

#7: Introduction Into Data Analysis at CBU MEG Lab

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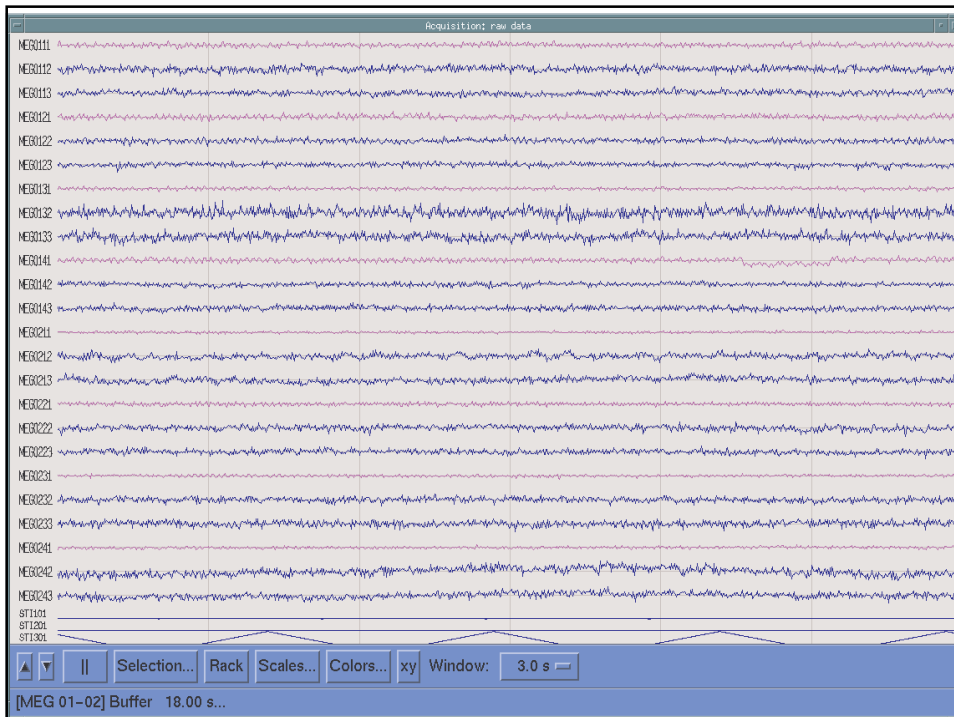
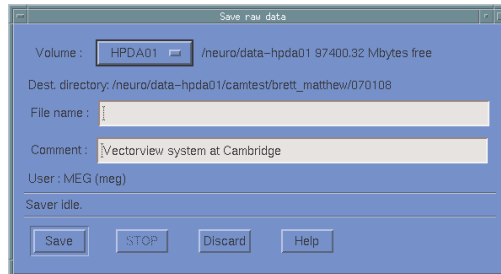


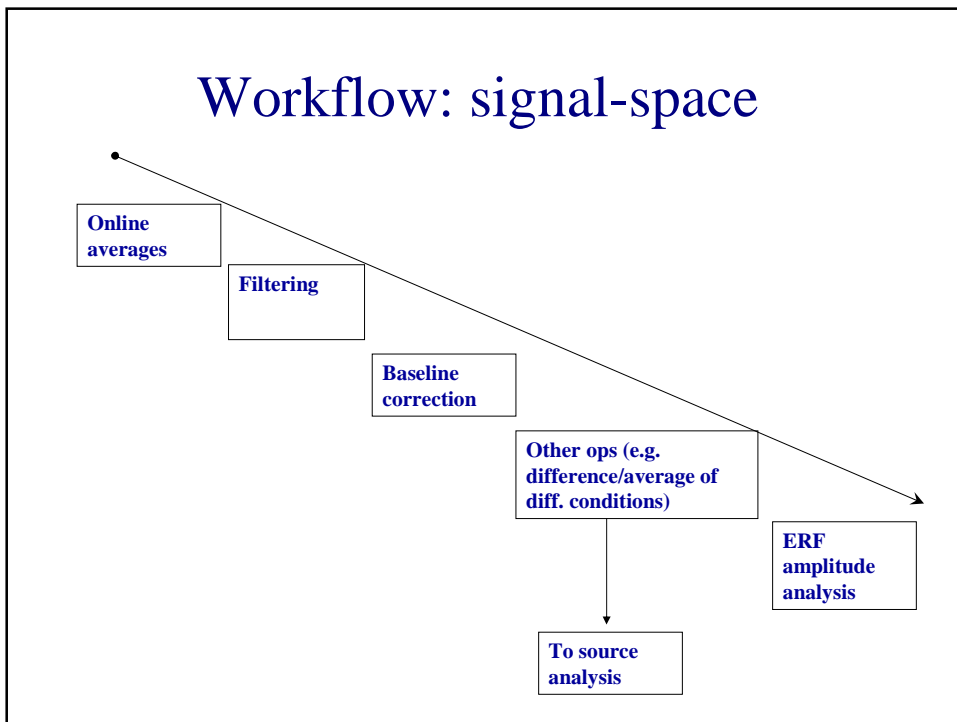
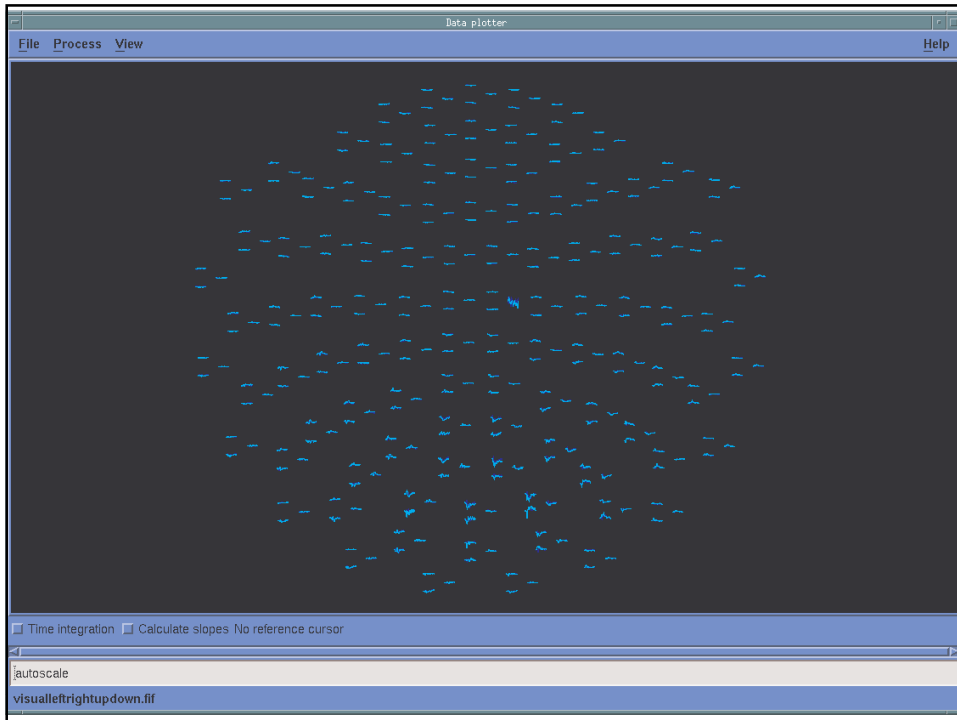
The main points

- Workflow
- Practicalities: the what and where
- Continuous data pre-processing
- Data viewing
- ERFs
- Dipoles
- Dipoles-MRI integration
- Other programs

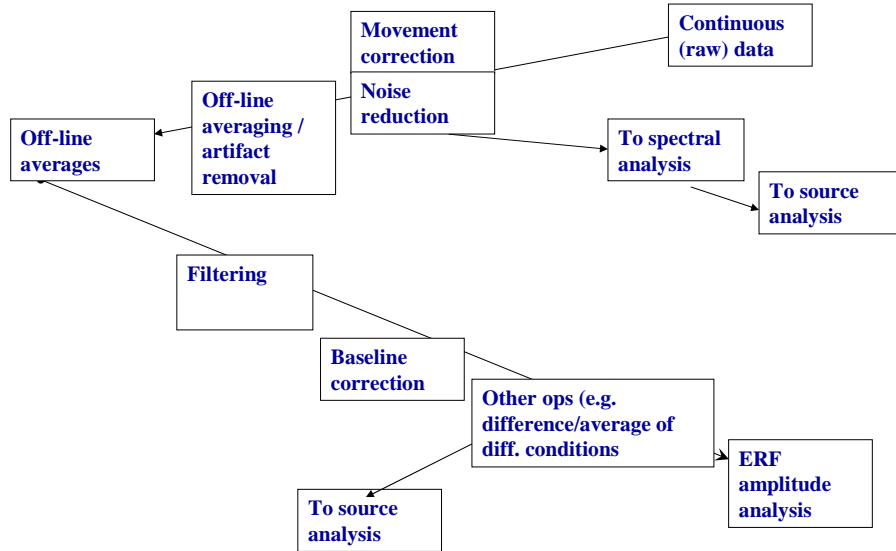
Two main data types

- Save averages
- AND raw data: two separate fiff-files

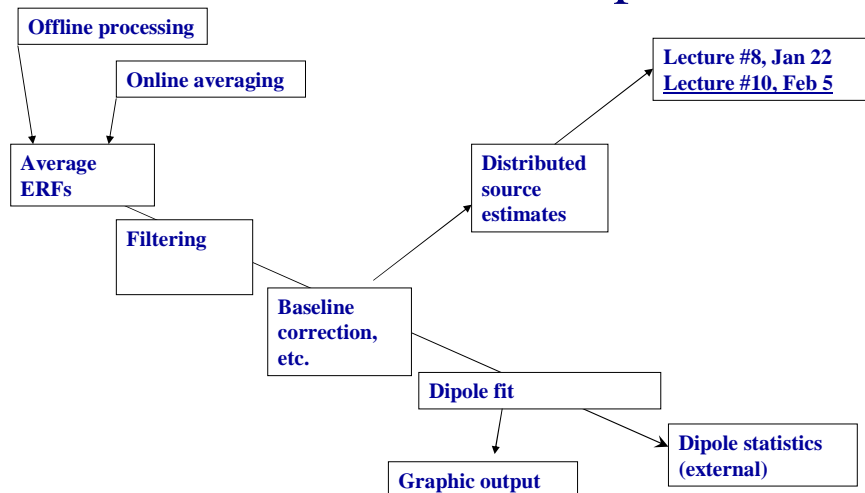




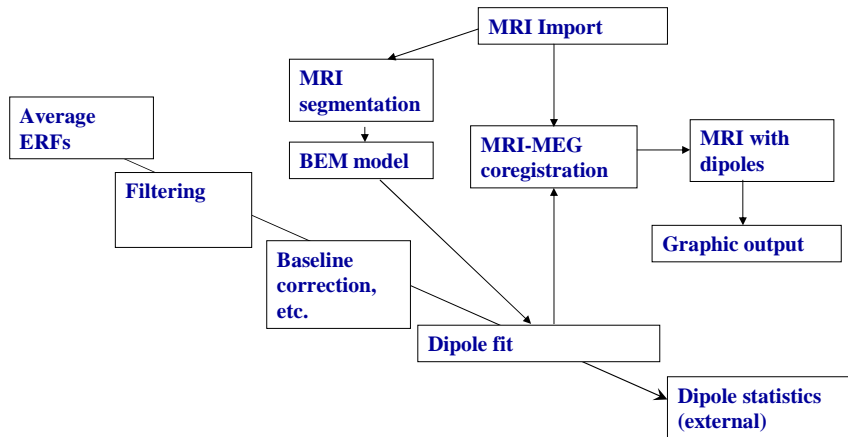
Workflow: signal-space



Workflow: source-space



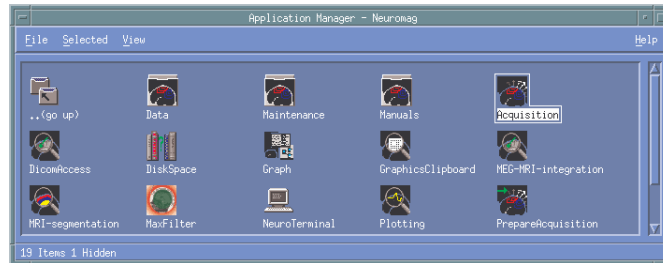
Workflow: source-space



Practicalities: data stream

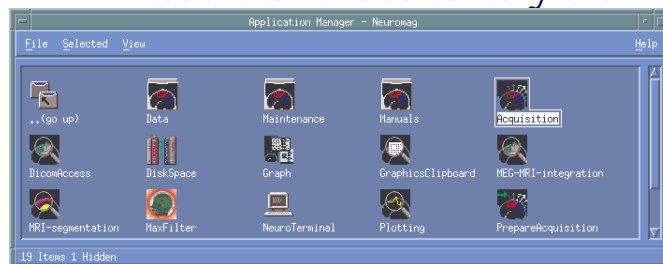
- Data recorded to /megdata/
Subdirectories, e.g.
 - /megdata/cbu = CBU
 - /megdata/cbu_csl = CSL data collected at CBU
- Access to these is read-only
- Data private to institutions
- For further processing, cp your data to your imaging space, /imaging/user-name for CBU

Practicalities: analysis



- On HPs: /opt/neuromag/bin/vue
- Continuous data access: Graph (Signal Processor), Maxfilter
- Movement correction, noise reduction: MaxFilter
- Averaging: Graph, MaxFilter

Practicalities: analysis



- Averaged ERFs: Plotter
- Source analysis: Xfit (dipoles), MCE (min norm)
- MRI:
 - DicomAccess – import into fiff format
 - MRILab – MEG-MRI integration
 - MRI-segmentation

Pre-processing of continuous data

- Viewing continuous data: Graph
- Powerful, can be used for batching
- Can control other applications
- The only tool for spectral/freq analysis in Neuromag package
- Complicated, user-unfriendly
- No linux version

Pre-processing of continuous data

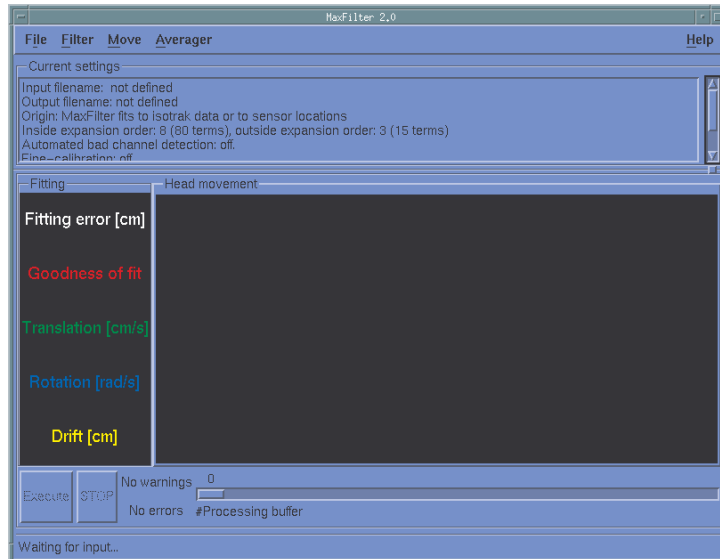
MaxFilter:

- SSS noise reduction techniques
- Movement correction
- Offline averaging

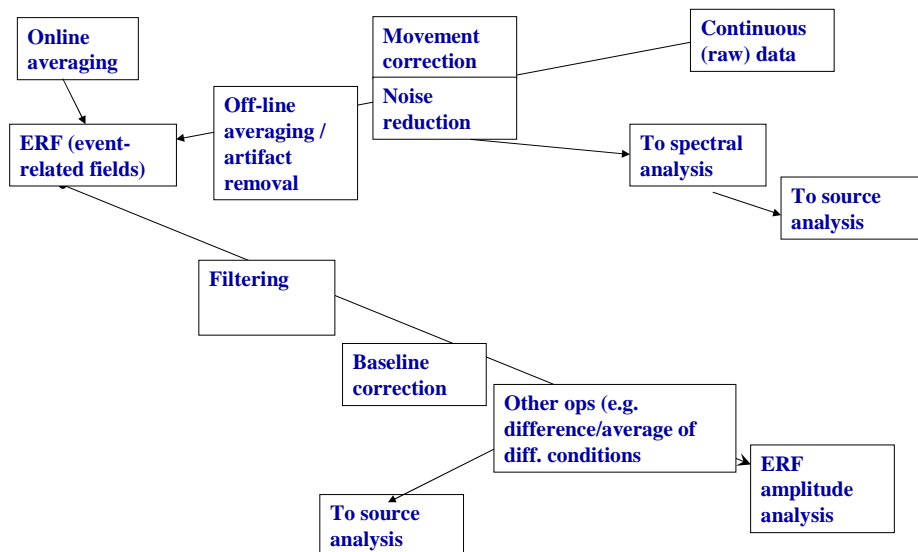
- Simple GUI
- Command line interface

- Averaging requires online averaging parameters to be pre-set

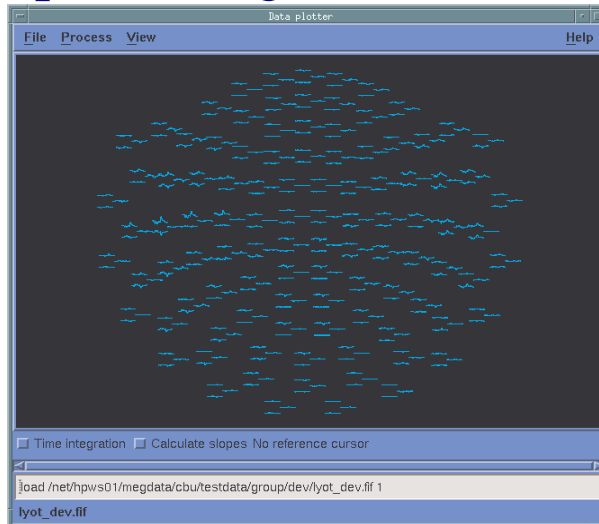
MaxFilter



Workflow: signal-space



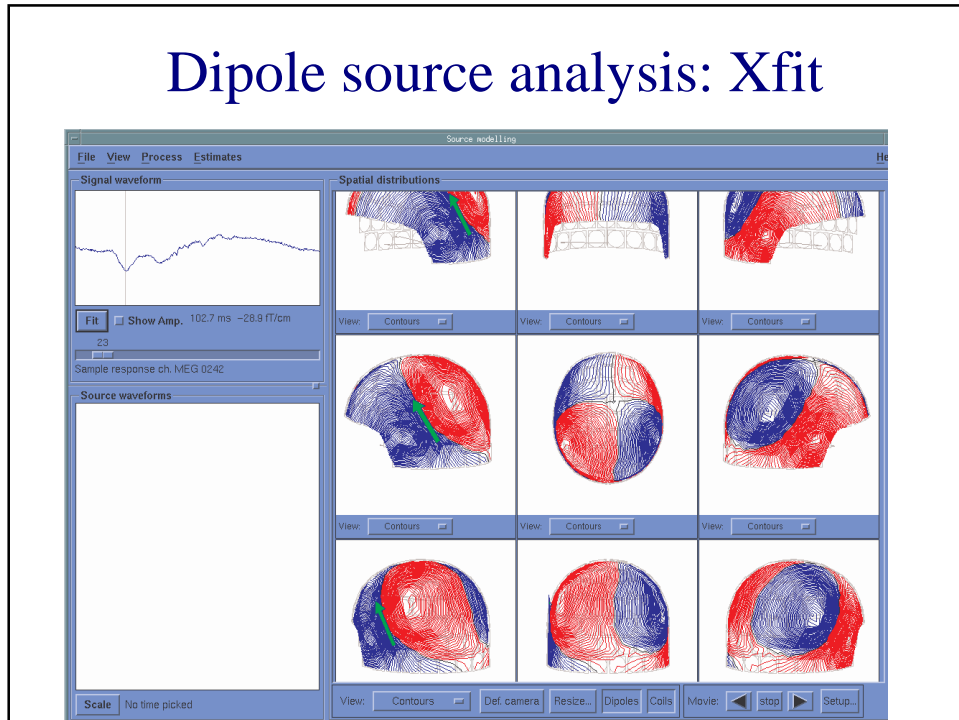
ERF viewing and basic processing: X-Plotter



ERF viewing and basic processing: X-Plotter

- Checking ERFs/bad channels
- Filtering, baseline correction
- Basic operation: subtraction, grand-average ERFs, etc.
- Peak detection and extraction for statistics
- Graphical output

Dipole source analysis: Xfit



Dipole source analysis: Xfit

- Field maps
- Single and multiple dipoles
- Re-iterated dipole fitting for a time range
- Dipole selection
- Link to MRI module
- Dipole info output for reporting/statistics
- Graphics

MRI utilities

- DicomAccess – to convert dicomm slices into fiff-format readable by Neuromag software



DicomAccess



MRI utilities

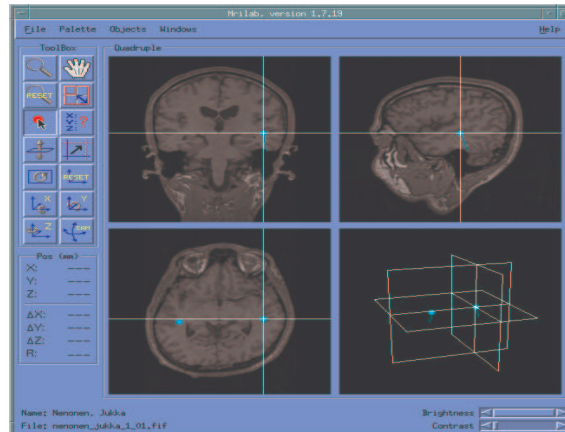
- MRILab –MEG-MRI integration



MEG-MRI integration

- Load MRI set (fiff-file)
- Co-register landmarks and or additional points
- Fit sphere (if not using BEM) and use in Xfit for head origin model
- Send dipoles from Xfit into MRILab

MEG-MRI integration



Other utilities

Neuromag:

- Minimum-norm current estimates
 - lectures by Olaf Hauk on 22.01 and
 - F. Pulvermuller on 5.02 (MCE)
- MRI Segmentation (seglab)
- Report composer
- Command-line utilities (MaxFilter, format conversion, access to fiff contents, etc.)

Other utilities

- CURRY
- BESA
- ASA
- MNE
- SPM5

Matlab access

- Fieldtrip:
<http://www.ru.nl/fcdonders/fieldtrip/>
- 4D toolbox:
<http://neuro.hut.fi/~tanzer/d4d/>
- FIFF Access:
<http://www.kolumbus.fi/kuutela/programs/meg-pd/>

HP, Linux

```
>ls /group/erp/bin/neuromag
```

```
>4Dtools-v1.1 meg-pd-1.2
```

Matlab access

- FIFF Access (Linux, HP):

```
>ls /group/erp/bin/neuromag/meg-pd-1.2
```

```
loadbdip.m  loadtri.m  rawdata.m  writeraw.m
```

```
badchans.m  loadfif.m  megmodel.m  savefif.m
```

```
chaninfo.m  loadmri.m  projmat.m  savemri.m
```

```
channames.m  loadtrans.m  rawchannels.m
```

```
subjno.m
```

- fiff2ascii conversion utility on HP machines
- Open-source Matlab tool for data access in progress

Next week

Lecture 8

**Dr Olaf Hauk: Introduction and overview
to EEG/MEG distributed source
information**

22.1.2007

