# Order-restricted one-sample comparison

#### import os

```
if "KERAS_BACKEND" not in os.environ:
    # set this to "torch", "tensorflow", or "jax"
    os.environ["KERAS_BACKEND"] = "jax"
import matplotlib.pyplot as plt
import numpy as np
import bayesflow as bf
import keras
```

INFO:bayesflow:Using backend 'jax'

The models are essentially the same as in the previous example on one-sample comparison, the difference is in the alternative model, where instead of having a Cauchy prior on the  $\delta$ , we restrict it to only negative values.

$$\begin{split} \mathcal{M}_{0} &: \delta = 0 \\ \mathcal{M}_{1} &: \delta \sim \operatorname{Cauchy}(0, 1)_{T(-\infty, 0)} \\ & \sigma \sim \operatorname{HalfCauchy}(0, 1) \\ & \mu = \delta \sigma \\ & x_{i} \sim \operatorname{Normal}(\mu, \sigma) \end{split}$$
(1)

#### Simulator

We need to define two simulator: one that represents the null hypothesis that  $\delta = 0$ , and one that represents the alternative hypothesis that  $\delta < 0$ . Then, we wrap them in a ModelComparisonSimulator, that will sample from either of them randomly.

We will also amortize over different sample sizes. Here we do this by randomly sampling values between 10 and 100. In the simulators, we will make sure that the output is always of length 100 (maximum sample size); the elements in the array whose index exceeds the actual sample size are filled with zeros. To make it easier for the networks to summarise such data, we will also create a binary indicator variable **observed**, which is one when the element in **x** is filled with an actual value, and zero otherwise.

```
max_n=100
def context():
    return dict(n=np.random.randint(10, max_n))
def prior_nuisance():
    sigma = np.random.standard_cauchy()
    sigma = np.abs(sigma)
    return dict(sigma=sigma)
def prior_null():
    return dict(delta=0)
def prior_alternative():
    delta = np.random.standard_cauchy()
    delta = - np.abs(delta)
    return dict(delta=delta)
def likelihood(sigma, delta, n):
    mu = sigma * delta
    x = np.zeros(max_n)
    x[:n] = np.random.normal(loc=mu, scale=sigma, size=n)
    observed = np.zeros(max_n)
    observed[:n] = 1
    return dict(x=x, observed=observed)
```

simulator\_null = bf.make\_simulator([context, prior\_nuisance, prior\_null, likelihood])
simulator\_alt = bf.make\_simulator([context, prior\_nuisance, prior\_alternative, likelihood])

simulator = bf.simulators.ModelComparisonSimulator([simulator\_null, simulator\_alt])

#### Approximator

The sample size is passed into the inference network directly, and the observations in x and the observed indicator are passed into a summary network first.

```
adapter=(
    bf.Adapter()
    .as_set(["x", "observed"])
    .rename("n", "classifier_conditions")
    .concatenate(["x", "observed"], into="summary_variables")
    .drop(["delta", "sigma"])
    )
```

Model comparison needs a classifier network to predict the posterior model probabilities. Here, we define a simple multi-layer perceptron to do this task.

```
inference_network = keras.Sequential([
    keras.layers.Dense(32, activation="gelu")
    for _ in range(6)
])
```

Then we wrap everything together.

```
approximator=bf.approximators.ModelComparisonApproximator(
    num_models=2,
    classifier_network=inference_network,
    summary_network=bf.networks.DeepSet(
        summary_dim=4,
        mlp_widths_equivariant=(32, 32),
        mlp_widths_invariant_inner=(32, 32),
        mlp_widths_invariant_outer=(32, 32),
        mlp_widths_invariant_last=(32, 32)
    ),
    adapter=adapter
)
```

#### Training

Here we will do offline traing. First, we will define the number of epochs, and the simulation budget (number of batches times the batch size). We also define an optimizer with a cosine decay schedule.

```
epochs=30
batches=20
batch_size=512
schedule=keras.optimizers.schedules.CosineDecay(1e-4, decay_steps=epochs*batches)
optimizer=keras.optimizers.Adam(schedule)
approximator.compile(optimizer)
```

Now we can prepare our training dataset.

train\_data=simulator.sample(batches\*batch\_size)

```
train_data=bf.datasets.OfflineDataset(
    data=train_data,
    batch_size=batch_size,
    adapter=adapter)
```

```
history=approximator.fit(
    dataset=train_data,
    epochs=epochs,
    num_batches=batches,
    batch_size=batch_size)
```

## f=bf.diagnostics.plots.loss(history=history)



### Validation

test\_data=simulator.sample(1000)

```
true_models = test_data["model_indices"]
pred_models = approximator.predict(conditions=test_data)
f=bf.diagnostics.plots.mc_calibration(
    pred_models=pred_models,
    true_models=true_models,
    model_names=[r"$\mathcal{M}_0$",r"$\mathcal{M}_1$"],
)
```

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```
normalize="true"
```

```
)
```

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#### Inference

winter=np.array([-0.05,0.41,0.17,-0.13,0.00,-0.05,0.00,0.17,0.29,0.04,0.21,0.08,0.37,0.17,0.0 0.17,0.17,0.33,0.04,0.04,0.04,0.00,0.21,0.13,0.25,-0.05,0.29,0.42,-0.05,0.12,0.04,0.29 summer=np.array([0.00,0.38,-0.12,0.12,0.25,0.12,0.13,0.37,0.00,0.50,0.00,0.00,-0.13,-0.37,-0 0.00,0.00,0.00,0.00,0.25,0.13,-0.25,-0.38,-0.13,-0.25,0.00,0.00,-0.12,0.25,0.00,0.50,0 n = len(winter)

```
x = np.zeros(max_n)
x[:n] = winter-summer
observed = np.zeros(max_n)
observed[:n] = 1
```

inference\_data = dict(

```
n = np.array([[n]]),
x = x[np.newaxis],
observed = observed[np.newaxis])
```

pred\_models = approximator.predict(conditions=inference\_data)[0]

```
pred_models[0]/pred_models[1]
```

7.3406186

Lee, M. D., & Wagenmakers, E.-J. (2013). Bayesian Cognitive Modeling: A Practical Course. Cambridge University Press.