```
In []: import numpy as np
    from MCMC_aux import get_model
    import Parameters as par
    import Models as mod
    import Kernels as ker
    import GP_Likelihood as gp
    from MCMC import run_MCMC as run
    import plotting as plot
    from saving import save
    import auxiliary as aux
```

2813.2308006313287 16620.585819951895

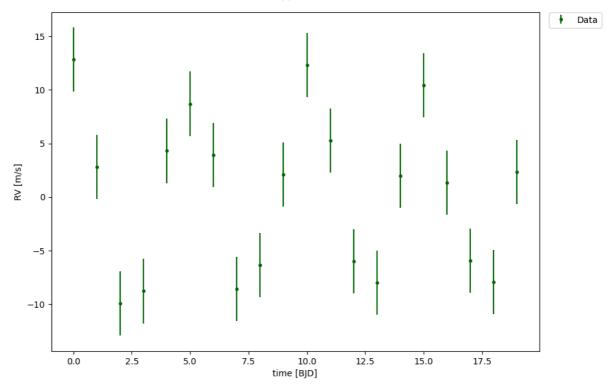
Creating fake data to work with

A cosine with a small jitter term is created as a fake set of data, this will act as the activity to model our kernel from

```
In []: # time array with 20 values
   time = np.arange(0,20,1)
   # set up the amplitude and period of the cosine
A = 10.
P = 5.
err = []
# set up a random jitter to add to the data
for i in time:
        err.append(np.random.uniform(-3,3))
# generate the rvs and errors
rv = A*np.cos(time*((2*np.pi)/P))+err
rv_err = np.ones_like(rv)*3
```

data_plot function will take the time, rv data, and rv errors and plot a scatter graph of the data. Axis labels, legend, and saving can all be controlled from the function inputs.

```
In [ ]: plot.data_plot(time = time, rv = rv, y_err = rv_err)
```



Creating the kernel

A kernel must be created using the par_create function, this will take only the name of the kernel and return an empty dictionary of hyperparameters to be filled out. This dictionary can be printed to view the hyperparamer names.

Currently available kernels along with their hyperparameter names can be viewed by running PrintKernelList:

Hyperparameters must then be assigned to each dictionary key using the parameter function. This function will also take errors and vary as inputs which will be used for the mcmc.

```
In []: # assign values to the dictionary
    hparam["gp_amp"] = par.parameter(value = 10.)
    hparam["gp_per"] = par.parameter(value = 5.)
    # printing now prints the filled dictionary
    print(hparam)
```

```
{'gp_amp': Parameter object: value = 10.0, error=2.0 (vary = True)
, 'gp_per': Parameter object: value = 5.0, error=1.0 (vary = True)
}
```

Priors should then be created by assigning the pri_create function to each prior and appeding it to a list of priors. The pri_create function takes the parameter name, the prior name, and the prior parameters as inputs which must be inputted in the correct form, this form can be viewed by running the PRINTPRIORDER function.

```
In [ ]: # view the correct form of prior parameter inputs
    par.PRINTPRIORDER()
```

Gaussian: List should take the form [mu, sigma] where all values are floats or ints

Jeffery: List should take the form [minval, maxval] where all values are flo ats or ints

Modified Jeffery: List should take the form [minval, maxval, kneeval] where all values are floats or ints

Uniform: List should take the form [minval, maxval] where all values are flo ats or ints

```
In []: # create empty prior list
    prior_list = []
    # uniform parameters used here so prior parameters inputted as [minval, maxv
    pri_amp = par.pri_create("gp_amp", "Uniform", [5.,15.])
    # then append the prior to the list
    prior_list.append(pri_amp)
    pri_per = par.pri_create("gp_per", "Uniform", [0.,10.])
    prior_list.append(pri_per)
    # print the list of all the priors
    print(prior_list)
```

```
[('gp_amp', 'Uniform', {'minval': 5.0, 'maxval': 15.0}), ('gp_per', 'Unifor
m', {'minval': 0.0, 'maxval': 10.0})]
```

Obtaining LogL and GP values

The GPLikelihood class should be defined and run with the time data, the rv data, the rv errors, the hyperparameters, and the kernel name. This allows the GPLikelihood.LogL function to be run with the prior_list which returns the initial log likelihood of the GP model.

In order to return the y values and errors of the GP model, a predicted x array must first be defined which should be smoother and longer than the initial time array, in this case it begins at -1 and ends at 21 with intervals of 0.1 which is around 10 times more data points than the initial time array. This must be then inputted into the GPLikelihood.predict function to return the y values and the errors of the GP.

```
In []: # GPLikelihood class called as loglik, run with the current inputs
loglik = gp.GPLikelihood(time, rv, rv_err, hparam, "Cosine")
# LogL obtained by running loglik.LogL with the prior_list as the only input
logL = loglik.LogL(prior_list)
# xpred is smoother and longer than time
xpred = np.arange(min(time)-1, max(time)+1, 0.1)
# GP_y and GP_err are arrays of the GP y values and errors of the same lengt
GP_y, GP_err = loglik.predict(xpred)
print('Initial Log Likelihood =', logL)
```

Initial Log Likelihood = -47.97958115218286

Plotting the GP

The GP y values could be manually plotted against xpred once obtained in the previous step however the GP_plot function allows for an alternative faster way of plotting. This function takes the time array, the rv data, the hyperparameters, the kernel name, and the rv errors and returns a plot of the data with the GP model plotted over it. Xpred, axis labels, residuals, legend, and saving can all be controlled by the function inputs.

