



# Functionality

Extractor → how to read RAT root files/ntuples?



fill

- Fit parameters – array of values for parameter

Systematics:

- Smear
- Scale
- Shift

DBIsotope → calculate double beta decay rates

Statistic e.g. chi\_squared

config  
Which dimensions?  
store.load  
Load from HDF5

Spectra

Core data structure – numpy.ndarray

- Spectra methods:
- shrink – cuts
  - scale – adjust rate/amplitude

Spectra – modified  
+ other Spectra

Fitting/Limit Setting

results

- Spectra methods:
- sum – number of events
  - output :
  - plot
    - plot\_projection
    - spectral\_plot
  - store
  - dump – save to HDF5

- Plots:
- Chi squared
  - SystAnalyser
- Table of limits
- Table of best fit values