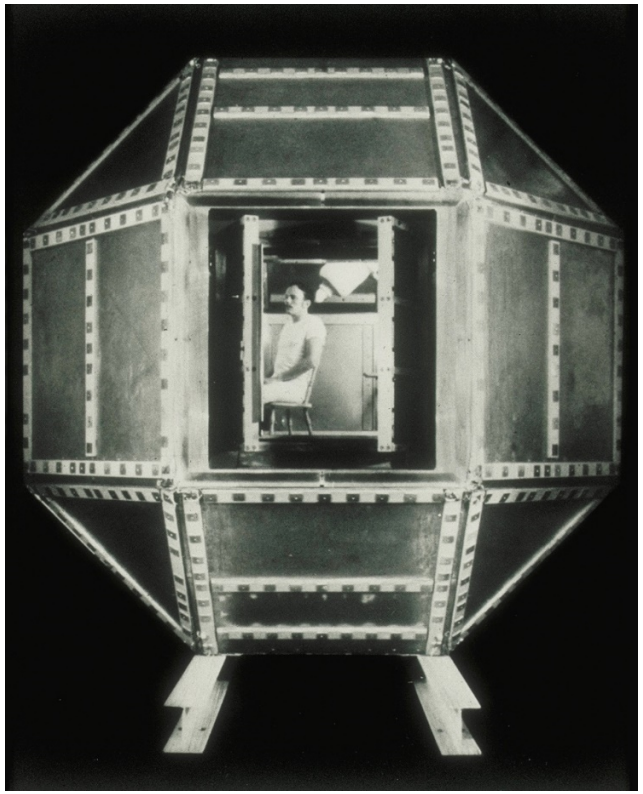
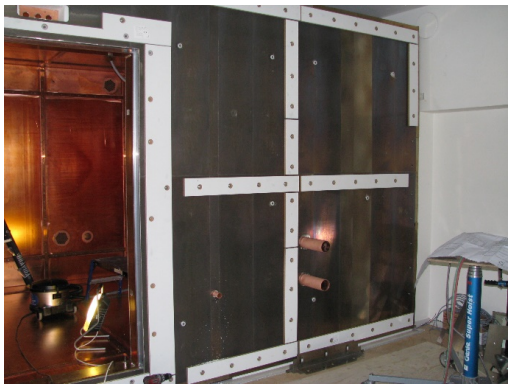


Passive Shielding

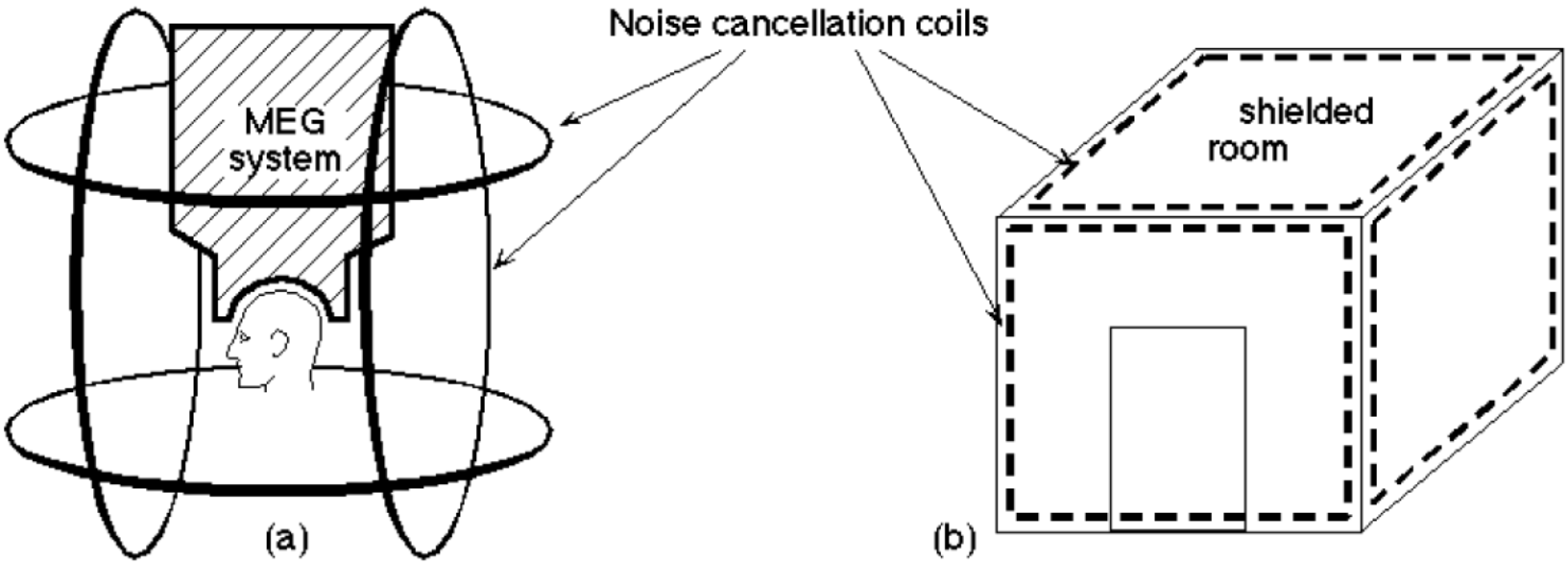


The magnetically shielded room built by David Cohen at MIT's Francis Bitter National Magnet Laboratory in 1969.

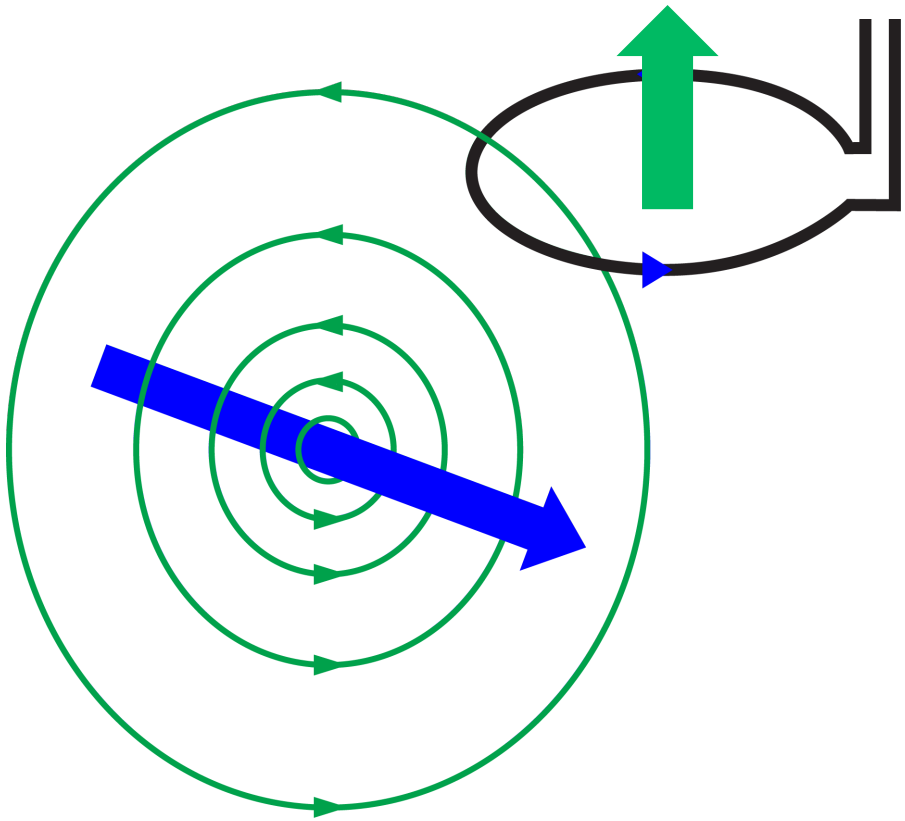


The magnetically shielded room at NatMEG in Stockholm (2013).

Active Shielding



Magnetometer



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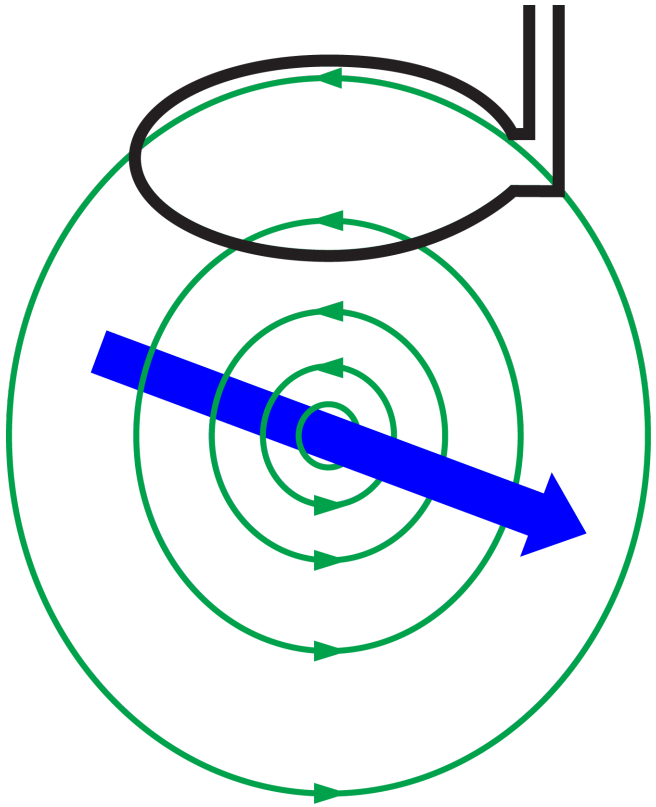

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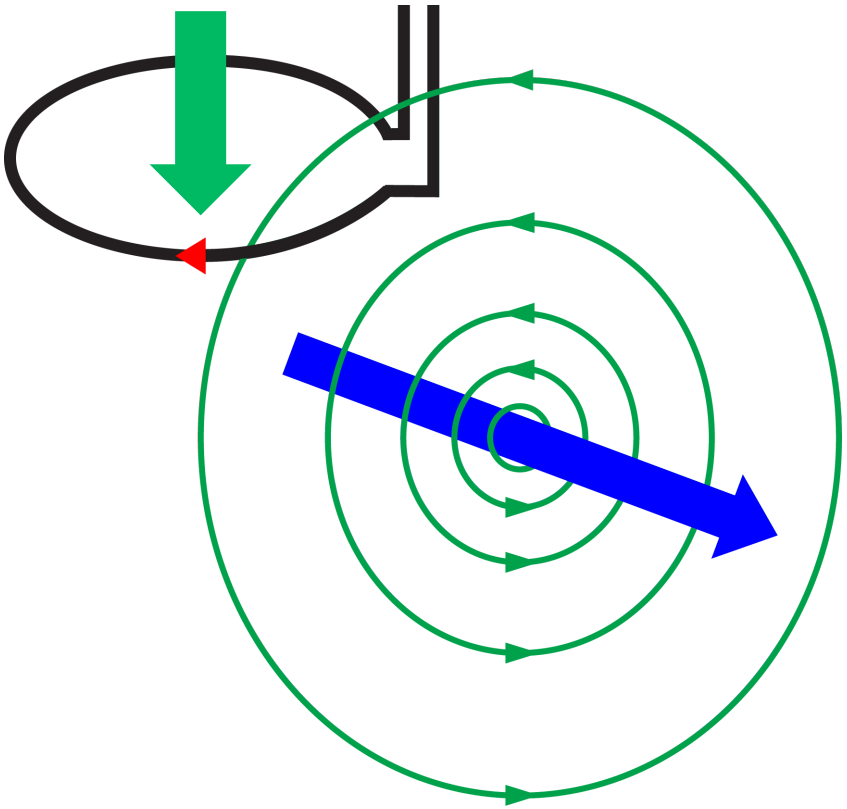
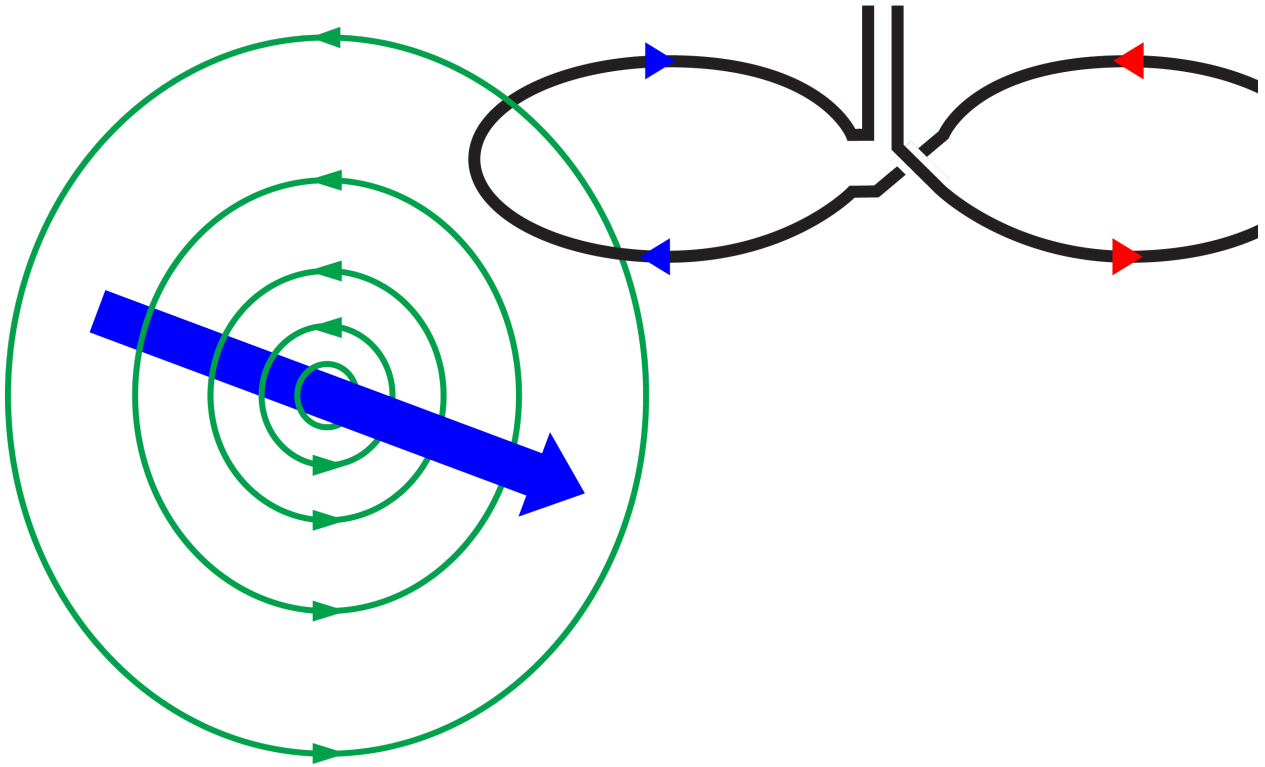
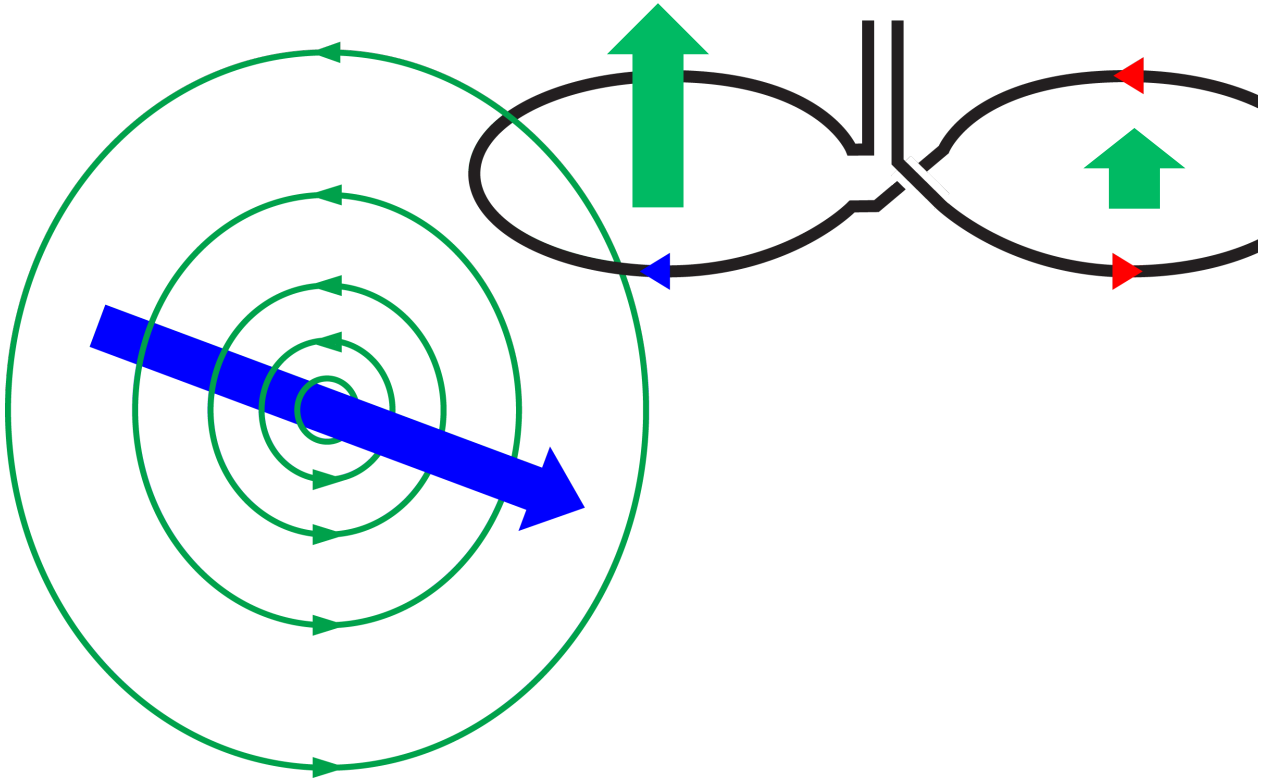




Planar gradiometer



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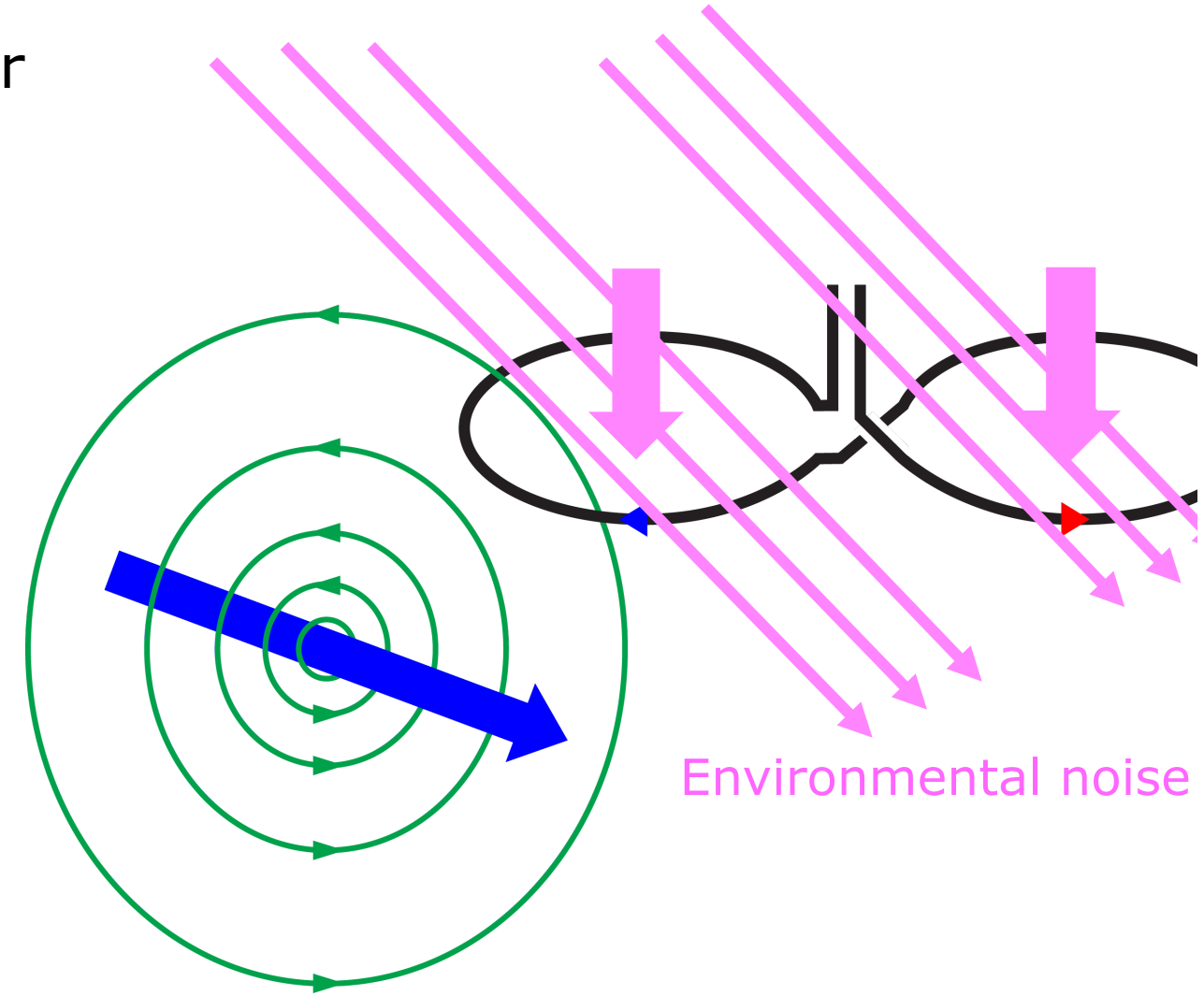

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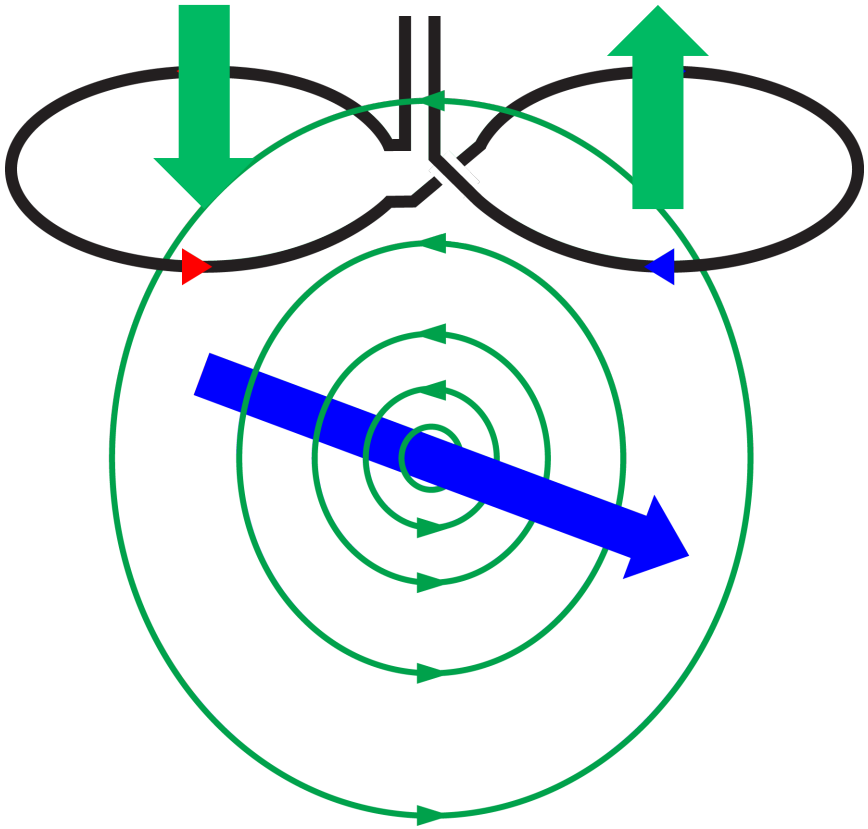
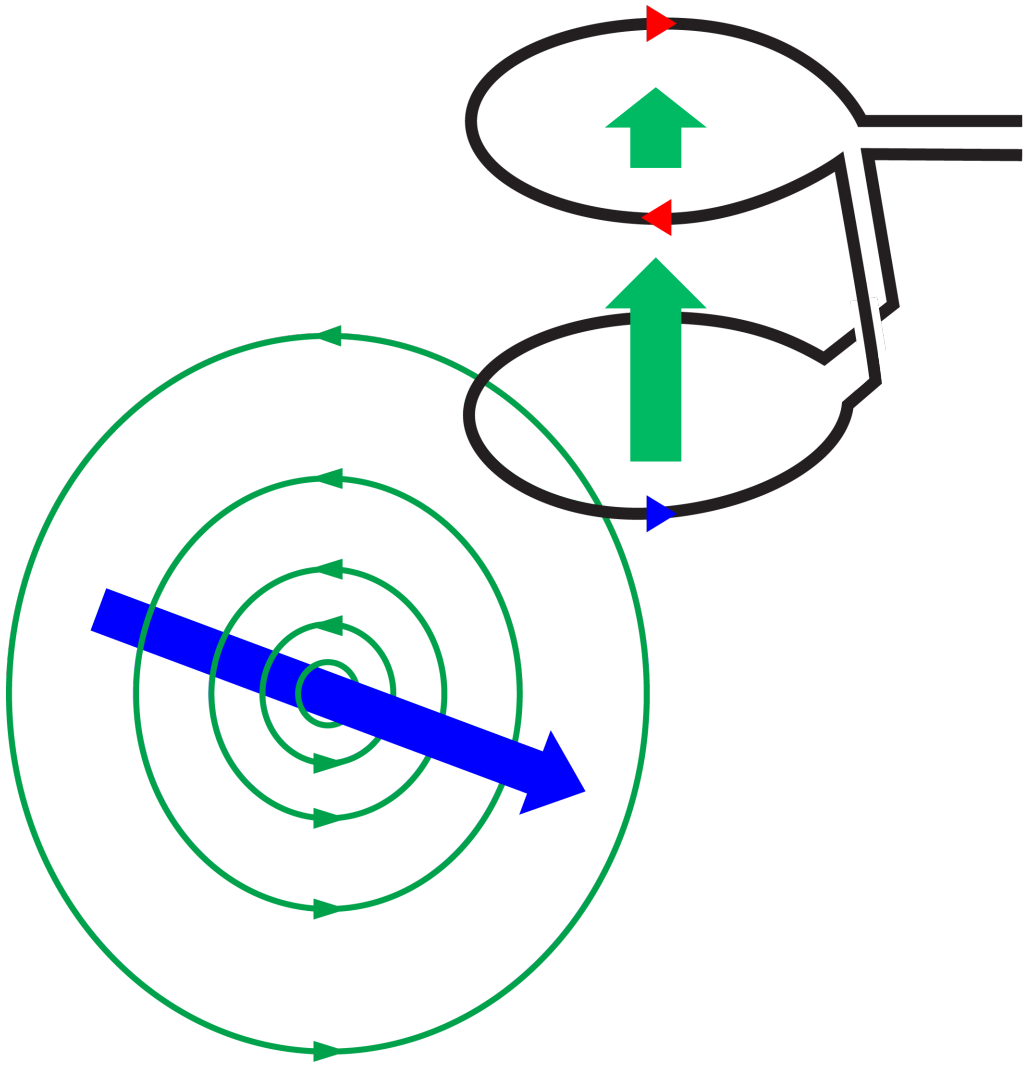


Axial gradiometer



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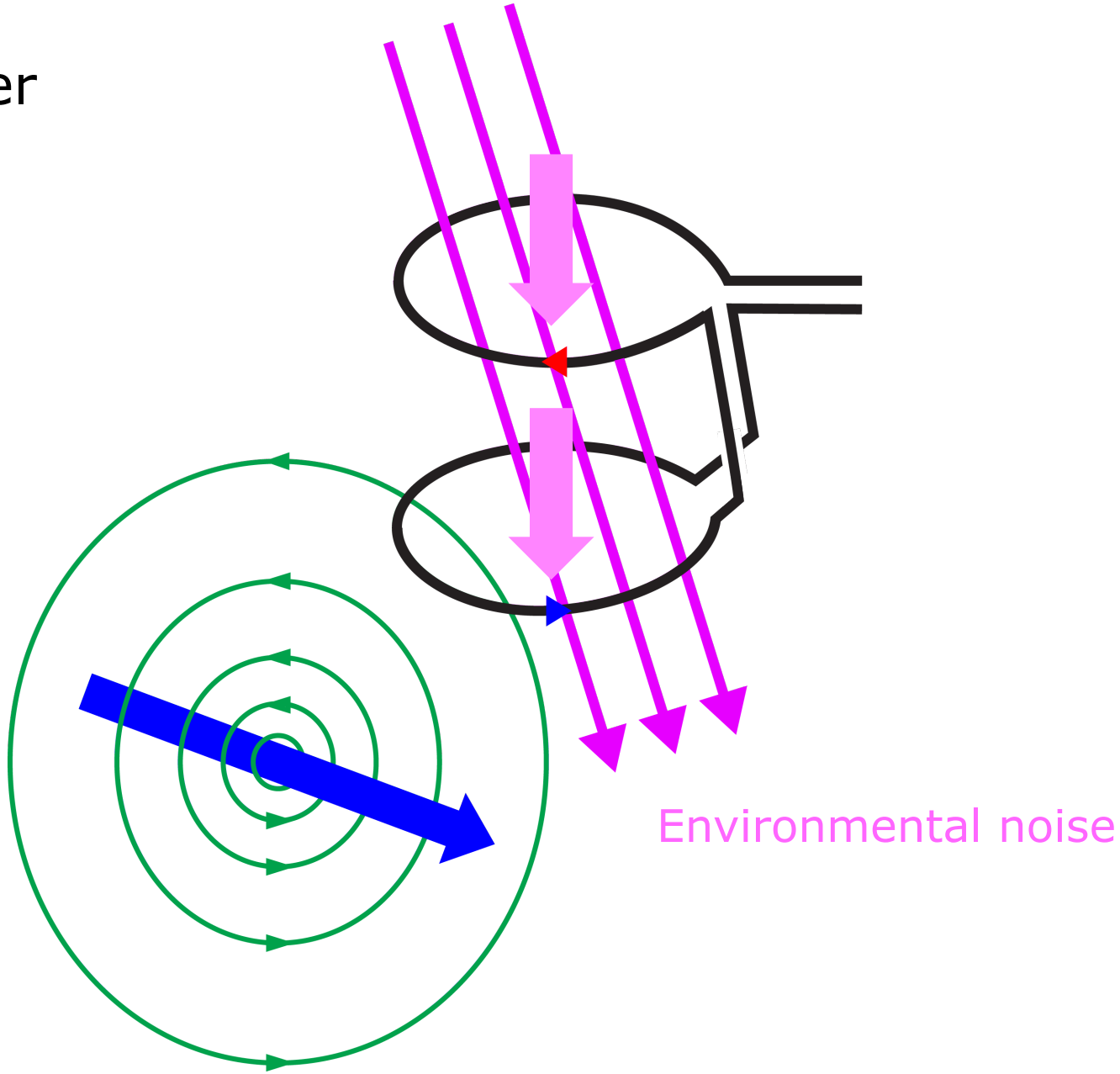
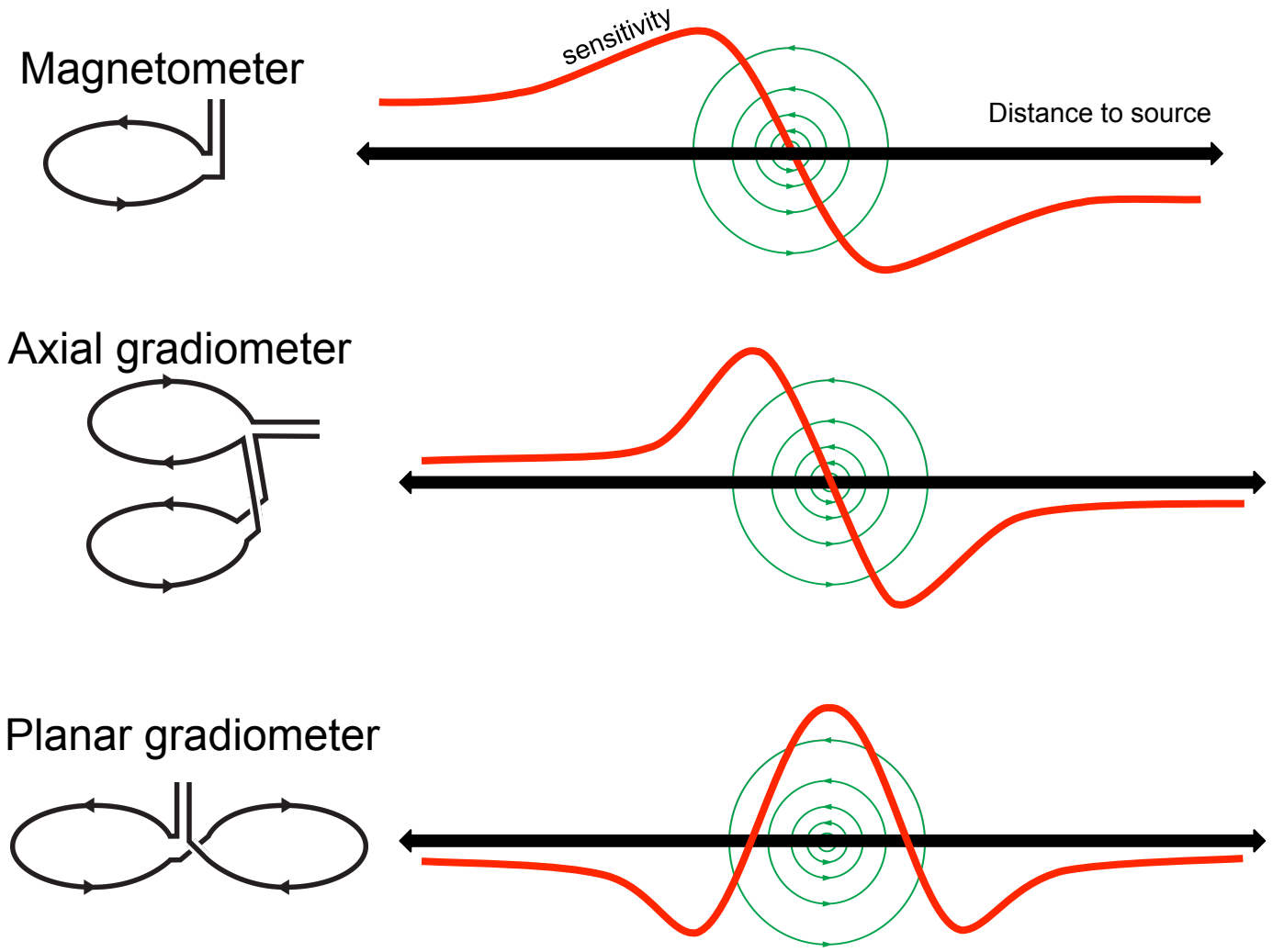


Different sensitivity profiles



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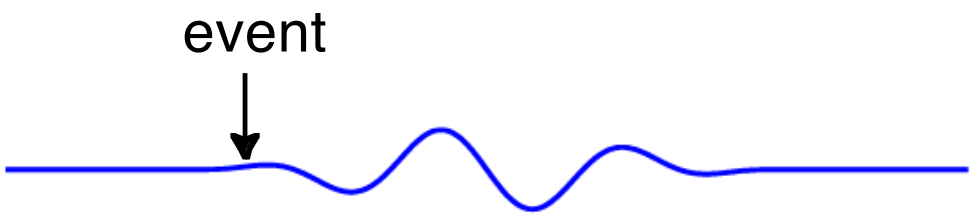
Analyzing those signals with FieldTrip

Background on the FieldTrip toolbox

Evoked activity



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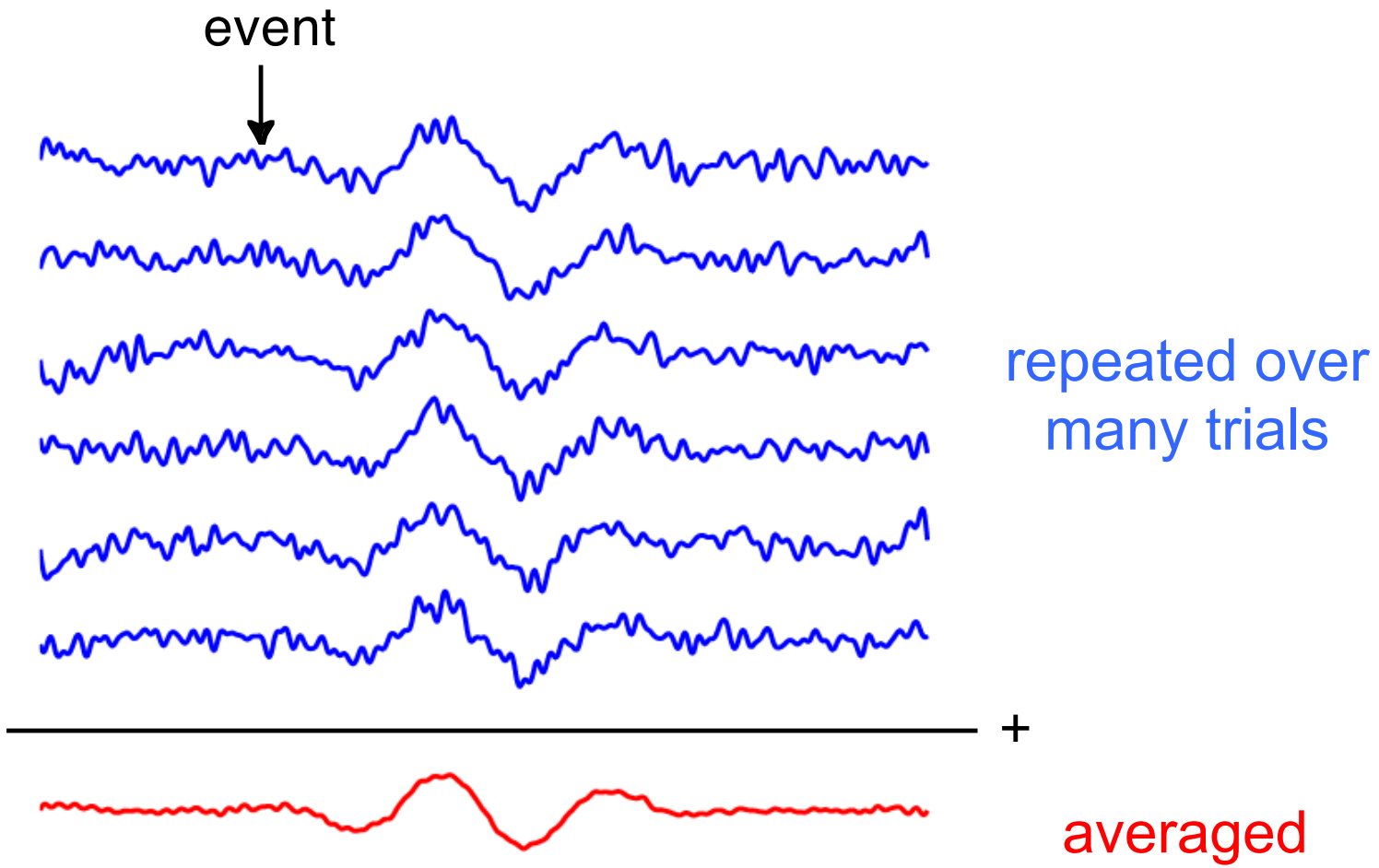
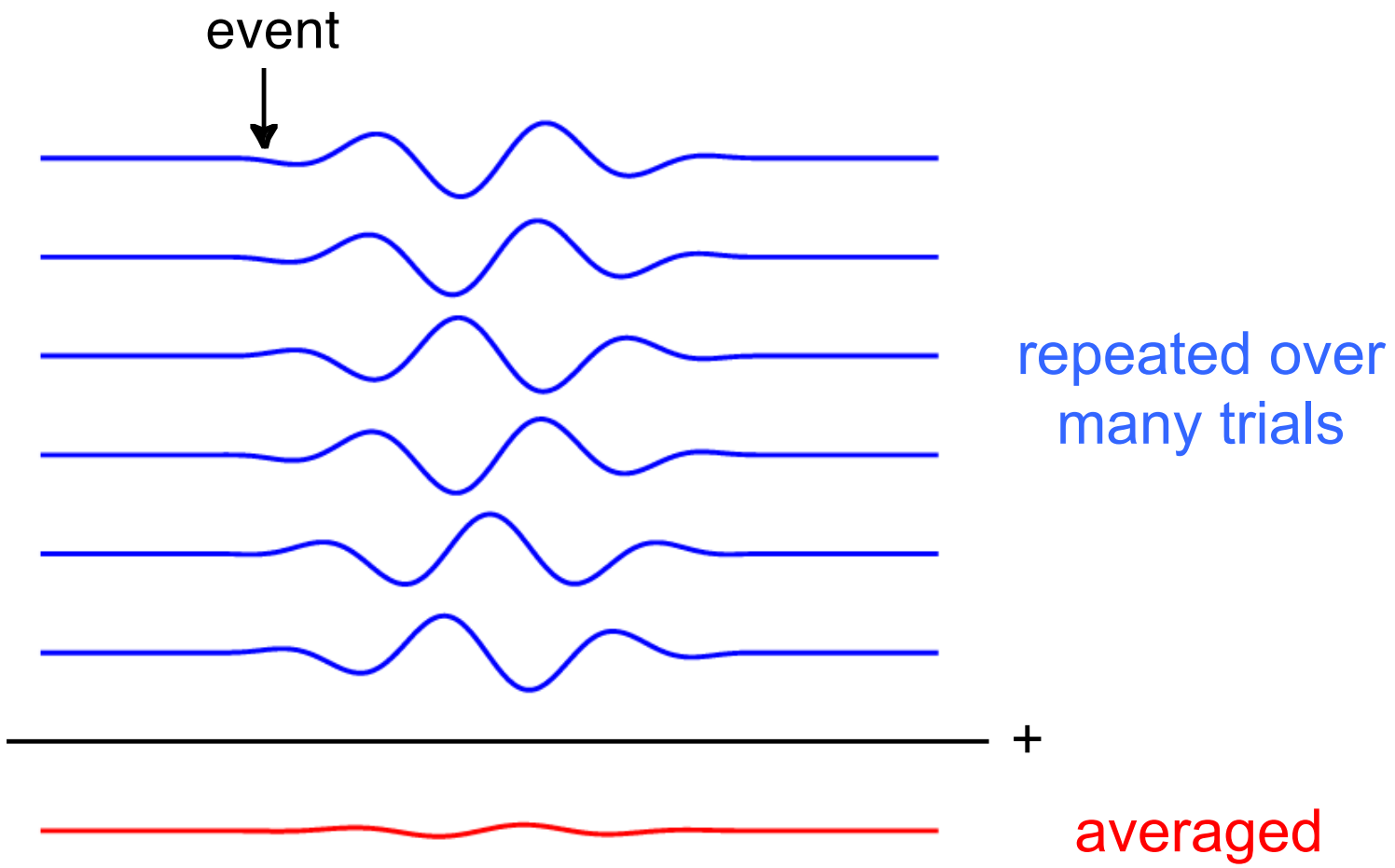


Induced activity



M/EEG signal characteristics considered during analysis

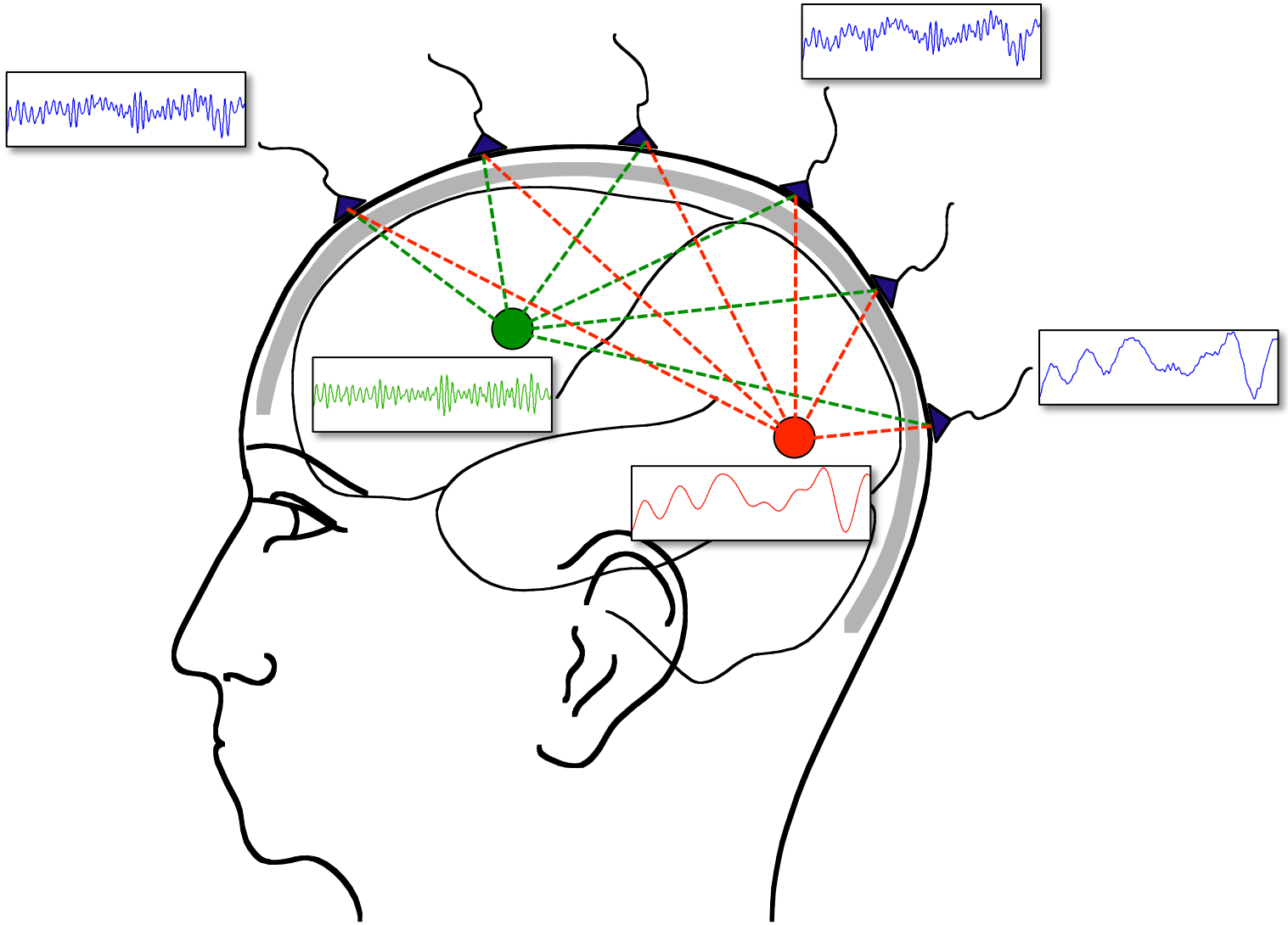
timecourse of activity
-> ERP

spectral characteristics
-> power spectrum

temporal changes in power
-> time-frequency response (TFR)

spatial distribution of activity over the head
-> source reconstruction

Superposition of source activity



Separating activity of different sources (and noise)

- Use the temporal aspects of the data at the channel level
  - ERF latencies
  - ERF difference waves
  - Filtering the time-series
  - Spectral decomposition
- Use the spatial aspects of the data
  - Volume conduction model of head
  - Estimate source model parameters

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Some FieldTrip basics

dataout = functionname(cfg, datain, …)

functionname(cf

dataout = functi

the “cfg” argument is a configuration structure, e.g.

```
cfg.channel = {‘C3’, C4’, ‘F3’, ‘F4’}
cfg.foilim = [1 70]
```

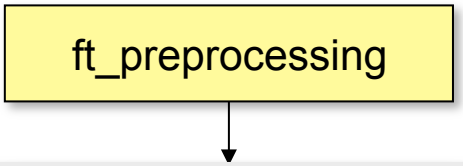
FieldTrip v.s. default Matlab

dataout = functionname(cfg, datain, …)

cfg.key1 = value1
cfg.key2 = value2

dataout = functionname(datain, ‘key1’, ‘value1’, …)

Using functions in an analysis protocol



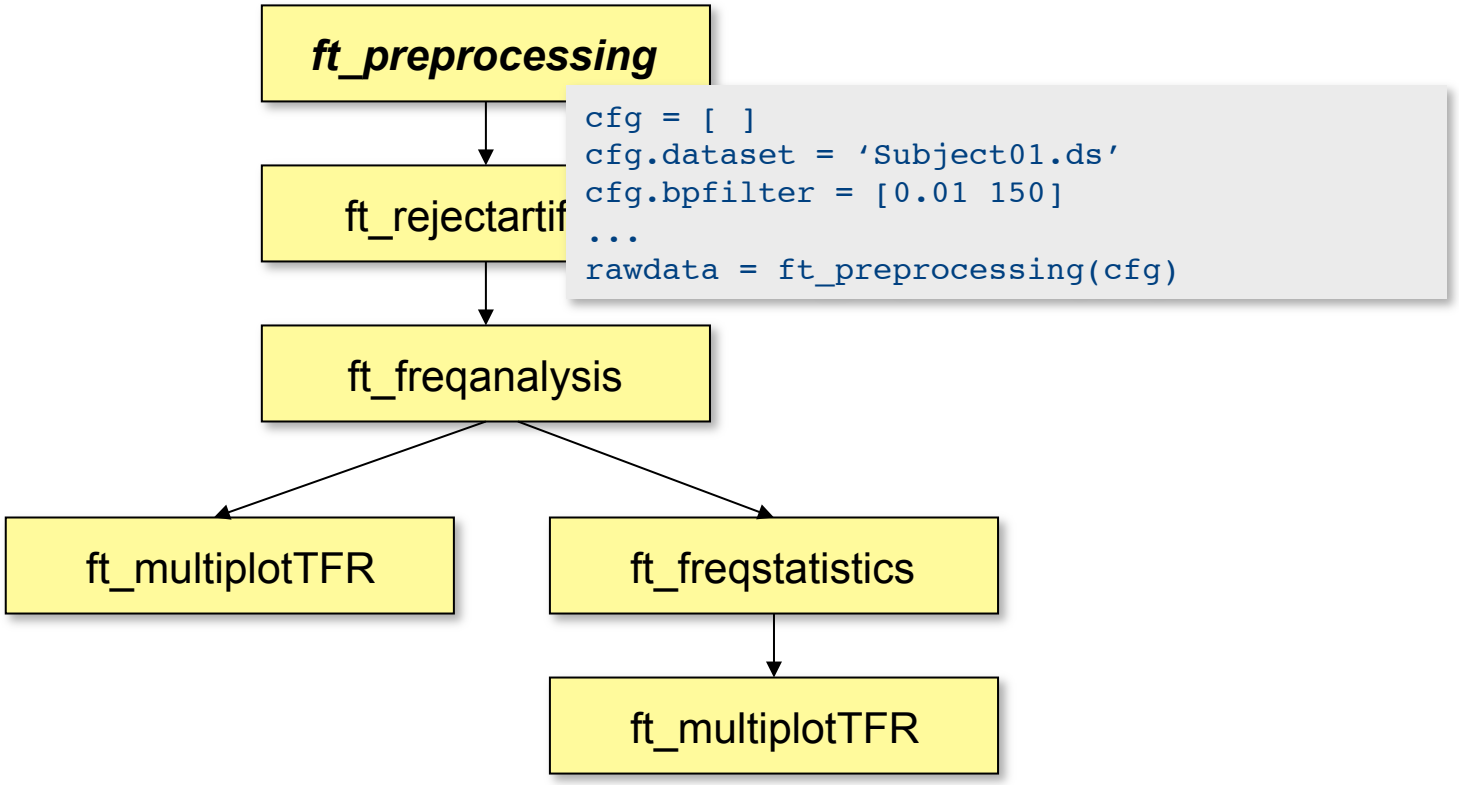

```
FT_PREPROCESSING reads MEG and/or EEG data according to user-specified
trials and applies several user-specified preprocessing steps to the
signals.

Use as
  [data] = ft_preprocessing(cfg)
or
  [data] = ft_preprocessing(cfg, data)

The first input argument "cfg" is the configuration structure, which
contains all details for the dataset filenames, trials and the
preprocessing options. You can only do preprocessing after defining the
segments of data to be read from the file (i.e. the trials), which is for
example done based on the occurence of a trigger in the data.

...
```

Using functions in an analysis protocol



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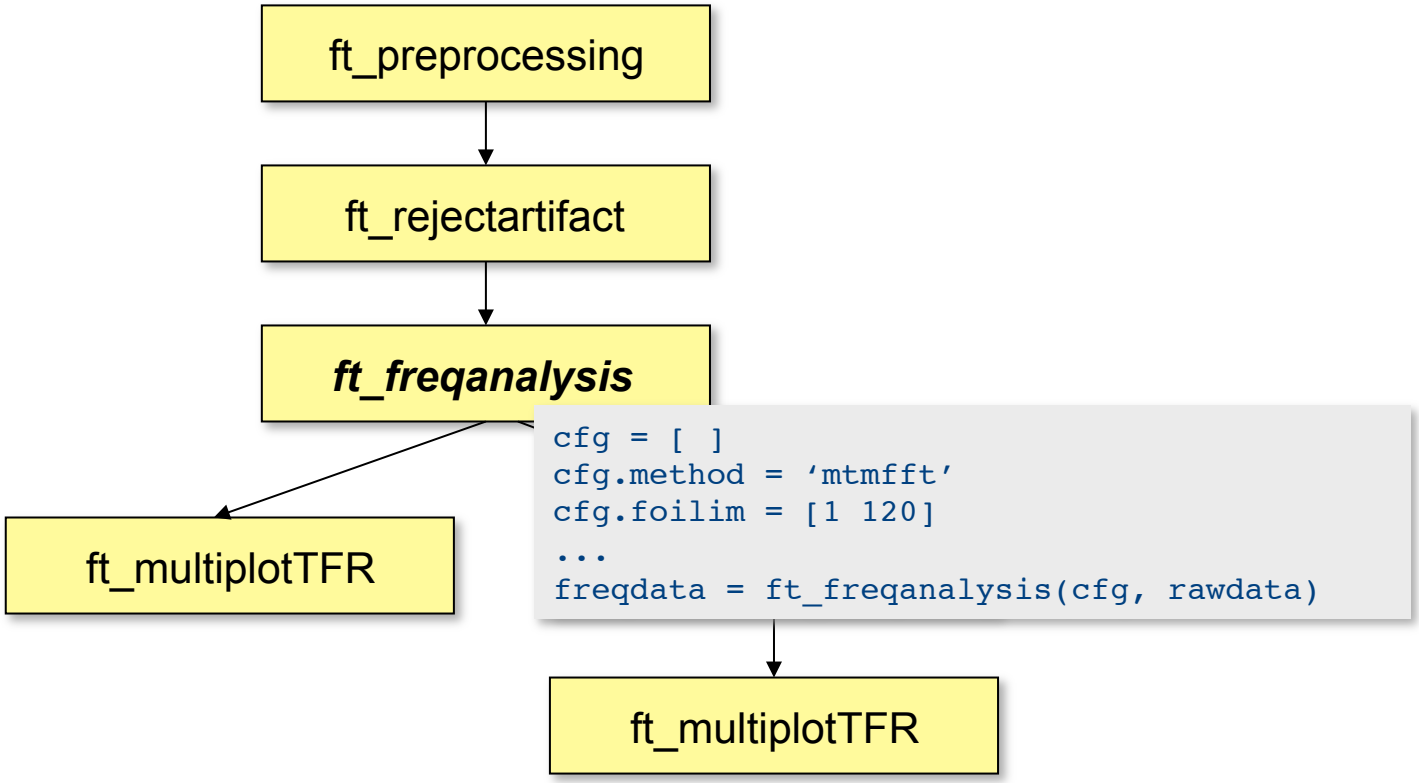


Raw data structure

```
rawData =
        label: {151x1 cell}
        trial: {1x87 cell}
         time: {1x87 cell}
          hdr: [1x1 struct]
          cfg: [1x1 struct]
```

Event related response

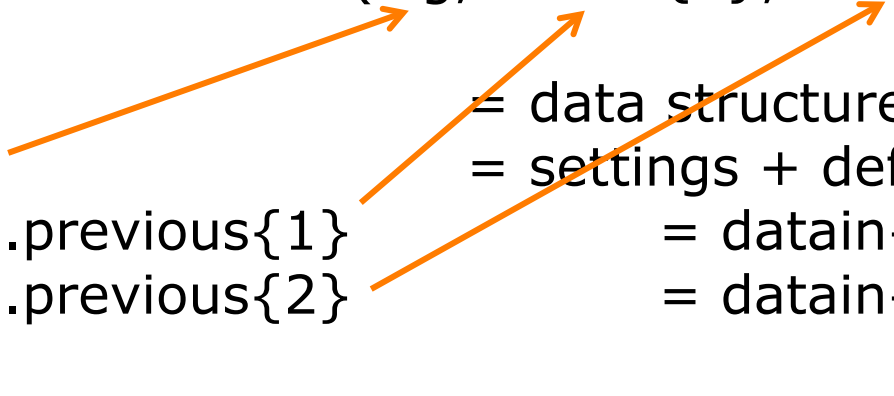
```
erpData =
        label: {151x1 cell}
          avg: [151x900 double]
          var: [151x900 double]
         time: [1x900 double]
       dimord: 'chan_time'
          cfg: [1x1 struct]
```

Keeping track of your analysis

input cfg structure specifies parameters

output cfg structure keeps history

dataout = functionname(cfg, datain{1}, datain{2},…)



details of computations are kept with data

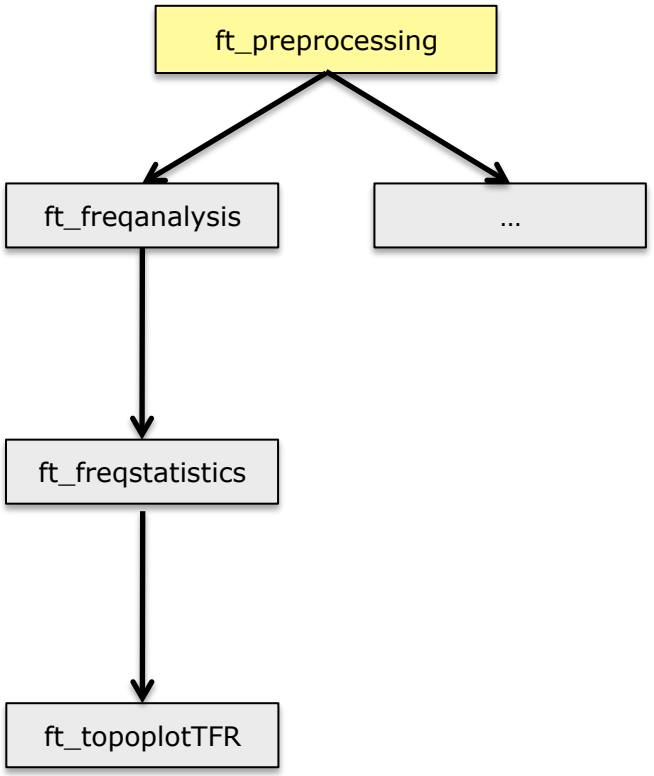
previous data is not kept, but can be reconstructed using cfg.previous.previous…

Example use in scripts

```
cfg = []
cfg.dataset = 'Subject01.ds'
cfg.bpfilter = [0.01 150]
...
rawdata = ft_preprocessing(cfg)
```

```
cfg = []
cfg.method = 'mtmfft'
cfg.foilim = [1 120]
...
freqdata = ft_freqanalysis(cfg, rawdata)
```

```
cfg = []
cfg.method = 'montecarlo'
cfg.statistic = 'indepsamplesT'
cfg.design = [1 2 1 2 2 1 2 1 1 2 ... ]
...
freqstat = ft_freqstatistics(cfg, freqdata)
```



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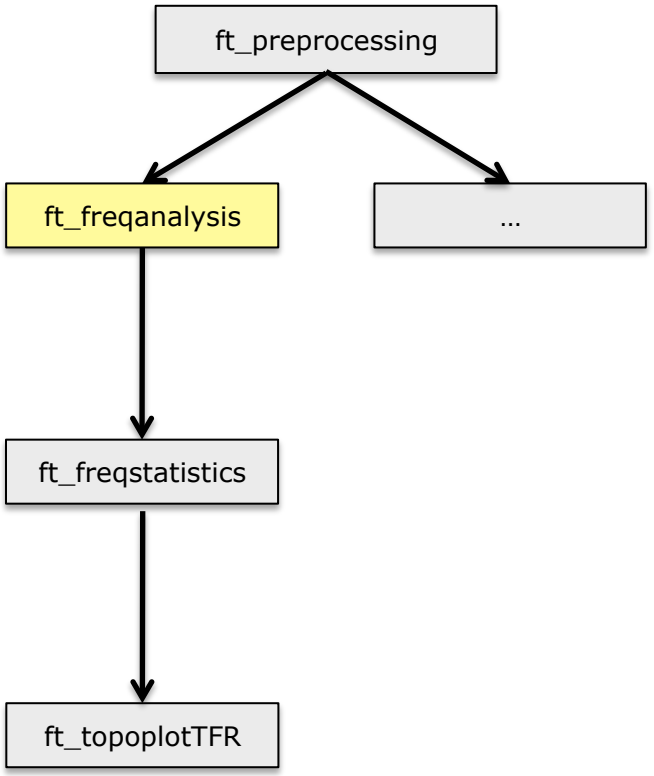

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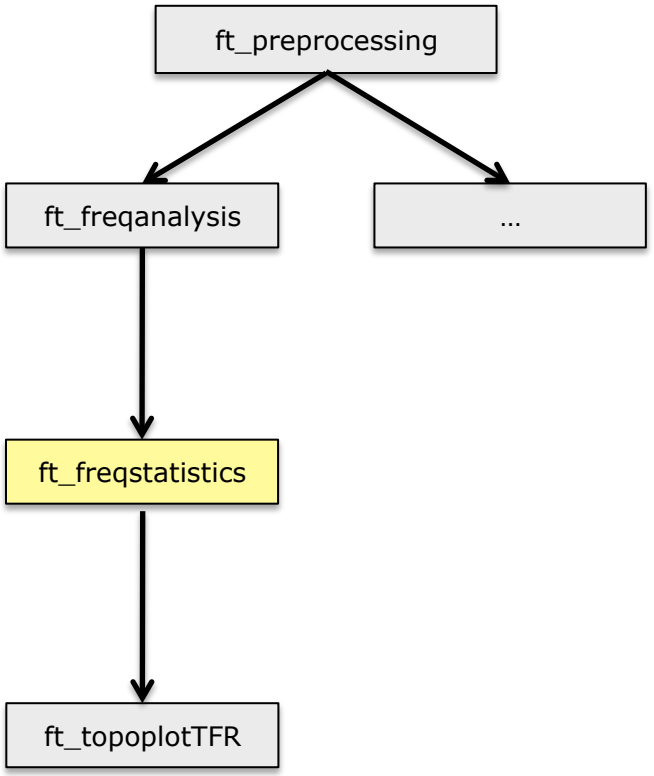

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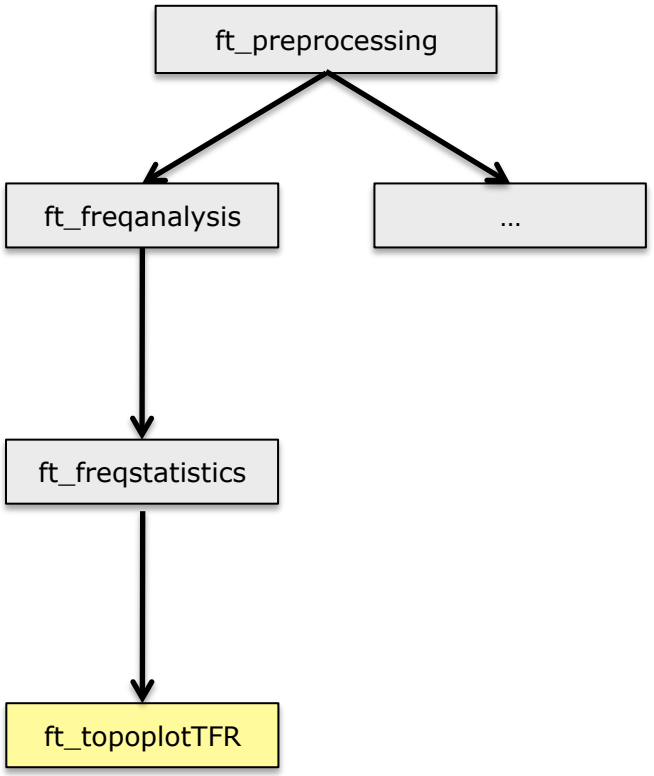

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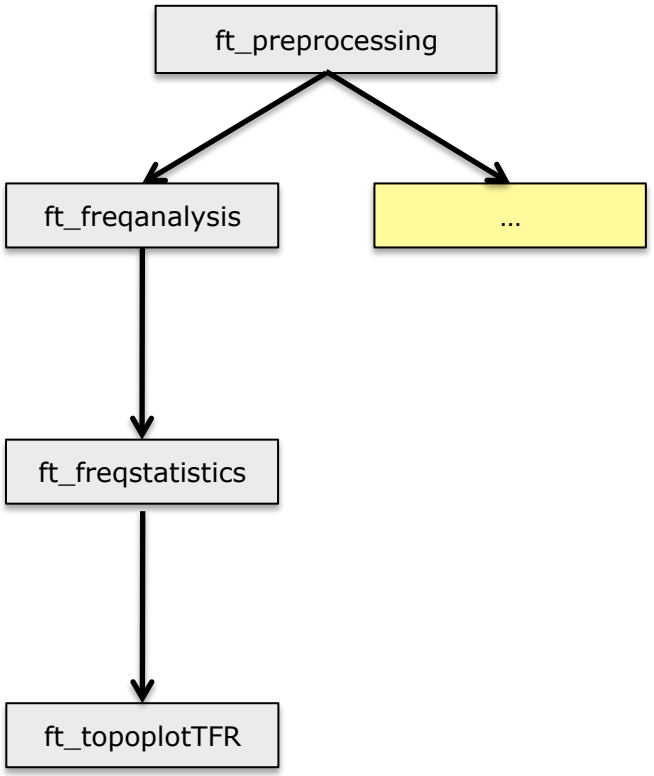


Example use in scripts

```
subj = {'S01.ds', 'S02.ds', …}
trig = [1 3 7 9]

for s=1:nsubj
for c=1:ncond

  cfg = []
  cfg.dataset  = subj{s}
  cfg.trigger  = trig(c)
  rawdata{s,c} = ft_preprocessing(cfg)

  cfg = []
  cfg.method    = 'mtmfft'
  cfg.foilim    = [1 120]
  freqdata{s,c} = ft_freqanalysis(cfg, rawdata{s,c})

end
end
```

Example use in scripts

```
subj = {'S01.ds', 'S02.ds', …}
trig = [1 3 7 9]

for s=1:nsubj
for c=1:ncond

  cfg = []
  cfg.dataset  = subj{s}
  cfg.trigger  = trig(c)
  rawdata      = ft_preprocessing(cfg)


  filename = sprintf('raw%s_%d.mat', subj{s}, trig(c));
  save(filename, 'rawdata')


end
end
```

Example use in distributed computing – with qsub

```
subj = {'S01.ds', 'S02.ds', …}
trig = [1 3 7 9]

for s=1:nsubj
for c=1:ncond

  cfgA{s,c} = []
  cfgA{s,c}.dataset    = subj{s}
  cfgA{s,c}.trigger    = trig(c)
  cfgA{s,c}.outputfile = sprintf('raw%s_%d.mat', subj{s}, trig(c))

  cfgB{s,c} = []
  cfgB{s,c}.dataset    = subj{s}
  cfgB{s,c}.trigger    = trig(c)
  cfgB{s,c}.inputfile  = sprintf('raw%s_%d.mat', subj{s}, trig(c));
  cfgB{s,c}.outputfile = sprintf('freq%s_%d.mat', subj{s}, trig(c));

end
end

qsubcellfun(@ft_preprocessing, cfgA)
qsubcellfun(@ft_freqanalysis,  cfgB)
```

Example use in distributed computing – with dfeval

```
subj = {'S01.ds', 'S02.ds', …}
trig = [1 3 7 9]

for s=1:nsubj
for c=1:ncond

  cfgA{s,c} = []
  cfgA{s,c}.dataset    = subj{s}
  cfgA{s,c}.trigger    = trig(c)
  cfgA{s,c}.outputfile = sprintf('raw%s_%d.mat', subj{s}, trig(c))

  cfgB{s,c} = []
  cfgB{s,c}.dataset    = subj{s}
  cfgB{s,c}.trigger    = trig(c)
  cfgB{s,c}.inputfile  = sprintf('raw%s_%d.mat', subj{s}, trig(c));
  cfgB{s,c}.outputfile = sprintf('freq%s_%d.mat', subj{s}, trig(c));

end
end

dfeval(@ft_preprocessing, cfgA)
dfeval(@ft_freqanalysis,  cfgB)
```

FieldTrip is a toolbox

the data and the separate functions are in the hands of the end-users

the scripts depend on the data properties, available memory and programming skills and style

scripts correspond to analysis protocols

- scripts can be reviewed by supervisors
- scripts are often shared with colleagues

Finding your way around in the FieldTrip toolbox

Matlab

help functionname

edit functionname

Website

http://www.fieldtriptoolbox.org

Email discussion list

Expertise in your local group

Talk outline

Recording signals from the brain

Temporal and spatial features of the signals

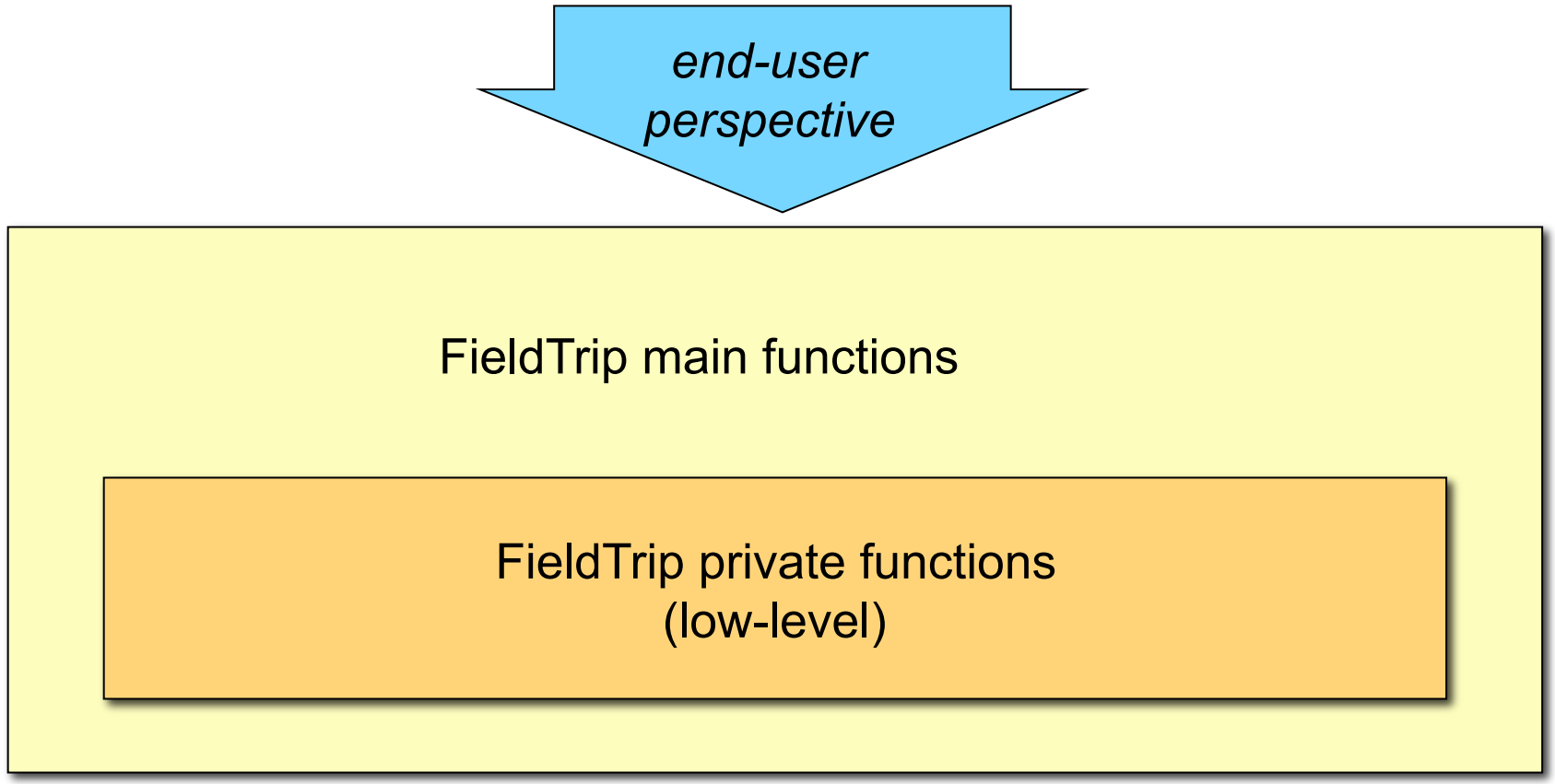
Analyzing those signals with FieldTrip

Background on the FieldTrip toolbox

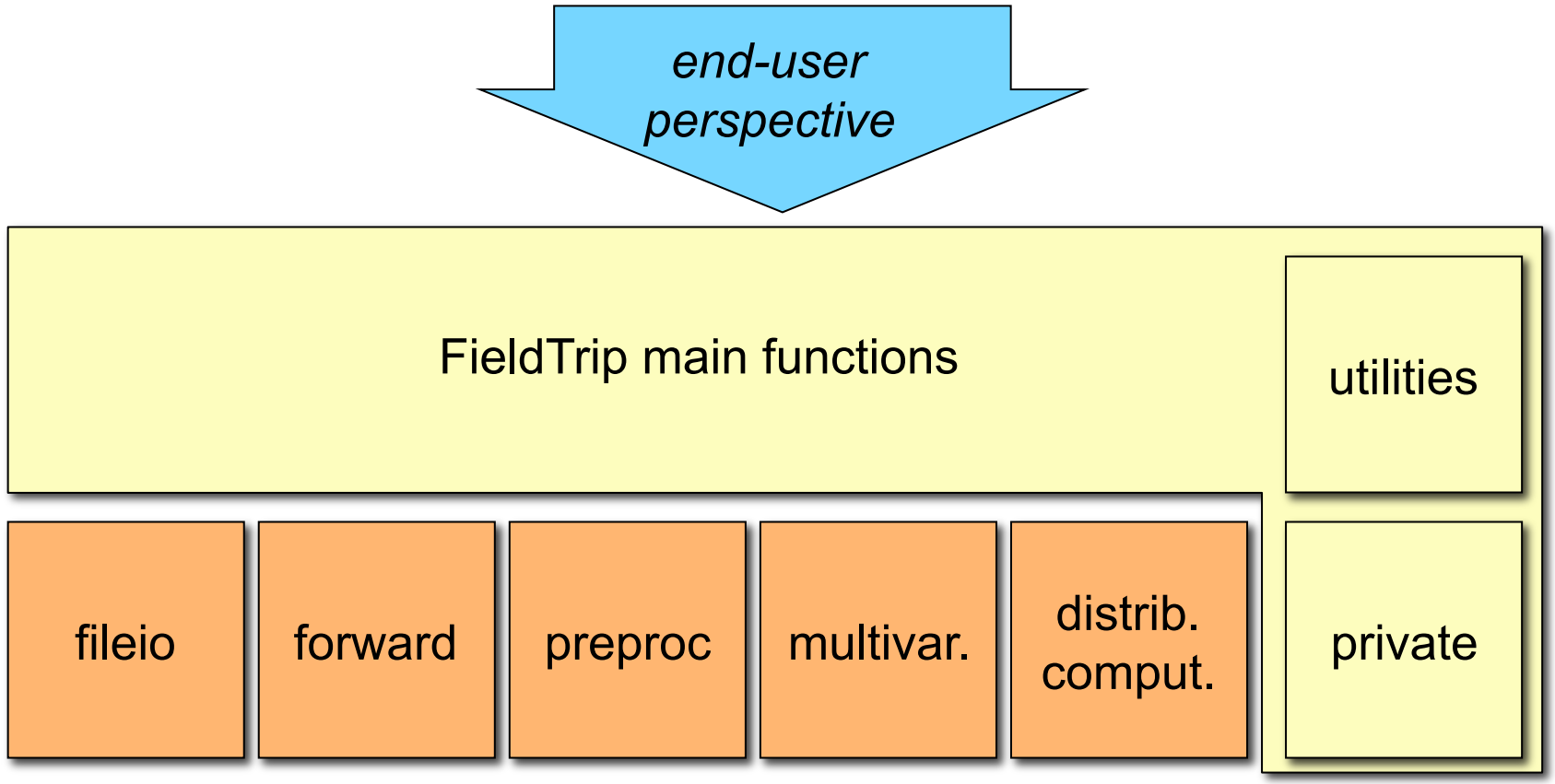
Who is the audience?

- experimental neuroscientists
  - no graphical user interface
  - more dedicated and ambitious researchers
- developers of other software packages
  - SPM
  - EEGLAB
  - BESA
  - BCI2000
  - SIMBIO
  - OpenMEEG
- developers of analysis methods

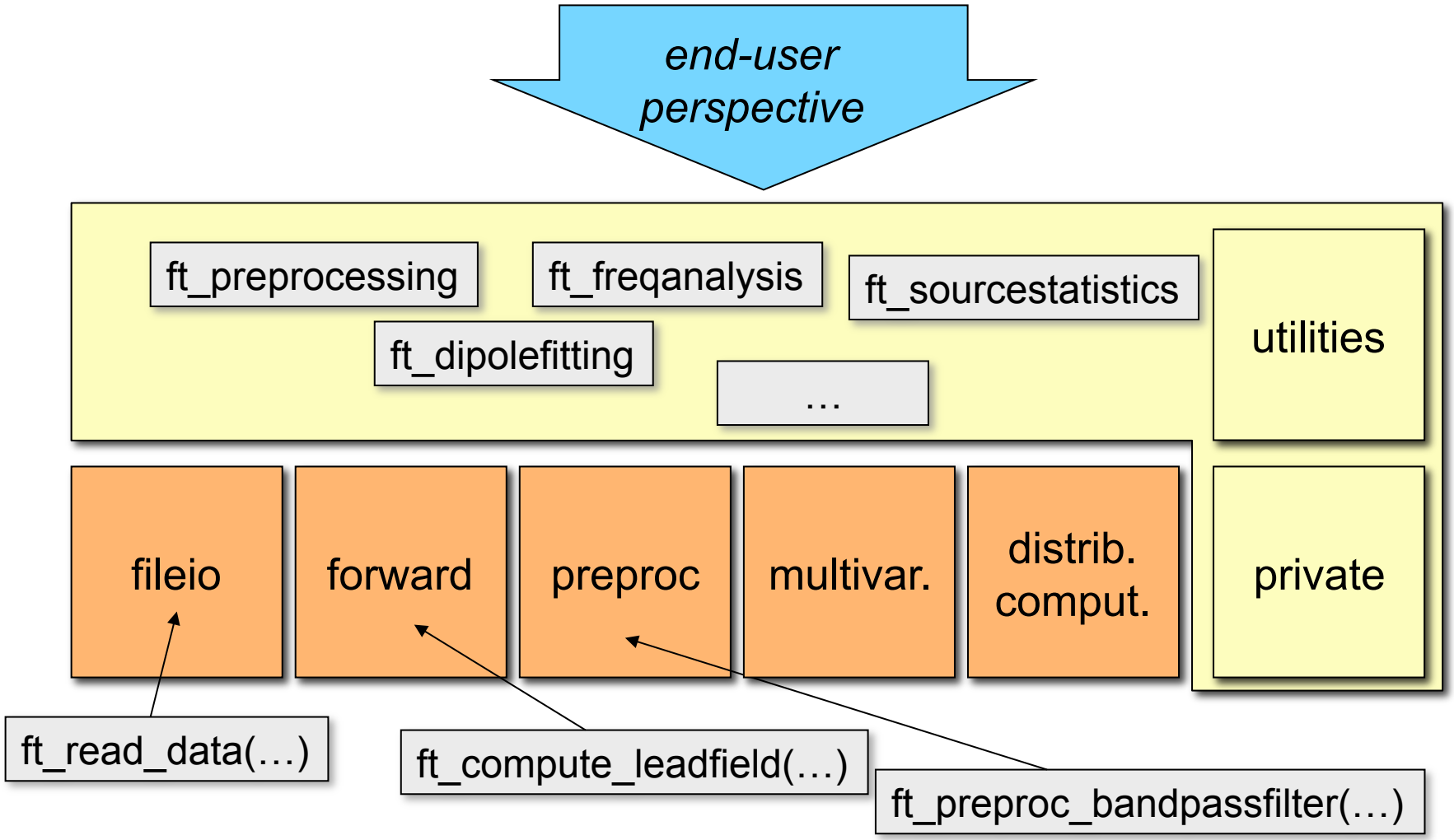
FieldTrip toolbox structure - at a glance



FieldTrip toolbox structure - a closer look



FieldTrip toolbox structure - a closer look



Summary

- Recording signals from the brain
- Temporal and spatial features of the signals
- Analyzing those signals with FieldTrip
- Background on the FieldTrip project

Hands-on

- Selecting pieces of data
- Preprocessing
- Averaging
- Plotting