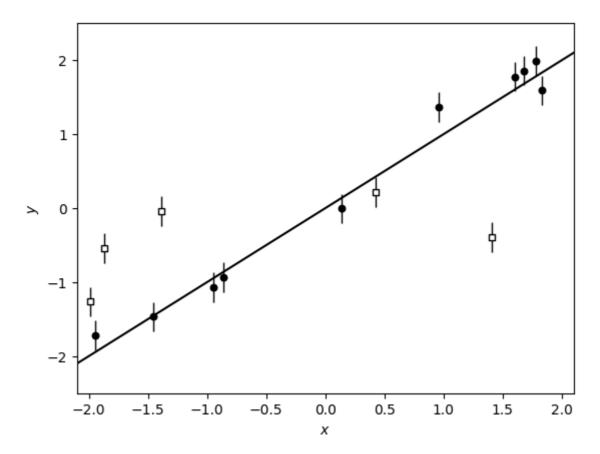
setup and data

```
In [3]: import numpy as np
        import matplotlib.pyplot as plt
In [4]: # We'll choose the parameters of our synthetic data.
        # The outlier probability will be 80%:
        true_frac = 0.8
        # The linear model has unit slope and zero intercept:
        true_params = [1.0, 0.0]
        # The outliers are drawn from a Gaussian with zero mean and unit variance
        true_outliers = [0.0, 1.0]
In [5]: # For reproducibility, let's set the random number seed and generate the
        np.random.seed(12)
        x = np.sort(np.random.uniform(-2, 2, 15))
        yerr = 0.2 * np.ones_like(x)
        y = true_params[0] * x + true_params[1] + yerr * np.random.randn(len(x))
        # Those points are all drawn from the correct model so let's replace some
        # them with outliers.
        m_bkg = np.random.rand(len(x)) > true_frac
        y[m bkg] = true outliers[0]
        y[m_bkg] += np.sqrt(true_outliers[1] + yerr[m_bkg] ** 2) * np.random.rand
        # Then save the *true* line.
        x0 = np.linspace(-2.1, 2.1, 200)
        y0 = np.dot(np.vander(x0, 2), true_params)
In [6]: def plot_data():
            plt.errorbar(x, y, yerr=yerr, fmt=",k", ms=0, capsize=0, lw=1, zorder
            plt.scatter(x[m_bkg], y[m_bkg], marker="s", s=22, c="w", edgecolor="k
            plt.scatter(
                x[\sim m_bkg], y[\sim m_bkg], marker="o", s=22, c="k", zorder=1000, label
            plt.plot(x0, y0, color="k", lw=1.5)
            plt.xlabel("$x$")
            plt.ylabel("$y$")
            plt.ylim(-2.5, 2.5)
            plt.xlim(-2.1, 2.1)
In [7]: plot_data()
```



numpyro: simple model, with and without treatment of outliers

```
In [8]: import jax
import jax.numpy as jnp

import numpyro
from numpyro import distributions as dist, infer
from numpyro_ext.distributions import MixtureGeneral
```

model formulation -->

numpyro.set_host_device_count(2)

```
In [10]:
         def linear_model(x, yerr, y=None):
             # These are the parameters that we're fitting and we're required to d
             # priors using distributions from the numpyro.distributions module.
             theta = numpyro.sample("theta", dist.Uniform(-0.5 * jnp.pi, 0.5 * jnp
             b_perp = numpyro.sample("b_perp", dist.Normal(0, 1))
             # Transformed parameters (and other things!) can be tracked during sa
             # "deterministics" as follows:
             m = numpyro.deterministic("m", jnp.tan(theta))
             b = numpyro.deterministic("b", b_perp / jnp.cos(theta))
             # Then we specify the sampling distribution for the data, or the like
             # Here we're using a numpyro.plate to indicate that the data are inde
             # isn't actually necessary here and we could have equivalently omitte
             # the Normal distribution can already handle vector-valued inputs. Bu
             # get into the habit of using plates because some inference algorithm
             # can take advantage of knowing this structure.
```

In [9]:

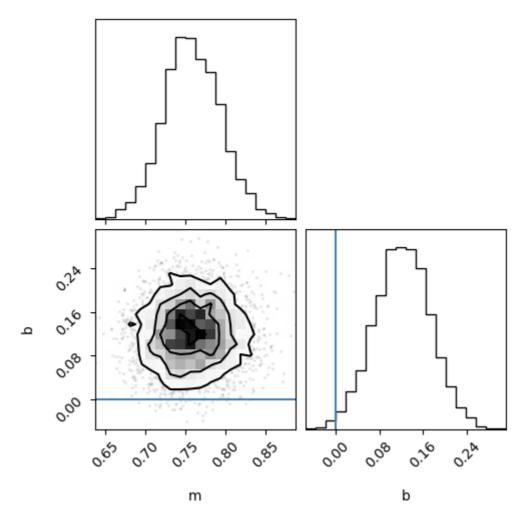
```
with numpyro.plate("data", len(x)):
                 numpyro.sample("y", dist.Normal(m * x + b, yerr), obs=y)
In [11]: # Using the model above, we can now sample from the posterior distribution
         # U-Turn Sampler (NUTS).
         sampler1 = infer.MCMC(
             infer.NUTS(linear_model),
             num_warmup=2000,
             num_samples=2000,
             num_chains=2,
             progress bar=True,
         %time sampler1.run(jax.random.PRNGKey(0), x, yerr, y=y)
                       | 0/4000 [00:00<?, ?it/s]
          0%|
                       | 0/4000 [00:00<?, ?it/s]
        CPU times: user 1.8 s, sys: 46.8 ms, total: 1.85 s
        Wall time: 1.87 s
In [13]: def linear mixture model(x, yerr, y=None):
             # Our "foreground" model is identical to the one we used previously:
             # parameterized by "theta" and "b_perp". Note that we don't wrap the
             # sampling distribution in a `numpyro.sample` here because we're goin
             # use it in the mixture distribution below.
             theta = numpyro.sample("theta", dist.Uniform(-0.5 * jnp.pi, 0.5 * jnp
             b_perp = numpyro.sample("b_perp", dist.Normal(0.0, 1.0))
             m = numpyro.deterministic("m", jnp.tan(theta))
             b = numpyro.deterministic("b", b_perp / jnp.cos(theta))
             fg_dist = dist.Normal(m * x + b, yerr)
             # Our outlier model is a Gaussian where we're fitting for the zero an
             # standard deviation.
             bg_mean = numpyro.sample("bg_mean", dist.Normal(0.0, 1.0))
             bg_sigma = numpyro.sample("bg_sigma", dist.HalfNormal(3.0))
             bg_dist = dist.Normal(bg_mean, jnp.sqrt(bg_sigma**2 + yerr**2))
             # We use a `Catagorical` distribution to define the outlier probabili
             # fit for the parameter `Q` which specifies the probability that any
             # individual point is a member of the foreground model. Therefore, th
             # "prior" outlier probability is `1 - Q`.
             Q = numpyro.sample("Q", dist.Uniform(0.0, 1.0))
             mix = dist.Categorical(probs=jnp.array([Q, 1.0 - Q]))
             # As with the previous model, the use of a `plate` here is optional,
             # let's do it anyways.
             with numpyro.plate("data", len(x)):
                 # The `numpyro.distributions` module doesn't yet implement a mixt
                 # distribution that is flexible enough for our use cases, so we'l
                 # one that I implemented in the `numpyro-ext` package. (This is s
                 # a lie: the `MixtureSameFamily` distribution in NumPyro _would_
                 # here, but not in our next example, so bear with me!)
                 numpyro.sample("obs", MixtureGeneral(mix, [fg_dist, bg_dist]), ob
In [14]: # Using the model above, we can now sample from the posterior distribution
         # U-Turn Sampler (NUTS).
         sampler2 = infer.MCMC(
             infer.NUTS(linear_mixture_model),
             num_warmup=2000,
             num_samples=2000,
             num_chains=2,
```

model evaluation -->

```
In [15]: import arviz as az
import corner
```

```
inf_data1 = az.from_numpyro(sampler1)
corner.corner(inf_data1, var_names=["m", "b"], truths=true_params);
az.summary(inf_data1)
```

| Out[16]: | | mean | sd | hdi_3% | hdi_97% | mcse_mean | mcse_sd | ess_bulk | ess_tai |
|----------|--------|-------|-------|--------|---------|-----------|---------|----------|---------|
| | b | 0.122 | 0.052 | 0.018 | 0.214 | 0.001 | 0.001 | 3891.0 | 2685.0 |
| | b_perp | 0.098 | 0.041 | 0.013 | 0.169 | 0.001 | 0.000 | 3894.0 | 2742.0 |
| | m | 0.758 | 0.035 | 0.691 | 0.826 | 0.001 | 0.000 | 3461.0 | 2719.(|
| | theta | 0.648 | 0.023 | 0.608 | 0.693 | 0.000 | 0.000 | 34610 | 2719 (|

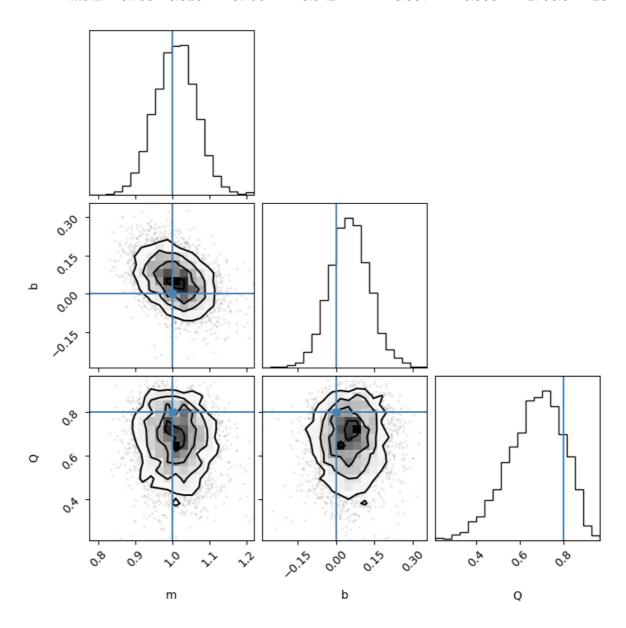


```
In [17]: inf_data2 = az.from_numpyro(sampler2)
    corner.corner(
```

```
inf_data2,
    var_names=["m", "b", "Q"],
    truths={
        "m": true_params[0],
        "b": true_params[1],
        "Q": true_frac,
     },
);
az.summary(inf_data2)
```

Out[17]:

| | mean | sd | hdi_3% | hdi_97% | mcse_mean | mcse_sd | ess_bulk | ess_ |
|----------|--------|-------|--------|---------|-----------|---------|----------|------|
| Q | 0.668 | 0.127 | 0.427 | 0.886 | 0.002 | 0.002 | 3513.0 | 290 |
| b | 0.052 | 0.078 | -0.097 | 0.198 | 0.001 | 0.001 | 2954.0 | 266 |
| b_perp | 0.037 | 0.056 | -0.069 | 0.140 | 0.001 | 0.001 | 2912.0 | 27 |
| bg_mean | -0.412 | 0.403 | -1.170 | 0.341 | 0.009 | 0.006 | 2264.0 | 22 |
| bg_sigma | 0.776 | 0.512 | 0.021 | 1.657 | 0.011 | 0.008 | 2195.0 | 15: |
| m | 1.008 | 0.056 | 0.897 | 1.106 | 0.001 | 0.001 | 2738.0 | 25 |
| theta | 0.789 | 0.028 | 0.738 | 0.842 | 0.001 | 0.000 | 2738.0 | 25 |



pymc: redo same models, with and without numpyro api

```
In [18]: import pymc as pm
         import numpy as np
         from pymc import Uniform, Normal, HalfNormal, Model, Mixture, Categorical
         model formulation -->
         simple model, no outlier handling
In [19]: with Model() as model:
             theta = Uniform("theta", -0.5*np.pi, 0.5*np.pi)
             b_perp = Normal("b_perp", 0, sigma=1)
             # transformed params
             m = pm.Deterministic("m", np.tan(theta))
             b = pm.Deterministic("b", b_perp / np.cos(theta))
             # likelihood
             likelihood = Normal("y", mu=m*x+b, sigma=yerr, observed=y)
             # inference
             %time idata1 = sample(2000, chains=2)
                                               100.00% [6000/6000 00:00<00:00
       Sampling 2 chains, 0 divergences]
        CPU times: user 1.46 s, sys: 179 ms, total: 1.64 s
        Wall time: 2.39 s
         ... and now with jax
In [20]: with Model() as model:
             theta = Uniform("theta", -0.5*np.pi, 0.5*np.pi)
             b_perp = Normal("b_perp", 0, sigma=1)
             # transformed params
             m = pm.Deterministic("m", np.tan(theta))
             b = pm.Deterministic("b", b_perp / np.cos(theta))
             # likelihood
             likelihood = Normal("y", mu=m*x+b, sigma=yerr, observed=y)
             # inference
             %time idata2 = sampling_jax.sample_numpyro_nuts(2000, chains=2)
        Compiling...
        Compilation time = 0:00:00.280433
        Sampling...
                        | 0/3000 [00:00<?, ?it/s]
          0%|
          0%|
                        | 0/3000 [00:00<?, ?it/s]
```

Sampling time = 0:00:01.486291

Transforming variables...

Transformation time = 0:00:00.079975

CPU times: user 1.82 s, sys: 54 ms, total: 1.87 s

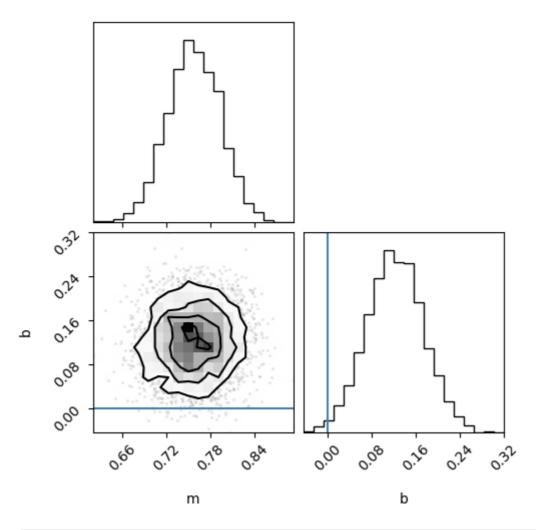
Wall time: 1.87 s

model evaluation -->

In [21]: corner.corner(idata1, var_names=["m", "b"], truths=true_params);
az.summary(idata1, var_names=["b_perp", "theta", "m", "b"])

| | F 7 | |
|-------|-----------------------|---|
| Out | 171 | = |
| U U L | $L \subseteq \perp J$ | |

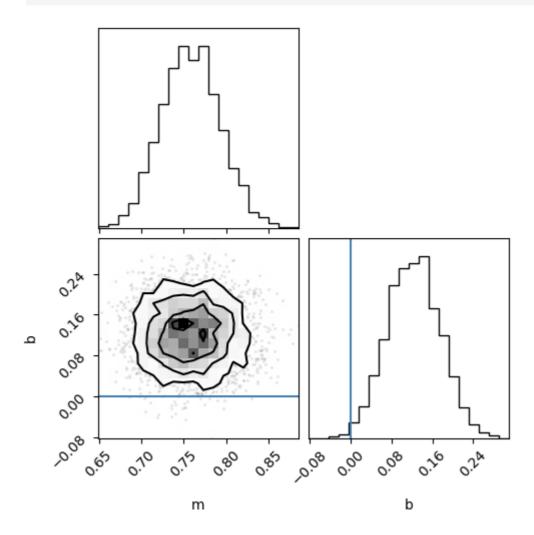
| | mean | sd | hdi_3% | hdi_97% | mcse_mean | mcse_sd | ess_bulk | ess_tai |
|--------|-------|-------|--------|---------|-----------|---------|----------|---------|
| b_perp | 0.098 | 0.041 | 0.020 | 0.174 | 0.001 | 0.000 | 4486.0 | 2971.(|
| theta | 0.648 | 0.023 | 0.606 | 0.690 | 0.000 | 0.000 | 4471.0 | 3046.0 |
| m | 0.757 | 0.036 | 0.693 | 0.825 | 0.001 | 0.000 | 4471.0 | 3046.0 |
| b | 0.123 | 0.051 | 0.028 | 0.222 | 0.001 | 0.001 | 4491.0 | 2992.0 |



In [22]: corner.corner(idata2, var_names=["m", "b"], truths=true_params);
az.summary(idata2, var_names=["b_perp", "theta", "m", "b"])

Out[22]:

| | | mean | sd | hdi_3% | hdi_97% | mcse_mean | mcse_sd | ess_bulk | ess_tai |
|--|--------|-------|-------|--------|---------|-----------|---------|----------|---------|
| | b_perp | 0.098 | 0.042 | 0.021 | 0.174 | 0.001 | 0.001 | 3133.0 | 2385.0 |
| | theta | 0.648 | 0.022 | 0.608 | 0.690 | 0.000 | 0.000 | 3119.0 | 2506.0 |
| | m | 0.758 | 0.035 | 0.696 | 0.825 | 0.001 | 0.000 | 3119.0 | 2506.0 |
| | b | 0.124 | 0.052 | 0.025 | 0.217 | 0.001 | 0.001 | 3130.0 | 2491.0 |



model formulation -->

model with some outlier handling

```
In [23]:
    with Model() as model:
        theta = Uniform("theta", -0.5*np.pi, 0.5*np.pi)
        b_perp = Normal("b_perp", 0.0, sigma=1.0)
        m = pm.Deterministic("m", pm.math.tan(theta))
        b = pm.Deterministic("b", b_perp / pm.math.cos(theta))
        fg_dist = Normal.dist(m*x*+b, yerr)

        bg_mean = Normal("bg_mean", 0.0, 1.0)
        bg_sigma = HalfNormal("bg_sigma", 3.0)
        bg_dist = Normal.dist(bg_mean, pm.math.sqrt(bg_sigma**2 + yerr**2))

        Q = Uniform("Q", 0.0, 1.0)
        mix = pm.Dirichlet("mix", pm.math.stack([Q, 1.0-Q]))

# likelihood
```

```
likelihood = Mixture("likelihood", w=mix, comp_dists=[fg_dist, bg_dis
# inference
%time idata3 = sample(2000, chains=2)
```

100.00% [6000/6000 00:03<00:00

Sampling 2 chains, 4 divergences]

```
CPU times: user 4.83 s, sys: 314 ms, total: 5.14 s Wall time: 8.78 s
```

```
In [24]:
with Model() as model:
    theta = Uniform("theta", -0.5*np.pi, 0.5*np.pi)
    b_perp = Normal("b_perp", 0.0, sigma=1.0)
    m = pm.Deterministic("m", pm.math.tan(theta))
    b = pm.Deterministic("b", b_perp / pm.math.cos(theta))
    fg_dist = Normal.dist(m*x+b, yerr)

bg_mean = Normal("bg_mean", 0.0, 1.0)
    bg_sigma = HalfNormal("bg_sigma", 3.0)
    bg_dist = Normal.dist(bg_mean, pm.math.sqrt(bg_sigma**2 + yerr**2))

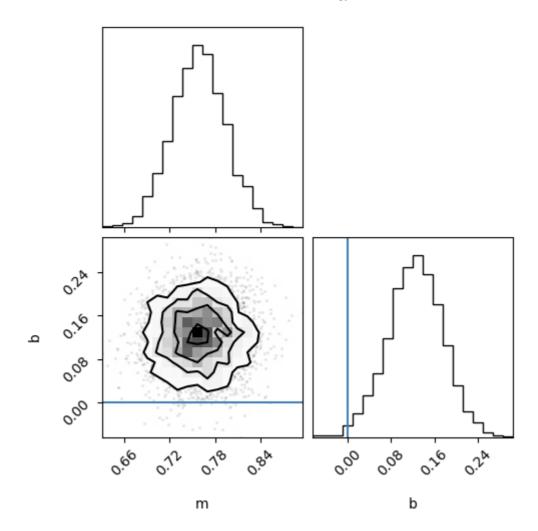
Q = Uniform("Q", 0.0, 1.0)
    mix = pm.Dirichlet("mix", pm.math.stack([Q, 1.0-Q]))

# likelihood
likelihood = Mixture("likelihood", w=mix, comp_dists=[fg_dist, bg_dis", inference
    %time idata4 = sampling_jax.sample_numpyro_nuts(2000, chains=2)
```

model evaluation -->

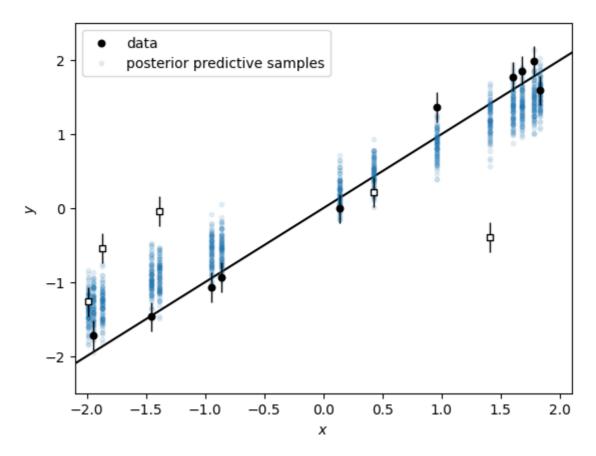
```
In [34]: corner.corner(idata3, var_names=["m", "b"], truths=true_params);
az.summary(idata3, var_names=["b_perp", "theta", "m", "b"])
```

Out[34]: sd hdi_3% hdi_97% mcse_mean mcse_sd ess_bulk ess_tai mean **b_perp** 0.099 0.041 0.020 0.176 0.001 0.001 2783.0 2377.0 **theta** 0.649 0.023 0.605 0.690 0.000 0.000 3319.0 2452.0 **m** 0.759 0.036 0.692 0.825 0.001 0.000 3319.0 2452.0 0.125 0.051 0.024 0.220 0.001 0.001 2768.0 2622.0



```
In []: corner.corner(idata4, var_names=["m", "b"], truths=true_params);
az.summary(idata4, var_names=["b_perp", "theta", "m", "b"])
```

simulate



In []: