

```
In [4]: import PALpulsarInit
import bayesutils as bu
import os, sys, glob
import numpy as np
import h5py as h5
import matplotlib.pyplot as plt
```

Getting Data into the HDF5 File Format

```
In [9]: # run the help function for PALpulsarInit
!python PALpulsarInit.py -h
```

Usage: PALpulsarInit.py [options]

Options:

```
-h, --help                show this help message and exit
--parDir=PARDIR           Full path to par files (required)
--timDir=TIMDIR           Full path to tim files (required)
--noiseDir=NOISEDIR       Full path to noise files
--outFile=OUTFILE         Full path to output filename (required)
--distFile=DISTFILE       Full path to pulsar distance file
--DMOFF=DMOFF             Turn on DMMODEL fitting
```

```
In [6]: # first lets just get the actual data into the file
!python PALpulsarInit.py --parDir './mdc_data/partim/open1/' --timDir './mdc_data/partim/open1/'
```

WARNING [TIM1]: Please place MODE flags in the parameter file

WARNING: duplicated warnings have been suppressed.

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469438580184544.000000

good=36028797018963968.000000

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469439262882464.000000

good=36028797018963968.000000

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469439263602336.000000

good=36028797018963968.000000

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469439263200448.000000

good=36028797018963968.000000

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469439263027584.000000

good=36028797018963968.000000

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469439263120992.000000

good=36028797018963968.000000

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469439263195232.000000

good=36028797018963968.000000

ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit

wmax=251469439262550016.000000

```

good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439263196128.000000
good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439262733792.000000
good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439263117696.000000
good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439262842176.000000
good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439262812448.000000
good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439263197280.000000
good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439262880416.000000
good=36028797018963968.000000
*****
ERROR [TKfit.C:176] Warning: wmax very large. Precision issues likely to break fit
wmax=251469439263134272.000000
good=36028797018963968.000000

```

```

In [10]: # now lets read in the file
pfile = h5.File('./mdc_data/open1_demo.hdf5')

```

```

In [13]: # we can look at which pulsars are in the data file
pfile['Data']['Pulsars'].keys()

```

```

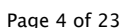
Out[13]: [u'J0030+0451',
u'J0218+4232',
u'J0437-4715',
u'J0613-0200',
u'J0621+1002',
u'J0711-6830',
u'J0751+1807',
u'J0900-3144',
u'J1012+5307',
u'J1022+1001',
u'J1024-0719',
u'J1045-4509',
u'J1455-3330',
u'J1600-3053',
u'J1603-7202',
u'J1640+2224',
u'J1643-1224',
u'J1713+0747',
u'J1730-2304',
u'J1732-5049',
u'J1738+0333',
u'J1741+1351',
u'J1744-1134',

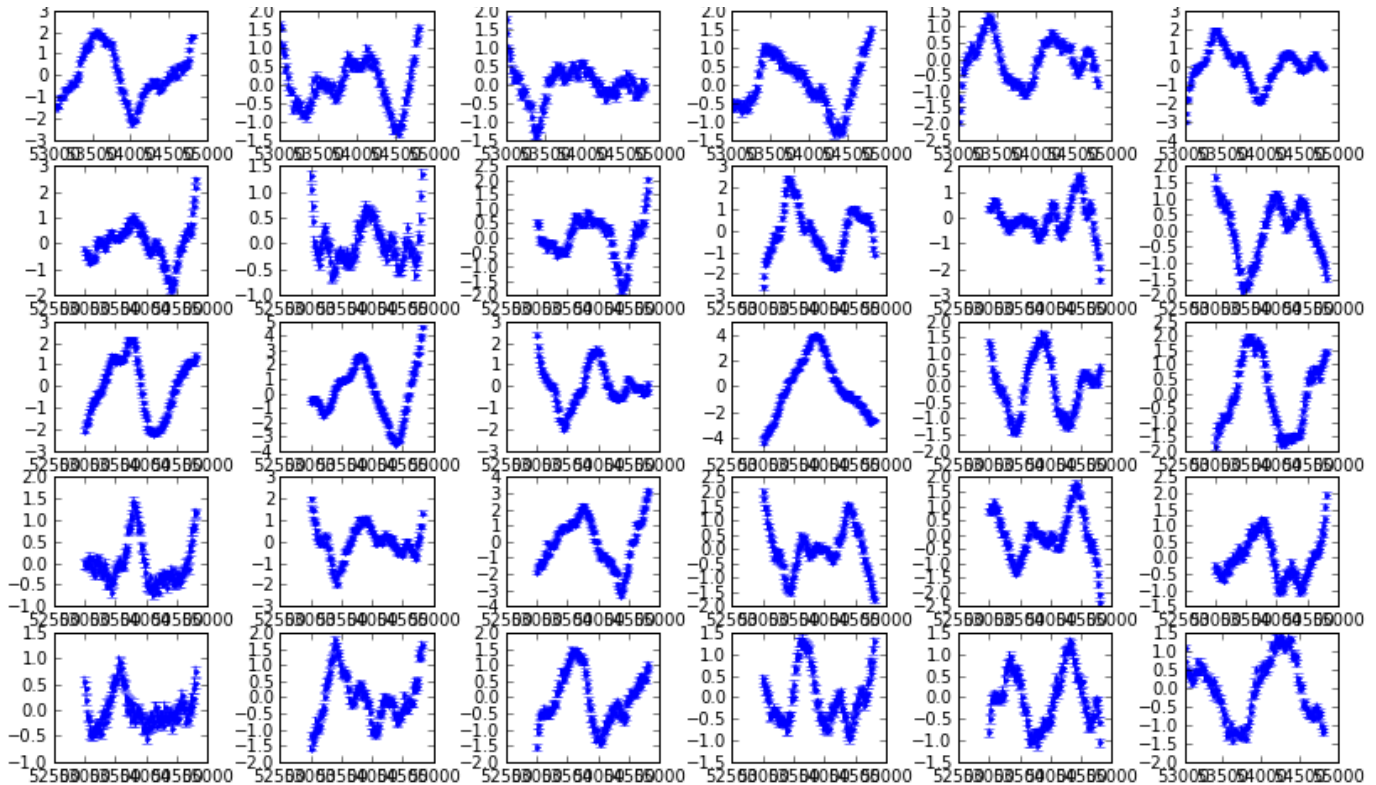
```

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```
In [21]: # this pulsar class works in a similar way but instead of using keys the datasets are just elements
figure(figsize=(11.5, 8))
for ct, p in enumerate(psr):
    plt.subplot(6, 6, ct+1)
    plt.errorbar(p.toas/86400, p.res*1e6, p.err*1e6, fmt='.')

plt.tight_layout(h pad=0.1, w pad=0.1)
```





Running Noise Estimation

```
In [22]: # at the moment I have a MultiNest and an MCMC implementation of the Hybrid Time-Frequency Method
!python testing/lentati_single_pulsar_noise.py -h
```

```
usage: lentati_single_pulsar_noise.py [-h] --h5File H5FILE [--outDir OUTDIR] --pulsar PNAME
                                     [--nmodes NModes] [--powerlaw] [--fc] [--broken] [--
single]
```

Run Lentati style noise estimation

optional arguments:

```
-h, --help            show this help message and exit
--h5File H5FILE       Full path to hdf5 file containing PTA data
--outDir OUTDIR       Full path to output directory (default = ./)
--pulsar PNAME        name of pulsar to use
--nmodes NModes       number of fourier modes to use (default=10)
--powerlaw            Use power law model (default = False)
--fc                  Use power law model with cross over frequency (default = False)
--broken              Use power law with two spectral indices and a cross over frequency (default =
False)
--single              Have one frequency and amplitude free to look for single frequency source
(default = False)
```

```
In [24]: # run one of the MDC pulsars as an example
!python testing/lentati_single_pulsar_noise.py --h5File ./mdc_data/open1_demo.hdf5 --outDir demo_
```

```
Reading in HDF5 file
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
Parameterizing Power spectrum coefficients by a power law
*****
MultiNest v3.0
```

```

*****
Copyright Farhan Feroz & Mike Hobson
Release Jun 2013

no. of live points = 500
dimensionality = 4
running in constant efficiency mode
*****
Starting MultiNest
generating live points
live points generated, starting sampling
Acceptance Rate: 0.994575
Replacements: 550
Total Samples: 553
Nested Sampling ln(Z): -440.906081

```

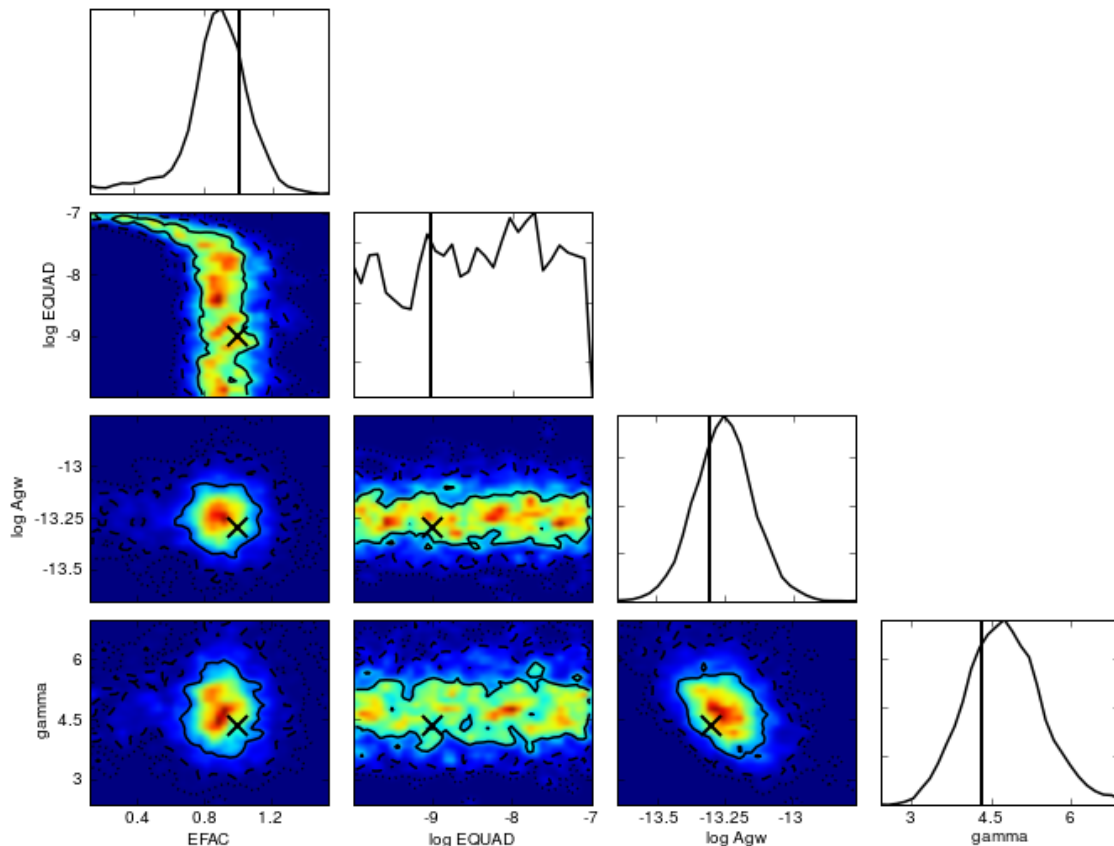
```

In [31]: # now take a look at the results
d = np.loadtxt('demo_noise_results/powerlaw/J0030+0451/testpost_equal_weights.dat')

# since this is the open MDC we know the "true" values [efac, log equad, log Agw, gw power spectral index]
inj = [1.0, -9, np.log10(5e-14), 4.333]
label = ['EFAC', 'log EQUAD', 'log Agw', 'gamma']

# call the triplot function from bayesutils to take a look at the 1 and 2 dimensional marginalized
# posteriors
bu.triplot(d[:, :-1], inj=inj, labels=label)

```



```

In [32]: # we can also do this with the model-independent method
!python testing/lentati_single_pulsar_noise.py --h5File ./mdc_data/open1_demo.hdf5 --outDir demo_

```

```

Reading in HDF5 file
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
Parameterizing Power spectrum coefficients by 10 independent coefficients

```



```

*****
MultiNest v3.0
Copyright Farhan Feroz & Mike Hobson
Release Jun 2013

no. of live points = 500
dimensionality = 12
running in constant efficiency mode
*****
Starting MultiNest
generating live points
live points generated, starting sampling
Acceptance Rate: 0.994575
Replacements: 550
Total Samples: 553
Nested Sampling ln(Z): 1044.474952

```

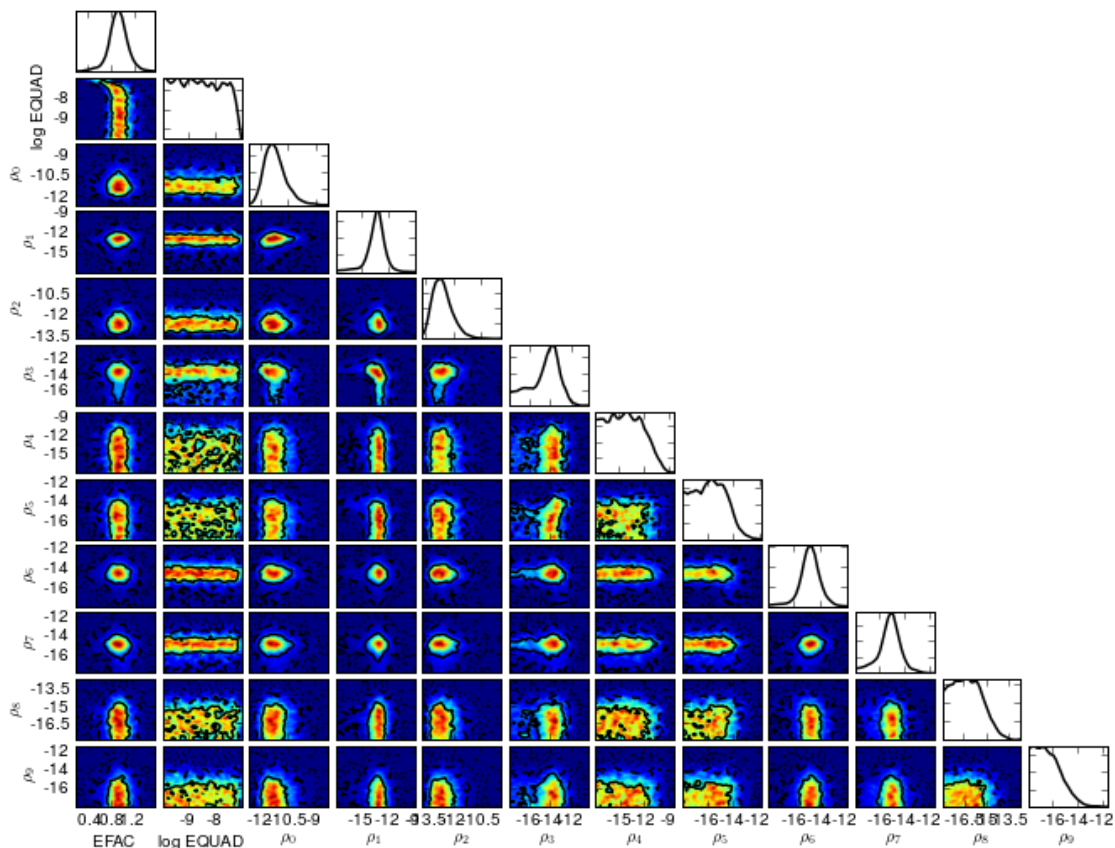
```

In [37]: # again we can make a triangle plot (it works for arbitrary size)
# now take a look at the results
d = np.loadtxt('demo_noise_results/independent/J0030+0451/testpost_equal_weights.dat')

# since this is the open MDC we know the "true" values
label = ['EFAC', 'log EQUAD']
for ii in range(d.shape[1]-3):
    label.append(r'$\rho_{%s}$'%(int(ii)))

bu.triplot(d[:,0:-1], labels=label)

```



```

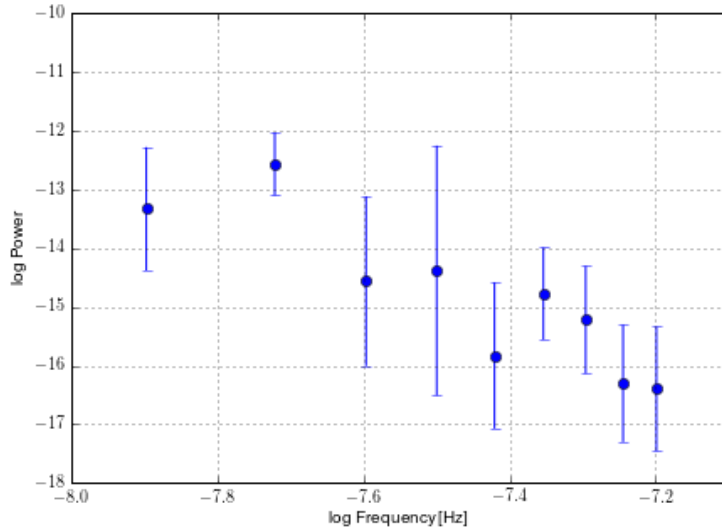
In [42]: # we can also plot the power spectrum
nmodes = 10
T = 5 * 3.16e7 # 5 year data span
f = np.linspace(1/T, nmodes/T, nmodes) # nyquist sampling

```

```
# log of power spectrum coefficients from output
lp = np.mean(d[:,2:-1], axis=0)
err = np.std(d[:,2:-1], axis=0)

# make plot
plt.figure(figsize=(7,5))
plt.errorbar(np.log10(f), lp, err, fmt='o')
plt.grid()
plt.xlim(-8.0, -7.1)
plt.xlabel('log Frequency [Hz]')
plt.ylabel('log Power')
```

Out[42]: <matplotlib.text.Text at 0x108f11ad0>



Getting Noise Values into HDF5 file

```
In [46]: # at the moment I don't have any nice scripts to make out noise files
# after we have done noise estimation we need to make nosise par files
# for each pulsar. Currently PAL only supports the powerlaw format but
# this will be changed soon as I'm going to change the inner workings
# of PALpulsarInit.

# an example of a noise par file is
!cat ./mdc_data/open_mdcl_noise_files/J0030+0451.noise

# here the amplitude is in GW units gamma is the power spectral index, equad is in seconds and ef
```

```
Pname J0030+0451
Amp 5e-14
gam 4.33
equad 0
efac 1
```

```
In [48]: # PALpulsarInit will recognize these keywords and store the noise values as well as the inverse c
# matrix term G (G^T C G)^-1 G^T for use in certain detection codes like the F-statistic

# for now we will need to delete our original hdf5 file in order to add the noise parameters
!rm ./mdc_data/open1_demo.hdf5

# run PALpulsarInit again
```



```
!python PALpulsarInit.py --parDir './mdc_data/partim/open1/' --timDir './mdc_data/partim/open1/'
J0030+0451.par J0030+0451.tim J0030+0451.noise
J0218+4232.par J0218+4232.tim J0218+4232.noise
J0437-4715.par J0437-4715.tim J0437-4715.noise
J0613-0200.par J0613-0200.tim J0613-0200.noise
J0621+1002.par J0621+1002.tim J0621+1002.noise
J0711-6830.par J0711-6830.tim J0711-6830.noise
J0751+1807.par J0751+1807.tim J0751+1807.noise
J0900-3144.par J0900-3144.tim J0900-3144.noise
J1012+5307.par J1012+5307.tim J1012+5307.noise
J1022+1001.par J1022+1001.tim J1022+1001.noise
J1024-0719.par J1024-0719.tim J1024-0719.noise
J1045-4509.par J1045-4509.tim J1045-4509.noise
J1455-3330.par J1455-3330.tim J1455-3330.noise
J1600-3053.par J1600-3053.tim J1600-3053.noise
J1603-7202.par J1603-7202.tim J1603-7202.noise
J1640+2224.par J1640+2224.tim J1640+2224.noise
J1643-1224.par J1643-1224.tim J1643-1224.noise
J1713+0747.par J1713+0747.tim J1713+0747.noise
J1730-2304.par J1730-2304.tim J1730-2304.noise
J1732-5049.par J1732-5049.tim J1732-5049.noise
```

Optimal Statistic and Cross Correlation Statistic

In [49]: *# run the help for the Optimal Statistic*

```
!python PALOptimalStatistic.py -h
```

Usage: PALOptimalStatistic.py [options]

Run the Optimal-statistic stochastic BG code as
Chamberlin, Creighton, Demorest et al (2013)

defined in

Options:

```
-h, --help            show this help message and exit
--h5File=H5FILE       Full path to hdf5 file containing PTA data
--outDir=OUTDIR        Full path to output directory (default = ./)
--spectralIndex=GAM    Power spectral index of stochastic background (default = 4.3333 (SMBHBs))
```

In [50]: *# run it on open MDC 1*

```
!python PALOptimalStatistic.py --h5File './mdc_data/open1_demo.hdf5'
```

Reading in HDF5 file

```
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
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WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
```

```
A_gw^2 = 2.41138648053e-27
std. dev. = 1.82299549806e-28
SNR = 13.2276052415
SNR of 13.2276052415 is above threshold!
```

```
In [54]: # run the help for the cross correlation function basically the same as the Optimal Statistic
!python PALCrossCorrelationStatistic.py -h

Usage: PALCrossCorrelationStatistic.py [options]

Run the Cross Correlation statistic defined in
Demorest et al. (2012)

Options:
  -h, --help                show this help message and exit
  --h5File=H5FILE           Full path to hdf5 file containing PTA data
  --outDir=OUTDIR           Full path to output directory (default = ./)
  --spectralIndex=GAM       Power spectral index of stochastic background (default = 4.3333 (SMBHBs))
```

[illegible]

```

WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
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WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
Computing Overlap Reduction Function Values
Running Cross correlation Statistic on 36 Pulsars
Results of Search

```

```

-----

A_gw^2 = 2.4113864814e-27
std. dev. = 1.82299549809e-28
Reduced Chi-squared = 0.933491011365
2-sigma upper limit based on chi-squared fit is A_gw < 5.20695934025e-14

```

```

In [57]: # get Agw
Agw = np.sqrt(2.4113864814e-27)    # same as Optimal Statistic!
print Agw

4.91058701318e-14

```

Simulating Data

```

In [59]: # run help for PALSimulation
!python PALSimulation.py -h

Usage: PALSimulation.py [options]

Simulate Fake Data (Under Construction)

Options:

```

```
In [63]: # first lets simulate a single source with SNR=15 and just white noise for the pulsars in open1_c
!python PALSimulation.py --h5File './mdc_data/open1_demo.hdf5' --single --snr 15 --gwchirpmass 2e
```

```
In [66]: # lets read the data into the pulsar class and take a look at the residuals
pfile = h5.File('./mdc_data/open1_demo_single.hdf5')

pulsargroup = pfile['Data']['Pulsars']

psr = [PALpulsarInit.pulsar(pulsargroup[key], addGmatrix=True, addNoise=True) for key in pulsargr

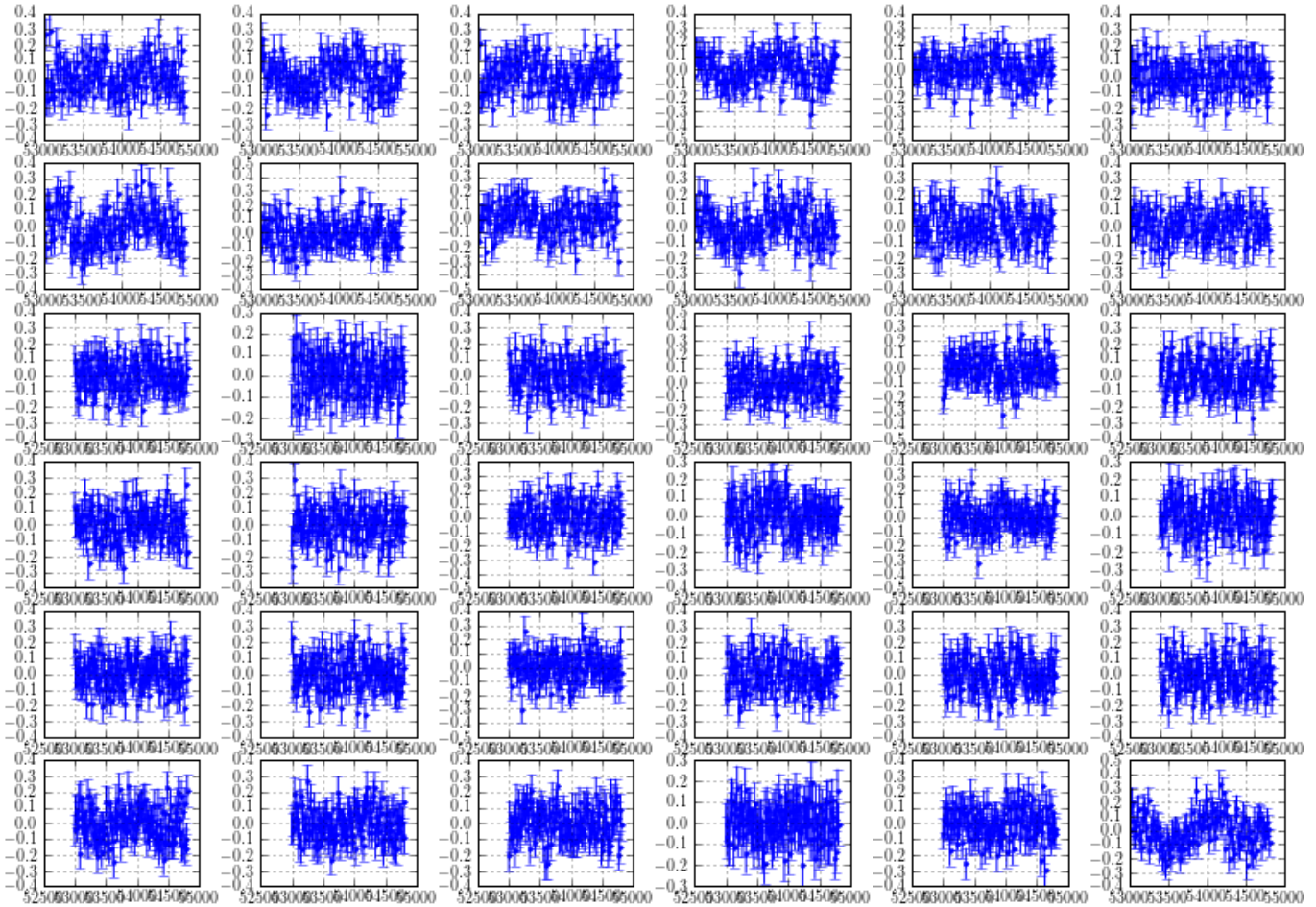
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
```

[illegible]


```
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
```

```
In [68]: # plot it
figure(figsize=(11.5, 8))
for ct, p in enumerate(psr):
    plt.subplot(6, 6, ct+1)
    plt.errorbar(p.toas/86400, p.res*1e6, p.err*1e6, fmt='. ')
    plt.grid()

plt.tight_layout(h_pad=0.1, w_pad=0.1)
```



```
In [70]: # now lets simulate a stochastic background with Agw = 5e-15 with white noise
!python PALSimulation.py --h5File './mdc_data/open1_demo.hdf5' --outFile './mdc_data/open1_demo_c
```

[illegible]

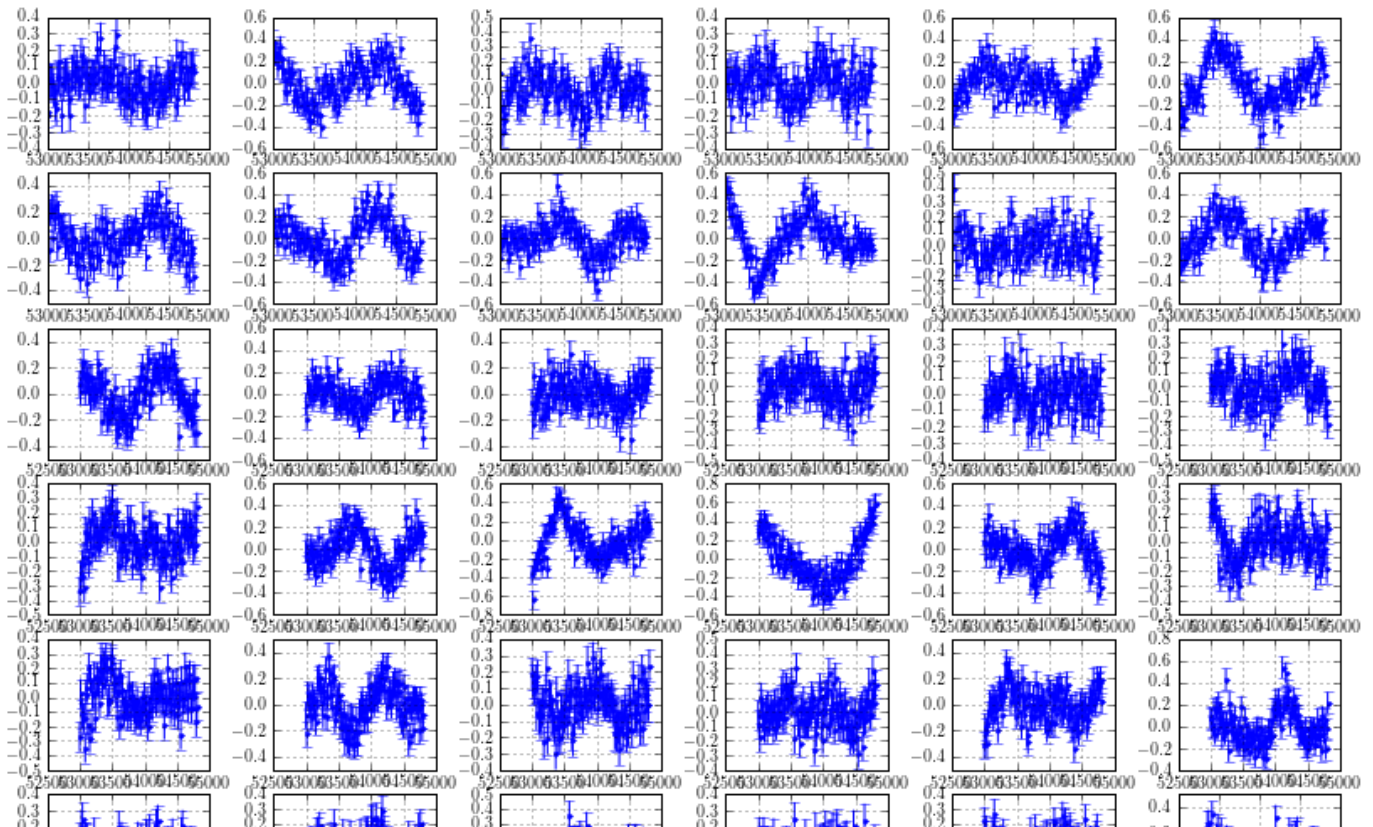
[illegible]

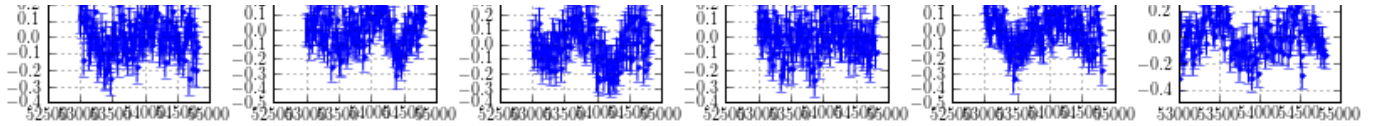
[illegible]

```
In [72]: # and plot it
```

```
figure(figsize=(11.5, 8))
for ct, p in enumerate(psr):
    plt.subplot(6, 6, ct+1)
    plt.errorbar(p.toas/86400, p.res*1e6, p.err*1e6, fmt='.')
    plt.grid()

plt.tight_layout(h_pad=0.1, w_pad=0.1)
```





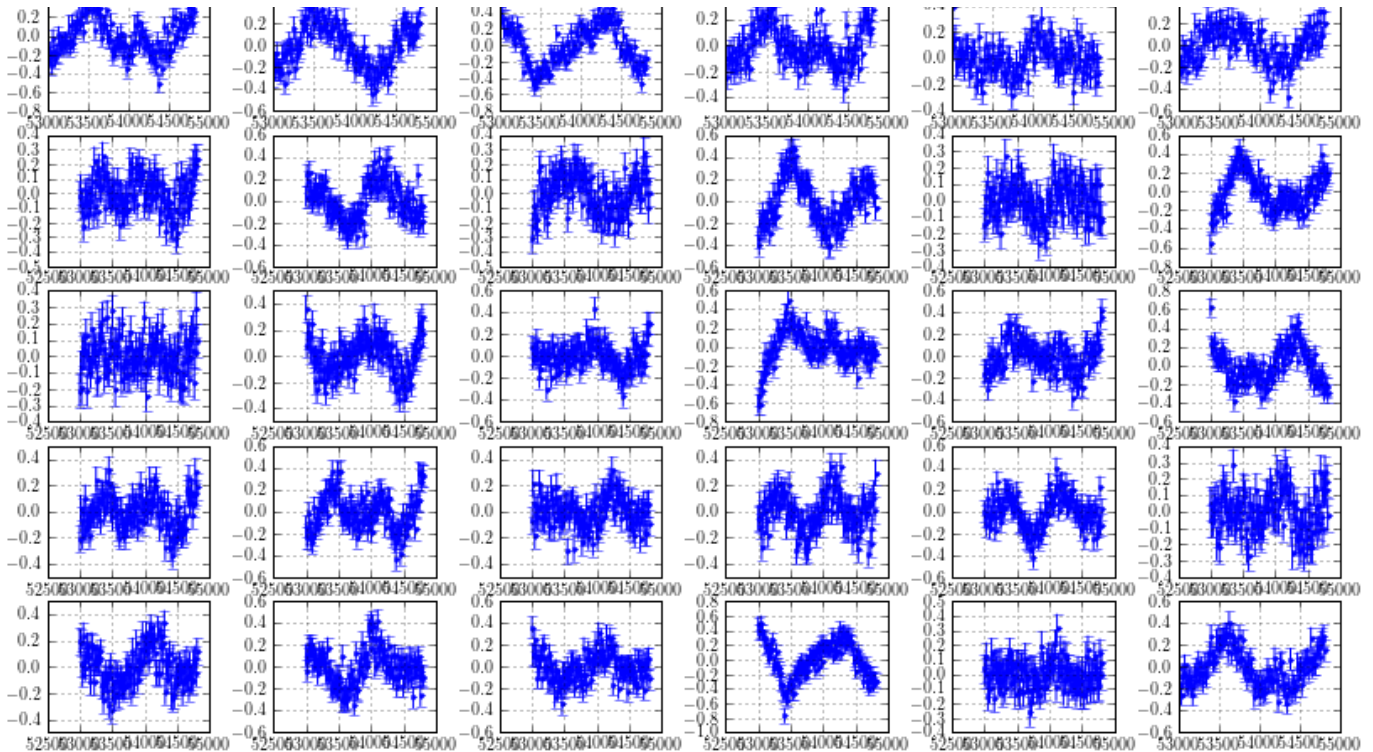
```

Saving file to ./mdc_data/open1_demo_both.hdf5
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
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WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc

```

[illegible]

```
In [75]: # and plot it
```

\mathcal{F}_p -Statistic

```
In [76]: # run the help function
!python PALFstatistic.py -h
```

Usage: PALFstatistic.py [options]

Run F-statistic search as defined in Ellis, Siemens, Creighton (2012)

Options:

-h, --help	show this help message and exit
--h5File=H5FILE	Full path to hdf5 file containing PTA data
--outDir=OUTDIR	Full path to output directory (default = ./)
--runFpStat	Option to run Incoherent Fp Statistic (default = True)
--runFeStat	Option to run Earth term Fe Statistic (default = False)
--fhigh=FHIGH	Highest frequency to search (default = 5e-7 Hz)
--nfreqs=NFREQS	Number of frequencies to search (default = 200)
--logsample	Sample in log frequency (default = False)
--best=BEST	Only use best pulsars based on weighted rms (default = 0, use all)

```
In [77]: # we will run the Fp statistic on our simulated single source dataset
# normally we will have to do a noise estimation but we know that there
# is only white noise and PALSimulation used white noise by default
!python PALFstatistic.py --h5File ./mdc_data/open1_demo_single.hdf5 --logsample
```

Reading in HDF5 file

```
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
WARNING: No distance error info, using sigma_d = 0.1 kpc
WARNING: No distance info, using d = 1 kpc
```

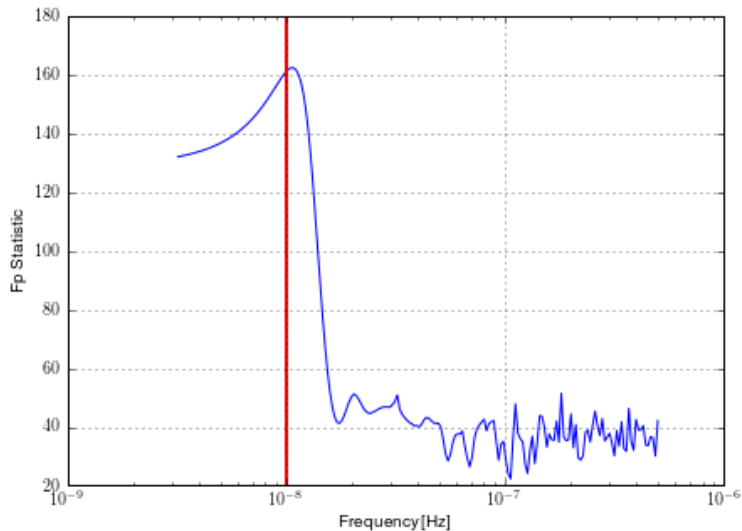
[illegible]

```
WARNING: No distance error info, using sigma_d = 0.1 kpc
Beginning Fp Search with 36 pulsars, with frequency range 3.20433591498e-09 -- 5e-07
Done Search. Computing False Alarm Probability
Writing results to file ../open1_demo_single.txt
```

```
In [79]: # read in the output file
d = np.loadtxt('open1_demo_single.txt')
```

```
In [87]: # plot the Fp statistic vs frequency
plt.figure(figsize=(7,5))
plt.semilogx(d[:,0], d[:,1])
plt.grid()
plt.xlabel('Frequency [Hz]')
plt.ylabel('Fp Statistic')
plt.axvline(1e-8, lw=2, color='r')
```

```
Out[87]: <matplotlib.lines.Line2D at 0x1092abd10>
```



MCMC Single Source Search for Non-Evolving Sources

```
In [91]: # run help function
!python testing/modelIndependent_time_domain_ss.py -h

usage: modelIndependent_time_domain_ss.py [-h] --h5File H5FILE [--outDir OUTDIR] [--best BEST]
                                           [--block BLOCK] [--scale SCALE] [--ntemps NTEMPS]
                                           [--nprocs NPROCS]
```

Run time domain single source MCMC

optional arguments:

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-h, --help            show this help message and exit
--h5File H5FILE       Full path to hdf5 file containing PTA data
--outDir OUTDIR       Full path to output directory (default = ./)
--best BEST           Only use best pulsars based on weighted rms (default = 0, use all)
--block BLOCK         How many parameters to update at each iteration (default = 0, use all)
--scale SCALE         Scale factor on jump covariance matrix (default = 1, native scaling)
--ntemps NTEMPS       Number of parallel temperature chains to run (default = 1)
--nprocs NPROCS       Number of processors to use with parallel tempering (default = 1)
```

```
In [*]: # run it
```


[illegible]