

```
In [1]: %matplotlib inline
import numpy as np
import matplotlib.pyplot as plt
plt.style.use('ggplot')
import pandas as pd
import statsmodels.api as sm
import pystan
```

第1章 p.18,19

混合正規分布

$$p(x|w) = (1 - a)\mathcal{N}(x) + a\mathcal{N}(x - b)$$

$$\mathcal{N}(x) = \frac{1}{\sqrt{2\pi}} \exp\left(-\frac{x^2}{2}\right)$$

$$W = \{(a, b) \mid 0 \leq a \leq 1, |b| \leq 5\}$$

$$\varphi(w) = \frac{1}{10}$$

真の分布

$$q(x) = p(x|a_0, b_0)$$

$$(a_0, b_0) = (0.5, 3.0), (0.5, 1.0), (0.5, 0.5)$$

サンプル数

$$n = 100$$

```
In [2]: def gen_mixed_gaussian(a0, b0, n, seed=None):
        if seed:
            np.random.seed(seed)
        bs = np.random.choice(2, size=n, p=[1.0-a0, a0])
        return np.vectorize(
            lambda b: np.random.normal(loc=b0) if b else np.random.normal(
            )(bs)
```

$n = 100$ のサンプルを作成する

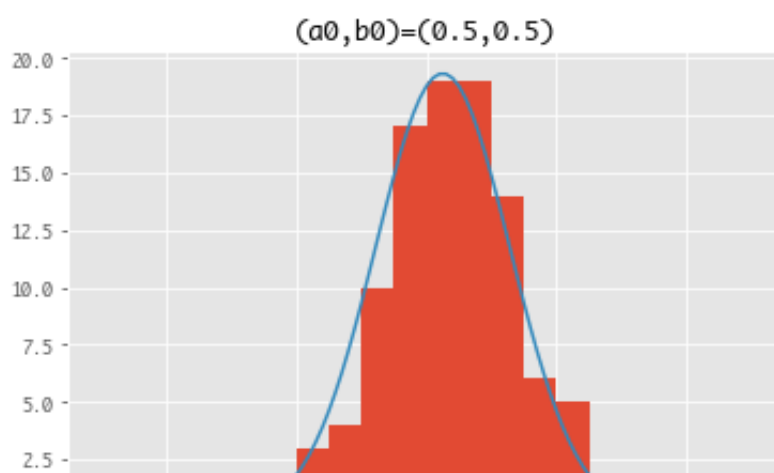
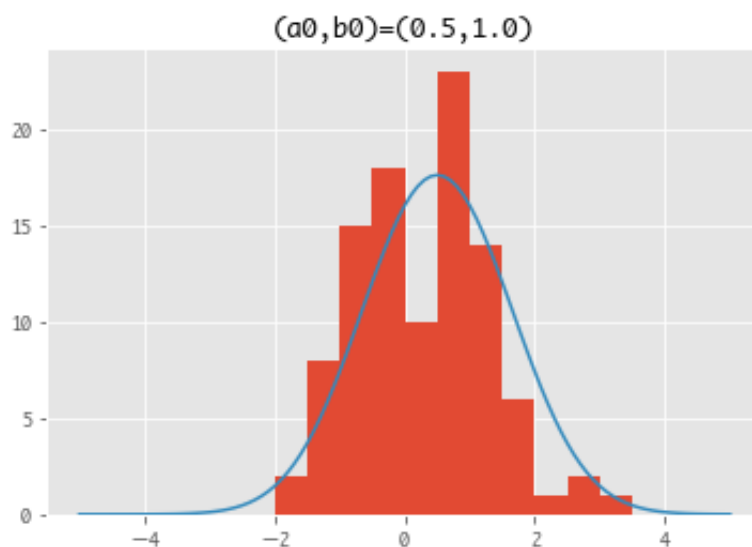
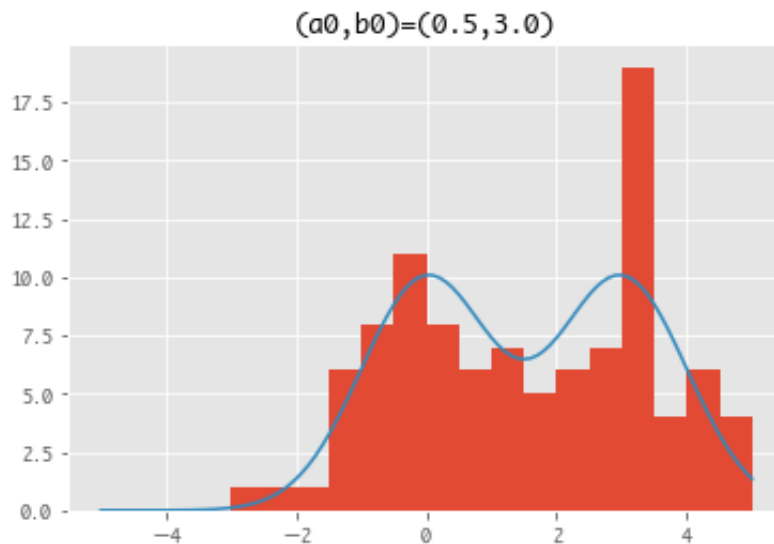
```
In [3]: np.random.seed(seed=123)
sample1 = gen_mixed_gaussian(0.5, 3.0, 100)
sample2 = gen_mixed_gaussian(0.5, 1.0, 100)
sample3 = gen_mixed_gaussian(0.5, 0.5, 100)
```

```
In [4]: from scipy.stats import norm
xs = np.linspace(-5, 5, 101)
def mg(a0, b0, n):
    return np.vectorize( \
        lambda x: ((1-a0)*norm.pdf(x) + a0*norm.pdf(x-b0))*n
    )
```

プロットしてみる

```
In [5]: plt.hist(sample1, bins=20, range=(-5.0, 5.0))
```

```
plt.hist(sample1, bins=20, range=(-5.0, 5.0),
plt.plot(xs, mg(0.5, 3.0, 50)(xs))
plt.title('(a0,b0)=(0.5,3.0)')
plt.show()
plt.hist(sample2, bins=20, range=(-5.0, 5.0))
plt.plot(xs, mg(0.5, 1.0, 50)(xs))
plt.title('(a0,b0)=(0.5,1.0)')
plt.show()
plt.hist(sample3, bins=20, range=(-5.0, 5.0))
plt.plot(xs, mg(0.5, 0.5, 50)(xs))
plt.title('(a0,b0)=(0.5,0.5)')
plt.show()
```





混合正規分布で当てはめするが、精度はでない。

```
In [6]: from sklearn.mixture import GaussianMixture
gm1 = GaussianMixture(n_components=2, covariance_type='spherical')
gm1.fit([[x] for x in sample1])
print(gm1.weights_)
print(gm1.means_)
print(gm1.covariances_)
```

```
[ 0.51323679  0.48676321]
[[ 3.14557991]
 [-0.15162801]]
[ 0.76946106  0.92362133]
```

```
In [7]: gm2 = GaussianMixture(n_components=2, covariance_type='spherical')
gm2.fit([[x] for x in sample2])
print(gm2.weights_)
print(gm2.means_)
print(gm2.covariances_)
```

```
[ 0.4115779  0.5884221]
[[-0.55853118]
 [ 0.91471121]]
[ 0.36718427  0.64473308]
```

```
In [8]: gm3 = GaussianMixture(n_components=2, covariance_type='spherical')
gm3.fit([[x] for x in sample3])
print(gm3.weights_)
print(gm3.means_)
print(gm3.covariances_)
```

```
[ 0.45176392  0.54823608]
[[-0.25324619]
 [ 0.83559722]]
[ 0.87647464  0.69496225]
```

PyStanで混合正規分布を推定してみる

```
In [9]: model = """
data {
  int<lower=1> N;
  vector[N] Y;
}
parameters {
  real <lower=0, upper=1> a;
  real <lower=-5, upper=5> b;
}
model {
  for(n in 1:N) {
    target += log_sum_exp(
      log(a) + normal_lpdf(Y[n] | 0, 1),
      loglm(a) + normal_lpdf(Y[n] | b, 1)
    );
  }
}
"""
```

```
In [10]: data1 = {'N':100, 'Y':sample1.tolist()}
```

```
In [11]: fit1 = pystan.stan(model_code=model, data=data1, iter=1000, chains=4)
print(fit1)
```

INFO:pystan:COMPILING THE C++ CODE FOR MODEL anon_model_6bdc421632bfad283cf5b68dafff39e6 NOW.

Inference for Stan model: anon_model_6bdc421632bfad283cf5b68dafff39e6.

4 chains, each with iter=1000; warmup=500; thin=1;

post-warmup draws per chain=500, total post-warmup draws=2000.

	mean	se_mean	sd	2.5%	25%	50%	75%	97.5%	n_eff
Rhat									
a	0.49	1.5e-3	0.06	0.39	0.45	0.49	0.53	0.6	1463
1.0									
b	3.15	4.2e-3	0.16	2.81	3.03	3.15	3.26	3.45	1509
1.0									
lp__	-195.7	0.04	1.04	-198.5	-196.2	-195.4	-195.0	-194.7	859
1.0									

Samples were drawn using NUTS at Thu Jun 8 18:04:35 2017.

For each parameter, n_eff is a crude measure of effective sample size,

and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat=1).

```
In [12]: data2 = {'N':100, 'Y':sample2.tolist()}
fit2 = pystan.stan(model_code=model, data=data2, iter=1000, chains=4)
print(fit2)
```

INFO:pystan:COMPILING THE C++ CODE FOR MODEL anon_model_6bdc421632bfad283cf5b68daffff39e6 NOW.

Inference for Stan model: anon_model_6bdc421632bfad283cf5b68daffff39e6.

4 chains, each with iter=1000; warmup=500; thin=1;
post-warmup draws per chain=500, total post-warmup draws=2000.

	mean	se_mean	sd	2.5%	25%	50%	75%	97.5%	n_eff
Rhat									
a	0.53	0.01	0.27	0.02	0.31	0.57	0.75	0.94	354
1.02									
b	0.76	0.02	0.48	0.14	0.42	0.67	1.01	1.96	543
1.01									
lp__	-146.4	0.07	1.3	-150.0	-146.9	-146.0	-145.4	-145.1	335
1.0									

Samples were drawn using NUTS at Thu Jun 8 18:04:56 2017.

For each parameter, n_eff is a crude measure of effective sample size,
and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat=1).

```
In [13]: data3 = {'N':100, 'Y':sample3.tolist()}
fit3 = pystan.stan(model_code=model, data=data3, iter=1000, chains=4)
print(fit3)
```

INFO:pystan:COMPILING THE C++ CODE FOR MODEL anon_model_6bdc421632bfad283cf5b68daffff39e6 NOW.

Inference for Stan model: anon_model_6bdc421632bfad283cf5b68daffff39e6.

4 chains, each with iter=1000; warmup=500; thin=1;
post-warmup draws per chain=500, total post-warmup draws=2000.

	mean	se_mean	sd	2.5%	25%	50%	75%	97.5%	n_eff
Rhat									
a	0.44	9.5e-3	0.23	0.03	0.26	0.45	0.61	0.84	567
1.01									
b	0.64	0.01	0.28	0.24	0.44	0.59	0.81	1.33	397
1.01									
lp__	-146.9	0.04	1.1	-149.8	-147.3	-146.6	-146.1	-145.8	640
1.01									

Samples were drawn using NUTS at Thu Jun 8 18:05:17 2017.

For each parameter, n_eff is a crude measure of effective sample size,
and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat=1).

事後分布の可視化

```
In [14]: from scipy.misc import logsumexp
def lp(a,b,Y):
    lp = 0
    for y in Y:
        lp += logsumexp([np.log(1-a)-y*y/2, np.log(a)-(y-b)*(y-b)/2])
    return lp
```

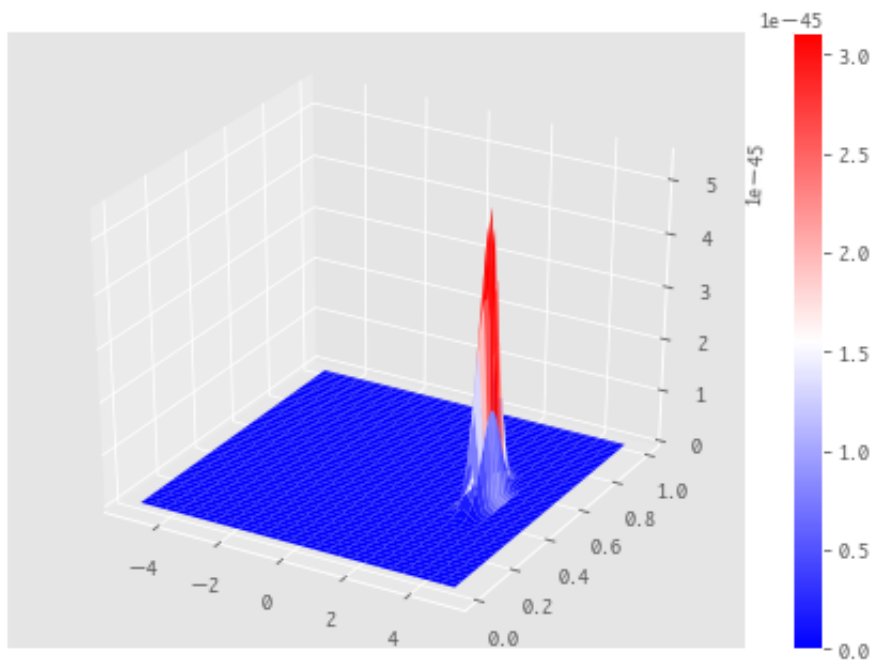
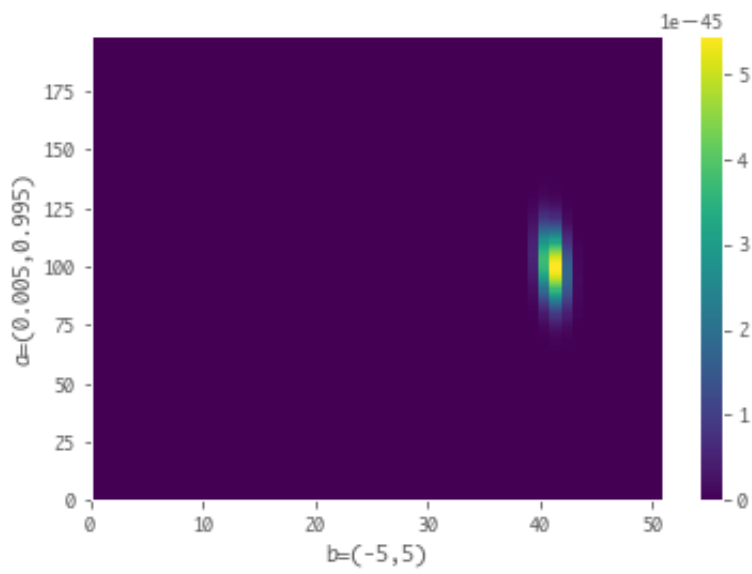
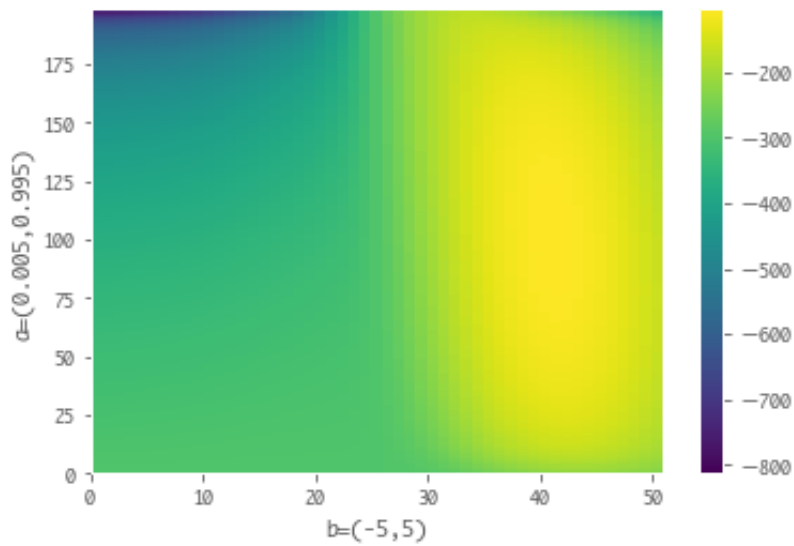
```
In [15]: from mpl_toolkits.mplot3d import Axes3D

def draw_graphs(sample):
    Z = np.array([[lp(a,b,sample) for b in np.linspace(-5, 5, 51)] for
    fig, ax = plt.subplots()
    im = ax.pcolor(Z)
    fig.colorbar(im, ax=ax)
    plt.xlabel('b=(-5,5)')
    plt.ylabel('a=(0.005,0.995)')
    plt.show()

    Ze = np.exp(Z)
    fig, ax = plt.subplots()
    im = ax.pcolor(Ze)
    fig.colorbar(im, ax=ax)
    plt.xlabel('b=(-5,5)')
    plt.ylabel('a=(0.005,0.995)')
    plt.show()

    fig = plt.figure()
    ax = Axes3D(fig)
    X, Y = np.meshgrid(np.linspace(-5, 5, 51), np.linspace(0.005, 0.995, 51))
    surf=ax.plot_surface(X,Y,Ze,cmap='bwr', linewidth=0)
    fig.colorbar(surf)
    plt.show()
```

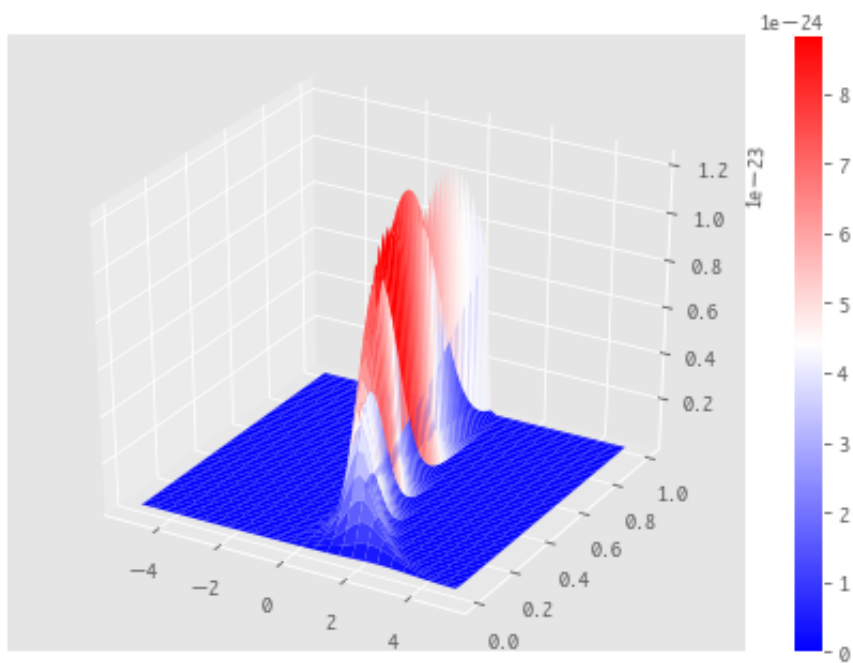
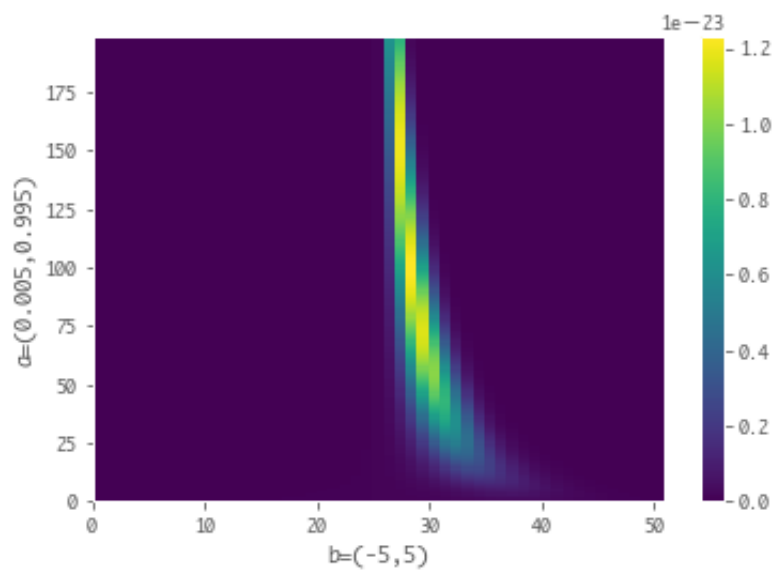
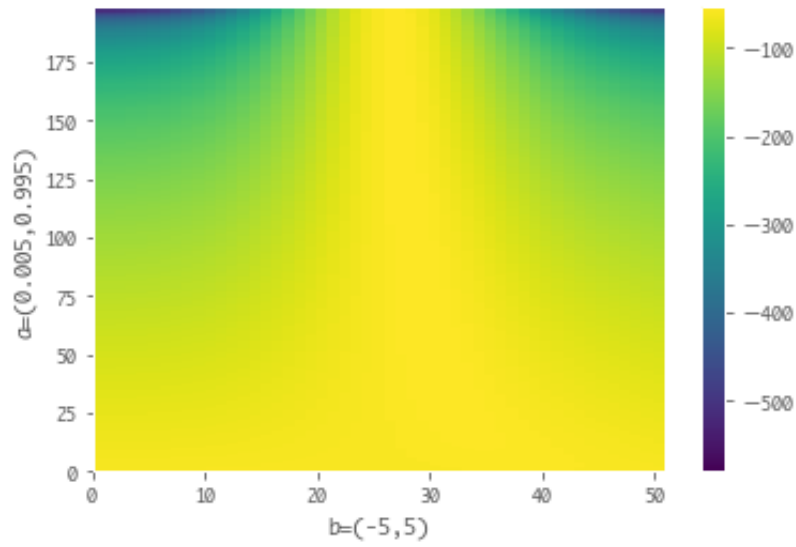
```
In [16]: draw_graphs(sample1)
```



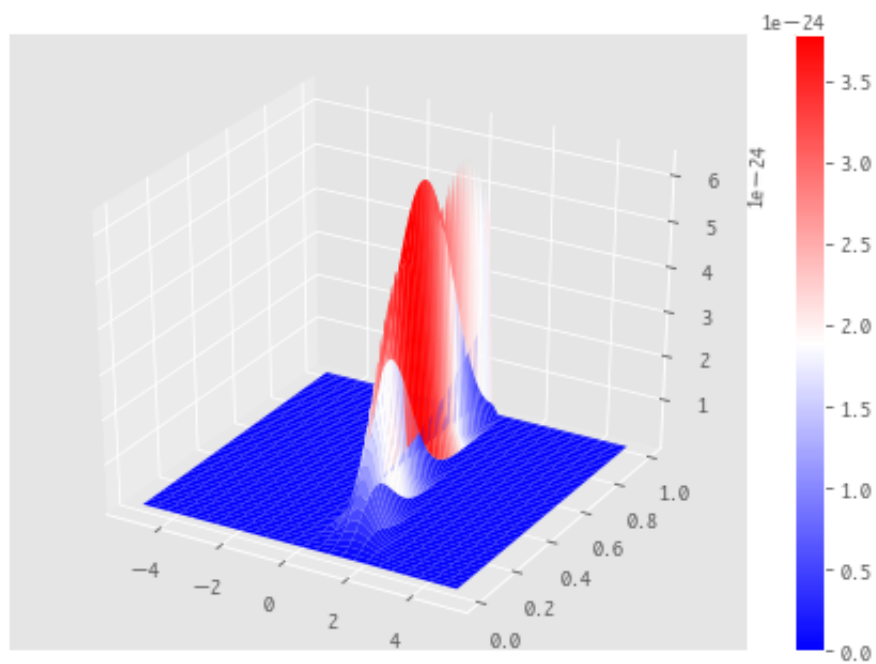
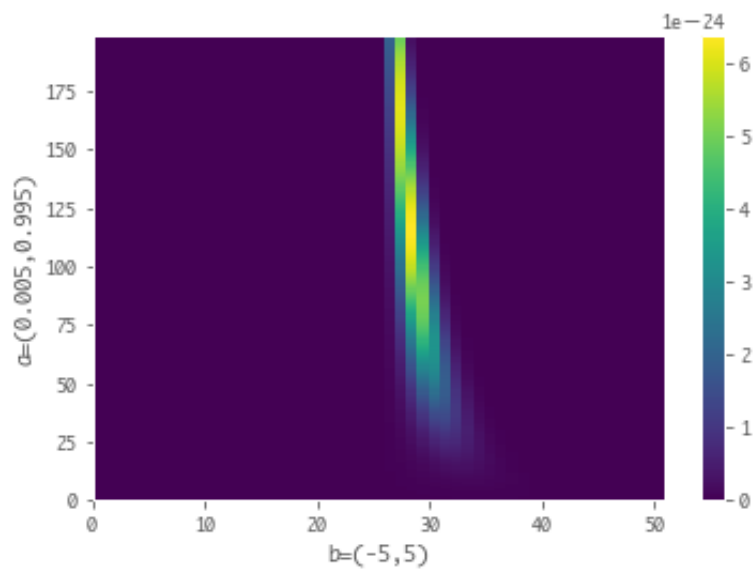
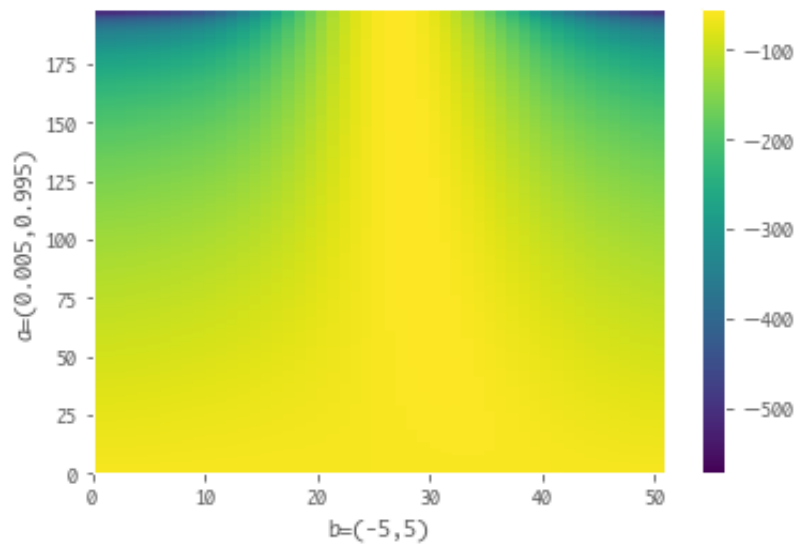
$$p(x|w) = (1 - a)\mathcal{N}(x) + a\mathcal{N}(x - b)$$

$a \simeq 0$ だと **b** の値は関係なくなる。

```
In [17]: draw_graphs(sample2)
```




```
In [18]: draw_graphs(sample3)
```



```
In [ ]:
```

In []: