

Rare-event simulation: Code demo PyMC3

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```
[1]: import numpy as np
import pandas as pd
import pymc3 as pm

%config InlineBackend.figure_format = 'retina'
import matplotlib.pyplot as plt
import seaborn as sns
sns.set()
```

```
[2]: import sys
print("Python version:", sys.version)
print("Numpy version:", np.__version__)
print("PyMC3 version:", pm.__version__)
```

```
Python version: 3.7.7 (default, Mar 23 2020, 23:19:08) [MSC v.1916 64 bit
(AMD64)]
Numpy version: 1.18.1
PyMC3 version: 3.8
```

```
[3]: df = pd.read_csv("intervals.csv")
```

```
[4]: df.head()
```

```
[4]:
```

| | EL | ER | SL | SR |
|---|----|-----------|------|-----------|
| 0 | 0 | 45.999306 | 49.0 | 49.999306 |
| 1 | 14 | 43.999306 | 44.0 | 54.999306 |
| 2 | 39 | 49.375000 | 53.0 | 53.500000 |
| 3 | 0 | 35.999306 | 35.0 | 35.999306 |
| 4 | 0 | 43.999306 | 0.0 | 43.999306 |

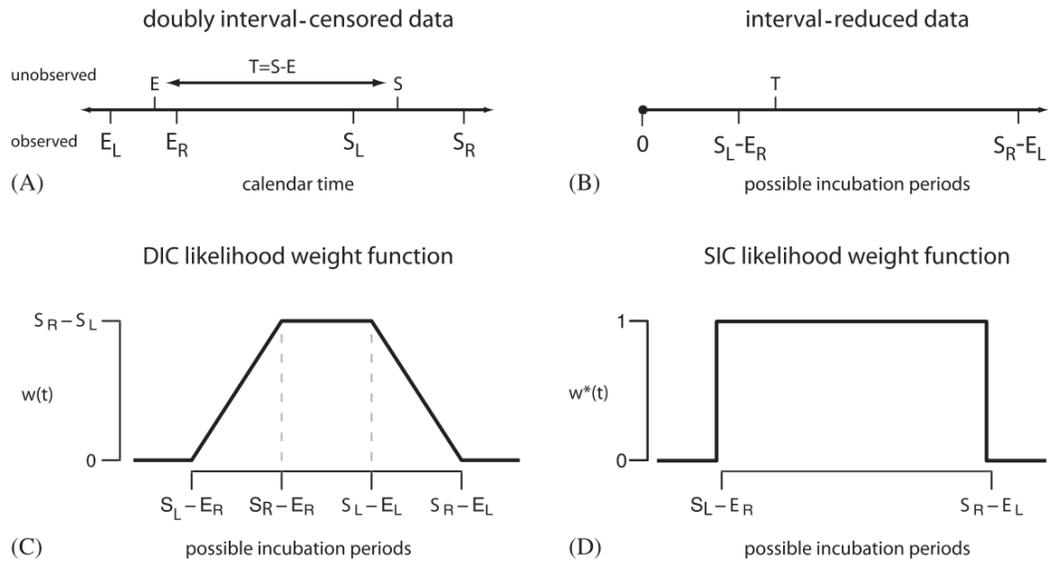
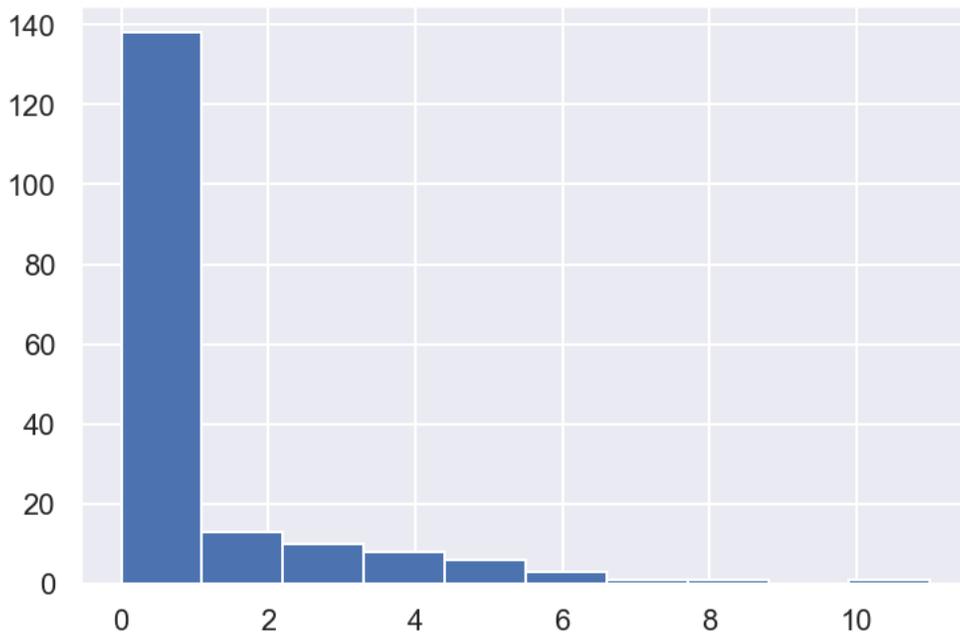


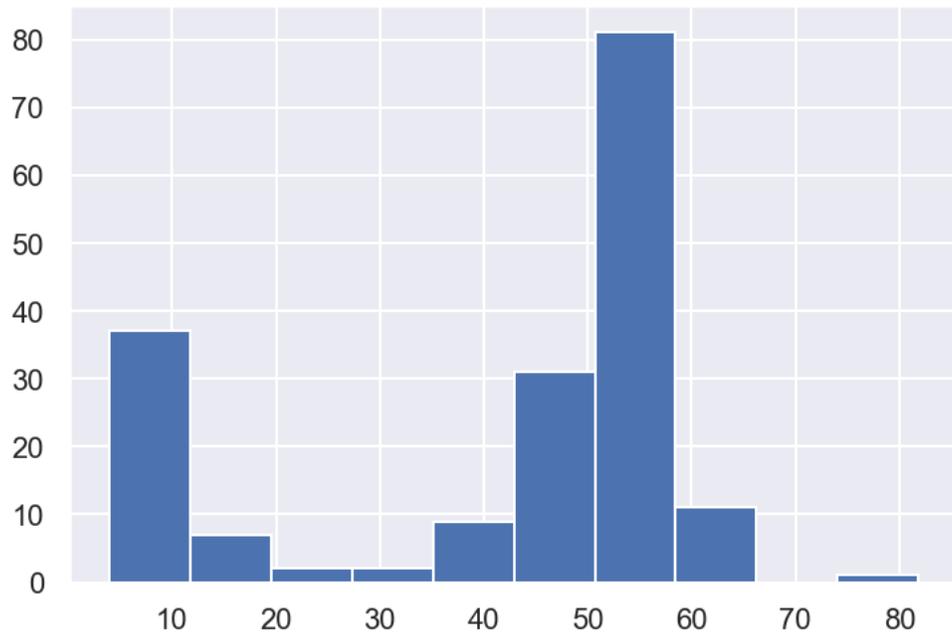
Figure 1 from Reich et al. (2009), *Estimating incubation period distributions with coarse data*.

```
[5]: Tmin = np.array(np.maximum(df["SL"]-df["ER"], 0))
      Tmax = np.array(df["SR"]-df["EL"])
```

```
[6]: plt.hist(Tmin);
```



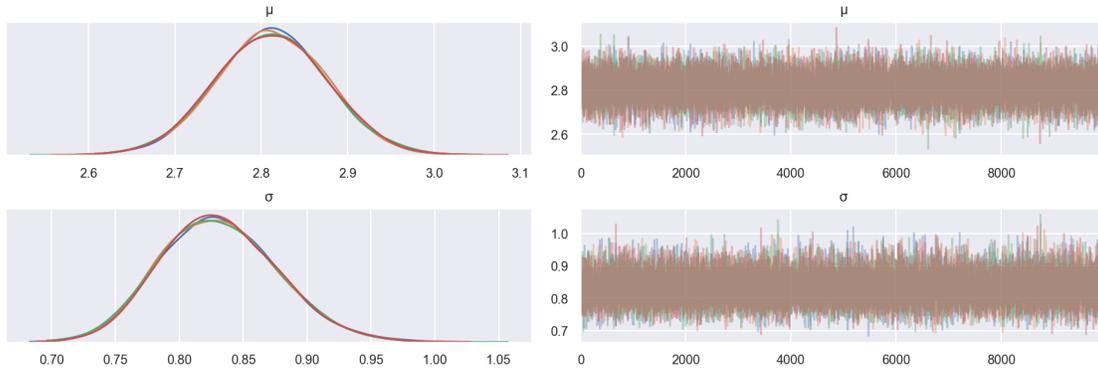
```
[7]: plt.hist(Tmax);
```



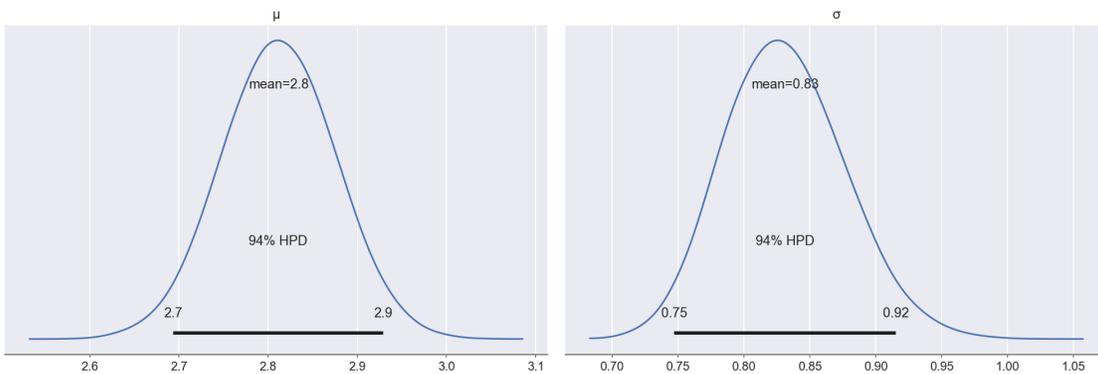
```
[8]: xs = np.linspace(0, 75, 500)
     ys = np.zeros(len(xs))

     nCases = len(Tmin)
     for i in range(nCases):
         ys[(xs >= Tmin[i]) & (xs <= Tmax[i])] += 1 / (nCases *
         ↪(Tmax[i]-Tmin[i]))

     plt.plot(xs, ys);
```

```
[12]: pm.plot_posterior(trace);
```



This is assuming we have observations $\mathbf{T} = (T_1, \dots, T_n)$ which gives us a likelihood of

$$L(\mu, \sigma | \mathbf{T}) = \prod_{i=1}^n f(T_i; \mu, \sigma)$$

where $f(x; \mu, \sigma)$ is the p.d.f. of the $\text{LogNormal}(\mu, \sigma^2)$ distribution.

However we don't have observations, just intervals. Say each unobserved period fell into $T_i \in [T_i^-, T_i^+]$. Our likelihood becomes

$$L(\mu, \sigma | \mathbf{T}^-, \mathbf{T}^+) = \prod_{i=1}^n [F(T_i^+; \mu, \sigma) - F(T_i^-; \mu, \sigma)]$$

where $F(x; \mu, \sigma)$ is the c.d.f. of the $\text{LogNormal}(\mu, \sigma^2)$ distribution.

```
[13]: import theano.tensor as tt
```

```
# Taken from PyMC3's pymc3/distributions/dist_math.py file
```

```

# starting at line 346.
def zvalue(x, sigma, mu):
    """
    Calculate the z-value for a normal distribution.
    """
    return (x - mu) / sigma

# Taken from PyMC3's pymc3/distributions/continuous.py file
# starting at line 1849.
def cdf(x, mu, sigma):
    """
    Compute the log of the cumulative distribution function for
    ↪Lognormal distribution
    at the specified value.

    Parameters
    -----
    x: numeric
        Value(s) for which log CDF is calculated. If the log CDF for
    ↪multiple
        values are desired the values must be provided in a numpy array
    ↪or theano tensor.

    Returns
    -----
    TensorVariable
    """
    z = zvalue(np.log(x), mu=mu, sigma=sigma)

    return tt.switch(
        tt.lt(z, -1.0),
        tt.erfcx(-z / tt.sqrt(2.)) / 2. * np.exp(-tt.sqr(z) / 2),
        tt.erfc(-z / tt.sqrt(2.)) / 2.
    )

```

With **Potential** we have to add log-terms to the likelihood. So

$$\log[L(\mu, \sigma | \mathbf{T}^-, \mathbf{T}^+)] = \sum_{i=1}^n \log[F(T_i^+; \mu, \sigma) - F(T_i^-; \mu, \sigma)].$$

```

[14]: %%time
with pm.Model() as model:
    mu = pm.Uniform('μ', lower=-25, upper=25)
    sigma = pm.Uniform('σ', lower=0, upper=25)
    pm.Potential('T', tt.sum(tt.log( cdf(Tmax, mu, sigma) - cdf(Tmin, mu, sigma) )))
    trace = pm.sample(10**5, step=pm.Metropolis())

```


[18]: 1.6333400315715902

[19]: `trace[" σ "].mean()`

[19]: 0.3841109415517976