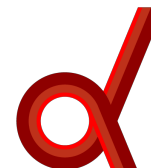


Radboud University



DONDERS
INSTITUTE



Introduction to EEG, MEG and analysis with the FieldTrip toolbox

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Donders Institute, Radboud University, Nijmegen, NL

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

What kind of signals are generated in the brain

How do we record those signals

Analyzing those signals with FieldTrip

Background on the FieldTrip toolbox

Talk outline

What kind of signals are generated in the brain

How do we record those signals

Analyzing those signals with FieldTrip

Background on the FieldTrip toolbox

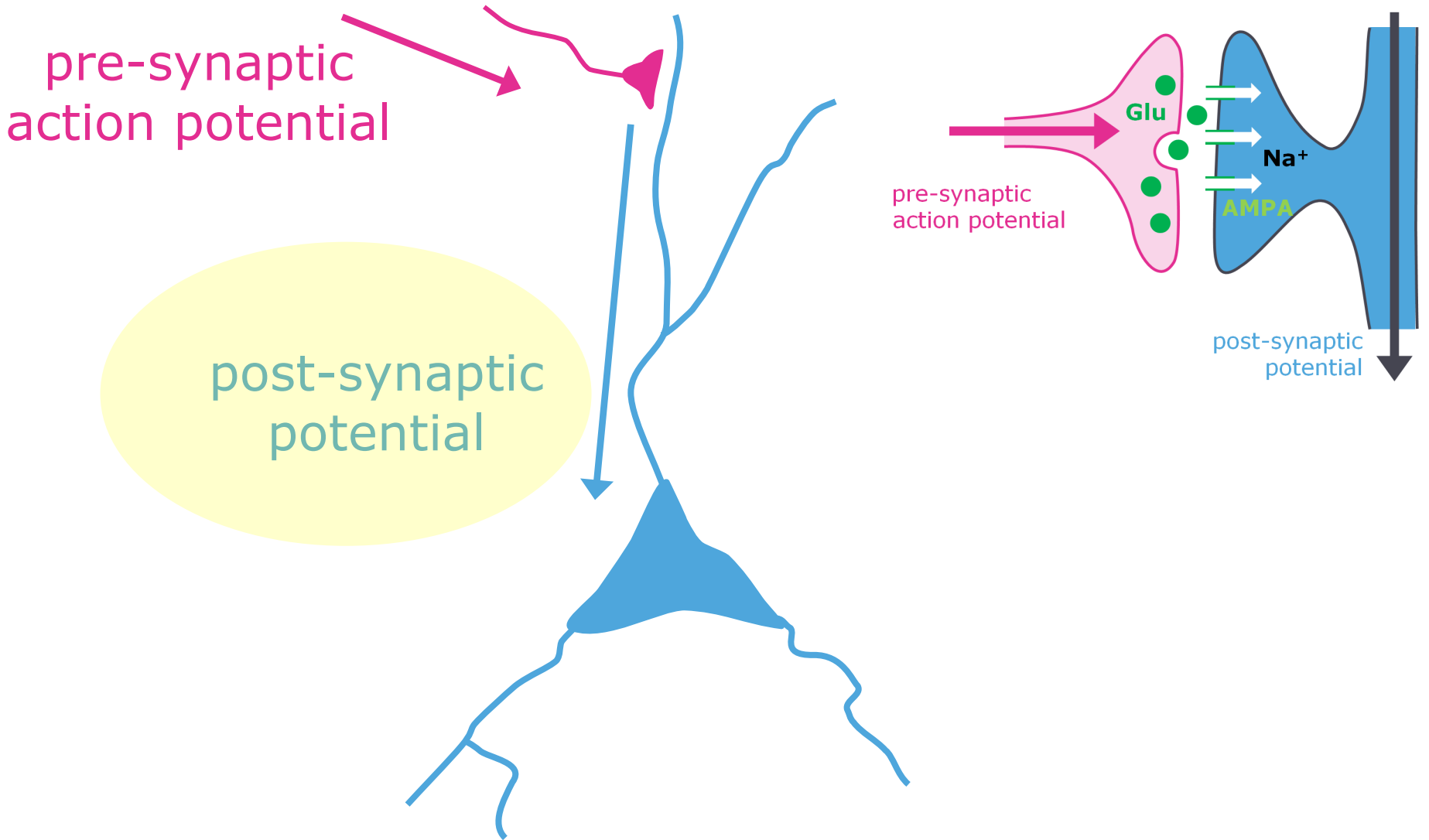
What kind of signals are generated in the brain

We measure the scalp potentials or field associated with post-synaptic potentials in pyramidal neurons

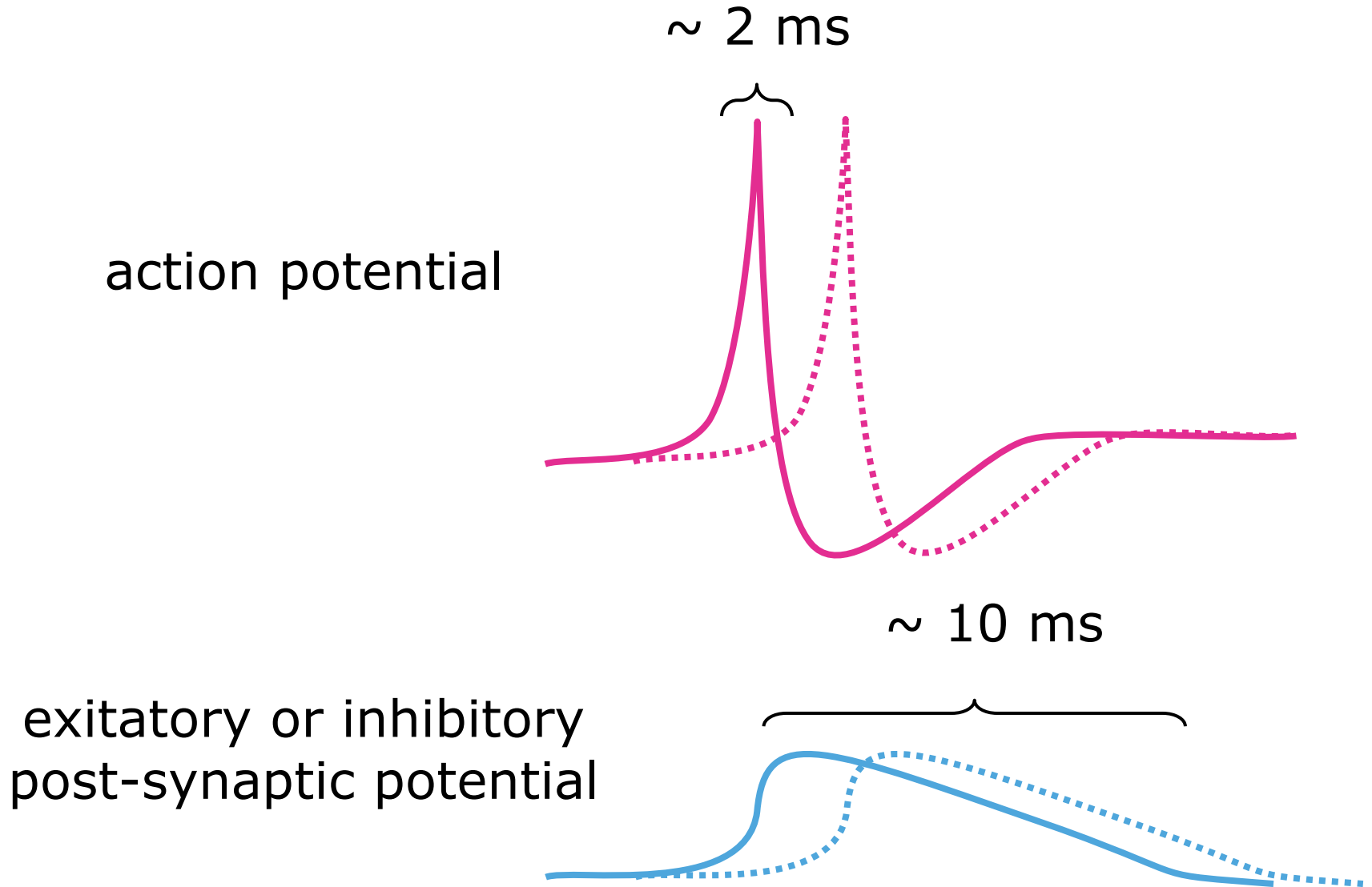
These PSPs represent the excitatory and inhibitory input that these neurons receive

Usually we study this neuronal input following the presentation of a stimulus or following a cognitive event

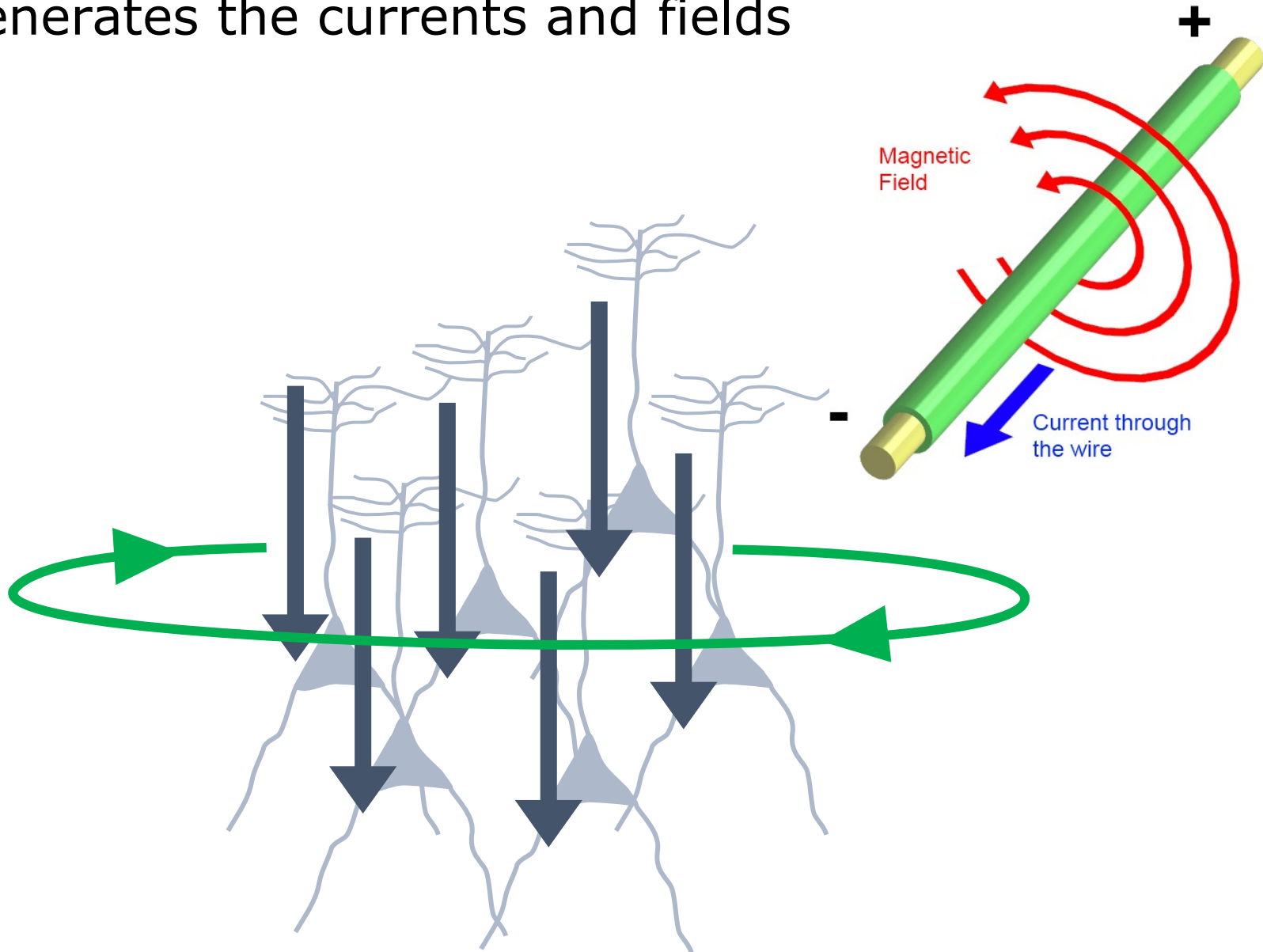
What produces the electric current and magnetic field?



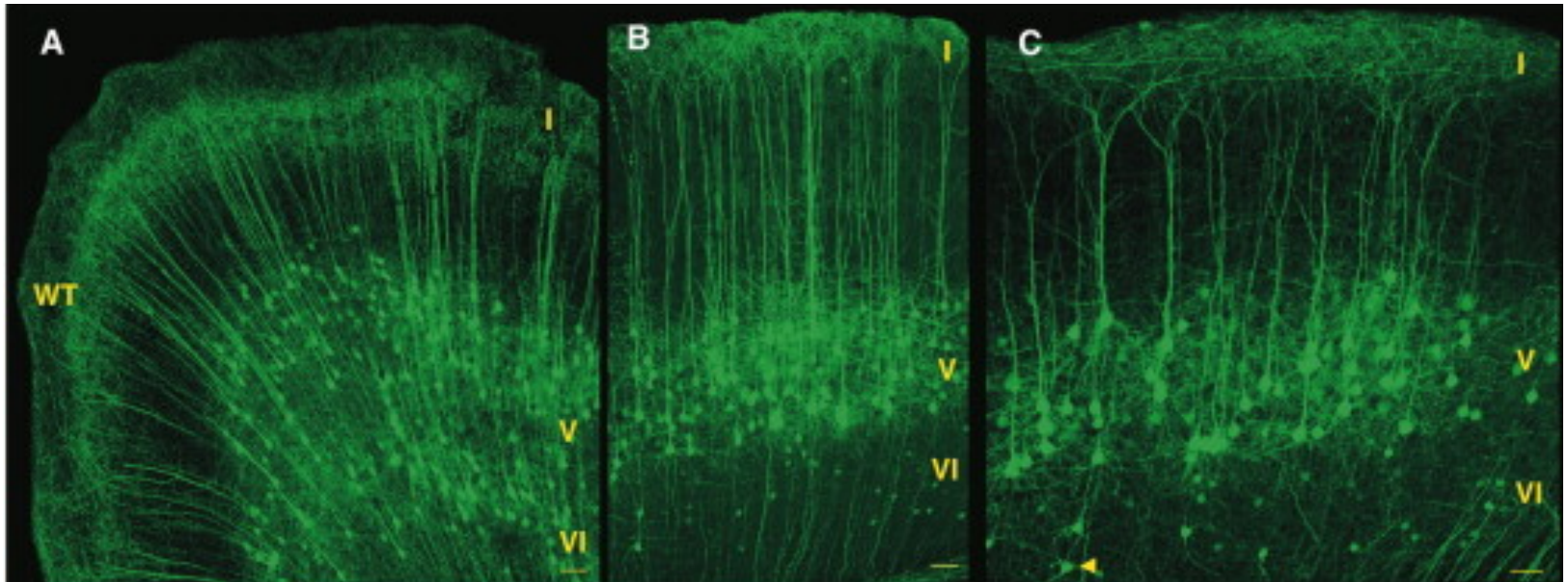
What produces the electric current and magnetic field



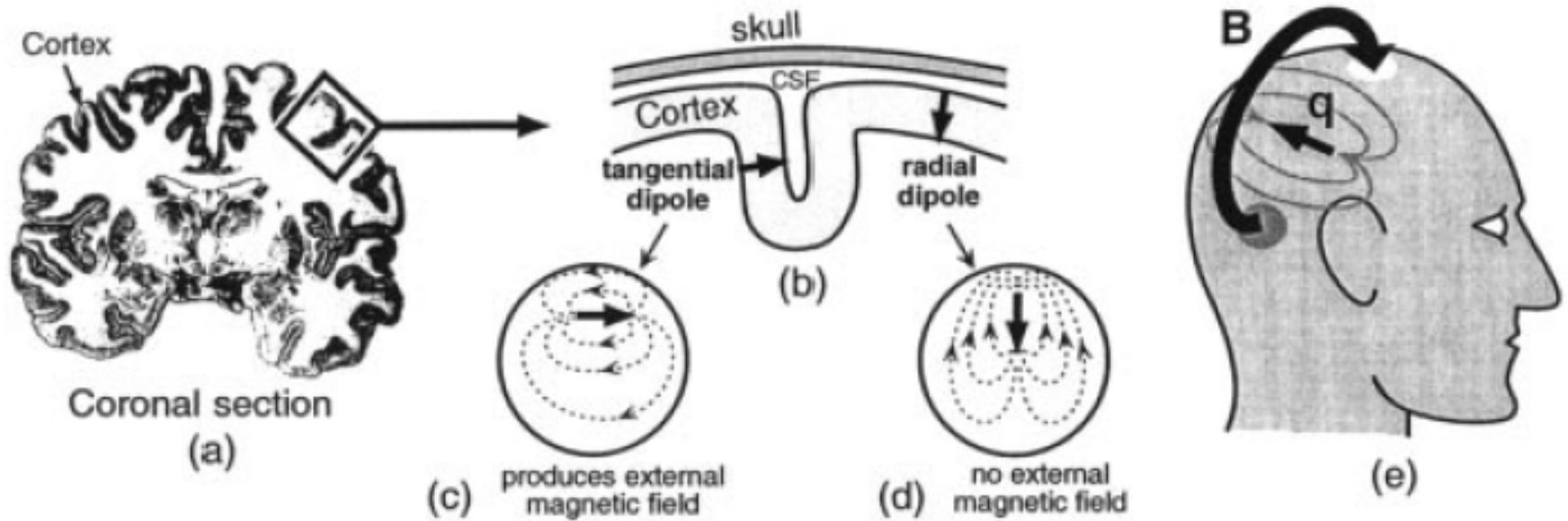
What generates the currents and fields



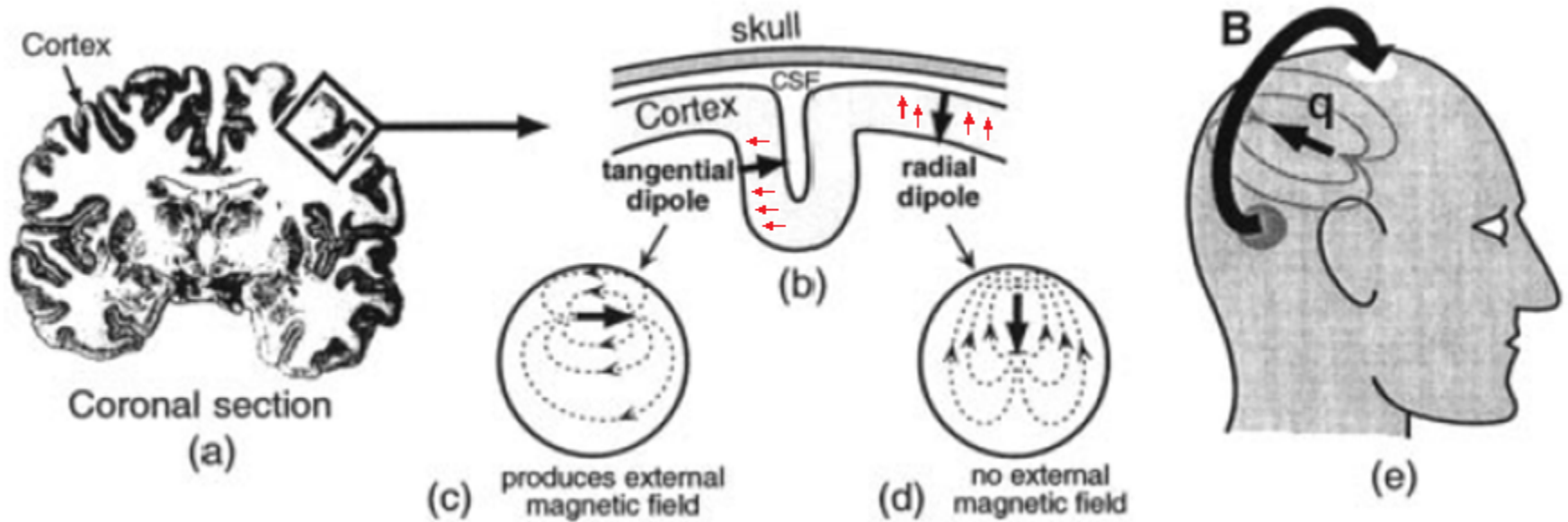
What generates the currents and fields



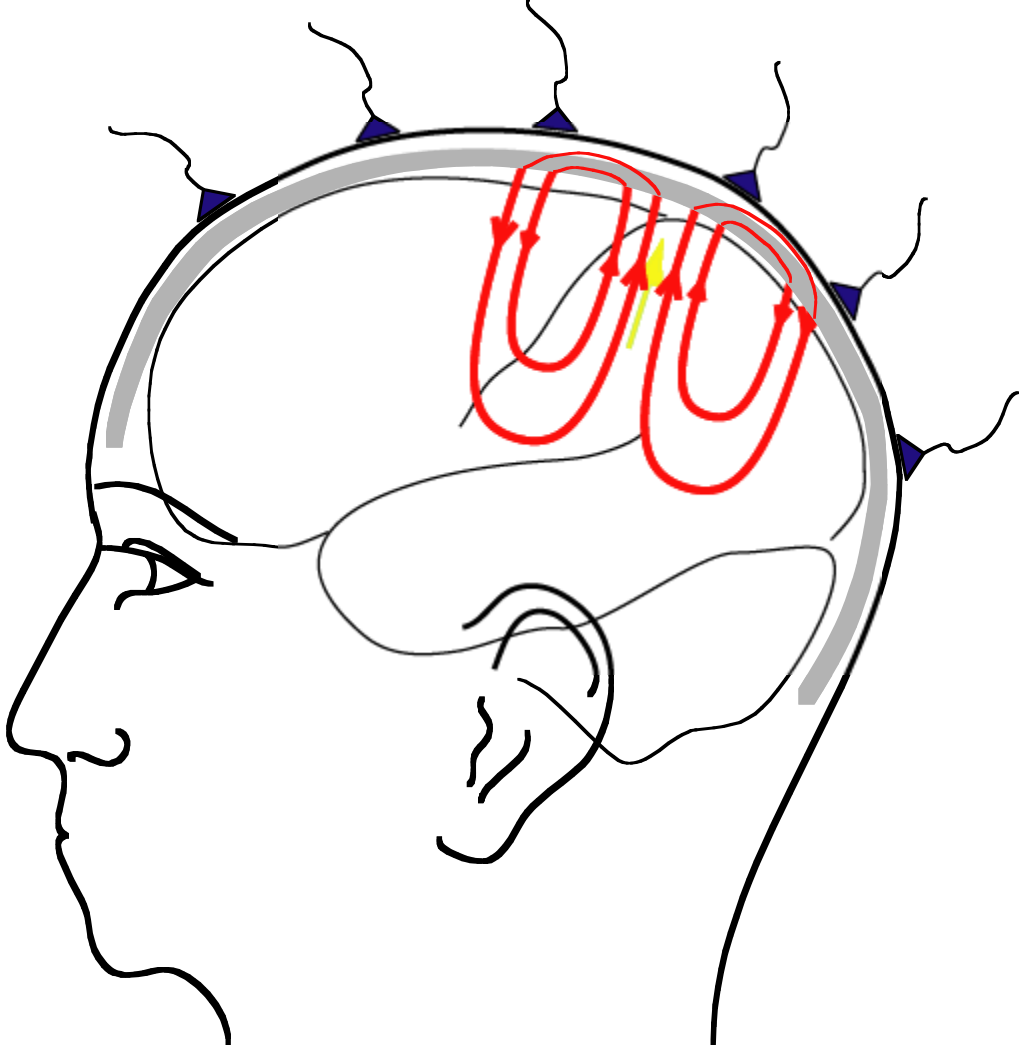
What generates the currents and fields



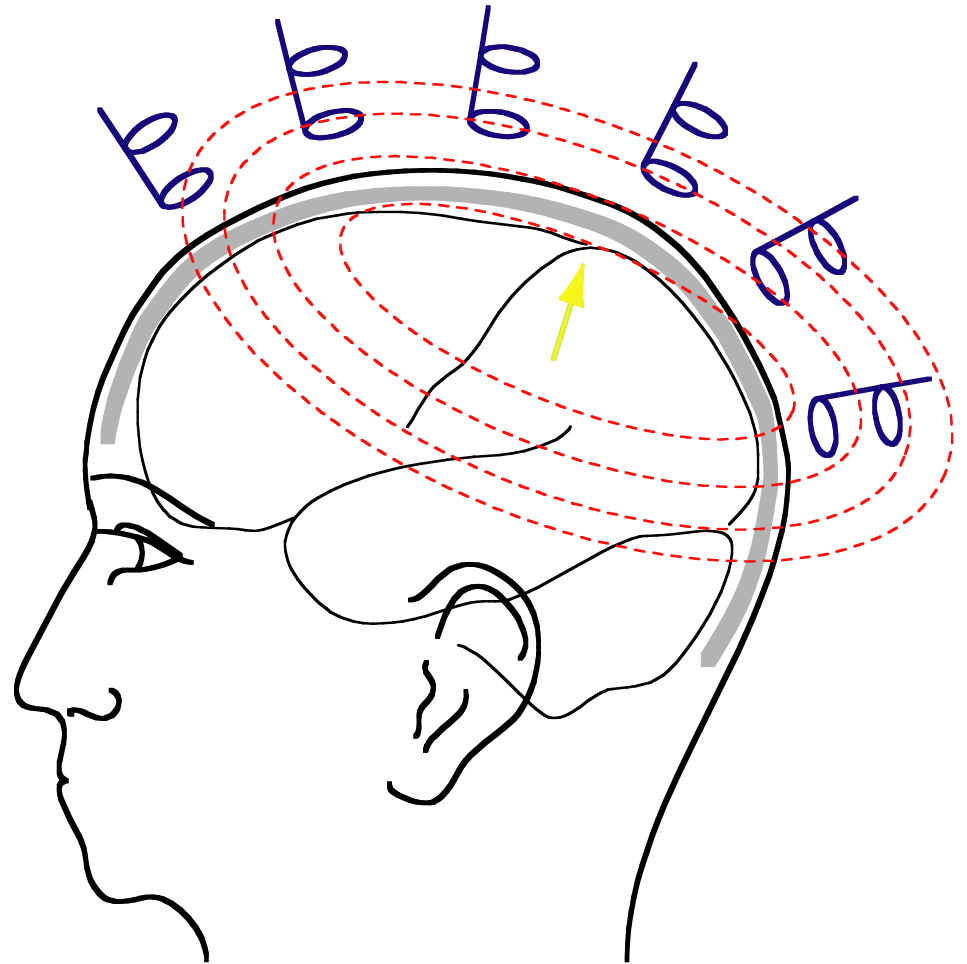
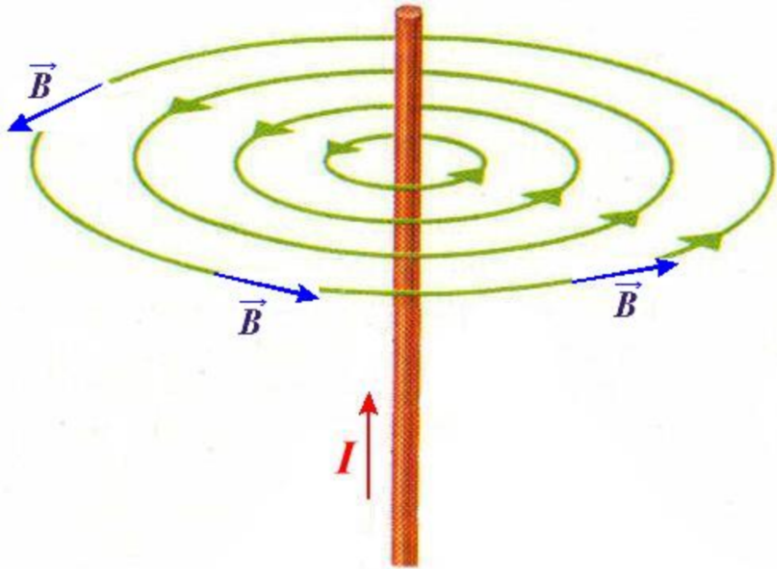
What generates the currents and fields



EEG volume conduction



Electric current \rightarrow magnetic field



Talk outline

What kind of signals are generated in the brain

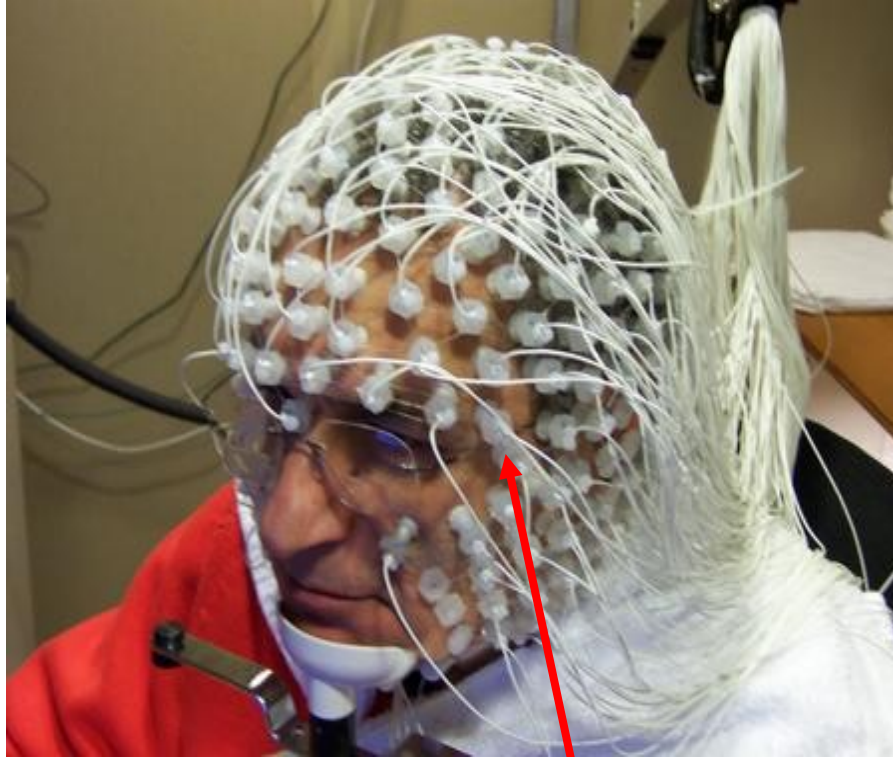
How do we record those signals

Analyzing those signals with FieldTrip

Background on the FieldTrip toolbox

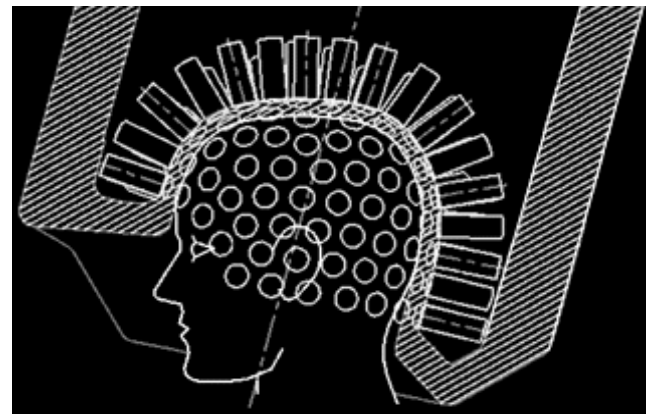
How can we measure brain signals (non-invasively)?

Electroencephalography (EEG)



10,000-1 Million
neurons

Magnetoencephalography (MEG)



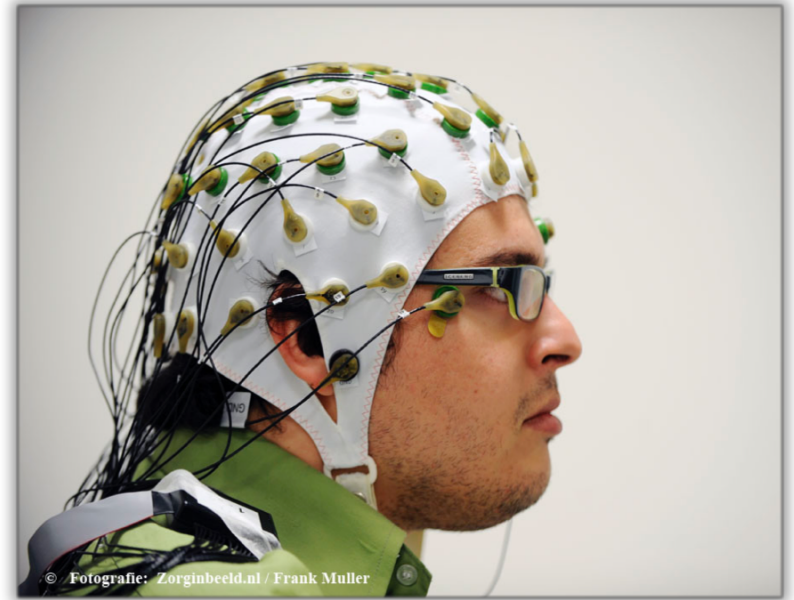
EEG apparatus

Sensors

- Conductivity and smearing
- Electrode positioning
- Referencing

Noise (environmental and internal)

- Shielding
- Movement (eyes, heart beat, muscle)



MEG acquisition system

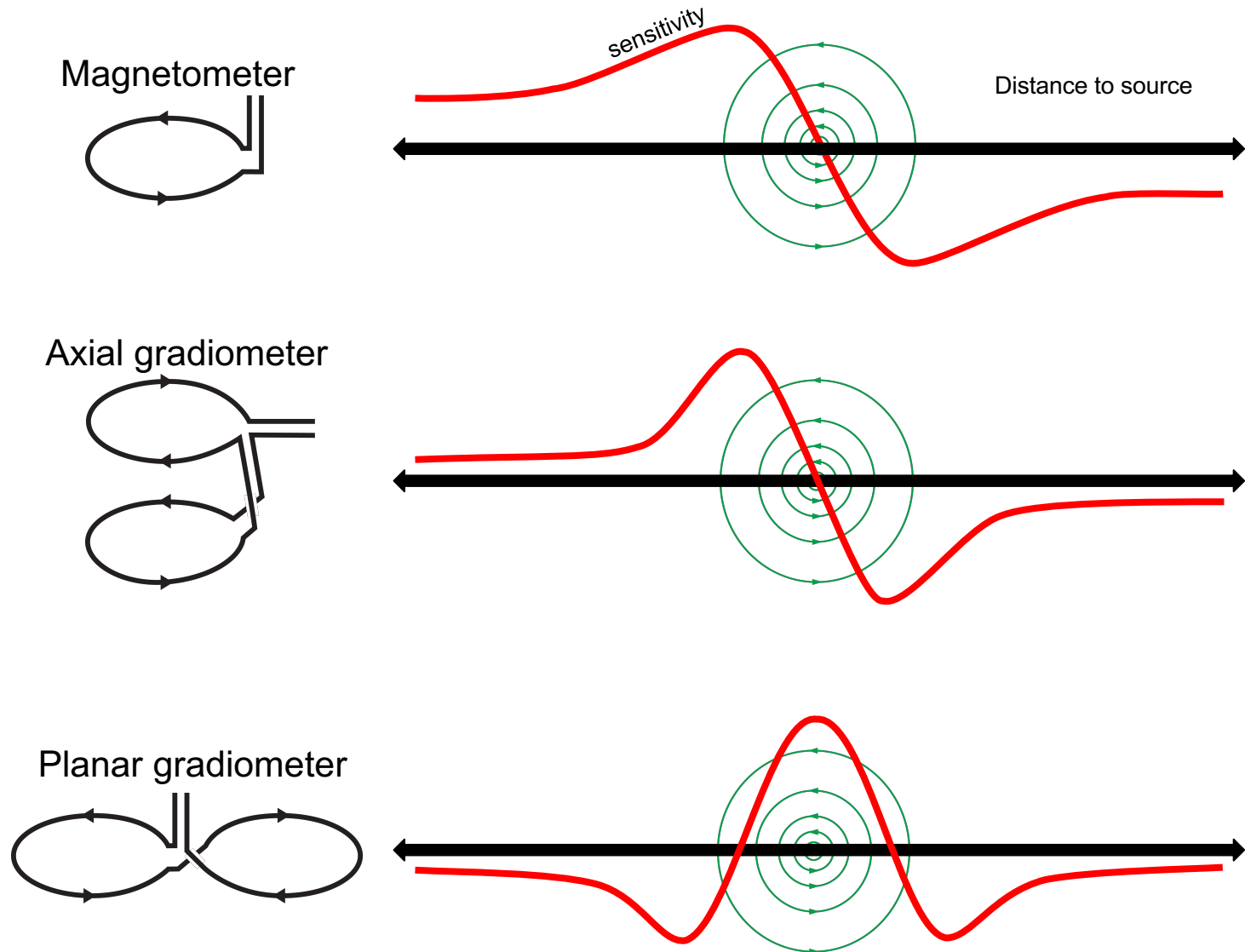
Magnetic detectors: magnetometers,
or planar/axial gradiometers

Whole head coverage: 100-300
sensors

Magnetically shielded room

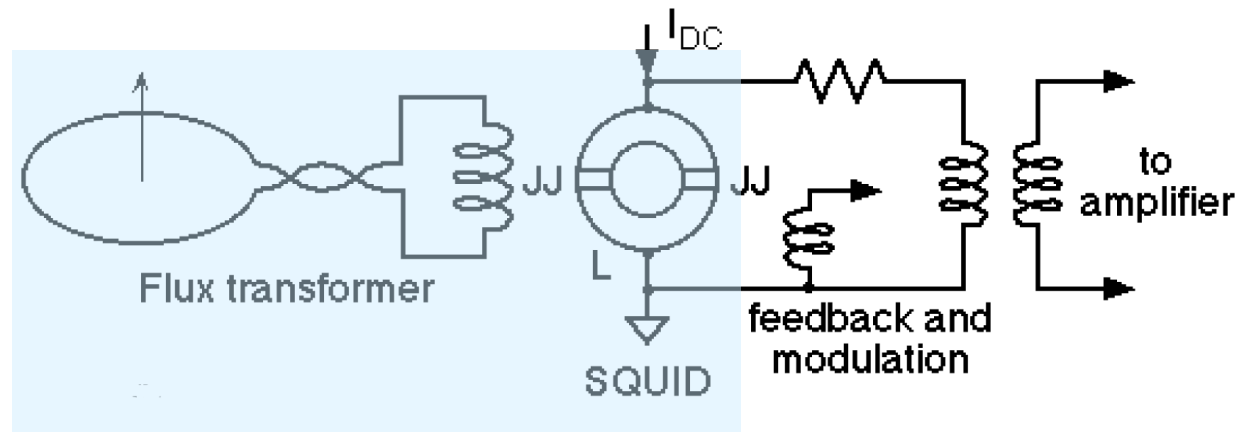
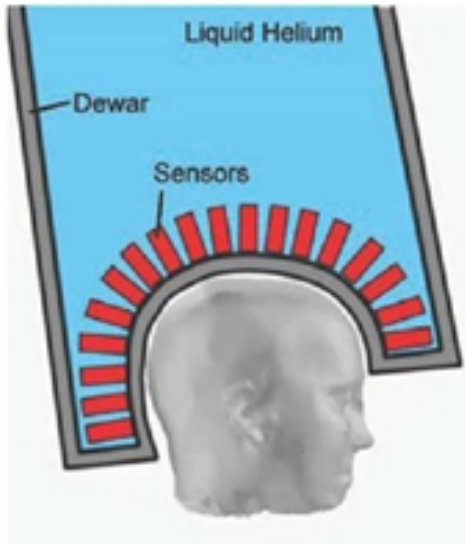
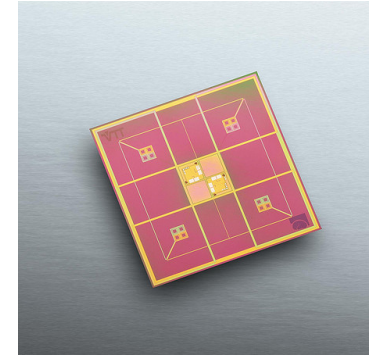


MEG sensor – sensitivity profile

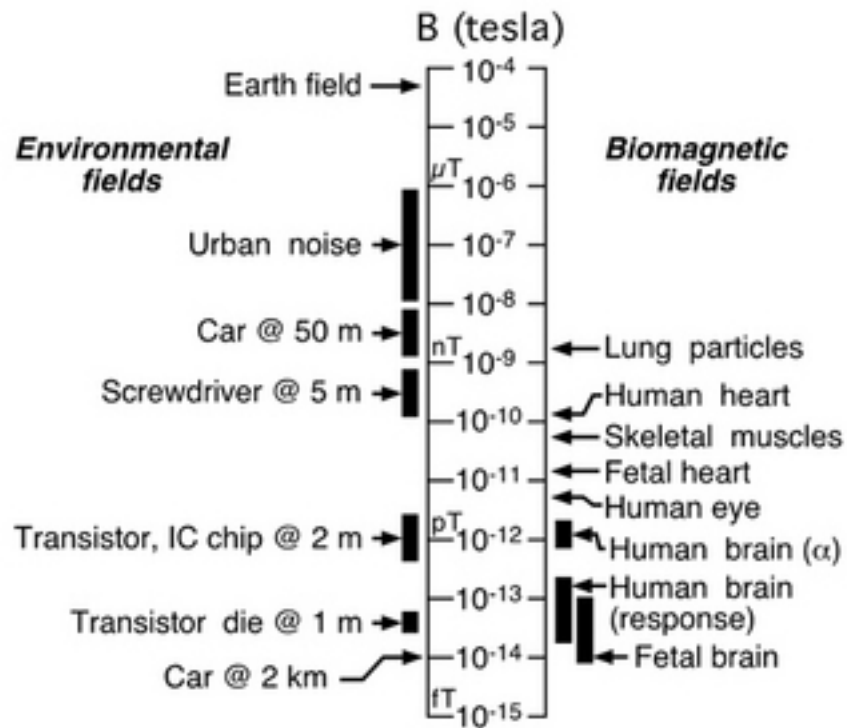


Magnetic field detectors

Superconducting **Q**uantum **I**nterference **D**evice



Technical challenges of MEG - Noise



Technical challenges of MEG

Requires sensitive magnetic detectors

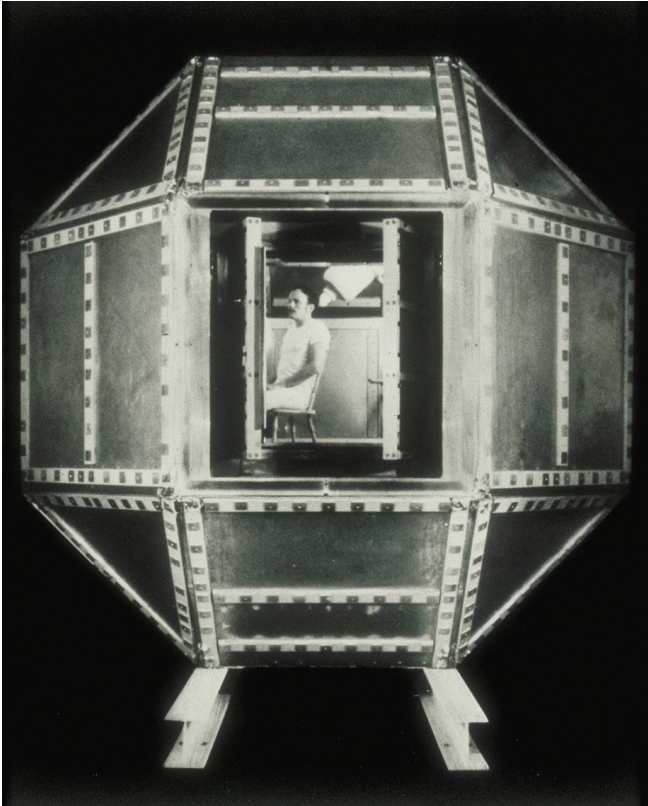
Deal with environmental noise

- shielding

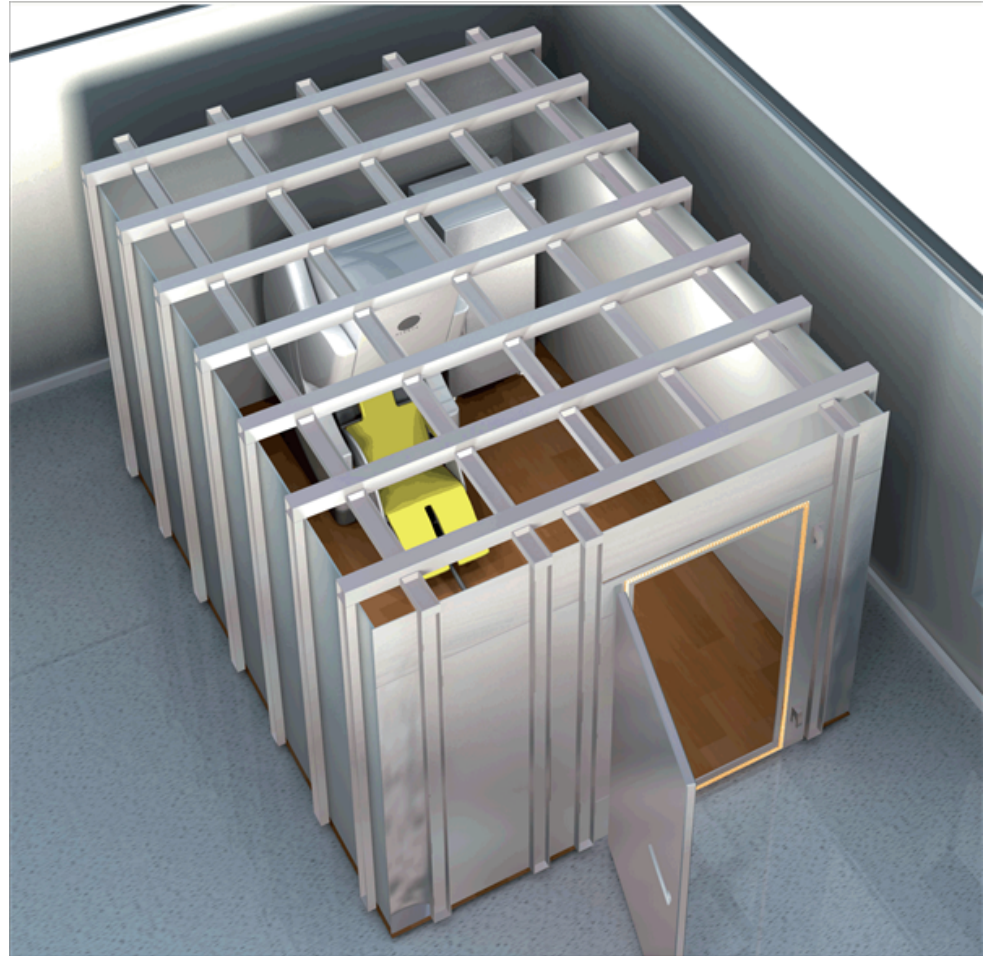
- sensor design

- reference sensors for noise subtraction

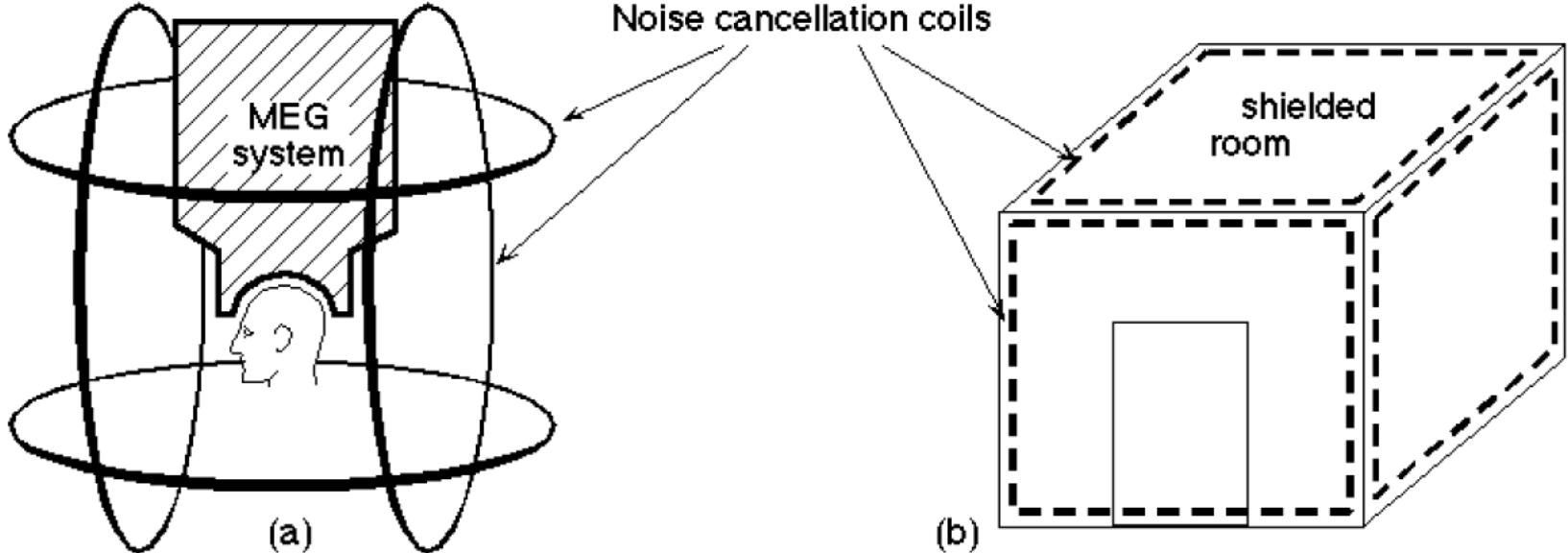
Shielding - passive



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



Shielding - active



EEG

vs.

MEG

- Direct measure of neuronal activity
- High temporal resolution

- Direct measure of neuronal activity
- High temporal resolution

EEG

vs.

MEG

- Direct measure of neuronal activity
- High temporal resolution

- Direct measure of neuronal activity
- High temporal resolution
- Only sensitive to neurons that are not perpendicular to the head

EEG

vs.

MEG

- Direct measure of neuronal activity
- High temporal resolution
- More sensitive to deep sources

- Direct measure of neuronal activity
- High temporal resolution
- Only sensitive to neurons that are not perpendicular to the head

EEG

vs.

MEG

- Direct measure of neuronal activity
- High temporal resolution
- More sensitive to deep sources
- Spatial smearing because of distortion by scalp and skull

- Direct measure of neuronal activity
- High temporal resolution
- Only sensitive to neurons that are not perpendicular to the head
- Sources can be localized better

EEG

vs.

MEG

- Direct measure of neuronal activity
- High temporal resolution



- Transportable

- A lot of preparation time needed

- Direct measure of neuronal activity
- High temporal resolution

- Only sensitive to neurons that are not perpendicular to the head
- Sources can be localized better

EEG

vs.

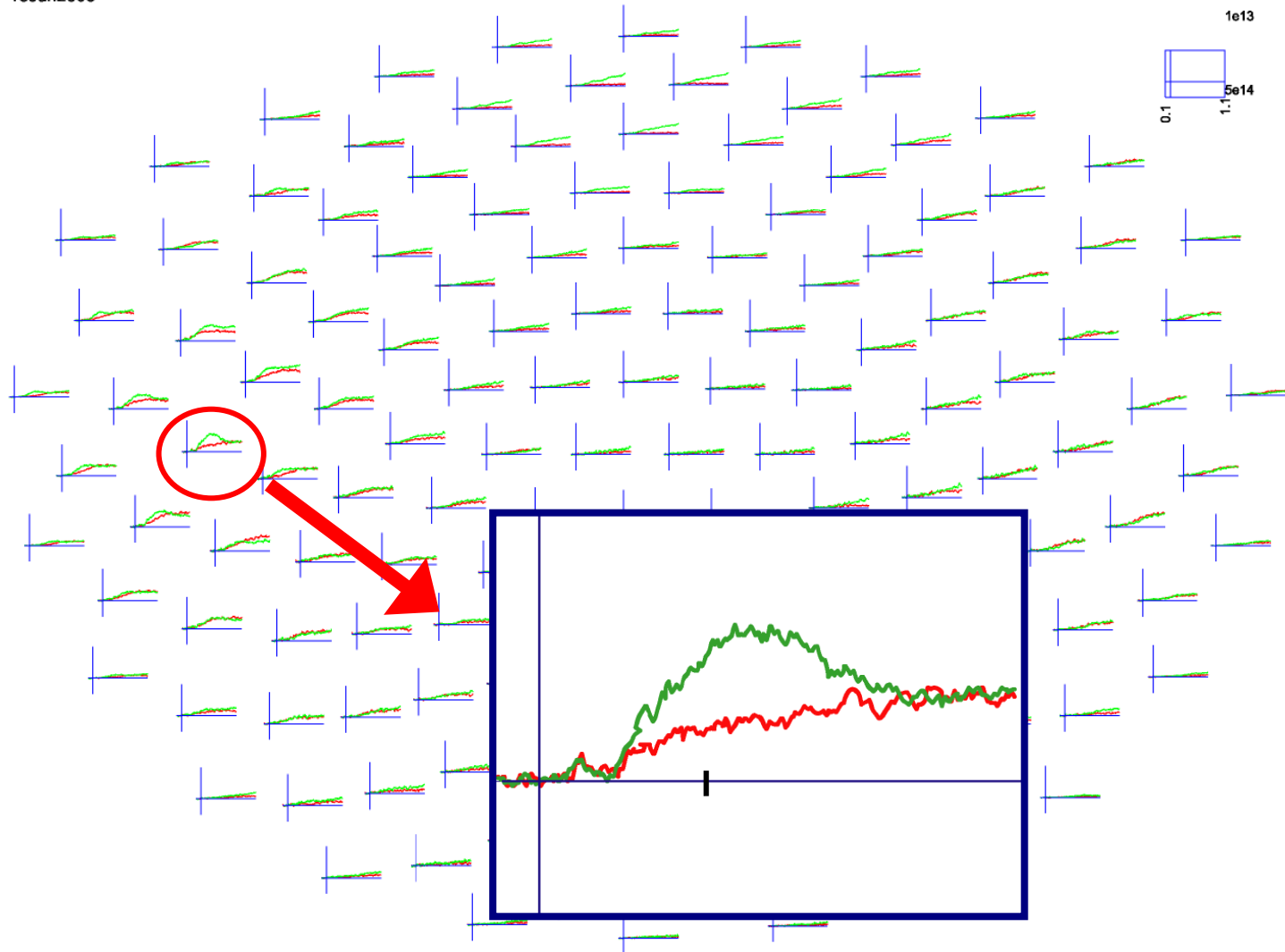
MEG

- Direct measure of neuronal activity
- High temporal resolution
- More sensitive to deep sources
- Spatial smearing because of distortion by scalp and skull
 - Transportable
- A lot of preparation time needed

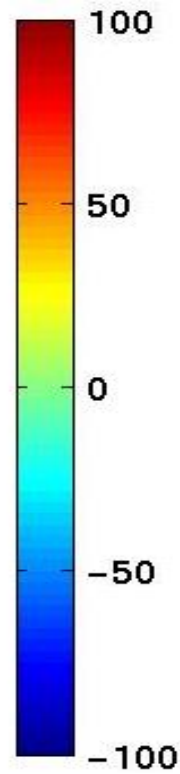
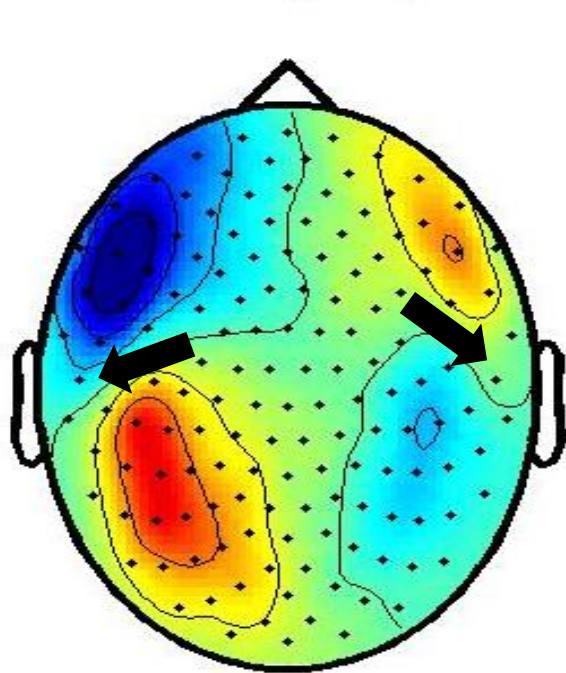
- Direct measure of neuronal activity
- High temporal resolution
- Only sensitive to neurons that are not perpendicular to the head
- Sources can be localized better
 - More expensive

N400 response in MEG

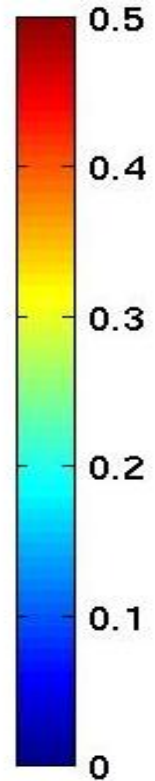
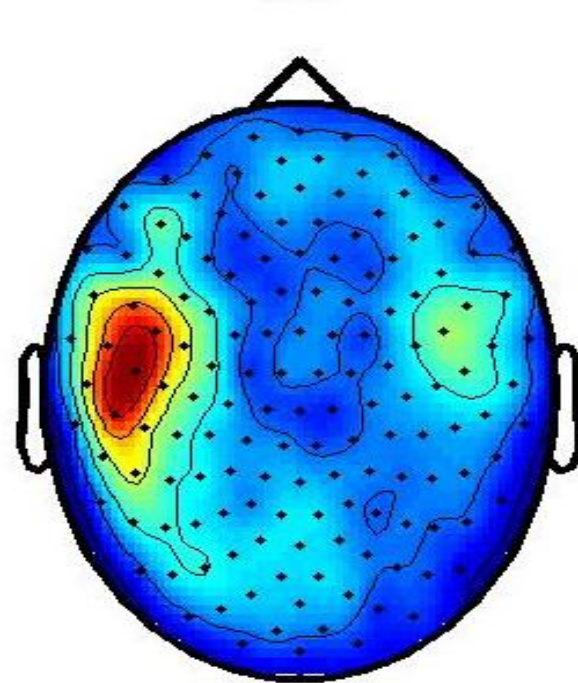
18Jun2003



N400 response - compared between MEG systems



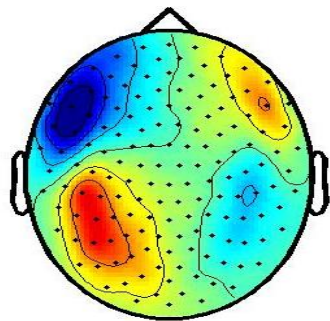
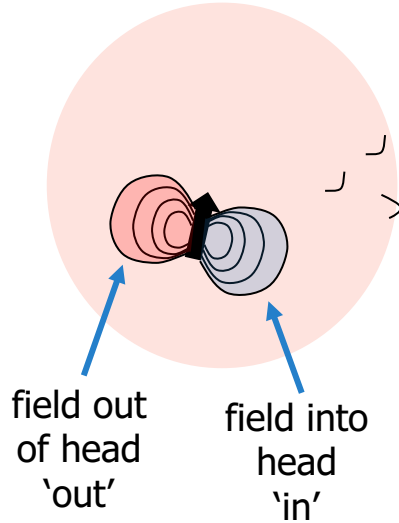
axial gradient



planar gradient

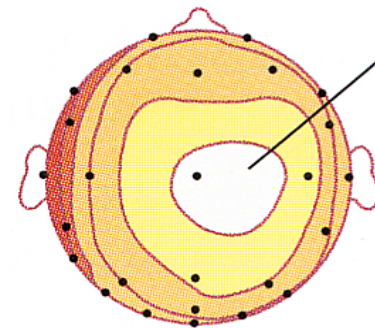
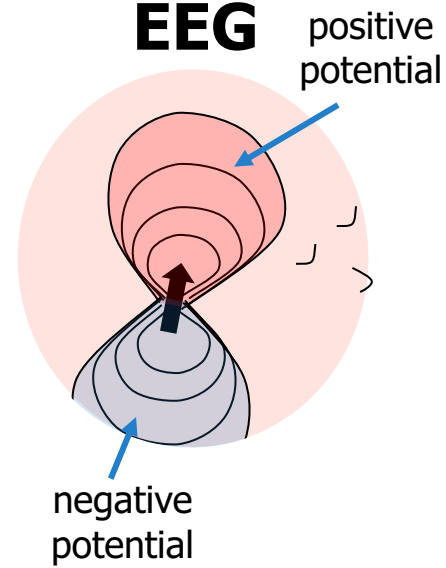
N400 response - EEG Compared to MEG

MEG



M400 field map

EEG



N400 field map

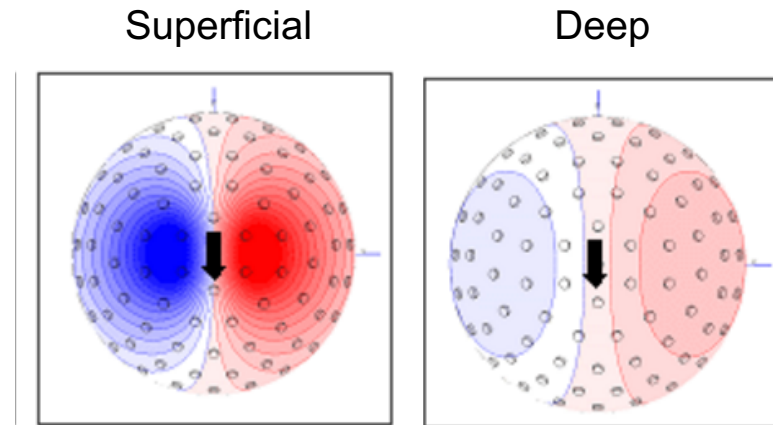
What affects the MEG/EEG signal?

Neural Current Depth

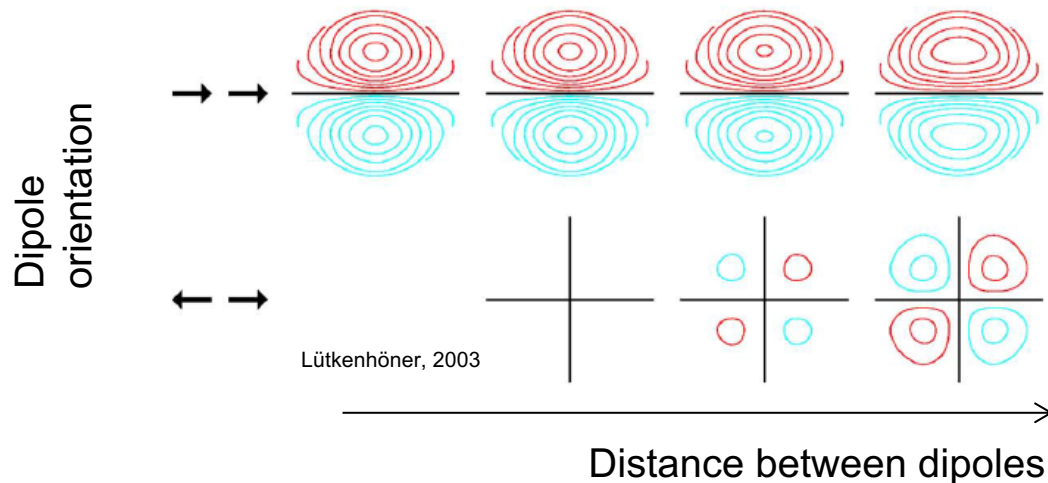
Deep source are difficult to measure

Sources close together can cancel out

- Perceived as one source
- Reduce signal amplitude



<http://imaging.mrc-cbu.cam.ac.uk/meg/IntroEEGMEG#eegrecordings>



Talk outline

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Background on the FieldTrip toolbox

M/EEG signal characteristics considered during analysis

timecourse of activity

-> ERP

spectral characteristics

-> power spectrum

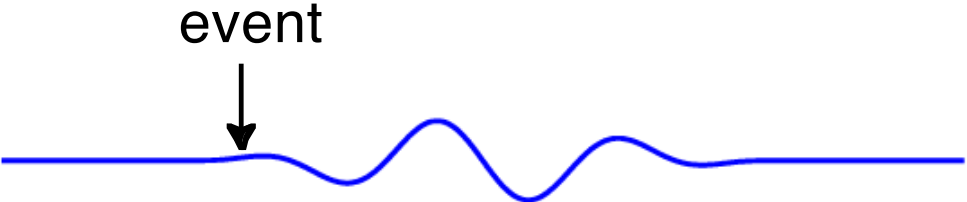
temporal changes in power

-> time-frequency response (TFR)

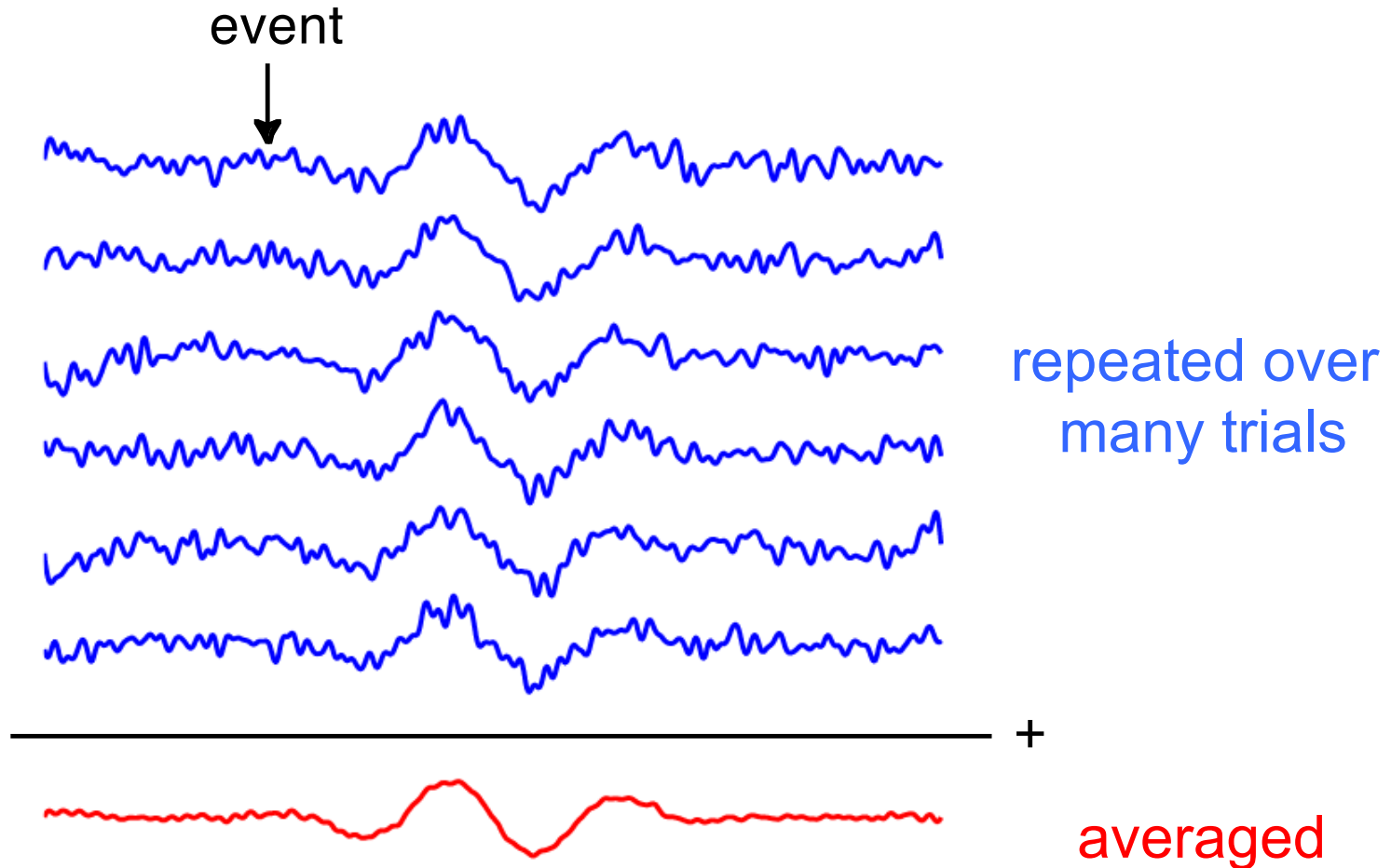
spatial distribution of activity over the head

-> source reconstruction

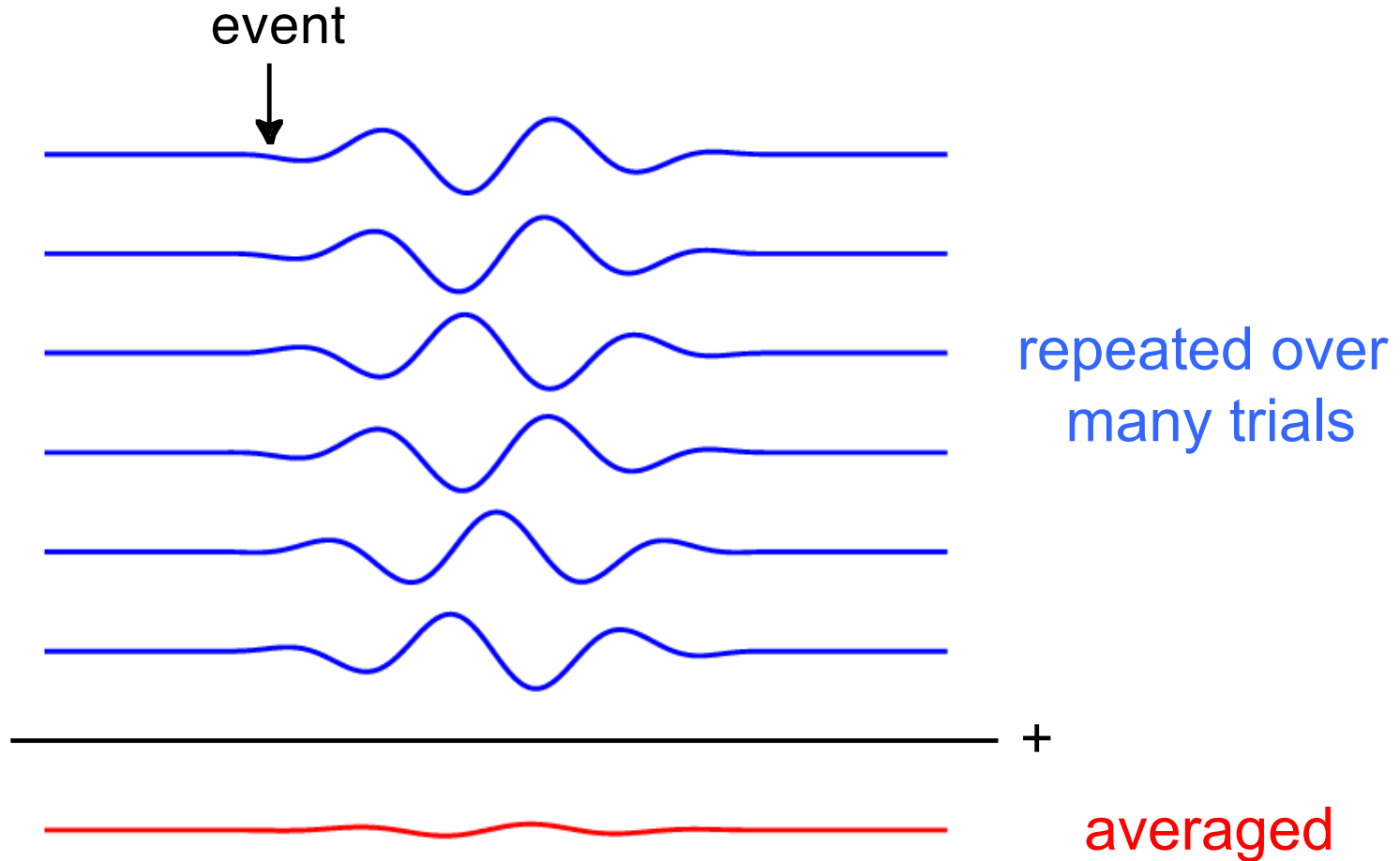
Evoked activity



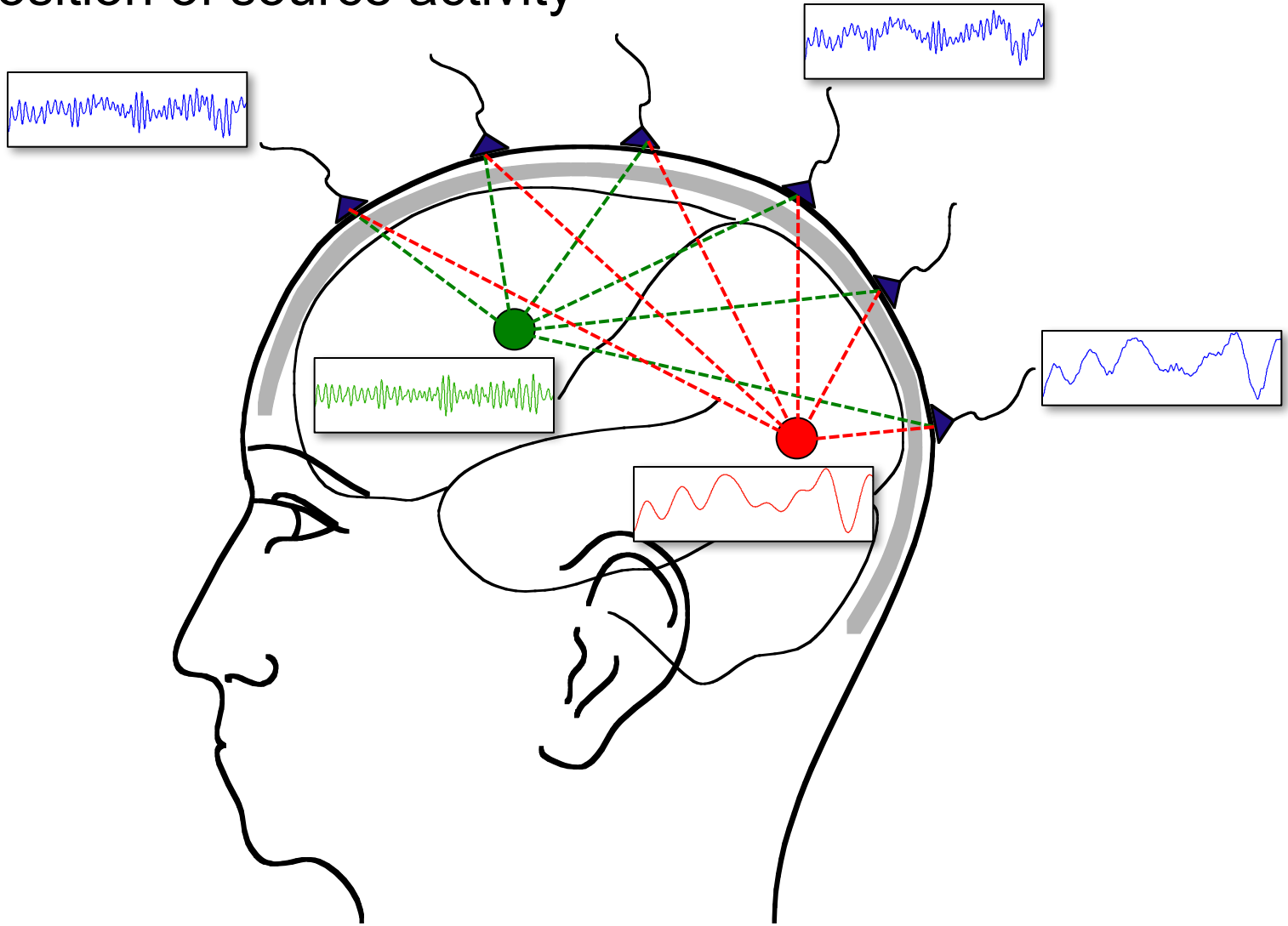
Evoked activity



Induced activity



Superposition of source activity



What is FieldTrip?

A MATLAB toolbox for electrophysiological analysis



Some FieldTrip basics

MATLAB default

```
dataout = functionname(datain, 'key1', 'value1', ...)
```

FieldTrip defaults

```
dataout = functionname(cfg, datain, ...)
```

```
functionname(cfg, datain, ...)
```

```
dataout = functionname(cfg)
```

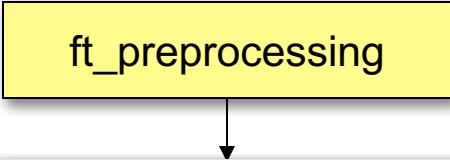
the "cfg" argument is a configuration structure, e.g.

```
cfg.channel = {'C3', 'C4', 'F3', 'F4'}
```

```
cfg.foilim = [1 70]
```

Using functions in an analysis protocol

ft_preprocessing



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)
```

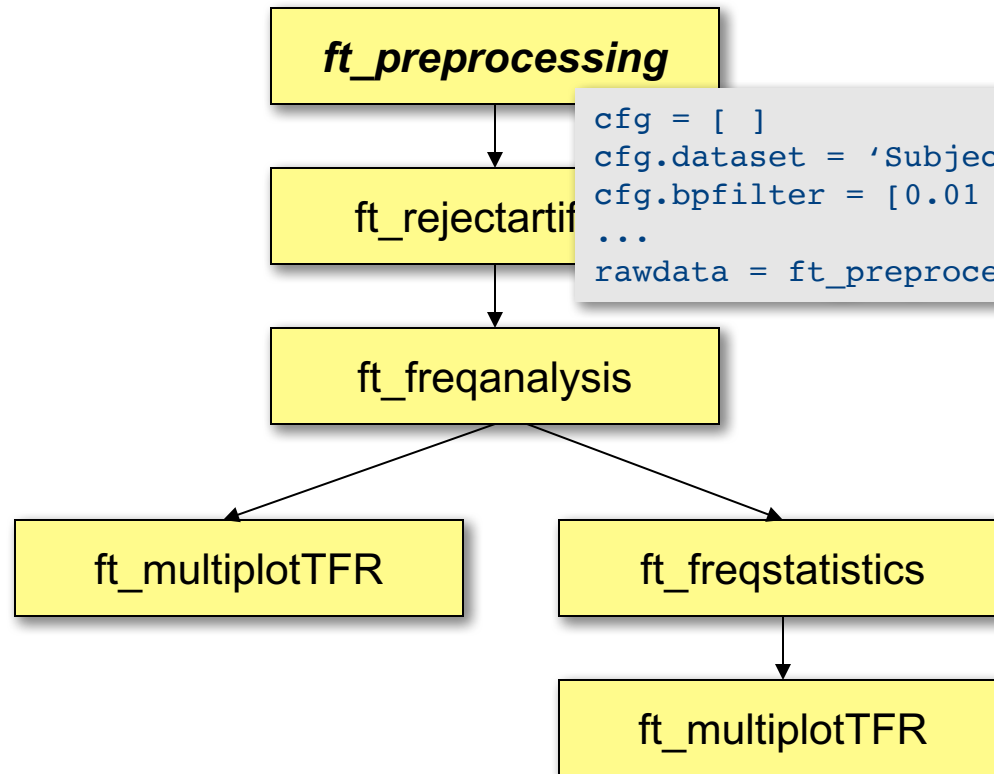
or

```
[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 occurrence of a trigger in the data.

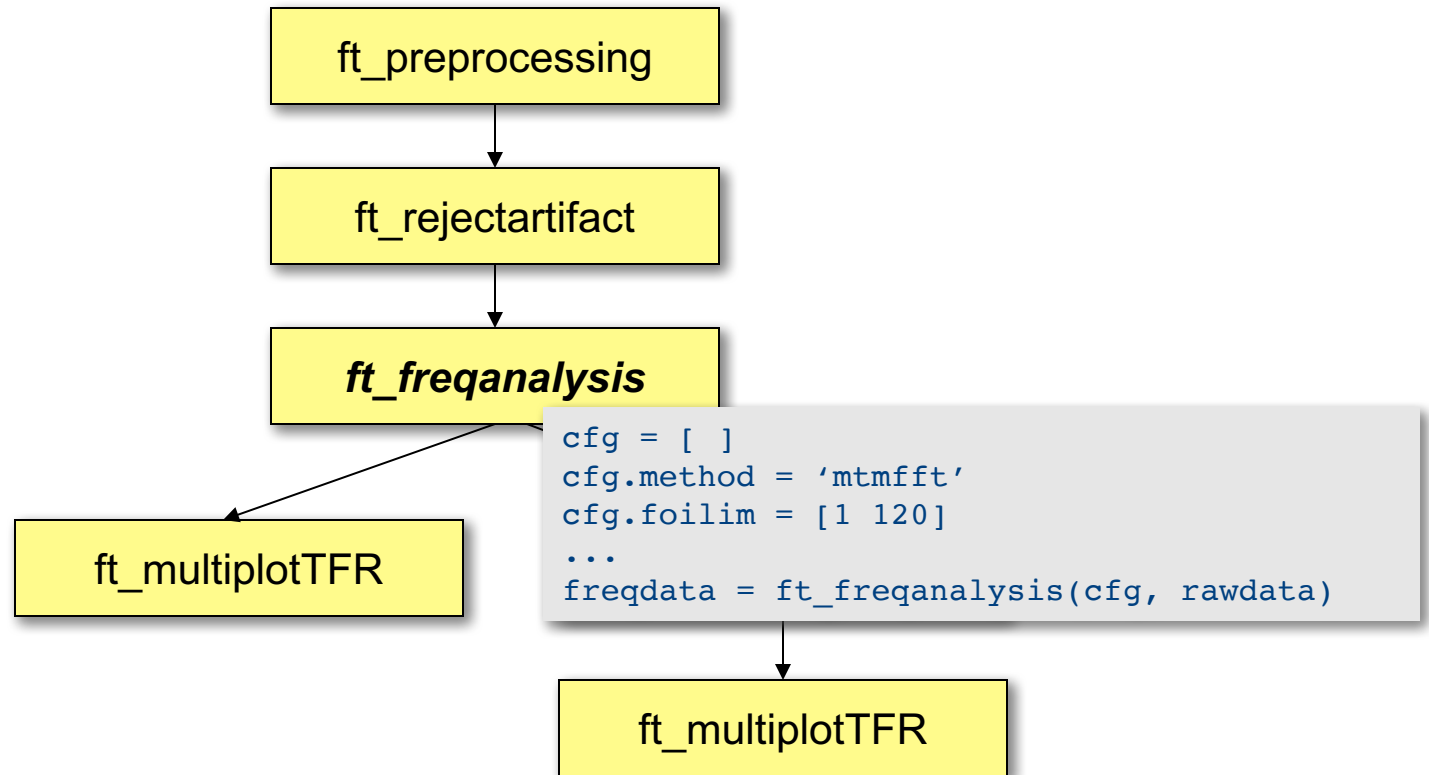
...

Using functions in an analysis protocol



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

Using functions in an analysis protocol



Raw data structure

```
rawData =  
    label: {151x1 cell}  
    trial: {1x80 cell}  
    time: {1x80 cell}  
fsample: 300  
    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},...)
```

dataout

dataout.cfg

dataout.cfg.previous{1}

dataout.cfg.previous{2}

...

= data structure with...

= settings + defaults

= datain{1}.cfg

= datain{2}.cfg

details of computations are kept with data

previous data is not kept, but can be reconstructed using

cfg.previous.previous...

Keeping track of your analysis

FieldTrip analysis pipeline, Tue 12-Apr-2016 16:30:38

file:///Users/roboos/Desktop/ERF_Stat_Letter_FacevsLetter.html

FieldTrip analysis pipeline, Tue 12-Apr-2016 16:30:38

```
graph TD; P1[ft_preprocessing] --> A[ft_appenddata]; P2[ft_preprocessing] --> A; P3[ft_preprocessing] --> A; P4[ft_preprocessing] --> A; A --> R1[ft_rejectvisual]; R1 --> R2[ft_rejectvisual]; R2 --> R3[ft_redefintrial]; R3 --> R4[ft_rejectartifact]; R4 --> R5[ft_componentanalysis]; R5 --> R6[ft_rejectcomponent]; R6 --> R7[ft_preprocessing]; R7 --> T1[ft_timelockanalysis]; R7 --> T2[ft_timelockanalysis]; T1 --> S[ft_timelockstatistics]; T2 --> S;
```

ft_preprocessing

User-specified configuration

```
cfg.dataset = '/home/fanny/Desktop/MEG_Emo_all/practi...';
cfg.trialfun = 'ft_trialfun_general';
cfg.trialdef.eventtype = 'STI101';
cfg.trialdef.eventvalue = [30 31 10 130 131 110];
cfg.trialdef.prestim = 0.7;
cfg.trialdef.poststim = 1.2;
cfg.callinfo.usercfg.dataset = '/home/fanny/Desktop/MEG_Emo_all/practi...';
cfg.callinfo.usercfg.trialf... = 'ft_trialfun_general';
cfg.callinfo.usercfg.triald... = 'STI101';
cfg.callinfo.usercfg.triald... = [30 31 10 130 131 110];
cfg.callinfo.usercfg.triald... = 0.7;
cfg.callinfo.usercfg.triald... = 1.2;
cfg.callinfo.usercfg.trackc... = 'off';
cfg.callinfo.usercfg.checkc... = 'loose';
cfg.callinfo.usercfg.checks... = 100000;
cfg.callinfo.usercfg.showca... = 'yes';
cfg.callinfo.usercfg.debug = 'no';
cfg.callinfo.usercfg.output... = 'overwrite';
cfg.callinfo.usercfg.trackc... = 'yes';
cfg.callinfo.usercfg.trackd... = 'no';
```

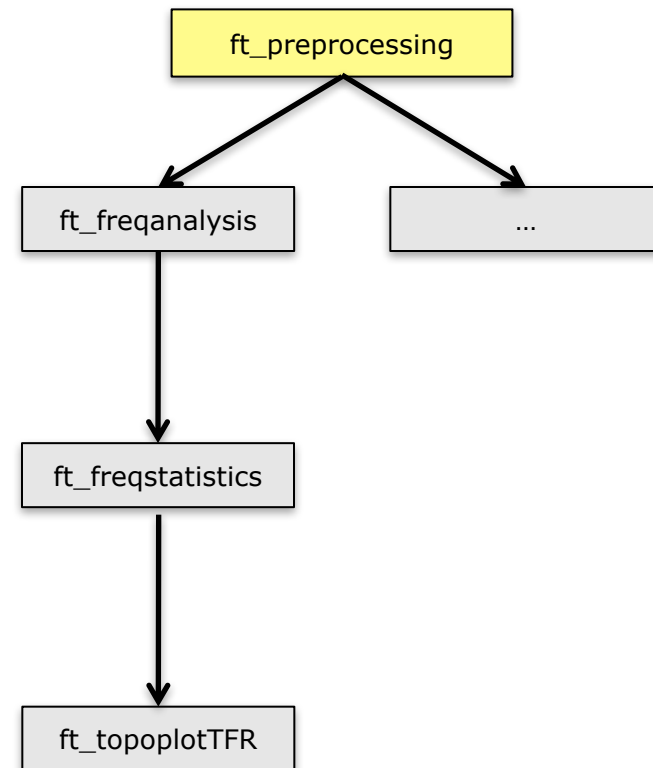
Pipeline HTML generated by roboos on Tue 12-Apr-2016 16:30:38.
Estimated total pipeline processing time: 17.1 minutes.

Example use in scripts

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

```
cfg = []  
cfg.method = 'mtmfft'  
cfg.foylim = [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)
```

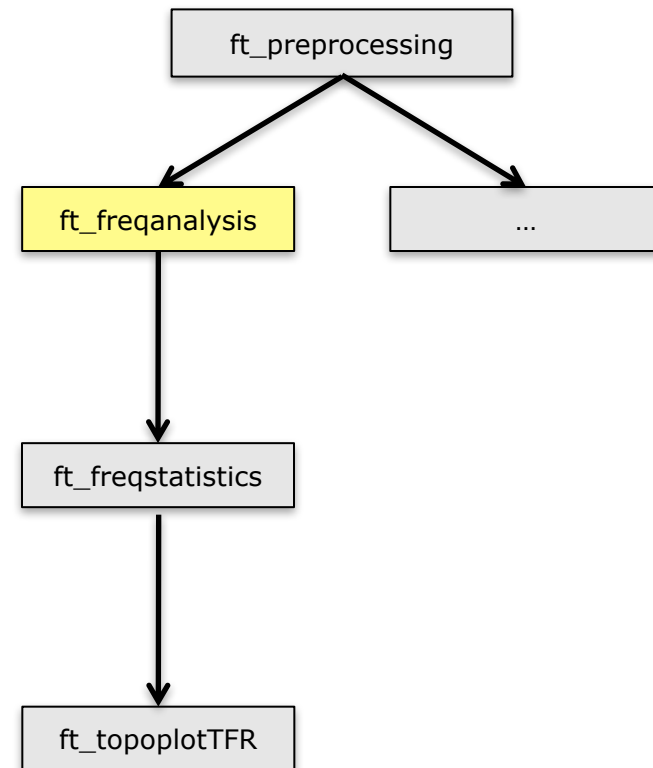


Example use in scripts

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

```
cfg = []
cfg.method = 'mtmfft'
cfg.foylim = [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)
```

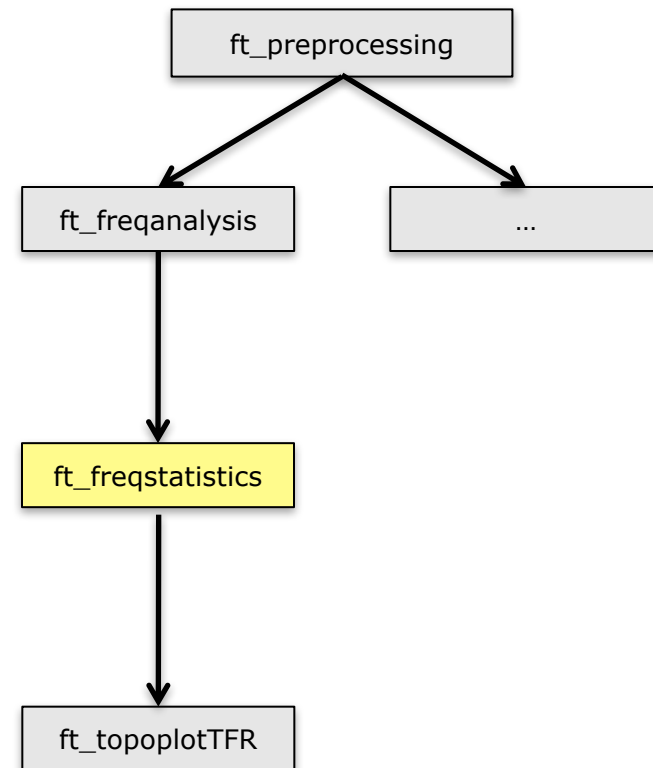


Example use in scripts

```
cfg = []  
cfg.dataset = 'Subject01.ds'  
cfg.bpfiler = [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)
```

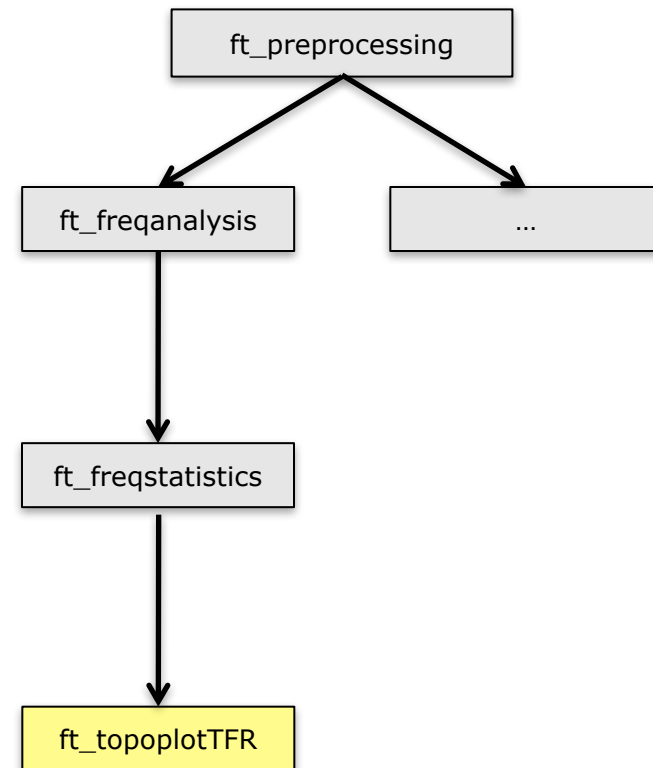


Example use in scripts

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

```
cfg = []  
cfg.method = 'mtmfft'  
cfg.foylim = [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)
```

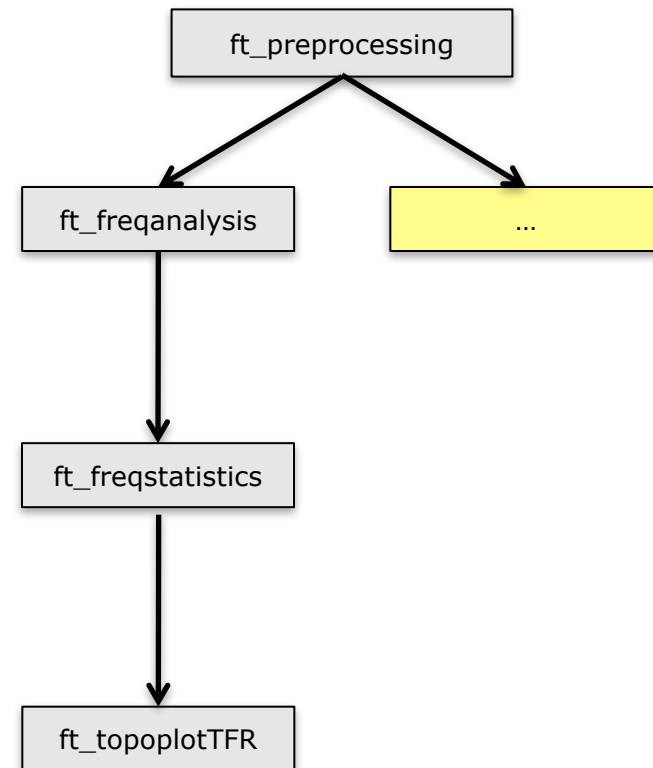


Example use in scripts

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

```
cfg = []  
cfg.method = 'mtmfft'  
cfg.foylim = [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)
```



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{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

```
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
your hands

the scripts depend on the data properties,
your computer and on your programming
skills and style

scripts correspond to analysis protocols

- scripts can be reviewed by supervisors

- scripts are often shared with colleagues

- scripts can be published/released

Finding your way around in the FieldTrip toolbox

Matlab

help functionname

edit functionname

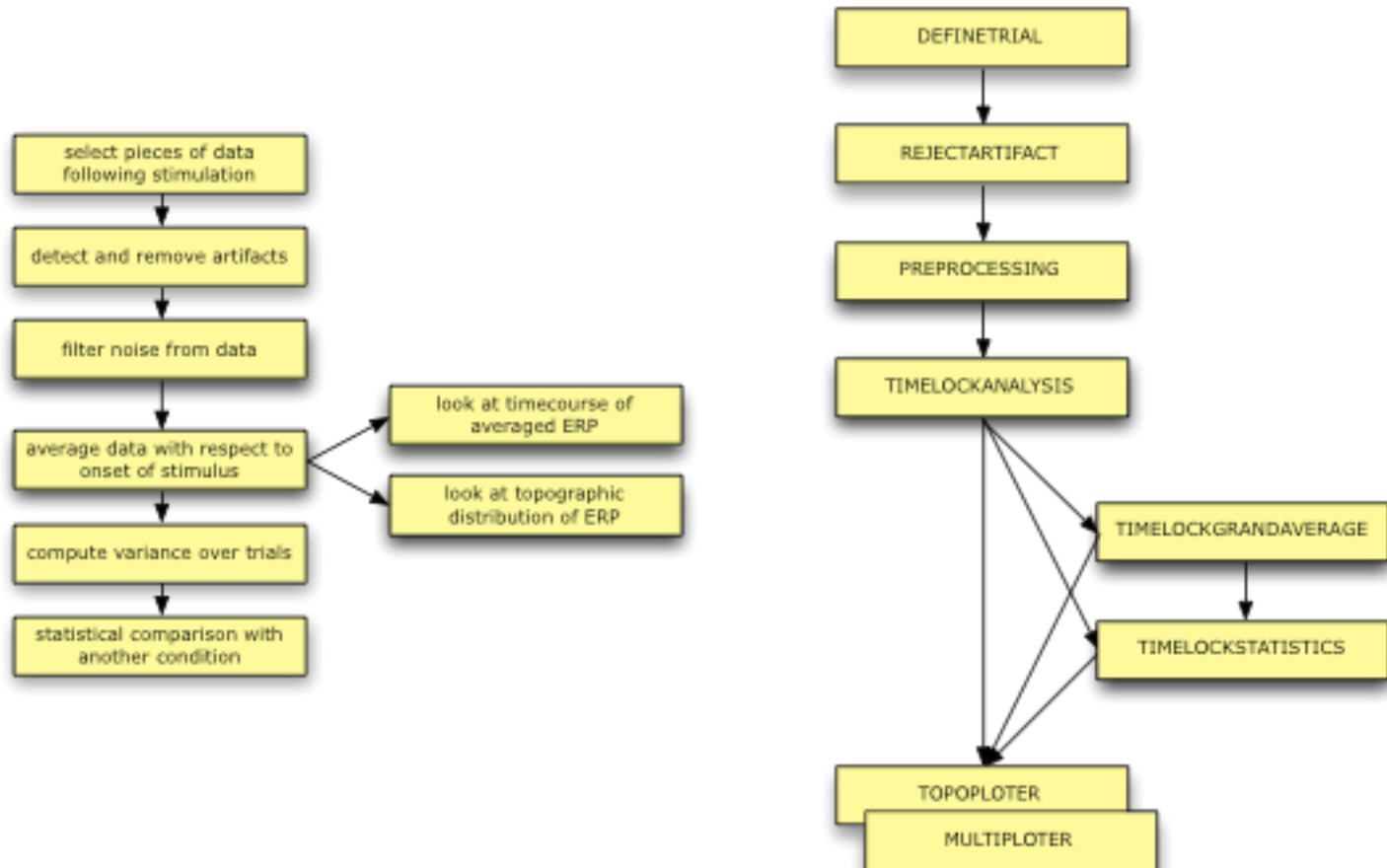
Website

<http://www.fieldtriptoolbox.org>

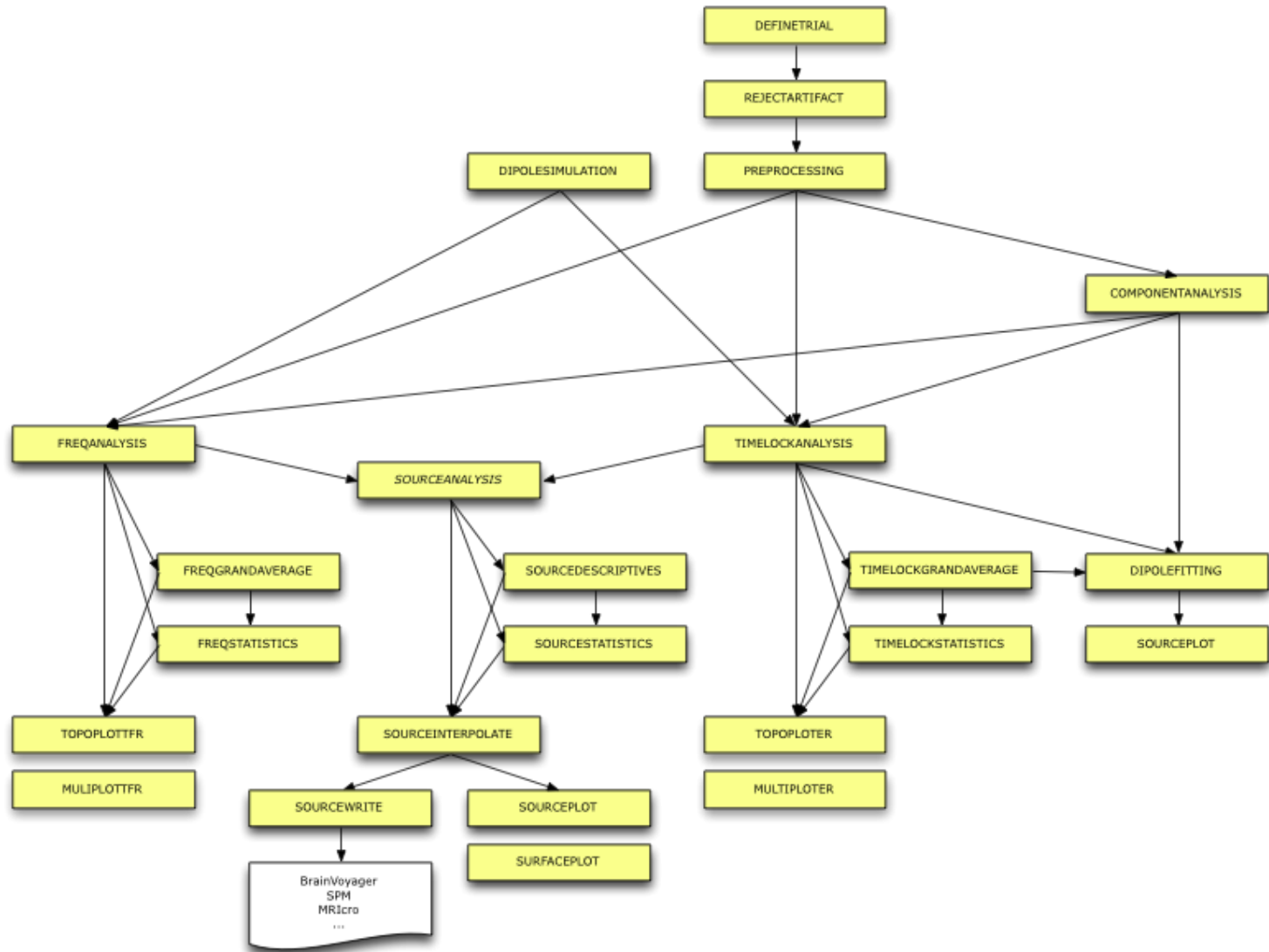
Email discussion list

Expertise in your local group

One-to-one mapping between analysis steps and toolbox functions



Overview of main functions



Talk outline

What kind of signals are generated in the brain

How do we record those 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

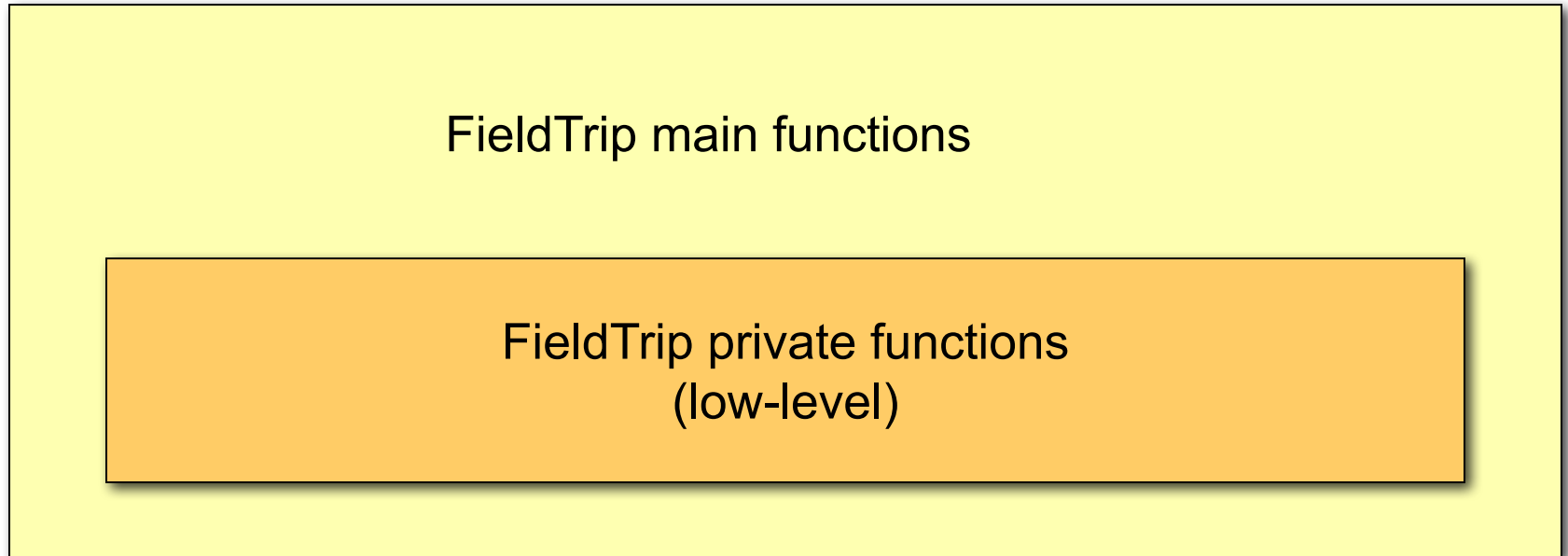
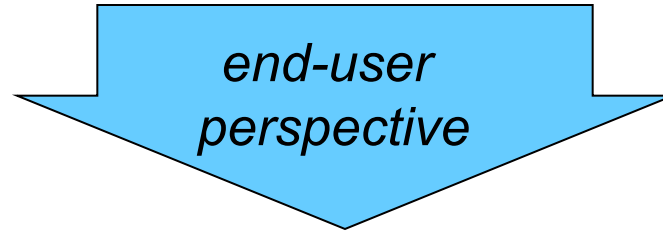
BCI2000

developers of analysis tools and methods

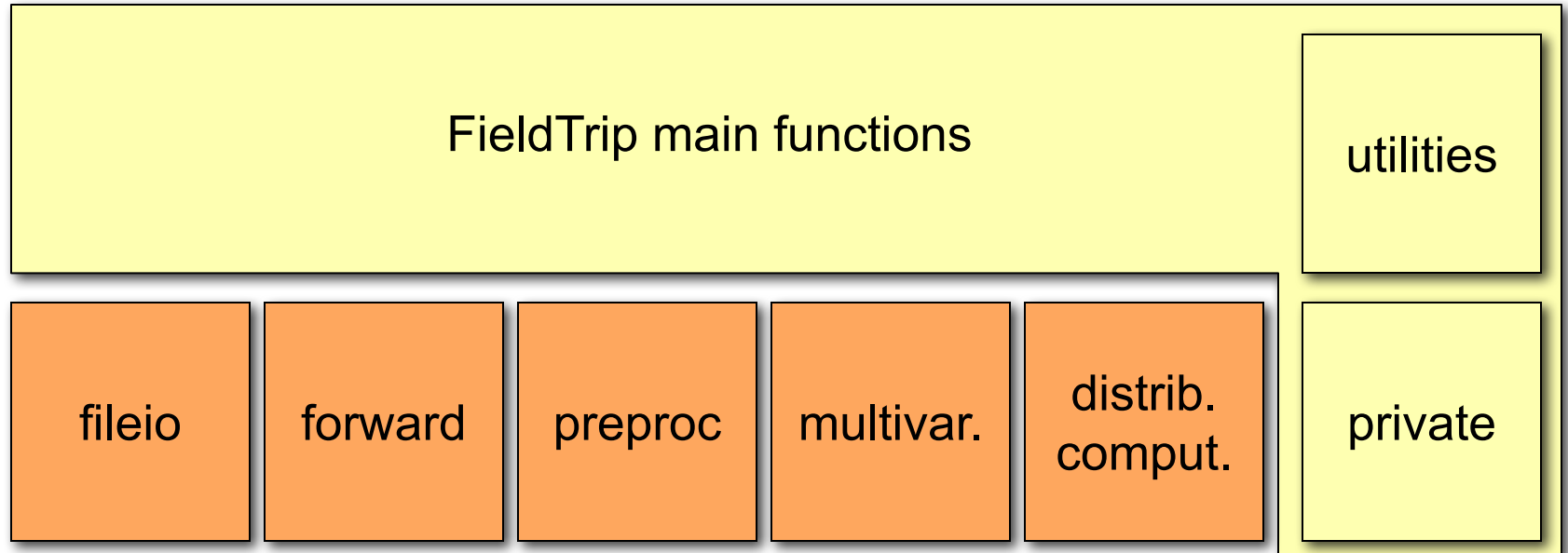
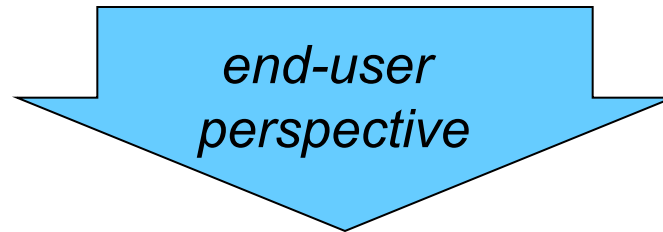
SIMBIO

OpenMEEG

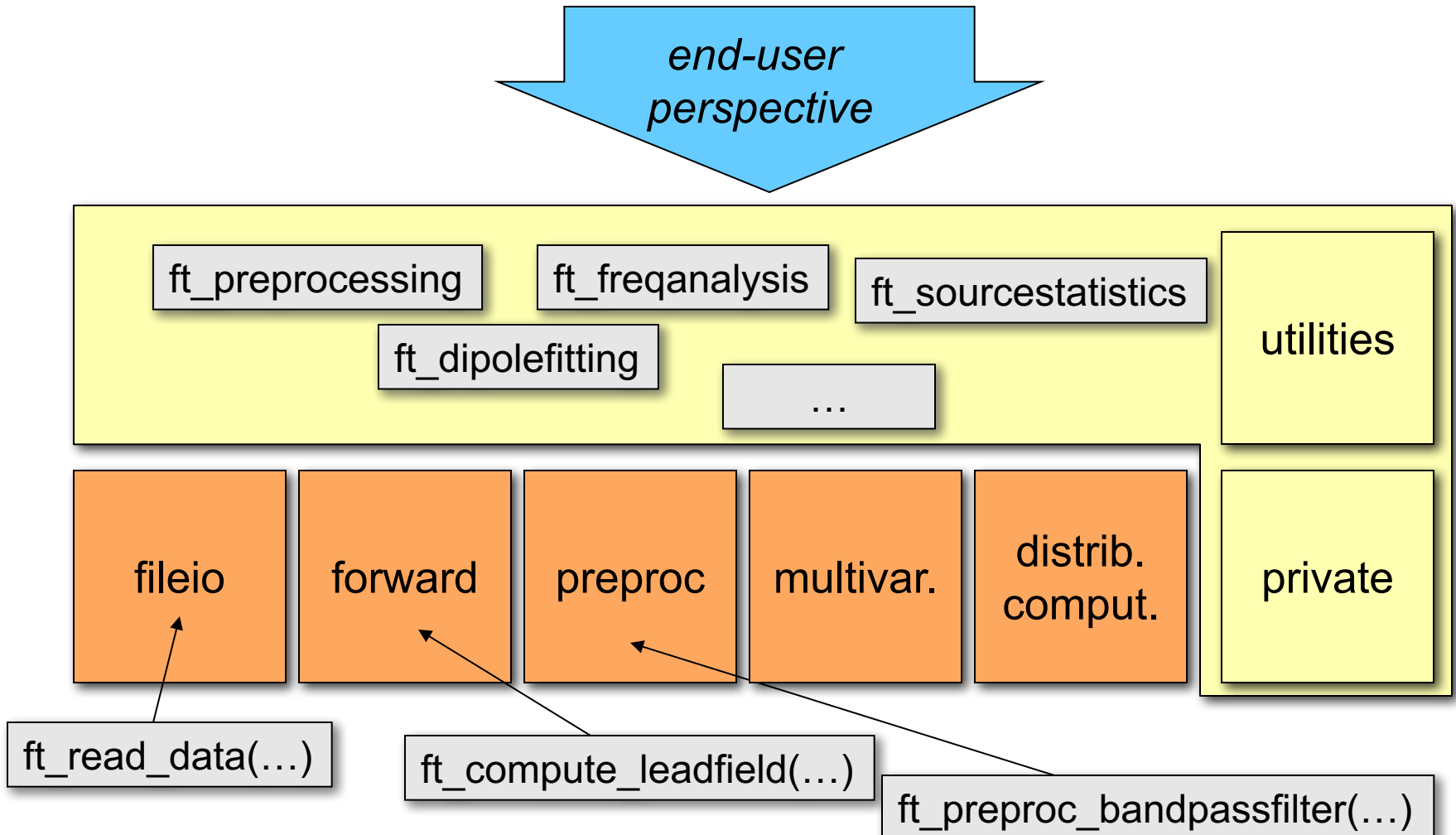
FieldTrip toolbox structure - at a glance



FieldTrip toolbox structure - a closer look



FieldTrip toolbox structure - a closer look



Summary

What kind of signals are generated in the brain

How do we record those signals

Analyzing those signals with FieldTrip

Background on the FieldTrip project

After lunch: hands-on

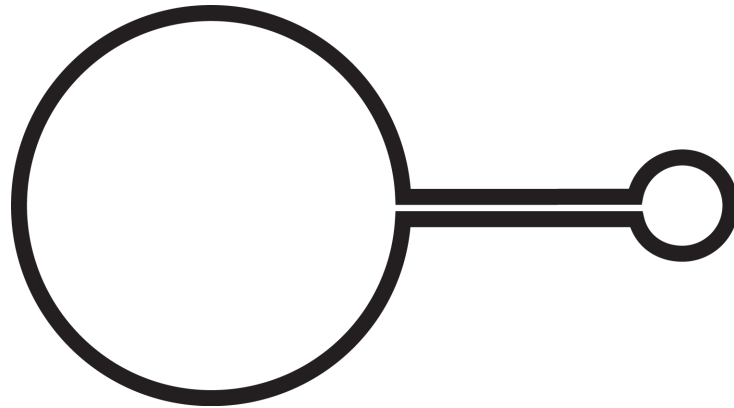
Selecting segments of data

Reading and preprocessing

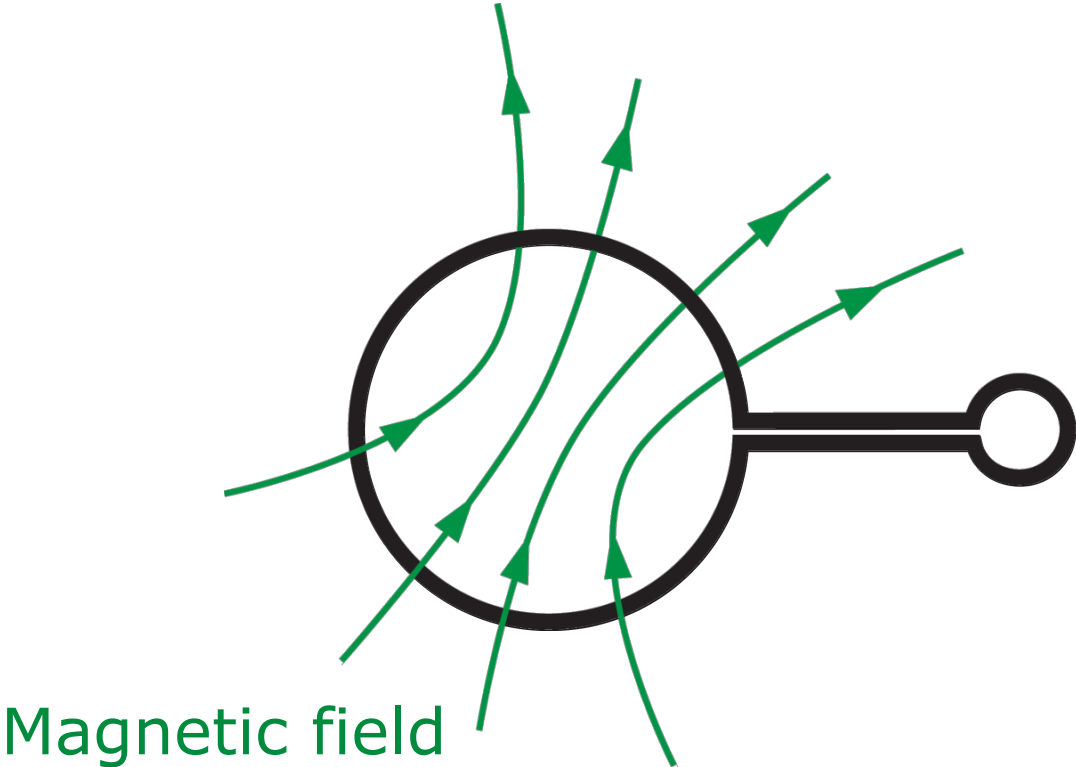
Averaging

Plotting

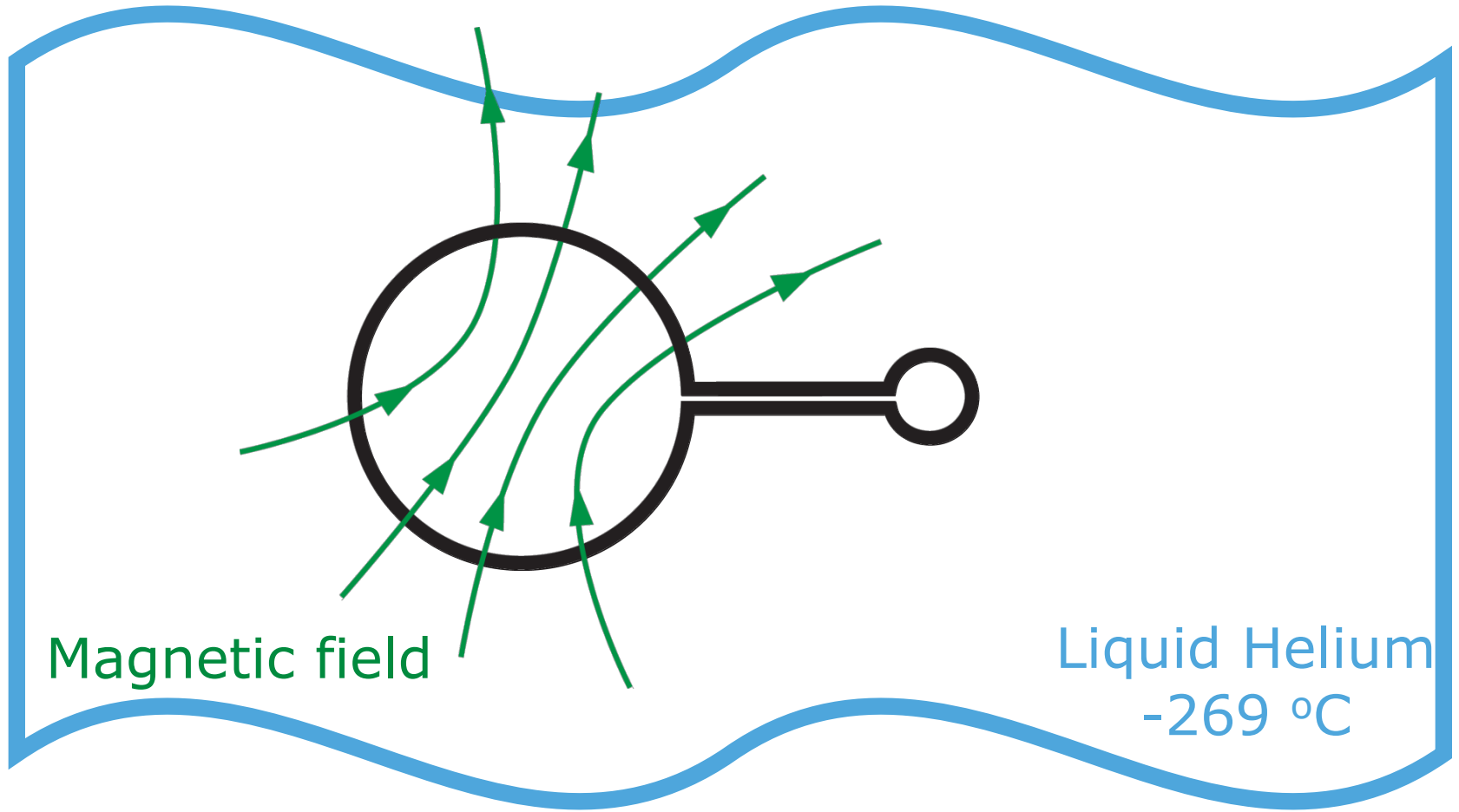
Recording small magnetic fields



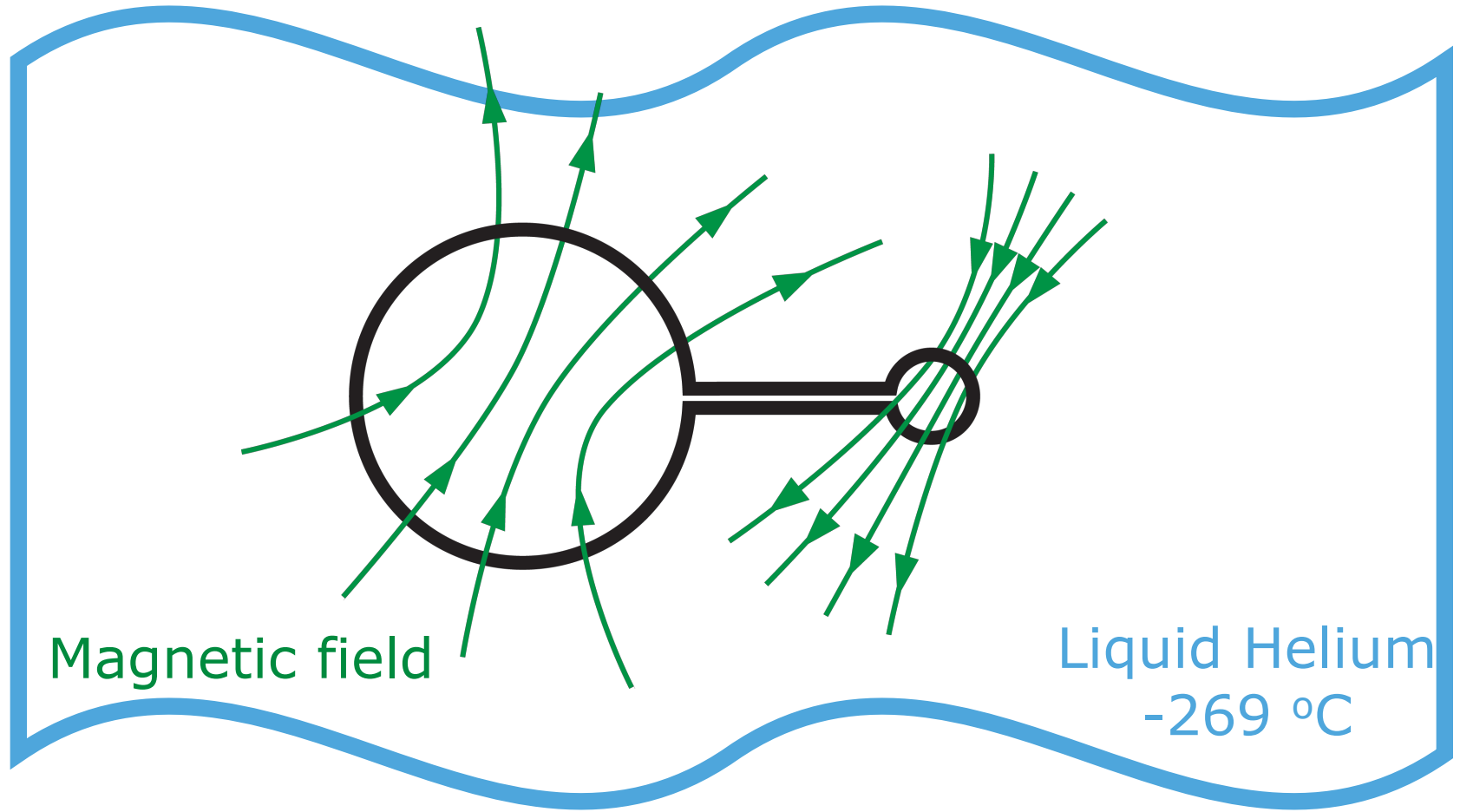
Recording small magnetic fields



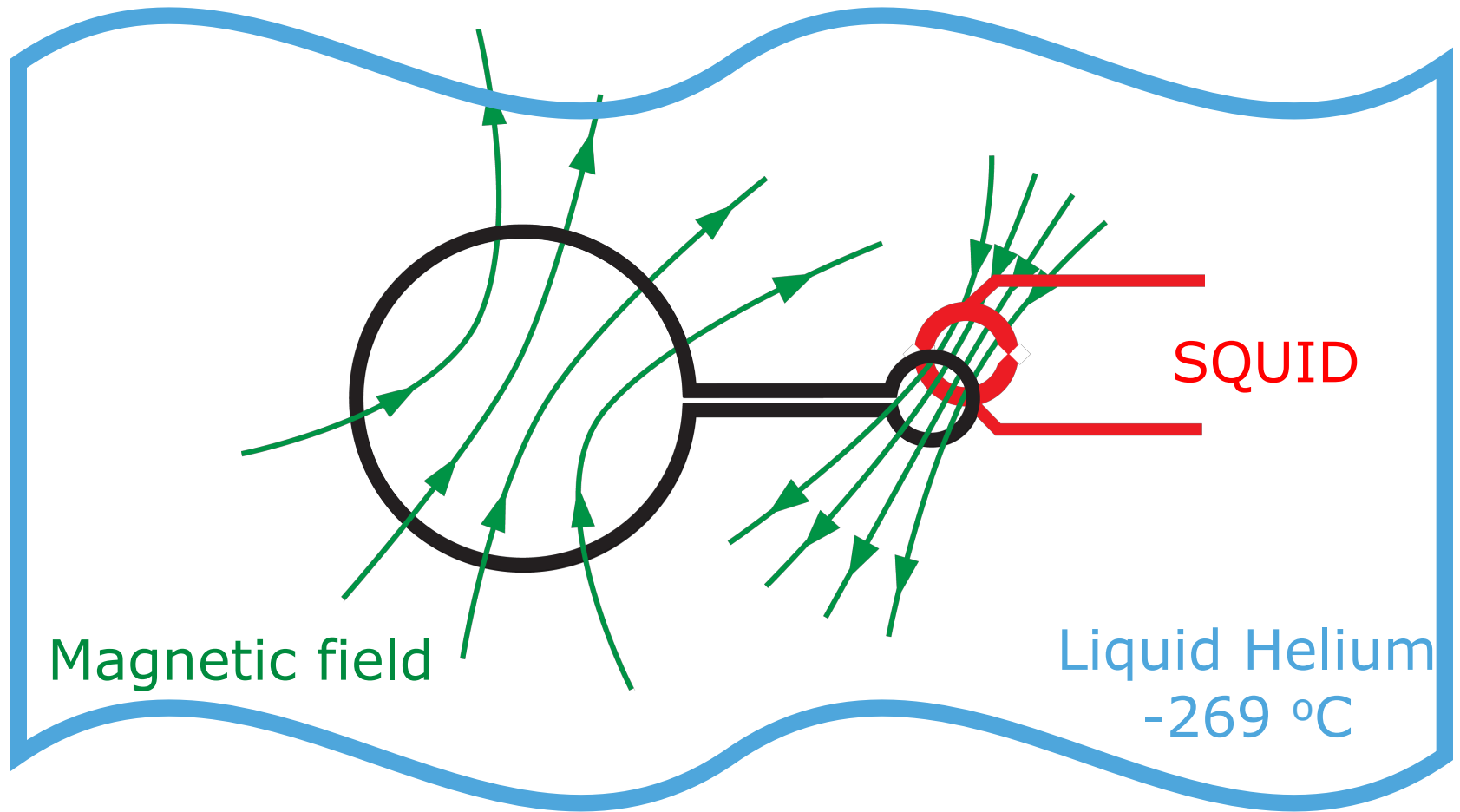
Recording small magnetic fields



Recording small magnetic fields

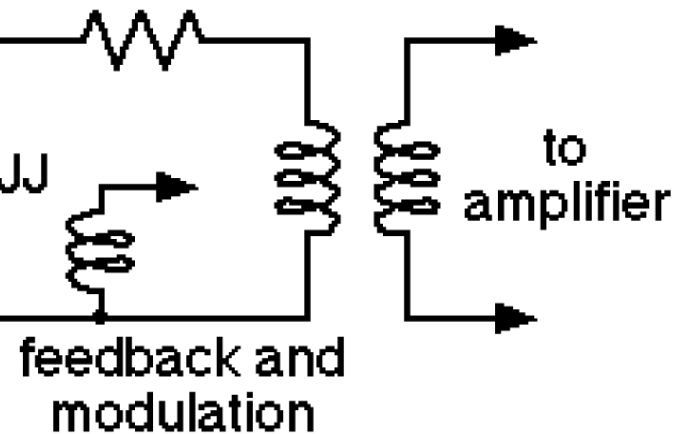
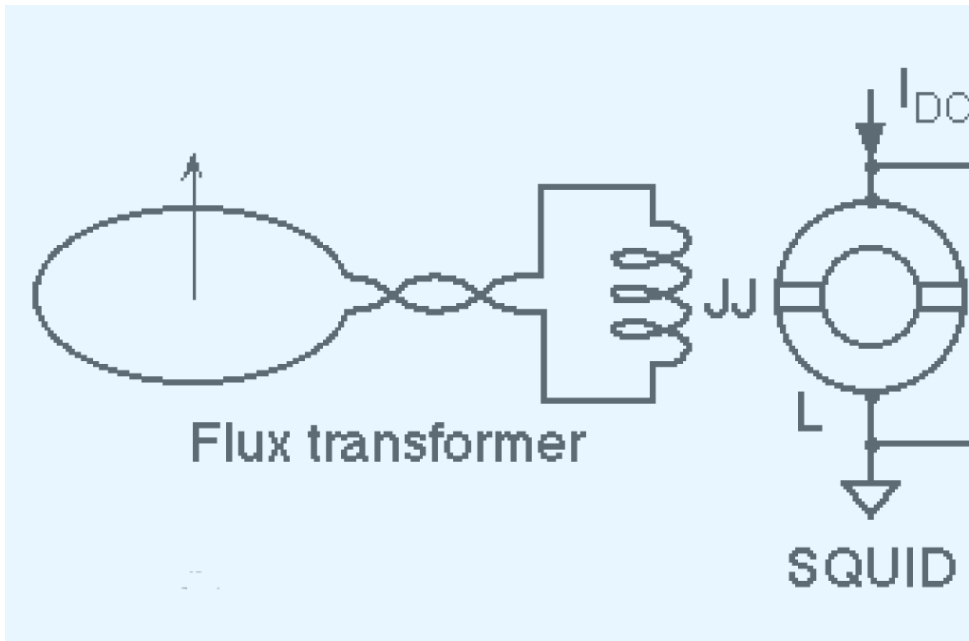
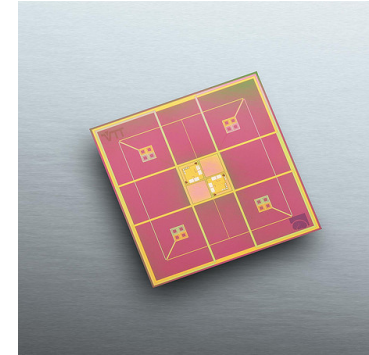


Recording small magnetic fields

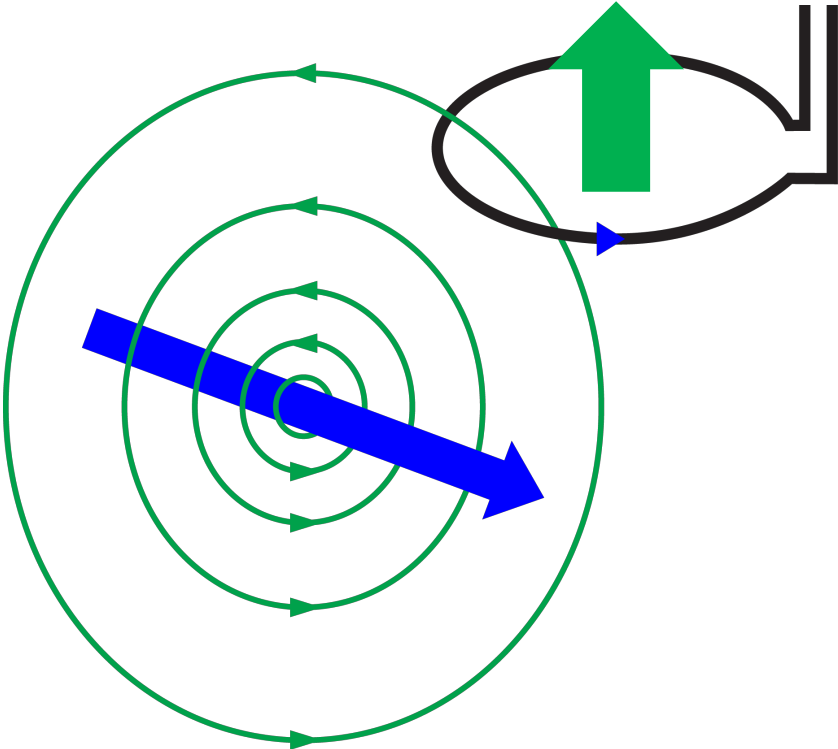


Magnetic field detectors

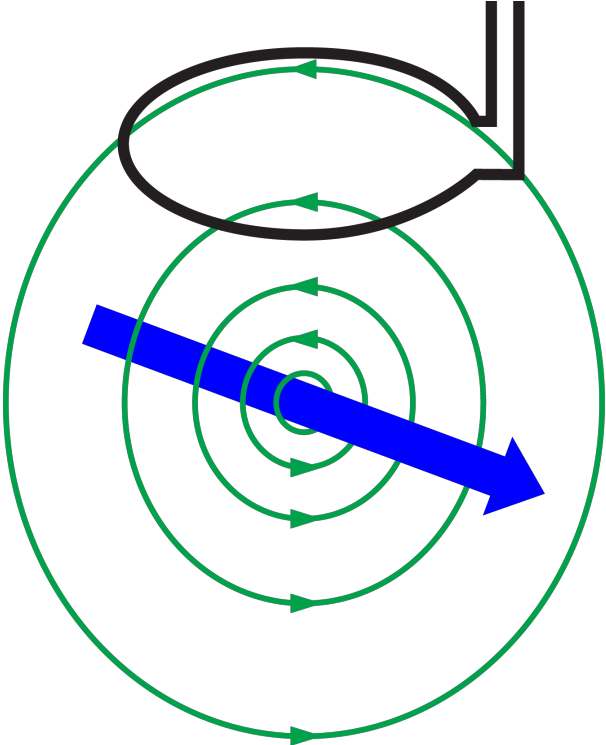
Superconducting **QU**antum **I**nterference **D**evice



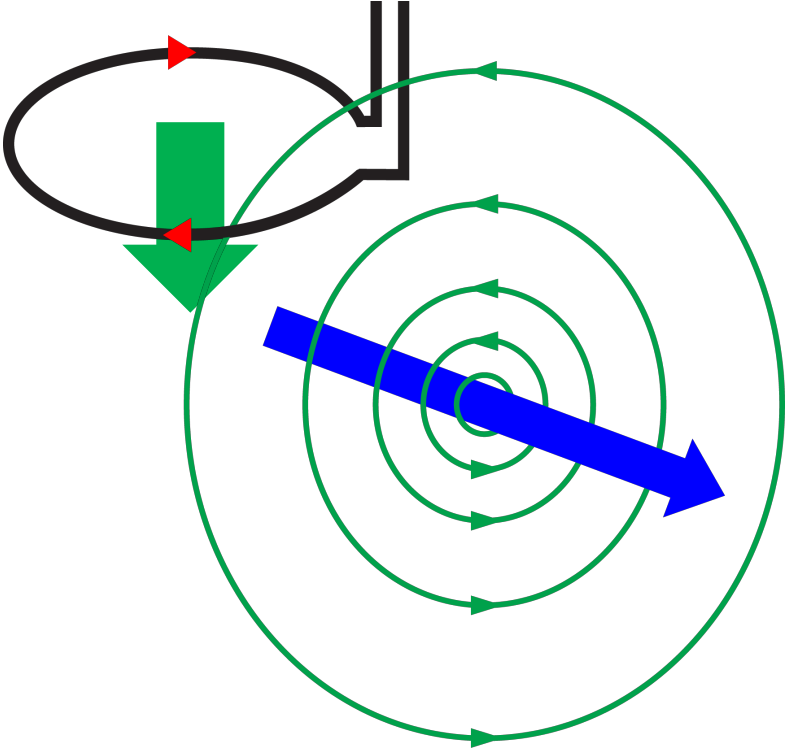
Magnetometer



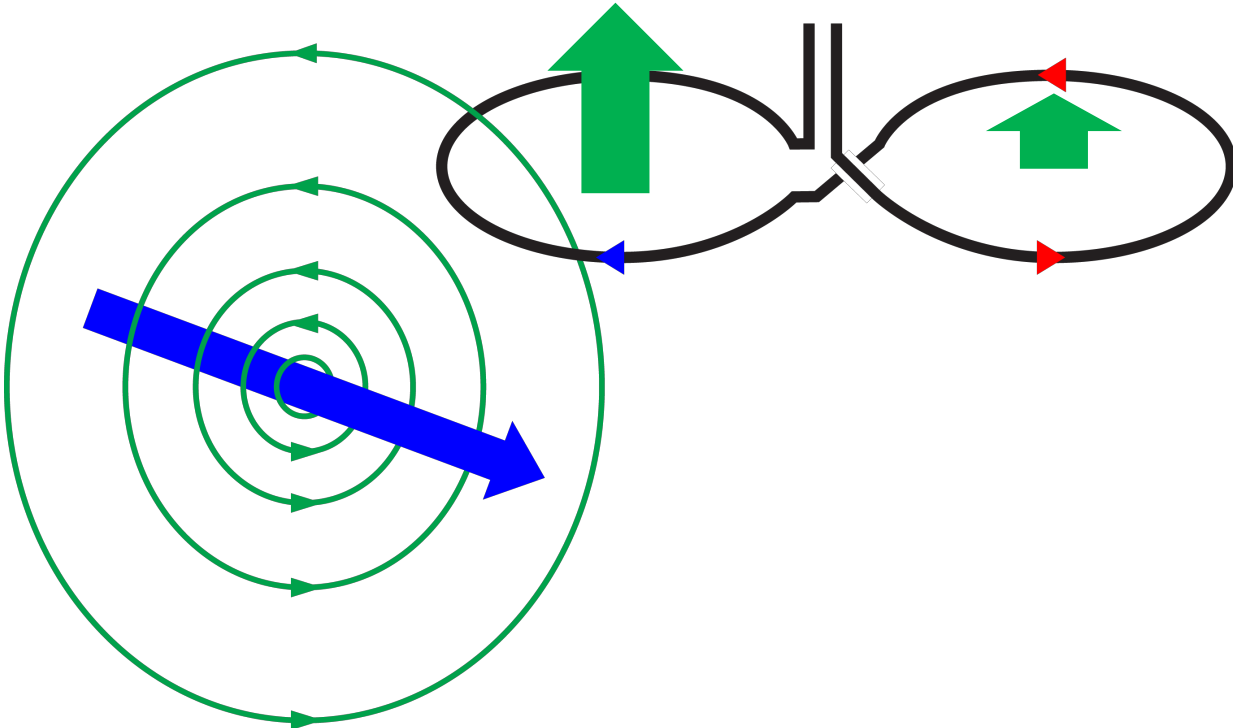
Magnetometer



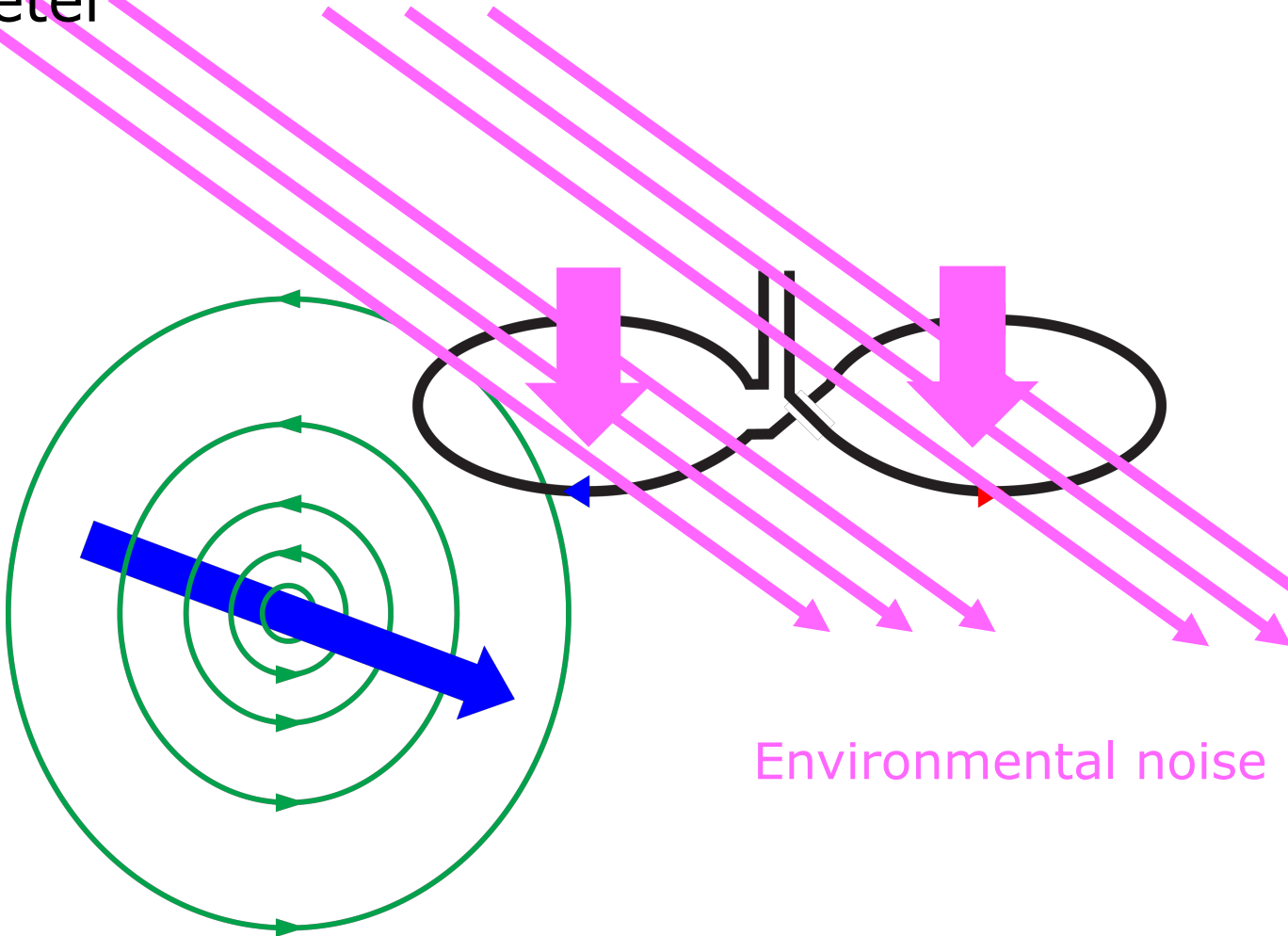
Magnetometer



Planar gradiometer

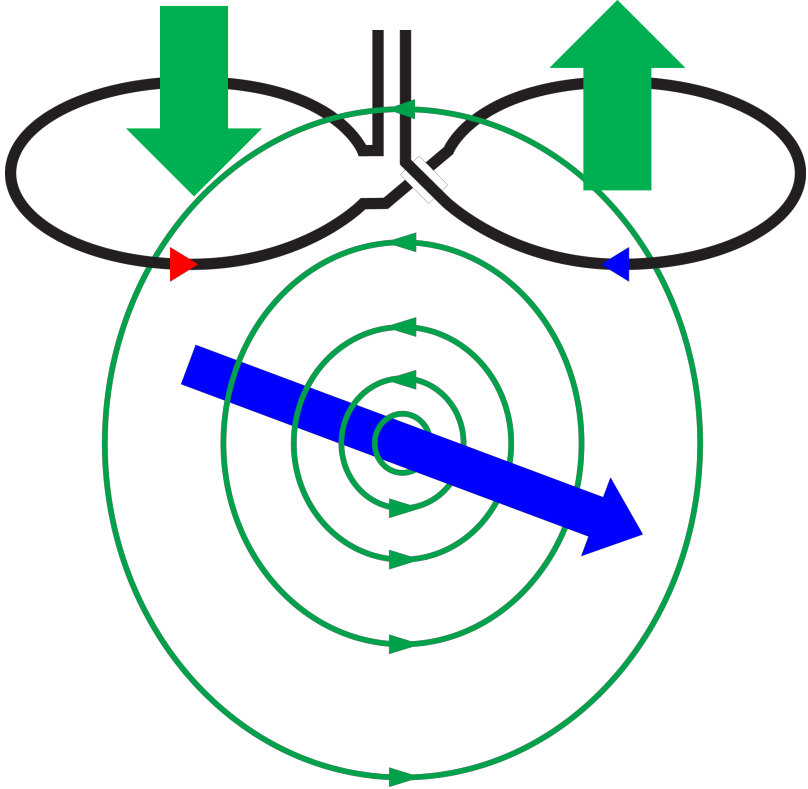


Planar gradiometer



Environmental noise

Planar gradiometer



Axial gradometer

