${\bf COMAP\ \, Nemo\ \, 0:\ \, COMAP\ \, Manchester} \\ {\bf Reduction\ \, Pipeline\ \, Documentation}$

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1 Overview

The Manchester COMAP data reduction pipeline is designed to take the uncalibrated, full resolution COMAP level 1 data and reduce it to wide 1 to 2 GHz bands that are calibrated using astronomical sources. We refer to the data calibrated using the vane and downsampled to 32 MHz channels as level 2 data. After this any further reductions are called level 3 data where astronomical calibration has been applied, and the data has been filtered of bad frequency channels and as many systematics as we can. Level 3 data is intended to be passed directly to a map-maker. There may be several level 3 datasets.

The final step of the Manchester COMAP pipeline is to filter out all observations with bad weather, poor noise statistics, or any other poor performance criteria. After the data have been filtered we can pass the selected good *level 3* data to the Destriping map-maker, resulting in a sky map, covariance map, and hit distribution map stored in FITS standard format with some predefined World Coordinate System (WCS).

A quick overview of the processing pipeline is:

- 1. All level 1 data is processed through several modules:
 - Measure the system temperature (T_{sys}) and gain (G) using the calibration vane.

[Calibration.CalculateVaneMeasurement]

- Calibrate the *level1* data using the *vane*. [Calibration.CreateLevel2Cont]
- Define edges of *scans* and store these. [Statistics.ScanEdges]
- Flag out any large spikes in the data exceeding 10σ of the noise. [Flagging.SigmaClip]
- Measure noise statistics per 32 MHz channel and fit 1/f statistics. [Statistics.FnoiseStats]
- 2. The resulting *level 2* data is then further processed into *level 3* data. There are several choices to make:
 - File lists of good data based on noise statistics.
 - Astronomical calibration to apply leave vane calibration? Tau A? Jupiter?

- Channel masks to apply, and account for how these change with time.
- Finally choose run name and create *level 3* reduction. [CreateLevel3.CreateLevel3]

2 Running the Pipeline

2.1 The Code

All the code you need for the pipeline are stored in the github repository found here: https://github.com/SharperJBCA/COMAPreduce. There are three modules:

- Analysis This contains all the data reduction pipeline for making level 3 data.
- MapMaking All the code for making maps.
- Tools Extra scripts that may be useful like coordinate transforms, stats not included in scipy, etc...

2.2 Installation

2.2.1 General Linux Machine

First you will need to install Python 3.X. To do this I recommend download the Anaconda distribution. Go to: https://www.anaconda.com/products/individual. This will provide you with a .sh file that from the command line you can install by:

> bash <filename>.sh

From here just follow the instructions, and make sure you install Anaconda somewhere on the local disks (e.g., /local/scratch/<username>). Now you will have a new command line tool call conda and you will be using this to setup your environment and install new Python packages. The first thing you must do after install conda is link the conda-forge (https://conda-forge.org/) repository, to do this run the command:

- > conda config --add channels conda-forge
- > conda config --set channel_priority struct

Next you will want to install the pipeline code. First you must clone the github repo to somewhere on your machine:

> git clone https://github.com/SharperJBCA/COMAPreduce.git .

This will create a directory called COMAPreduce, enter this. We will now create a new virtual environment that matches mine so that you can be sure we are running the code with all the same packages. To do this simply execute:

> conda create --name <env> --file requirements.txt

Where **<env>** is the name of the virtual environment you want to use. Whenever you **ssh** into the machine you will need to use the command:

> conda activate <env>

to ensure your system is setup correctly. Alternatively you can add this line to your rc file (e.g., .tcshrc in your home area).

Finally, you can now install the pipeline routines as a module into your current Python setup by simply invoking:

> python setup.py install in the COMAPreduce directory.

2.3 Running the Pipeline

Will update later...

2.4 Running the Map-Maker

To run the map-maker create a new working directory somewhere on your system. Next copy from the COMAPreduce/comancpipeline/MapMaking directory the following files/directorys to your working directory:

- run_destriper.py
- CreateFilelist.py
- ParameterFiles

Inside the ParameterFiles directory you will find a number of .ini files that are pre-setup for running the M31 data (fg4), the Galactic plane survey (GFields/fg6), and the Perseus data (fg7). These are called:

- ParameterFilesparameters_fg4.ini
- ParameterFilesparameters_fg7.ini
- ParameterFilesparameters_GFields.ini

First you will need to create filelists of processed *level 3* data files by invoking the CreateFilelists.py script:

> python CreateFilelists.py <source-name>

where <source-name> is fg4, fg7, GField, etc... Note: feel free to edit this file so you can optimise the file lists. The filelists will be stroed in a FileLists/directory.

Once you have a filelist ready you can invoke the map-maker using: > python run_destriper.py <parameter-file>

3 Old-Stuff

Requirements:

- Python 3.0 or higher.
- Compiled shared libraries of the FORTRAN version of the Starlink astronomical libraries (SLALIB) available from http://starlink.eao.hawaii.edu/starlink/2018ADownload.
- A copy of parallel ready H5Py. Installation instructions can be found here http://docs.h5py.org/en/stable/mpi.html. N.B.: If you are using an Anaconda packaged version of PYTHON installation you may need to remove the existing CONDA install of HDF5.
- The latest version of HEALPY, MPI4PY (either openMPI or MPICH work as backends), NUMPY, SCIPY, MATPLOTLIB, and ASTROPY.

To install the Manchester COMAP reduction pipeline:

- Clone/download the github repository found here: https://github.com/SharperJBCA/COMAPreduce.
- Enter the directory: cd COMAPreduce and run python setup.py install. If your SLALIB libaries are not in standard location you must define the environment variable: SLALIB_LIBS
- To run the COMAP pipeline make a new directory above COMAPreduce (e.g. cd ../ && mkdir runcomapreduce) and copy the RUN.PY, and .INI files there.

• The pipeline can then be run using the command: mpirun -n X python run.py -F FILELIST.list -P PARAMETERS.ini. FILELIST.list should contain a list of files with either just the filenames to be processed or the full path to files to be processed. PARAMETERS.ini will control the processing to be performed, details of which are described in Sections 4 and 5.

4 Usage

4.1 Parameter Files

There are several example parameter files already included:

- Ambload.ini This will calculate the $T_{\rm sys}$ and gain (e.g. volts per Kelvin) from ambient load stare observations.
- Downsample Ini This will downsample a data file in frequency by factor times and also check to see if any pointing needs to be added.
- FITJUPITER.INI This will fix the pointing, downsample, and calibrate a Jupiter observation to the ambient load. Then it will fit a Gaussian to the time ordered data to derive amplitude, pointing and beam width measurements. It will also produce a calibration scale in units of Janskys/Kelvin for every horn and frequency channel.

5 Classes

5.1 BaseClass.H5Data

Useful functions for defining new classes:

- getdset Retrieve a dataset, if it is not in memory load it.
- setdset Load a dataset into memory.
- resized et Resize a dataset by passing it a new array.
- updatedset Update a dataset values.
- getAttr Get an attribute (stored in output file)
- setAttr Set an attribute
- getextra Get a dataset from the extra outputs

- setextra Set an array to be assigned to extra outputs (must describe the shape of the array, e.g. which axis refers to horns, frequencies, etc...)
- resizeextra Change dimensions of an extra dataset.

N.B. Never write directly to the dset or extras attributes of the H5Data class.

Useful attributes/functions for MPI routines:

- splitType Axis type being split for MPI purposes (i.e., either Types._HORNS_, Types._SIDEBANDS_, Types._FREQUENCY_, Types._TIME_).
- selectType Axis type being explicity selected (i.e., as above)
- selectIndex Index being selected along selectType axis.
- splitFields Names of fields in COMAP data structure that will contain a split axis.
- selectFields Names of fields in COMAP data structure that will have a selected axis.
- hi/lo Dictionary, for each splitField, defining where in the larger structure this process is accessing data.
- ndims Dictionary containing dimensions of each dataset in memory for this process.
- fullFieldLengths Dictionary containing dimensions of each dataset in memory in total.
- getDataRange Function that returns how to split N values between M processes.