## **HW 29**

```
In [1]: # 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
from scipy.optimize import minimize
```

The file aer.txt contains data from the analysis of an enzyme reaction. Enzymes catalyze the rate of chemical reactions. In this data set, the dependent variable is the rate of the chemical reaction and the independent variable is the amount of enzyme.

The data should be fit to the four parameter model

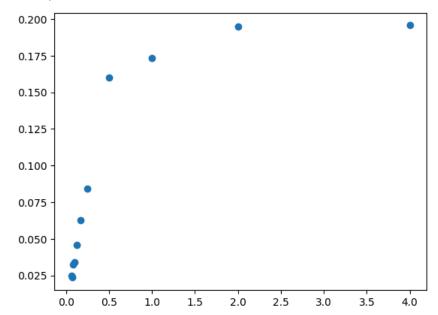
$$y_{\phi}(u) = \frac{\phi_1(u^2 + \phi_2 u)}{u^2 + \phi_3 u + \phi_4}.$$

The parameters should each satisfy  $\phi_i > 0$ , so we take as our parameter vector:

$$\theta = (\log \phi_1, \log \phi_2, \log \phi_3, \log \phi_4)'.$$

```
In [2]: data: pd.DataFrame = pd.read_csv("aer.txt", delimiter=r'\s+', header=None).values # type: ignore
enzy: np.ndarray = data[:, 0] # type: ignore
rate: np.ndarray = data[:, 1] # type: ignore
plt.scatter(enzy, rate)
```

Out[2]: <matplotlib.collections.PathCollection at 0x1192c5010>



```
In [3]:

def f(theta: np.ndarray, u: float):
    assert isinstance(theta, np.ndarray) and len(theta.shape)==1 and theta.shape[0]==4
    assert isinstance(u, float)
    th1, th2, th3, th4 = theta
    numerator = th1*(u**2+th2*u)
    denominator = u**2+th3*u+th4
    return numerator/denominator

def f_vec(theta: np.ndarray, u: np.ndarray):
    return np.array([f(theta, v) for v in u])
```

# NOTE: theta\_log = np.array([np.log(th) for th in theta])

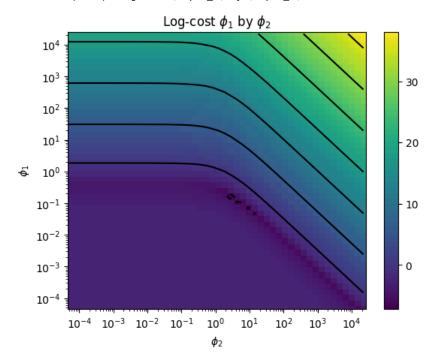
```
In [4]: # a. Which parameters could go to zero without resulting in a trivial model? Rewrite the model
        # for each of these cases.
        # thetal*theta2 is identifiable but theta1/theta2 is unidentifiable: theta2 is
        # similarly unidentifiable
        # All parameters except for thetal could go to zero without resulting in a trivial model.
        # In each case, associated values simplify to zero
        # drop th2
        # drop th3
        # drop th4
        # drop th2 & th3
        # drop th2 & th4
        # drop th3 & th4
        # drop th2 & th3 & th4
        # drop th2
        def f_drop2(theta: np.ndarray, u: float):
            th1, _{-}, th3, th4 = theta
            numerator = th1*u**2
            denominator = u**2+th3*u+th4
            return numerator/denominator
        # drop th3
        def f_drop3(theta: np.ndarray, u: float):
            th1, th2, \_, th4 = theta
            numerator = th1*(u**2+th2*u)
            denominator = u**2+th4
            return numerator/denominator
        # drop th4
        def f_drop4(theta: np.ndarray, u: float):
            th1, th2, th3, _{-} = theta
            numerator = th1*(u**2+th2*u)
            denominator = u**2+th3*u
            return numerator/denominator
        # drop th2 & th3
        def f_drop23(theta: np.ndarray, u: float):
            th1, _{-}, _{-}, th4 = theta
            numerator = th1*u**2
            denominator = u**2+th4
            return numerator/denominator
        # drop th2 & th4
        def f_drop24(theta: np.ndarray, u: float):
            th1, _, th3, _ = theta
            numerator = th1*u**2
            denominator = u**2+th3*u
            return numerator/denominator
        # drop th3 & th4
        def f_drop34(theta: np.ndarray, u: float):
            th1, th2, _, _ = theta
            numerator = th1*(u**2+th2*u)
            denominator = u**2
            return numerator/denominator
        # drop th2 & th3 & th4
        def f_drop234(theta: np.ndarray, u: float):
            th1, *_ = theta
            numerator = th1*u**2
            denominator = u**2
            return numerator/denominator
        simplified_callables_a = [f_drop2, f_drop3, f_drop4, f_drop23, f_drop24, f_drop34, f_drop234]
```

In [5]: # b. Consider what happens when  $\phi 2$  is much larger than  $\phi 1$ . What would the numerator simplify # to? Rewrite the model with this simplifying assumption.

```
# when \phi 2 is much larger than \phi 1, the numerator simplifies to th1*th2*u,
        # as if \phi1 is negligible/zero but not zero
        def f_2morethan1(theta: np.ndarray, u: float):
             th1, th2, th3, th4 = theta
             numerator = th1*th2*u
             denominator = u**2+th3*u+th4
             return numerator/denominator
        simplified_callables_b = [f_2morethan1]
In [6]: # c. Repeat part b for \phi3 and \phi4. Why did we ignore \phi1?
        # when \phi 3 is much larger than \phi 4, the denominator simplifies
        # as if \phi 4 is negligible/zero but not zero
        def f_3morethan4(theta: np.ndarray, u: float):
             th1, th2, th3, _{-} = theta
             numerator = th1*(u**2+th2*u)
             denominator = u**2+th3*u
             return numerator/denominator
        # when \phi 4 is much larger than \phi 3, the denominator simplifies
        # as if φ3 is negligible/zero but not zero
        def f_4morethan3(theta: np.ndarray, u: float):
             th1, th2, th3, th4 = theta
             numerator = th1*(u**2+th2*u)
             denominator = u**2+th3*u+th4
             return numerator/denominator
        # we ignore \phi1 because it is not directly related to \phi3 or \phi4--it would not be interesting
        # to do that comparison
        simplified_callables_c = [f_3morethan4, f_4morethan3]
In [7]: # save my functions
        simplified_callables = simplified_callables_a
        simplified_callables.extend(simplified_callables_b)
        simplified_callables.extend(simplified_callables_c)
In [8]: # d. For the full model, construct a cost surface running from -10 \le \log(\theta i) \le 10. Produce six
        # plots which slow slices of the cost surface with respect to pairs of the model parameters.
        def c(theta, enzy, rate):
             x_{hat} = f_{vec}(theta, enzy)
             diff = x_hat-rate
             norm_d = np.linalg.norm(diff, ord=2)
             return 0.5 * norm_d**2
        # our cost surface range is equivalent to e^-10 \le \theta i \le e^10
        log_theta_range = np.linspace(-10, 10, 50)
        theta_range = np.exp(log_theta_range)
        meshg = np.meshgrid(theta_range, theta_range)
        c_{surf_values} = np.zeros((50, 50, 50, 50))
        for i, th1 in enumerate(theta_range):
             for j, th2 in enumerate(theta_range):
                 for k, th3 in enumerate(theta_range):
                     for l, th4 in enumerate(theta_range):
                         theta = np.array([th1, th2, th3, th4]).flatten()
                         c_surf_values[i, j, k, l] = c(theta, enzy, rate)
        log_c_tensor = np.log(c_surf_values)
        # this is a 4-D tensor, so let's visualize slices
In [9]: # th1 x th2
        plt.contour(*meshg, log_c_tensor[:, :, 25, 25], colors="black")
        plt.pcolormesh(*meshg, log_c_tensor[:, :, 25, 25])
        plt.colorbar()
        plt.xscale("log")
        plt.yscale("log")
        plt.ylabel(r'$\phi_1$')
```

```
plt.xlabel(r'$\phi_2$')
plt.title(f"Log-cost {r'$\phi_1$'} by {r'$\phi_2$'}")
```

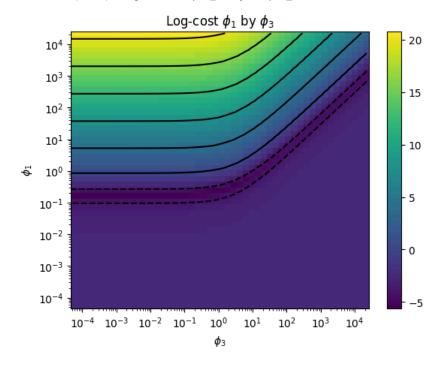
Out[9]: Text(0.5, 1.0, 'Log-cost \$\\phi\_1\$ by \$\\phi\_2\$')



```
In [10]: # th1 x th3

plt.contour(*meshg, log_c_tensor[:, 25, :, 25], colors="black")
plt.pcolormesh(*meshg, log_c_tensor[:, 25, :, 25])
plt.colorbar()
plt.xscale("log")
plt.yscale("log")
plt.ylabel(r'$\phi_1$')
plt.xlabel(r'$\phi_3$')
plt.xlabel(r'$\phi_3$')
plt.title(f"Log-cost {r'$\phi_1$'} by {r'$\phi_3$'}")
```

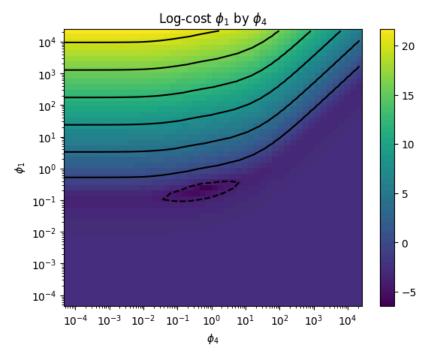
Out[10]: Text(0.5, 1.0, 'Log-cost \$\\phi\_1\$ by \$\\phi\_3\$')



```
In [11]: # th1 x th4

plt.contour(*meshg, log_c_tensor[:, 25, 25, :], colors="black")
plt.pcolormesh(*meshg, log_c_tensor[:, 25, 25, :])
plt.colorbar()
plt.xscale("log")
plt.yscale("log")
plt.yscale("log")
plt.ylabel(r'$\phi_1$')
plt.xlabel(r'$\phi_4$')
plt.title(f"Log_cost {r'$\phi_1$'} by {r'$\phi_4$'}")
```

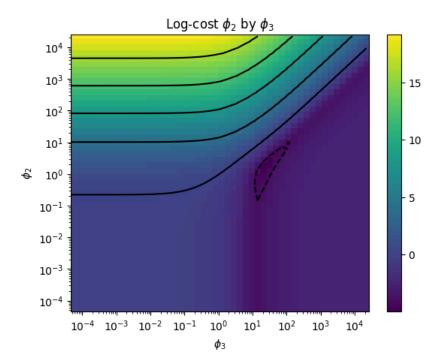
Out[11]: Text(0.5, 1.0, 'Log-cost \$\\phi\_1\$ by \$\\phi\_4\$')



```
In [12]: # th2 x th3

plt.contour(*meshg, log_c_tensor[25, :, :, 25], colors="black")
plt.pcolormesh(*meshg, log_c_tensor[25, :, :, 25])
plt.colorbar()
plt.xscale("log")
plt.yscale("log")
plt.ylabel(r'$\phi_2$')
plt.xlabel(r'$\phi_3$')
plt.xlabel(r'$\phi_3$')
plt.title(f"Log-cost {r'$\phi_2$'} by {r'$\phi_3$'}")
```

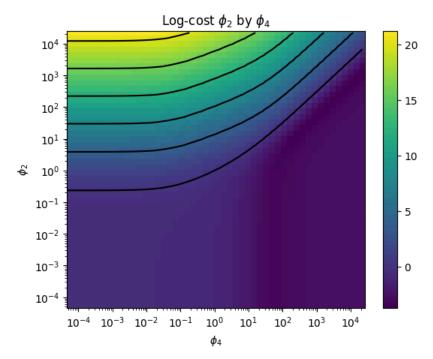
Out[12]: Text(0.5, 1.0, 'Log-cost \$\\phi\_2\$ by \$\\phi\_3\$')



```
In [13]: # th2 x th4

plt.contour(*meshg, log_c_tensor[25, :, 25, :], colors="black")
plt.pcolormesh(*meshg, log_c_tensor[25, :, 25, :])
plt.colorbar()
plt.xscale("log")
plt.yscale("log")
plt.yscale("log")
plt.ylabel(r'$\phi_2$')
plt.xlabel(r'$\phi_4$')
plt.title(f"Log-cost {r'$\phi_2$'} by {r'$\phi_4$'}")
```

Out[13]: Text(0.5, 1.0, 'Log-cost  $\phi_2$  by  $\phi_4$ ')

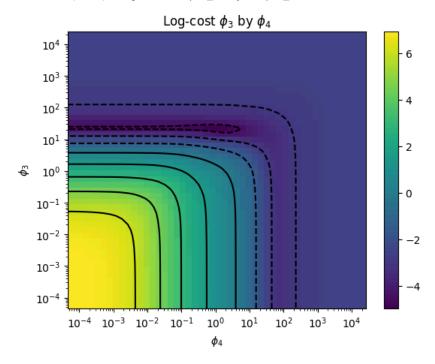


```
In [14]: # th3 x th4

plt.contour(*meshg, log_c_tensor[25, 25, :, :], colors="black")
plt.pcolormesh(*meshg, log_c_tensor[25, 25, :, :])
plt.colorbar()
plt.xscale("log")
```

```
plt.yscale("log")
plt.ylabel(r'$\phi_3$')
plt.xlabel(r'$\phi_4$')
plt.title(f"Log-cost {r'$\phi_3$'} by {r'$\phi_4$'}")
```

Out[14]: Text(0.5, 1.0, 'Log-cost \$\\phi\_3\$ by \$\\phi\_4\$')



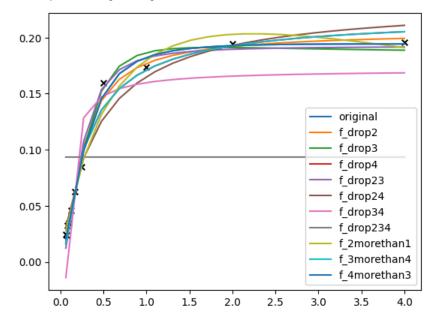
In [15]: # e. Identify regions on your cost surface plots which correspond to your simplified models

```
# simplified models:
          # drop th2 (lower left, th1-v-th2)
         # drop th3 (lower left, th1-v-th3; lower left, th2-v-th3)
# drop th4 (lower left, th1-v-th4)
         # drop th2 & th3 (bottom, th2-v-th3)
         # drop th2 & th4 (bottom, th2-v-th4)
          # drop th3 & th4 (bottom left, th3-v-th4)
          # drop th2 & th3 & th4 (bottom left of all plots except the last)
         \# \phi 2 >> \phi 1, \phi 1 -> 0 (lower left, th1-v-th2; bottom, th1-v-th3; bottom, th1-v-th4)
          # \phi 3 >> \phi 4, \phi 4 -> 0 (top left, th3-v-th4)
          # \phi 4 >> \phi 3, \phi 3 -> 0 (right, th3-v-th4)
In [16]: # f. Fit the full model to data.(Hint: At the best fit, I find that \phi 1 \approx 0.2.)
          fit = minimize(c, np.array([1, 1, 1, 1]), args=(enzy, rate))
          fit.x # this matches \phi1 \approx 0.2
Out[16]: array([0.19277051, 0.1922218 , 0.12335622, 0.13646131])
In [17]: # g. Repeat Part f for each of your simplified models.
          def custom_c(theta, enzy, rate, call):
              x_hat = np.array([call(theta, en) for en in enzy])
              diff = x_hat-rate
              norm_d = np.linalg.norm(diff, ord=2)
              return 0.5 * norm_d**2
          fits = {}
          for call in simplified_callables:
              simple_fit = minimize(custom_c, np.array([1, 1, 1, 1]), args=(enzy, rate, call))
              fits[call.__name__] = simple_fit.x
              print(f"For the simplified model '{call.__name__}', the fit parameters are {simple_fit.x}")
```

```
For the simplified model 'f_drop2', the fit parameters are [0.20603113 1.
                                                                                              0.12575824 0.0382867 1
        For the simplified model 'f_drop3', the fit parameters are [0.18305676 0.1607094 1.
                                                                                                          0.138236911
        For the simplified model 'f_drop4', the fit parameters are [ 0.21919701 -0.04144616 0.22761978 1.
        For the simplified model 'f_drop23', the fit parameters are [0.19230461 1. For the simplified model 'f_drop24', the fit parameters are [0.23237847 1.
                                                                                               0.40714765 1.
        For the simplified model 'f_{\rm d}rop34', the fit parameters are [ 0.1714469 \, -0.06759888 \, 1.
                                                                                                               1.
        For the simplified model 'f_drop234', the fit parameters are [0.09374545 1.
                                                                                                1.
        For the simplified model 'f_2morethan1', the fit parameters are [1.5582906 1.5582906 7.45169457 5.0266313
        For the simplified model 'f 3morethan4', the fit parameters are [ 0.21919701 -0.04144616 0.22761978 1.
        For the simplified model 'f_4morethan3', the fit parameters are [0.19277051 0.1922218 0.12335622 0.1364613
        1]
In [18]: # h. Plot all of your models on the same set of axes with the data. What does this plot tell you
         # about the identifiability of this model+parameter combination?
         x_plotting = np.linspace(min(enzy), max(enzy), 20)
         y_plotting = np.array([f(fit.x, en) for en in x_plotting])
         plt.plot(x_plotting, y_plotting, label="original")
         for call in simplified_callables:
             y_plotting = np.array([call(fits[call.__name__], en) for en in x_plotting])
             plt.plot(x_plotting, y_plotting, label=call.__name__)
         plt.scatter(enzy, rate, color='black', marker="x", s=30)
         plt.legend()
         # Regarding identifiability of this model+parameter combination, this plot shows that
         # there is a lot of signal aliased between parameters such that most can be dropped
         # or reduced to a negligible quantity without significant effect on the fit. In other
         # words, some parameters are unidentifiable. There are a couple exceptions like
         # dropping both parameters in the denominator, or dropping all parameters but thetal,
```

Out[18]: <matplotlib.legend.Legend at 0x1192c7a10>

# but otherwise all other models fit relatively well.



## 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