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#### HW 24-25

## **Imports**

```
import warnings

import jax
import jax.numpy as jnp
import numpy as np
import pandas as pd
# import PIL
# import scipy
# import sympy as sp
from matplotlib import pyplot as plt

warnings.filterwarnings("ignore")
```

### HW 24

In [2]: # In this problem, you will practice using automatic differentiation to construct the Fisher Information # Matrix of a non-linear system and using said matrix to make informed decisions about sampling.

Consider the model 
$$f(\vec{x}) = \begin{pmatrix} e^{-\theta_1 x_1} \tan \theta_2 x_2 \\ \cosh \theta_1 x_1 x_2 \end{pmatrix}$$

```
In [3]: # (a) Use automatic differentiation to create the Jacobian of this function. Note that your result
# will be a programmatic object, not something that you can write out by hand

def f_x(x, th):
    x: jnp.ndarray (2,)
    th: jnp.ndarray (2,)
    in: jnp.ndarray (2,)
    in: jnp.exp(-th[0]*x[0])*jnp.tan(th[1]*x[1]),
    jnp.cosh(th[0]*x[0])*x[1])
    return output

J = jax.jacrev(f_x, argnums=1)

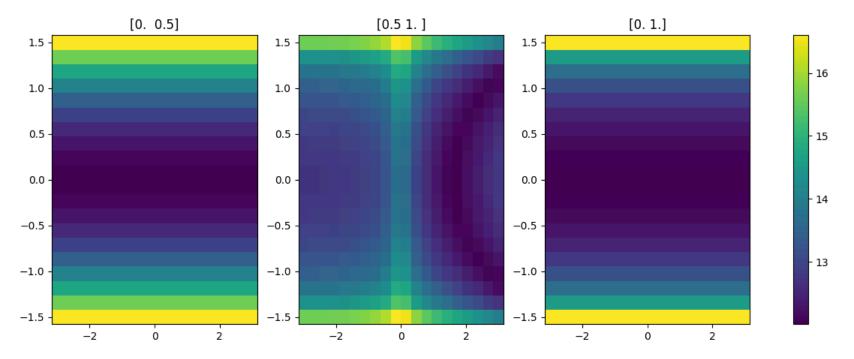
In [4]: # (b) The experiment that you want to test this model on has three possible sample points: 0, 0.5,
# and 1, but due to cost, you can only choose to sample at two points. Explore the behavior of
# your model by plotting the condition number of the Fisher Information Matrix as a function
```

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```
# of \theta. Use the following procedure:
# • Create three plots, one for each pair of possible sampling points: (0,0.5), (0.5,1), and (0,1)
# • Sample θ1 from (-3, 3)
# • Sample \theta2 from (-1.5, 1.5) to avoid singularities
# • Use a sparse grid to sample \thetai to keep computational time low (~ 20 samples from each \thetai)
# • Report the base-10 logarithm of the condition number
samples = jnp.array([[0, 0.5], [0.5, 1], [0, 1]])
th_1 = jnp.linspace(-3, 3, 20)
th_2 = jnp.linspace(-1.5, 1.5, 20)
fig, ax = plt.subplots(1, 3, figsize=(15,5))
for i, sample in enumerate(samples):
    ax[i].set_title(sample)
    m = jnp.zeros((3, 20, 20))
    for j, t1 in enumerate(th_1):
        for k, t2 in enumerate(th_2):
            ja = J(sample, jnp.array([t1, t2]))
            FIM = jnp.matmul(ja.T, ja)
            s = jnp.linalg.svd(FIM, compute_uv=False)
            s=s+1e-12 # from zulip
            cn = s[0]/s[-1]
            log_cn = jnp.log10(cn)
            m = m.at[:, j, k].set(jnp.array([t1, t2, log_cn]))
    pcm = ax[i].pcolormesh(*m)
fig.colorbar(pcm, ax=ax) # type: ignore
```

Out[4]: <matplotlib.colorbar.Colorbar at 0x118017b60>

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```
In [5]: # (c) What does a low condition number tell you about your model? Which of your plots has the
# greatest area with the lowest condition number? With that in mind, which set of sampling
# points is best for this model and why?

# A low condition number suggests that information in our model is evenly distributed across parameters,
# enabling more effective model fitting.
# The (0, 1) plot has the greatest area with the lowest condition number, so I recommend that set of sampling
# points because that some space will be easiest to traverse.
```

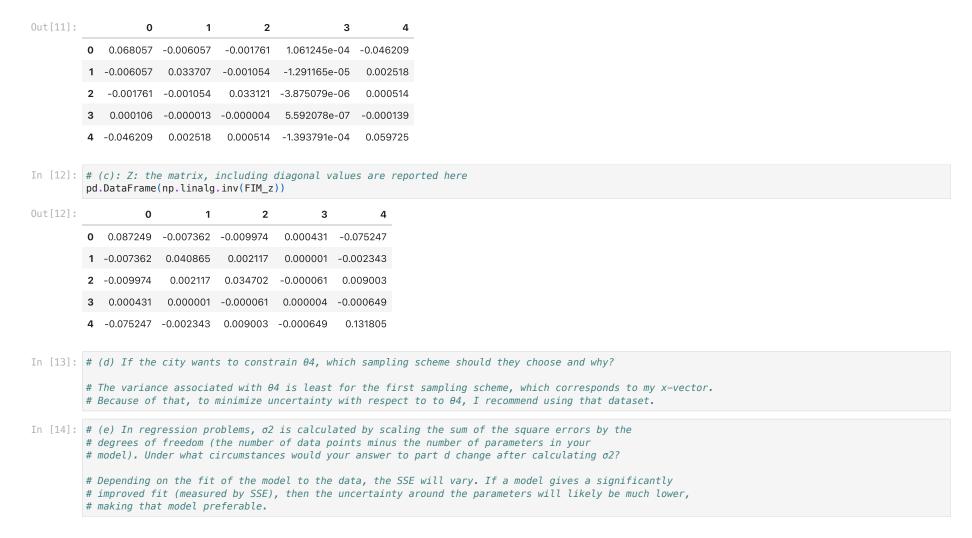
### **HW 25**

$$f(t_i) = \theta_1 \sin(\frac{2\pi}{365}t_i) + \theta_2 \sin(\frac{2\pi}{30}t_i) + \theta_3 \sin(\frac{2\pi}{7}t_i) + \theta_4 t_i + \theta_5$$

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```
In [7]: # (a) Construct the system matrix A for each set of points
         def design_mat(vec: np.ndarray) -> np.ndarray:
             """ adapted from HW 19 """
             m = np.array([
                 np.sin(2*np.pi*(1/365)*vec),
                 np.sin(2*np.pi*(1/30)*vec),
                 np.sin(2*np.pi*(1/7)*vec),
                 vec,
                 np.ones(shape=vec.shape)
             ]).T
             return m
         A_x = design_mat(x)
         A_y = design_mat(y)
         A_z = design_mat(z)
In [8]: # (b) Calculate the Fisher Information Matrix using the equation AT A for each set of points (Note
         # that we have not yet generated a fit, so we are setting \sigma-2 to 1 for simplicity's sake).
         FIM_x = np.matmul(A_x.T, A_x)
         FIM_y = np.matmul(A_y.T, A_y)
         FIM_z = np.matmul(A_z.T, A_z)
In [9]: # (c) Invert the Fisher Information Matrix to yield the covariance matrix. Report the diagonal
         # values for each covariance matrix, which are the uncertainties associated with each \theta value.
In [10]: # (c): X: the matrix, including diagonal values are reported here
         pd.DataFrame(np.linalg.inv(FIM_x))
Out[10]:
                                                     3
         0 0.039593 -0.000276 -0.000714 4.917626e-05 -0.017949
         1 -0.000276  0.033906  -0.000719  -1.916879e-06  0.000333
         2 -0.000714 -0.000719 0.033451 -5.016914e-06 0.001948
         3 0.000049 -0.000002 -0.000005 4.246699e-07 -0.000155
         4 -0.017949 0.000333 0.001948 -1.550002e-04 0.073245
In [11]: # (c): Y: the matrix, including diagonal values are reported here
         pd.DataFrame(np.linalg.inv(FIM_y))
```

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# Acknowledgment

Work in this repository and with associated assignments and projects may be adapted or copied from similar files used in my prior academic and industry work (e.g., using a LaTeX file or Dockerfile as a starting point). Those files and any other work in this repository may have been developed with the help of LLM's like ChatGPT. For example, to provide context, answer questions, refine writing, understand function call syntax, and assist with repetitive tasks. In these cases, deliverables and associated work reflect my best efforts to optimize my learning and demonstrate my capacity, while using available resources and LLM's to facilitate the process.

**ChatGPT Conversation**