## Lecture 11 Demo: Posterior predictive checks

```
In [5]: %matplotlib inline
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

import pymc3 as pm
from pymc3 import glm
import arviz as az
sns.set()
```

#### Create OK turbines data set

```
In [6]: turbines = pd.read_csv('turbines.csv')
# The "year" column contains how many years since the year 2000
turbines['year'] = turbines['p_year'] - 2000
turbines = turbines.drop('p_year', axis=1)
turbines.head()
```

```
Out[6]:
           t_state t_built t_cap year
         0
                        6 390.0
                                 -3.0
               ΑK
               ΑK
                        6 475.0
                                 -1.0
                           100.0
                                  0.0
               ΑK
                        1 1500.0
               ΑK
                                   1.0
               ΑK
                        1 100.0
                                   2.0
```

```
In [7]: # Turbines in Oklahoma
  ok_filter = (turbines.t_state == 'OK') & (turbines.year >= 0)
  ok_turbines = turbines[ok_filter].sort_values('year')
  ok_turbines["totals"] = np.cumsum(ok_turbines["t_built"])
  # Log-transform the counts, too
  ok_turbines["log_totals"] = np.log(ok_turbines["totals"])
```

# Draw posterior samples from Gaussian, Poisson and negative binomial models

```
In [8]: # Bayesian regression model using Gaussian likelihood (equivalent to OLS)

with pm.Model() as gaussian_model:
    # Specify glm and pass in data. This is similar to the code from
    # Lab 3 that created `theta = pm.Beta(...)`, etc., but using PyMC3's
    # GLM module sets everything up automaticaly.
    # The resulting linear model, its likelihood and
    # and all its parameters are automatically added to our model.
    glm.GLM.from_formula('log_totals ~ year', ok_turbines)
    # draw posterior samples using NUTS sampling
    gaussian_trace = pm.sample(1000, cores=2, target_accept=0.95)
# Not counting variable name changes, there are two differences
```

```
# between this cell and the version we did before: can you find them?
with pm.Model() as poisson model:
    glm.GLM.from_formula('totals ~ year', ok_turbines, family=glm.families.Poisson())
    # draw posterior samples using NUTS sampling
    poisson trace = pm.sample(1000, cores=2, target accept=0.95)
with pm.Model() as negbin model:
    glm.GLM.from_formula('totals ~ year', ok_turbines, family=glm.families.NegativeBinom
    # draw posterior samples using NUTS sampling
    negbin trace = pm.sample(1000, cores=2, target accept=0.95)
/tmp/ipykernel_24/11348518.py:11: FutureWarning: In v4.0, pm.sample will return an `arvi
z.InferenceData` object instead of a `MultiTrace` by default. You can pass return infere
ncedata=True or return inferencedata=False to be safe and silence this warning.
  gaussian_trace = pm.sample(1000, cores=2, target_accept=0.95)
Auto-assigning NUTS sampler...
Initializing NUTS using jitter+adapt_diag...
Multiprocess sampling (2 chains in 2 jobs)
NUTS: [sd, year, Intercept]
                                       100.00% [4000/4000 00:03<00:00 Sampling 2 chains, 0
divergences]
Sampling 2 chains for 1_000 tune and 1_000 draw iterations (2_000 + 2_000 draws total) t
ook 3 seconds.
/tmp/ipykernel_24/11348518.py:18: FutureWarning: In v4.0, pm.sample will return an `arvi
z.InferenceData` object instead of a `MultiTrace` by default. You can pass return infere
ncedata=True or return_inferencedata=False to be safe and silence this warning.
  poisson trace = pm.sample(1000, cores=2, target accept=0.95)
Auto-assigning NUTS sampler...
Initializing NUTS using jitter+adapt_diag...
Multiprocess sampling (2 chains in 2 jobs)
NUTS: [mu, year, Intercept]
                                       100.00% [4000/4000 00:06<00:00 Sampling 2 chains, 0
divergences]
Sampling 2 chains for 1 000 tune and 1 000 draw iterations (2 000 + 2 000 draws total) t
ook 7 seconds.
/tmp/ipykernel_24/11348518.py:24: FutureWarning: In v4.0, pm.sample will return an `arvi
z.InferenceData` object instead of a `MultiTrace` by default. You can pass return infere
ncedata=True or return inferencedata=False to be safe and silence this warning.
  negbin_trace = pm.sample(1000, cores=2, target_accept=0.95)
Auto-assigning NUTS sampler...
Initializing NUTS using jitter+adapt_diag...
Multiprocess sampling (2 chains in 2 jobs)
NUTS: [alpha, mu, year, Intercept]
```

100.00% [4000/4000 00:05<00:00 Sampling 2 chains, 0

divergences]

Sampling 2 chains for  $1\_000$  tune and  $1\_000$  draw iterations ( $2\_000 + 2\_000$  draws total) took 5 seconds.

### Credible intervals for coefficients

```
In [10]: |
         β_year_samples = negbin_trace['year']
         β_year_samples
Out[10]: array([0.22088704, 0.23019972, 0.23461718, ..., 0.24547829, 0.25445352,
                0.234716081)
In [11]: \beta_year_samples.shape
Out[11]: (2000,)
         plt.hist(β_year_samples, bins=20)
Out[12]: (array([ 3.,
                                3., 14., 34., 76., 120., 205., 262., 337., 335.,
                         3.,
                 240., 170., 97., 55., 19., 12.,
                                                        5.,
                                                              7.,
                                                                    3.]),
          array([0.11942477, 0.13156762, 0.14371048, 0.15585333, 0.16799618,
                 0.18013903, 0.19228188, 0.20442474, 0.21656759, 0.22871044,
                 0.24085329, 0.25299614, 0.26513899, 0.27728185, 0.2894247,
                 0.30156755, 0.3137104 , 0.32585325, 0.33799611, 0.35013896,
                 0.36228181]),
          <BarContainer object of 20 artists>)
          350
          300
          250
          200
          150
          100
           50
            0
                                                            0.30
                       0.15
                                    0.20
                                                0.25
                                                                        0.35
In [13]: |
         sorted\_samples = np.sort(\beta\_year\_samples)
         sorted samples
Out[13]: array([0.11942477, 0.12179607, 0.12432532, ..., 0.35316032, 0.36213128,
                0.36228181])
In [14]:
         0.95 * 2000
Out[14]: 1900.0
In [15]: sorted_samples[:1900]
Out[15]: array([0.11942477, 0.12179607, 0.12432532, ..., 0.28930384, 0.28933486,
                0.2895472 ])
```

```
In [16]: 0.288 - 0.138
Out[16]: 0.14999999999999997
In [17]: | sorted_samples[100:2000]
Out[17]: array([0.18868333, 0.18869092, 0.18881772, ..., 0.35316032, 0.36213128,
               0.36228181])
In [18]: 0.358 - 0.194
Out[18]: 0.1639999999999998
In [19]: sorted samples [40:1940]
Out[19]: array([0.17415211, 0.17441136, 0.17544495, ..., 0.29621778, 0.2966495,
               0.296964071)
In [20]: 0.295 - 0.181
Draw posterior predictive samples from Gaussian, Poisson and negative
         binomial models
         We use PyMC3's sample posterior predictive:
In [21]: with gaussian_model:
            gaussian ppc = pm.sample posterior predictive(
                gaussian_trace, var_names=["year", "Intercept", "y"]
                                              100.00% [2000/2000 00:01<00:00]
In [22]: with poisson_model:
            poisson_ppc = pm.sample_posterior_predictive(
                poisson_trace, var_names=["year", "Intercept", "y"]
             )
                                              100.00% [2000/2000 00:01<00:00]
In [23]: with negbin_model:
            negbin ppc = pm.sample posterior predictive(
                negbin_trace, var_names=["year", "Intercept", "y"]
             )
                                              100.00% [2000/2000 00:01<00:00]
In [33]: negbin_ppc
```

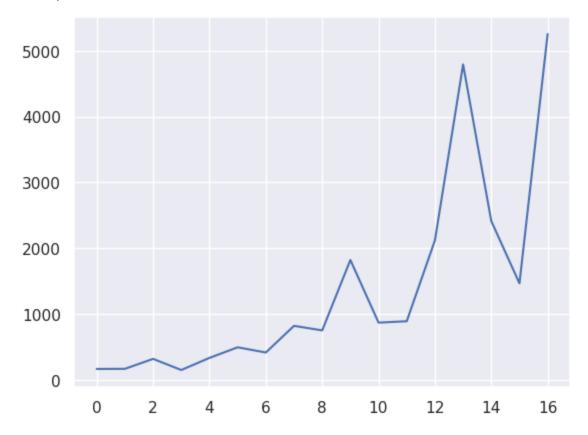
```
Out[33]: {'year': array([0.22088704, 0.23019972, 0.23461718, ..., 0.24547829, 0.25445352,
                 0.23471608]),
          'Intercept': array([4.51208039, 4.2686655 , 4.31509783, ..., 4.18148262, 4.04335691,
                 4.21057468]),
          'y': array([[ 205,
                                       310, ..., 4373, 10755, 13293],
                                108,
                           208,
                                  243, ..., 6419, 12672, 6433],
                     70,
                 [
                           152,
                                  255, ..., 7372, 3831, 7449],
                     80,
                 . . . ,
                 [ 103,
                            55,
                                   97, ...,
                                             3840, 4972,
                            54,
                                             2656, 15923,
                                                           5515],
                     78,
                                  168, ...,
                     43,
                            62,
                                  117, ...,
                                             5658, 9861, 10956]])}
```

```
In [34]: negbin_ppc['y'].shape
```

Out[34]: (2000, 17)

```
In [35]: plt.plot(negbin_ppc['y'][47, :])
```

Out[35]: [<matplotlib.lines.Line2D at 0x7f934c78deb0>]



In GLMs, we have the "average prediction"  $\bar{y}=$  InverseLinkFunction $(X\beta)$ , and then we have the actual predictions themselves, which are randomly distributed around this average, based on the chosen likelihood model.

For example, suppose year = 4, the coefficient  $\beta_{year}$  is 0.19, and the intercept is 4.8. Then the average prediction  $\bar{y}$  for that year will be  $\exp(4 \cdot 0.19 + 4.8)$ :

```
In [24]: np.exp(4 * 0.19 + 4.8)
Out[24]: 259.8228363229507
```

```
In []: np.exp(4 * 0.23 + 4.8)
```

or, about 260 turbines. The actual distribution for the number of turbines we actually observe will be a

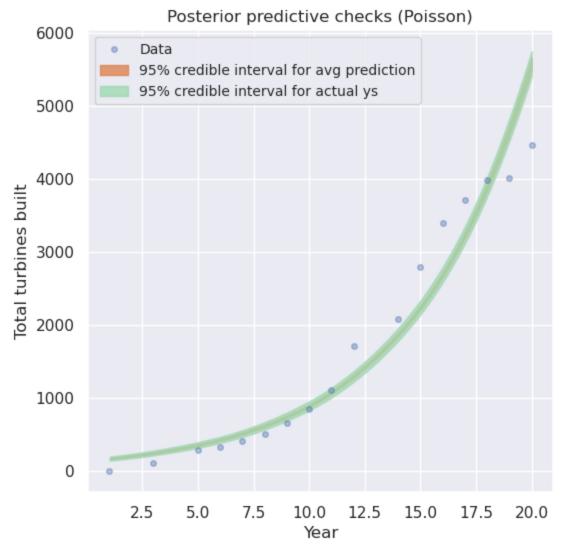
negative binomial with mean 259.8 (and variance estimated from the data).

Note that there is uncertainty in the average prediction too: the coefficient 0.19 in the example above could have been different. So, it's important to track our uncertainty at two levels:

- The uncertainty for the average prediction at each point (we'll also call this the 'regression output') reflects how uncertain we are about the coefficients, and what that means for the average prediction
- The uncertainty for the actual observed data reflects how much variation around the mean we should expect based on our data.

### Plot posterior predictive distribution for each model

```
In [25]: def compute_avg_prediction(ppc):
             linear_part = np.outer(ppc["year"], ok_turbines.year.values) + ppc["Intercept"][:,np
             return np.exp(linear_part)
In [27]: # Compute the posterior regression function (y hat values) for each model
         y_hat_poisson = compute_avg_prediction(poisson_ppc)
         y_hat_negbin = compute_avg_prediction(negbin_ppc)
         y_hat_gaussian = compute_avg_prediction(gaussian_ppc)
         y_hat_vals = [y_hat_poisson, y_hat_gaussian, y_hat_negbin]
In [28]: \_, ax = plt.subplots(figsize = (6,6))
         ax.plot(ok_turbines.year, ok_turbines.totals, "o", ms=4, alpha=0.4, label="Data")
         az.plot_hdi(
             ok turbines.year,
             y_hat_poisson,
             ax=ax,
             hdi prob=0.95,
             fill_kwargs={"alpha": 0.8, "label": "95% credible interval for avg prediction"},
         az.plot_hdi(
             ok_turbines.year,
             poisson_ppc["y"],
             ax=ax,
             hdi prob=0.95,
             fill_kwargs={"alpha": 0.8, "color": "#a1dab4", "label": "95% credible interval for a
         ax.set xlabel("Year")
         ax.set ylabel("Total turbines built")
         ax.set title("Posterior predictive checks (Poisson)")
         ax.legend(fontsize=10);
         /opt/conda/lib/python3.9/site-packages/arviz/plots/hdiplot.py:157: FutureWarning: hdi cu
         rrently interprets 2d data as (draw, shape) but this will change in a future release to
         (chain, draw) for coherence with other functions
           hdi_data = hdi(y, hdi_prob=hdi_prob, circular=circular, multimodal=False, **hdi_kwarg
         /opt/conda/lib/python3.9/site-packages/arviz/plots/hdiplot.py:157: FutureWarning: hdi cu
         rrently interprets 2d data as (draw, shape) but this will change in a future release to
         (chain, draw) for coherence with other functions
           hdi_data = hdi(y, hdi_prob=hdi_prob, circular=circular, multimodal=False, **hdi_kwarg
         s)
```



```
In [32]:
         _, ax = plt.subplots(1,3, figsize = (15,7.5), dpi=100)
         names = ["Poisson", "Gaussian", "Negative binomial"]
         for idx, ppc in enumerate([poisson_ppc, gaussian_ppc, negbin_ppc]):
             ax[idx].plot(ok_turbines.year, ok_turbines.totals, "o", ms=4, alpha=0.4, label="Data
             az.plot_hdi(
                 ok_turbines.year,
                 y_hat_vals[idx],
                 ax=ax[idx],
                 hdi_prob=0.95,
                 fill_kwargs={"alpha": 0.8, "label": "95% credible interval for avg prediction"},
             )
             if idx==1:
                 pp_y = np.exp(ppc["y"])
             else:
                 pp_y = ppc["y"]
             az.plot_hdi(
                 ok_turbines.year,
                 pp_y,
                 ax=ax[idx],
                 hdi_prob=0.95,
                 fill_kwargs={"alpha": 0.8, "color": "#a1dab4", "label": "95% credible interval f
             )
             ax[idx].set_xlabel("Year")
             ax[idx].set_ylabel("Total turbines built")
             ax[idx].set_title(names[idx])
             ax[idx].legend(fontsize=10)
```

ax[idx].set\_ylim(-100, 90000)
plt.tight\_layout()

