

Modifications to ucl-nir-analysis in order to include 31P-data

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The modified version of the synchronisation script requires the same folder structure as the original ucl-nirs-analysis script with an additional subfolder /phosphorus to every LWPXXX folder containing the 31P data sheets.

```
pigletdatadir
├── studyDir
│   ├── LWPXXX
│   │   ├── nirs
│   │   ├── systemic
│   │   └── phosphorus
│   │       └── LWPXXX_insult_analysis.xlsx
│   ├── LWPXXY
│   ├── LWPXXZ
│   ├── study_log.xlsx
│   ├── 31p_log.xlsx
│   └── ...
```

In order to read out the 31p data and synchronise it with the nirs/systemic data, the following modifications must be made to the folder containing the original ucl-nirs-analysis scripts:

- Copy the new file *phosAnalyseAndSync.m* to the original /func folder.
- Also in the original /func folder, replace the original *load31PData.m* with the new version

- Use the included files *DataAnalyse_31p.m* and *31p_log.xlsx* as templates and adjust them to the files you want to analyse and the folder topology on your PC.
- (If script runs very slowly: comment out lines 345-349 in */tarragona/Broadband_Tarragona.m*)

Running *DataAnalyse* will then create a variable *pigDataPTB* that has the same basic structure as *pigDataNTB* with additional rows of data for various MRS variables. This variable is stored in a file called *pigDataPTB* in the specified output folder.

Note that for the synchronisation NIRS and systemic data are down-sampled to the sampling time of the 31p-data (usually 1min).