

Problem 3

```
In [2]: import load_helper

import pandas as pd
import matplotlib.pyplot as plt
import numpy as np

import importlib

from torch import tensor
```

```
In [53]: fname = '3.daphne'
graph = load_helper.graph_helper(fname)
%cat 3.daphne

(let [data [1.1 2.1 2.0 1.9 0.0 -0.1 -0.05]
      likes (foreach 3 []
                     (let [mu (sample (normal 0.0 10.0))
                           sigma (sample (gamma 1.0 1.0))]
                       (normal mu sigma)))
      pi (sample (dirichlet [1.0 1.0 1.0]))
      z-prior (discrete pi)
      z (foreach 7 [y data]
                (let [z (sample z-prior)
                    _ (observe (get likes z) y)]
                  z))]
      (= (first z) (second z)))
```

```
In [16]: import bbvi
importlib.reload(bbvi)

from bbvi import graph_bbvi_algo12
```

```
In [34]: %%time
T=1000
L=100
lr=0.05
r, logW, sigma = bbvi.graph_bbvi_algo12(graph,T=T,L=L,lr=lr,
                                         do_log=False)
```

t=0, Q after step={'sample3': Gamma(concentration: 1.0499999523162842, rate: 0.54132479429245), 'sample1': Gamma(concentration: 1.0499999523162842, rate: 0.54132479429245), 'sample2': Normal(loc: -0.05000000074505806, scale: 9.999954223632812), 'sample4': Normal(loc: -0.04999999701976776, scale: 9.999954223632812), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 0.05000000074505806, scale: 9.999954223632812), 'sample5': Gamma(concentration: 0.949999988079071, rate: 0.54132479429245)}

t=100, Q after step={'sample3': Gamma(concentration: 1.7900134325027466, rate: 0.3037601709365845), 'sample1': Gamma(concentration: 1.8541914224624634, rate: 0.27189576625823975), 'sample2': Normal(loc: 0.1060582771897316, scale: 10.16331672668457), 'sample4': Normal(loc: 0.2834552824497223, scale: 10.029430389404297), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 0.4662581980228424, scale: 9.678585052490234), 'sample5': Gamma(concentration: 1.4942097663879395, rate: 0.3359958529472351)}

t=200, Q after step={'sample3': Gamma(concentration: 1.8218340873718262, rate: 0.288897842168808), 'sample1': Gamma(concentration: 2.0777390003204346, rate: 0.21965110301971436), 'sample2': Normal(loc: 0.428939551115036, scale: 9.960811614990234), 'sample4': Normal(loc: 0.5669648051261902, scale: 9.839775085449219), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 0.9046857953071594, scale: 9.574071884155273), 'sample5': Gamma(concentration: 1.7429133653640747, rate: 0.27125629782676697)}

t=300, Q after step={'sample3': Gamma(concentration: 1.8476452827453613, rate: 0.28053200244903564), 'sample1': Gamma(concentration: 2.1341969966888428, rate: 0.20382164418697357), 'sample2': Normal(loc: 0.4442814290523529, scale: 9.737642288208008), 'sample4': Normal(loc: 0.38011208176612854, scale: 9.466002464294434), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 0.8249368667602539, scale: 9.532282829284668), 'sample5': Gamma(concentration: 1.8130440711975098, rate: 0.2528989017009735)}

t=400, Q after step={'sample3': Gamma(concentration: 1.8373630046844482, rate: 0.29939261078834534), 'sample1': Gamma(concentration: 2.2015764713287354, rate: 0.17110460996627808), 'sample2': Normal(loc: 0.470374737320961, scale: 9.841106414794922), 'sample4': Normal(loc: 0.22893062233924866, scale: 9.439980506896973), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 1.2280818223953247, scale: 9.21605110168457), 'sample5': Gamma(concentration: 1.9015353918075562, rate: 0.19762399792671204)}

t=500, Q after step={'sample3': Gamma(concentration: 1.8658350706100464, rate: 0.2749086618423462), 'sample1': Gamma(concentration: 2.231902837753296, rate: 0.15848882496356964), 'sample2': Normal(loc: 0.47287917137145996, scale: 9.806499481201172), 'sample4': Normal(loc: 0.32633745670318604, scale: 9.028315544128418), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 1.039696216583252, scale: 9.088177680969238), 'sample5': Gamma(concentration: 1.9733400344848633, rate: 0.17851786315441132)}

t=600, Q after step={'sample3': Gamma(concentration: 1.8757519721984863, rate: 0.26896676421165466), 'sample1': Gamma(concentration: 2.243215322494507, rate: 0.15821991860866547), 'sample2': Normal(loc: 0.4916940927505493, scale: 9.686244010925293), 'sample4': Normal(loc: 0.3182411754312134, scale: 8.951693534851074), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 1.0337268114089966, scale: 9.07003116607666), 'sample5': Gamma(concentration: 1.9913407564163208, rate: 0.17566214501857758)}

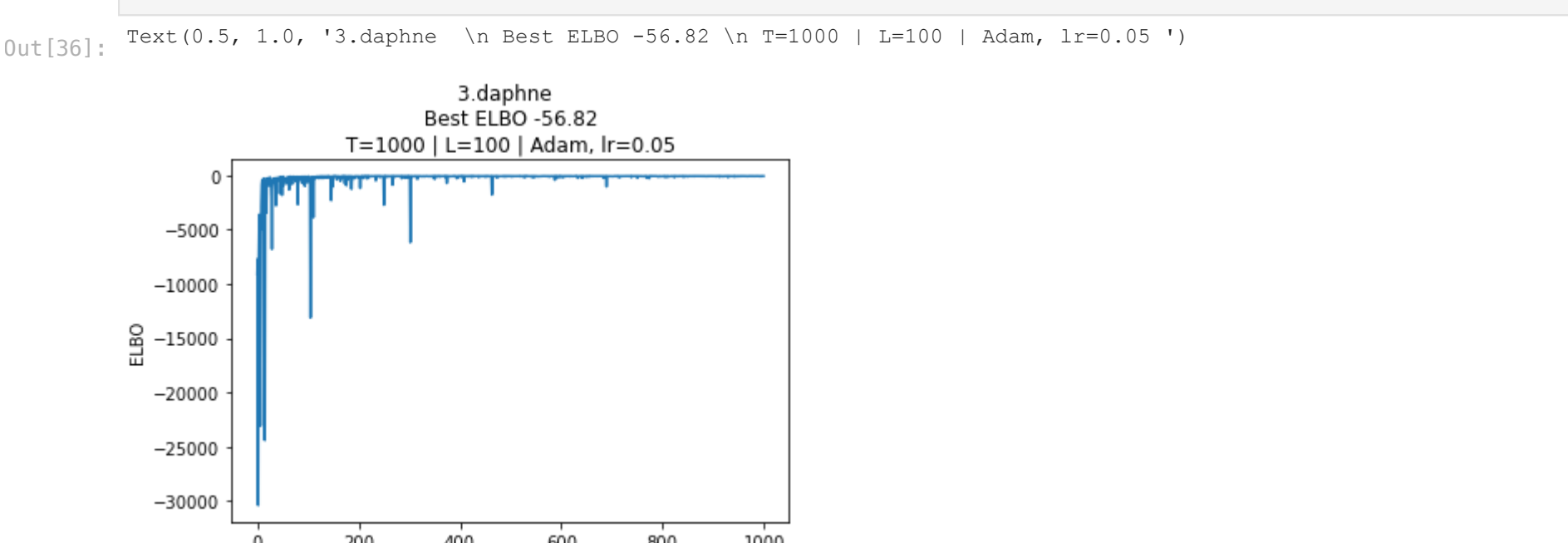
t=700, Q after step={'sample3': Gamma(concentration: 1.8960187435150146, rate: 0.2543301284313202), 'sample1': Gamma(concentration: 2.267752170562744, rate: 0.15137308835983276), 'sample2': Normal(loc: 0.381063848733902, scale: 9.35001277923584), 'sample4': Normal(loc: 0.40551134943962097, scale: 8.892863273620605), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 0.9871072769165039, scale: 9.003787994384766), 'sample5': Gamma(concentration: 2.0286083221435547, rate: 0.16498127579689026)}

t=800, Q after step={'sample3': Gamma(concentration: 1.9163165092468262, rate: 0.2426215559244156), 'sample1': Gamma(concentration: 2.2835867404937744, rate: 0.1482381969690323), 'sample2': Normal(loc: 0.3498328626155853, scale: 9.097940444946289), 'sample4': Normal(loc: 0.4434855282306671, scale: 8.850457191467285), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 1.0016863346099854, scale: 8.960394859313965), 'sample5': Gamma(concentration: 2.0418074131011963, rate: 0.16577433049678802)}

t=900, Q after step={'sample3': Gamma(concentration: 1.9285883903503418, rate: 0.23700419068336487), 'sample1': Gamma(concentration: 2.2909774780273438, rate: 0.15304233133792877), 'sample2': Normal(loc: 0.3641284108161926, scale: 9.029130935668945), 'sample4': Normal(loc: 0.443512886762619, scale: 8.797557830810547), 'sample6': Dirichlet(concentration: torch.Size([3])), 'sample11': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample13': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample19': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample15': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample9': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample7': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample17': Categorical(probs: torch.Size([3])), logits: torch.Size([3])), 'sample0': Normal(loc: 1.0233283042907715, scale: 8.92258358001709), 'sample5': Gamma(concentration: 2.0463082790374756, rate: 0.17054669559001923)}

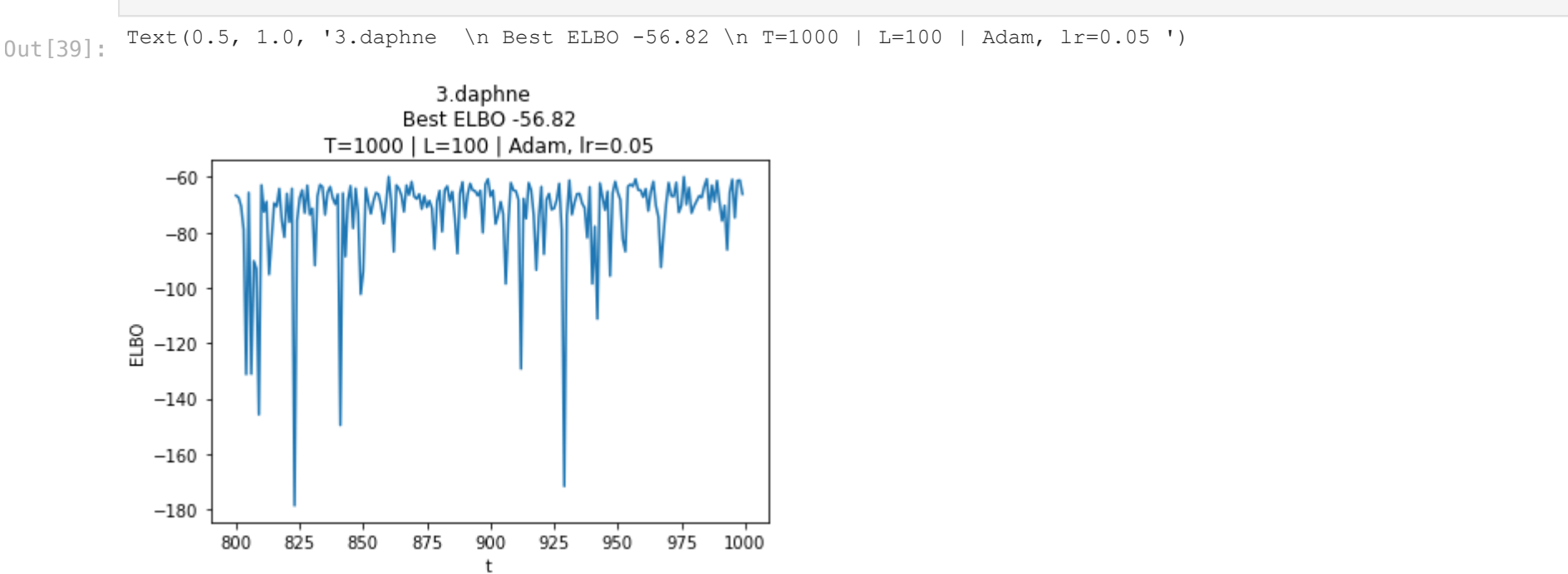
CPU times: user 48min 53s, sys: 6.19 s, total: 48min 59s
Wall time: 2h 13min 5s

```
In [36]: elbo = logW.mean(1)
pd.Series(elbo).plot()
plt.xlabel('t')
plt.ylabel('ELBO')
plt.title('{} \n Best ELBO {:.12f} \n T={} | L={} | Adam, lr={} '.format(fname,elbo.max(),T,L,lr))
```



```
In [37]: r_array = np.array(r)
```

```
In [39]: sr = pd.Series(elbo[-200:])
sr.index = np.arange(elbo.size-200,elbo.size)
sr.plot()
plt.xlabel('t')
plt.ylabel('ELBO')
plt.title('{} \n Best ELBO {:.12f} \n T={} | L={} | Adam, lr={} '.format(fname,elbo.max(),T,L,lr))
```



```
In [42]: probs = np.exp(logW)
probs /= probs.sum()
probs = probs.reshape(T,L)
posterior_r = (probs * r_array).sum(axis=0,1)
posterior_r2 = (probs * r_array**2).sum(axis=0,1)

std_r = np.sqrt(posterior_r2 - posterior_r**2)
```

```
In [51]: print('{} expected posterior z1==z2 {:.13f} | std posterior z1==z2 {:.13f}'.format(fname,posterior_r,std_r))

3.daphne expected posterior z1==z2 0.049 | std posterior z1==z2 0.215
```

Also write a paragraph or two about the mode-seeking behavior of VI on models like this that have internal symmetries. Note what the symmetries are and why they might cause problems with optimization.

This problem is a Gaussian mixture model. As discussed in office hours, the choice of $KL(p||q)$ instead of $KL(q||q)$ makes us snap to one mode, in this multimodal problem. See the inclusive divergence and exclusive divergence remarks on [Stefano Ermon's Probabilistic Graphical Models Winter class \(Stanford 2020-21\)](#). This GMM problem is multimodal, because there is no meaning to the cluster labels attached to the data points, they are just arbitrary labels. Stochastically, if some datapoints are relabelled, the cluster can shift in the sense that, the labels can swap between clusters. When this is happening in the algorithm, we are "jumping out" of a local minima, and "jumping into" another.