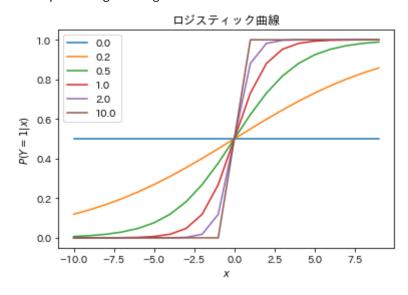
```
import copy
import japanize_matplotlib
%matplotlib inline
import matplotlib.pyplot as plt
import numpy as np
import scipy
from matplotlib.pyplot import imshow
from numpy.random import randn
from scipy import stats
import matplotlib.pyplot as plt
np.random.seed(42)
```

問題21 (b)

Out[3]: <matplotlib.legend.Legend at 0x7fd70cce2460>



問題22

```
N=100に対してp=2の場合
```

```
In [4]: 
N = 100

p = 2

X = np_random_randn(N, p)
```

```
X = np\_concatenate([np\_ones(N)\_reshape(N, 1), X], axis=1)
beta = np.random.randn(p+1)
y = np.zeros(N)
s = np.dot(X, beta)
prob = 1 / (1 + np.exp(s))
for i in range(N):
  if np.random.rand(1) > prob[i]:
    y[i] = 1
  else:
    y[i] = -1
beta
```

Out[4]: array([0.35778736, 0.56078453, 1.08305124])

```
In [5]:
         beta = np.inf
         gamma = np.random.randn(p + 1)
         while np.sum((beta - gamma) ** 2) > 0.001:
           beta = gamma.copy()
           s = np_{\bullet}dot(X, beta)
           v = np_exp(-s * y)
           u = y * v / (1 + v)
           w = v / (1 + v) ** 2
           z = s + u / w
           W = np.diag(w)
           gamma = np.dot(np.linalg.inv(X.T @ W @ X), np.dot(X.T @ W, z))
                                                                                ##
           print(gamma)
         beta
```

```
[-0.08319037 0.64260297 1.08943998]
       [0.46802113 0.75346512 0.56215532]
      [0.42157733 0.78281238 0.67917968]
      [0.42198408 0.78533849 0.68339308]
Out[5]: array([0.42157733, 0.78281238, 0.67917968])
      N=100に対してp=200の場合etaの各値が大きくなってoverflowが発生している
```

```
In [6]:
         N = 10
         p = 20
         X = np.random.randn(N, p)
         X = np.concatenate([np.ones(N).reshape(N, 1), X], axis=1)
         beta = np.random.randn(p+1)
         y = np_z zeros(N)
         s = np.dot(X, beta)
         prob = 1 / (1 + np.exp(s))
         for i in range(N):
           if np_random_rand(1) > prob[i]:
             y[i] = 1
           else:
             y[i] = -1
         beta = np.inf
         gamma = np_random_randn(p + 1)
         while np.sum((beta - gamma) ** 2) > 1e-10:
           beta = gamma.copy()
           s = np.dot(X, beta)
           v = np_exp(-s * y)
           u = y * v / (1 + v)
           w = v / (1 + v) ** 2
           z = s + u / (w + 1e-10) #divide by zero を防ぐ
           W = np.diag(w)
           gamma = np.dot(np.linalg.inv(X.T @ W @ X), np.dot(X.T @ W, z))
         # print(gamma)
         beta
```

```
<ipython-input-6-e171288a67a7>:19: RuntimeWarning: overflow encountered in exp v = np.exp(-s * y) <ipython-input-6-e171288a67a7>:20: RuntimeWarning: invalid value encountered in true_divide u = y * v / (1 + v) <ipython-input-6-e171288a67a7>:21: RuntimeWarning: invalid value encountered in true_divide w = v / (1 + v) ** 2
Out[6]: array([-455782., 11779920., -405240., -1183160., -2064004., 3737908., -721972., -3214680., -3661340., 1798388., 1121604., 1165528., -1881996., -1350324., 4851096., 381573., 7384400., 303920., -2144132., -1236972., 495416.])
```

問題24

```
In [7]: #Linuxでの実行(Windows, Google Colaboratory不可)
import numpy as np
import matplotlib.pyplot as plt
import glmnet_python
import sys
from cvglmnet import cvglmnet
from cvglmnetCoef import cvglmnetCoef
from cvglmnetPlot import cvglmnetPlot
```

```
In [8]:
         # Linuxマシンのフォルダに"breastcancer.csv"をおく
         x = np.loadtxt("../data/breastcancer.csv"
                 delimiter=",", skiprows=1, usecols=range(1000))
         y = np.loadtxt("../data/breastcancer.csv",
                 delimiter=",", skiprows=1, dtype="unicode", usecols=1000)
         n = len(y)
         yy = np.ones(n)
         for i in range(n):
           if y[i] == "control":
             yy[i] = 1
           else:
             yy[i] = -1
         fit1 = cvglmnet(x=x.copy(), y=yy.copy(), ptype="deviance", family="binomial")
         fit2 = cvglmnet(x=x.copy(), y=yy.copy(), ptype="class", family="binomial")
         beta = cvglmnetCoef(fit1)
         np.sum(beta != 0)
```

/opt/conda/lib/python3.9/site-packages/numpy/lib/scimath.py:288: RuntimeWarning: divide by zero encountered in log return nx.log(x)

Out[8]: 35

非ゼロ要素の個数は35

問題26

```
def soft_th(lam, x):
    return np.sign(x) * np.maximum(np.abs(x) - lam, np.zeros(1))

def linear_lasso(X, y, lam=0, beta=None):
    n, p = X.shape
    if beta is None:
        beta = np.zeros(p)
```

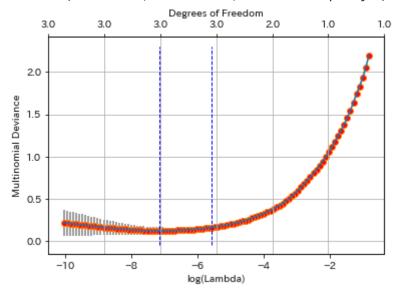
```
X, y, X_bar, X_sd, y_bar = centralize(X, y) #中心化(下記参照)
  eps = 1
  beta_old = copy.copy(beta)
  while eps > 0.00001: #このループの収束を待つ
    for j in range(p):
      r = y
      for k in range(p):
        if j != k:
           r = r - X[:, k] * beta[k]
      z = (np_*dot(r, X[:, j]) / n) / (np_*dot(X[:, j], X[:, j]) / n)
      beta[j] = soft\_th(lam, z)
    eps = np.linalg.norm(beta - beta_old, 2)
    beta_old = copy.copy(beta)
  beta = beta / X_sd # 各変数の係数を正規化前のものに戻す
  beta_0 = y_bar - np_dot(X_bar, beta)
  return beta, beta_0
def centralize(X0, y0, standardize=True):
  X = copy.copy(X0)
  y = copy_copy(y0)
  n, p = Xshape
  X_bar = np_zeros(p)
                           # Xの各列の平均
  X_sd = np_zeros(p)
                               #Xの各列の標準偏差
  for j in range(p):
    X_bar[j] = np_mean(X[:, j])
    X[:, j] = X[:, j] - X_bar[j]
                             #Xの各列の中心化
    X_sd[j] = np_std(X[:, j])
    if standardize is True:
      X[:, j] = X[:, j] / X_sd[j] # Xの各列の標準化
  if np.ndim(y) == 2:
    K = y_shape[1]
                               # yの平均
    y_bar = np_zeros(K)
    for k in range(K):
      y_bar[k] = np_mean(y[:, k])
      y[:, k] = y[:, k] - y_bar[k] #yの中心化
                         # yがベクトルの場合
  else:
    y_bar = np_mean(y)
    y = y - y_bar
  return X, y, X_bar, X_sd, y_bar
def W_linear_lasso(X, y, W, lam=0):
  n, p = X.shape
  X_bar = np_zeros(p)
  for k in range(p):
    X_bar[k] = np_sum(np_dot(W, X[:, k])) / np_sum(W)
    X[:, k] = X[:, k] - X_bar[k]
  y_bar = np_sum(np_dot(W, y)) / np_sum(W)
  y = y - y_bar
  L = np_linalq_cholesky(W) #
\# L = np.sqrt(W)
  u = np.dot(L, y)
  V = np.dot(L, X)
  beta, beta_0 = linear_lasso(V, u, lam)
  beta_0 = y_bar - np_dot(X_bar, beta)
  return beta_0, beta
def multi_lasso(X, y, lam):
  n, p = X.shape
  K = len(set(y))
  beta = np.ones((K, p))
  gamma = np_zeros((K, p))
  while np_linalg_norm(beta - gamma, 2) > 0.1:
    gamma = beta_copy()
    for k in range(K):
      r = 0
```

```
for h in range(K):
                   if k != h:
                     r = r + np.exp(np.dot(X, beta[h, :]))
                 v = np.exp(np.dot(X, beta[k, :])) / r
                 u = (y == k) - v / (1 + v)
                 w = v / (1 + v) ** 2
                 z = np_dot(X, beta[k, :]) + u / w
                 beta_0, beta_1 = W_{linear_lasso}(X[:, range(1, p)],
                                   z, np.diag(w), lam=lam)
                 beta[k, :] = np.block([beta_0, beta_1]).copy()
            for j in range(p):
               med = np_median(beta[:, j])
               for h in range(K):
                 beta[h, j] = beta[h, j] - med
            return beta
In [10]:
          # Linuxでの実行(Windows, Google Colaboratory不可)
          import numpy as np
          import matplotlib.pyplot as plt
          import glmnet_python
          from glmnet import glmnet
          import sys
          from cyglmnet import cyglmnet
          from cvglmnetCoef import cvglmnetCoef
          from cyglmnetPlot import cyglmnetPlot
In [11]:
          from sklearn.datasets import load_iris
          iris = load_iris()
          X = np.array(iris["data"])
          y = np.array(iris["target"], dtype="float64")
          cvfit3 = cvglmnet(x=X_copy(), y=y_copy(),
                    ptype="deviance", family="multinomial")
          lam_min = cvfit3["lambda_min"]
          beta = cvglmnetCoef(cvfit3)
          print(lam_min)
          print(beta)
         [0.00077805]
         [array([[ 9.2302786 ],
             [ 0.
                     1,
             [2.6568387],
             [-2.73371535],
             [-0.46547548]]), array([[5.62976947],
             [0.71170415],
             ١٥.
                    ],
             ۲٥.
                     ],
             [0.
                     ]]), array([[-12.78950282],
             [ 0.
                     ],
             [-2.20996807],
             [ 3.35592123],
             [ 7.69194562]])]
         /opt/conda/lib/python3.9/site-packages/numpy/lib/scimath.py:288: RuntimeWarning: divide by
         zero encountered in log
          return nx.log(x)
         /opt/conda/lib/python3.9/site-packages/glmnet_python-0.2.0-py3.9.egg/glmnet_python/cvm
         ultnet.py:72: RuntimeWarning: invalid value encountered in multiply
          ly = bigY*scipy.log(bigY)
In [12]:
          fig3 = plt.figure()
```

```
cvglmnetPlot(cvfit3)
fig3.savefig("img3.png")
```

/opt/conda/lib/python3.9/site-packages/glmnet_python-0.2.0-py3.9.egg/glmnet_python/cvgl mnetPlot.py:113: MatplotlibDeprecationWarning: Case-insensitive properties were deprecated in 3.3 and support will be removed two minor releases later

```
ax2.set(XLim=xlim1, XTicks = atdf, XTickLabels = prettydf)
```



```
In [13]:
          def table_count(m, u, v):
             n = u.shape[0]
             count = np.zeros([m, m])
             for i in range(n):
               count[int(u[i]), int(v[i])] += 1
             return(count)
          K = 3
          p = 5
          n = 150
          gamma = np.zeros((K, p))
          for k in range(K):
             for j in range(p):
               gamma[k, j] = np.sum(beta[k][j])
          v = np_zeros(n)
          for i in range(n):
             max_value = -np.inf
             for k in range(K):
               value = gamma[k, 0] + np.dot(gamma[k, range(1, p)], X[i, :]) #空欄
               if value > max_value:
                 v[i] = k
                 max_value = value
          table_count(3, y, v)
```

問題28

```
ln [14]:

def poisson_lasso(X, y, lam):
    p = X.shape[1] # plますべて1の列を含んでいる
    beta = np.random.randn(p)
    gamma = np.random.randn(p)
```

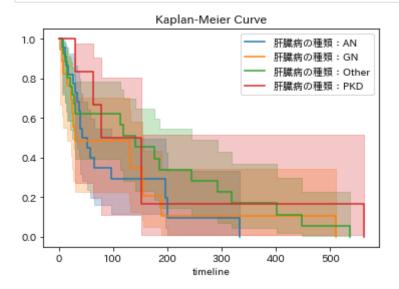
```
beta = gamma
               s = np.dot(X, beta)
                                            #デザイン行列と重みの積
               w = np_exp(s)
                                          #exp
               u = y - w
                                        #uの計算
               z = s + u / w
               gamma_0, gamma_1 = W_linear_lasso(X[:, range(1, p)],
                                  z, np.diag(w), lam)
               gamma = np.block([gamma_0, gamma_1]).copy()
               print(gamma)
             return gamma
In [15]:
          N = 100 # lambdaの値が小さいと発散して,推定値が出ないことがある。
          p = 3
          X = np.random.randn(N, p)
          X = np_{\text{-}}concatenate([np_{\text{-}}ones(N)_{\text{-}}reshape(N, 1), X], axis=1)
          beta = np.random.randn(p + 1)
          s = np.dot(X, beta)
          y = np.random.poisson(lam=np.exp(s))
          print(beta)
          [-0.04946371 0.67481949 -1.12272202 0.38240975]
In [16]:
          poisson_lasso(X,y,2)
          [ 2.21953756 -0.31371836 0.35286932 -0.
                                                           ]
          [ 1.64649324 -0.
                                0.
                                        0.
          [ 1.17841304 0.
                               -0.
                                        0.
                                              ]
          [1.02784931 0.
                               -0.
                                        0.
                                              ]
          [ 1.01530996 0.
                               -0.
                                        0.
                                              ]
          [ 1.01523068 0.
                               -0.
                                        0.
                                              ]
                                      , -0.
Out[16]: array([ 1.01523068, 0.
                                             , 0.
                                                      ])
         問題29
In [17]:
          import pandas as pd
          from lifelines import KaplanMeierFitter
In [18]:
          df = pd_read_csv("../data/kidney.csv")
          df_drop(df_columns[0], axis=1, inplace=True)
          df
               id time
                                          disease
Out[18]:
                        status
                                age sex
                                                    frail
           0
                1
                     8
                             1
                                 28
                                        1
                                             Other
                                                     2.3
           1
                1
                    16
                             1
                                 28
                                        1
                                             Other
                                                     2.3
           2
               2
                    23
                             1
                                 48
                                        2
                                               GN
                                                     1.9
           3
               2
                    13
                             0
                                 48
                                        2
                                               GN
                                                     1.9
           4
               3
                    22
                             1
                                 32
                                        1
                                             Other
                                                     1.2
                                  ...
                     ...
                             ...
                                       ...
                                                     ...
              36
                    16
                             0
                                 42
                                        2
                                             Other
                                                     0.7
          72
              37
                     6
                             0
                                 52
                                        2
                                              PKD
                                                     2.1
```

while np.sum((beta - gamma) ** 2) > 0.0001:

| | id | time | status | age | sex | disease | frail |
|----|----|------|--------|-----|-----|---------|-------|
| 73 | 37 | 78 | 1 | 52 | 2 | PKD | 2.1 |
| 74 | 38 | 63 | 1 | 60 | 1 | PKD | 1.2 |
| 75 | 38 | 8 | 0 | 60 | 1 | PKD | 1.2 |

76 rows × 7 columns

```
| kmf = KaplanMeierFitter() | ax = None | for name, group in df.groupby("disease"): | kmf.fit(group["time"], event_observed=group["status"], | label="肝臓病の種類:" + str(name)) | if ax is None: | ax = kmf.plot() | else: | ax = kmf.plot(ax=ax) | plt.title("Kaplan-Meier Curve") | plt.show()
```



問題32

(a)

```
In [20]:
```

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import glmnet_python
from glmnet import glmnet
from glmnetCoef import glmnetCoef
import sys
from cvglmnet import cvglmnet
from cvglmnetCoef import cvglmnetCoef
from cvglmnetPlot import cvglmnetPlot
import os
import rpy2.robjects as robjects
from rpy2.robjects import pandas2ri
from rpy2.robjects.packages import importr
from lifelines import KaplanMeierFitter
```

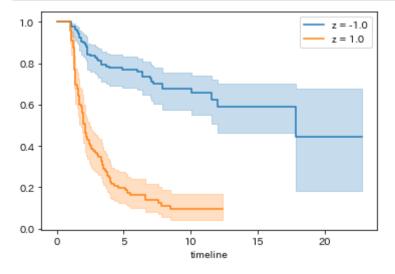
```
base = importr("base")
base_load("../data/LymphomaData.rda")
w = robjects_globalenv["patient.data"]
X = np_array(w[0]).T
y = np_array(w[1])
delta = np_array(w[2])
v = np_concatenate([y_reshape(240, 1), delta_reshape(240, 1)], axis=1)
fit = glmnet(x=X_copy(), y=v_copy(), family="cox")
beta = glmnetCoef(fit, s=np_float64([0.119787]))
print(np_sum(beta != 0))
```

Warning: Cox model has no intercept!

29個の要素が非ゼロとなっている

(b)

```
In [22]:
          z = np.sign(np.dot(X, beta))
          df2 = pd.DataFrame(
             np.concatenate(
               [y_reshape(240, 1), delta_reshape(240, 1), z], axis=1))
          df2.columns = ["time", "status", "sign"]
          fig = plt.figure()
          kmf = KaplanMeierFitter()
          ax = None
          for name, group in df2 groupby ("sign"):
             kmf_fit(group["time"], event_observed=group["status"],
                 label="z = " + str(name))
             if ax is None:
               ax = kmf.plot()
             else:
               ax = kmf_plot(ax=ax)
```



```
In [23]: #plt.title("Kaplan-Meier Curve") #fig.savefig("img7.png")
```

問題32

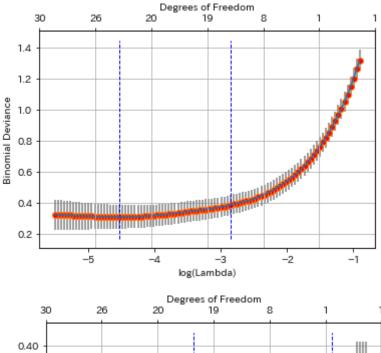
In [24]:

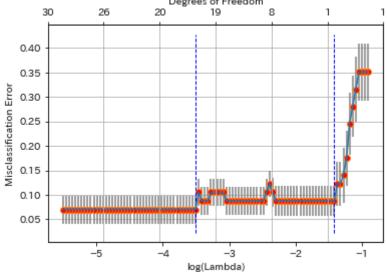
import scipy, importlib, pprint, matplotlib.pyplot as plt, warnings from glmnet import glmnet; from glmnetPlot import glmnetPlot from glmnetPrint import glmnetPrint; from glmnetCoef import glmnetCoef; from glmnetPredict in
from cvglmnet import cvglmnet; from cvglmnetCoef import cvglmnetCoef
from cvglmnetPlot import cvglmnetPlot; from cvglmnetPredict import cvglmnetPredict

```
In [25]:
          df = pd_read_csv("../data/leukemia_big.csv", names=range(72))_T
In [26]:
           df.head()
                              1
                                            2
                                                                        4
                                                                                       5
                                                                                                     6
Out[26]:
               0
                                                          3
             ALL
                  -1.533621735
                                -1.235672925 -0.333982875
                                                              0.488702121 -1.300893264 -1.682668245
          1 ALL -0.867609617
                                 -1.275500531
                                                0.375926519
                                                              0.444010978
                                                                            -1.229659818
                                                                                          -1.642071845
          2 ALL -0.433171907
                                 -1.184492247
                                               -0.459196002
                                                              0.436263539
                                                                            -1.325882367 -1.407263942
          3 ALL
                  -1.671903187 -1.596423985
                                               -1.422571436
                                                              0.193352945
                                                                            -1.818328843
                                                                                          -1.744469301
            ALL -1.187689367 -1.335255683 -0.797492909
                                                              0.235631518
                                                                             -1.311205961 -1.654380542
         5 rows × 7129 columns
In [27]:
           y = df.iloc[:,0].apply(lambda x: 0 if x == "ALL" else 1)
          X = df_{iloc}[:,1:]
In [28]:
          from sklearn.model_selection import train_test_split
In [29]:
          X_train, X_test, y_train, y_test = train_test_split(X,y,train_size=0.8, random_state=42, stratif
In [30]:
          X_train = X_train.values.astype('float')
          X_test = X_test.values.astype('float')
          y_train = y_train.values.astype('float')
          y_test = y_test.values.astype('float')
In [31]:
           # Linuxマシンのフォルダに"breastcancer.csv"をおく
          fit1 = cvglmnet(x=X_train_copy(), y=y_train_copy(), ptype="deviance", family="binomial")
          fit2 = cvglmnet(x=X_train_copy(), y=y_train_copy(), ptype="class", family="binomial")
In [32]:
           fig = plt.figure()
           cvglmnetPlot(fit1)
           fig2 = plt.figure()
           cvglmnetPlot(fit2)
          /opt/conda/lib/python3.9/site-packages/glmnet_python-0.2.0-py3.9.egg/glmnet_python/cvgl
          mnetPlot.py:113: MatplotlibDeprecationWarning: Case-insensitive properties were deprecated in
```

3.3 and support will be removed two minor releases later ax2.set(XLim=xlim1, XTicks = atdf, XTickLabels = prettydf)
/opt/conda/lib/python3.9/site-packages/glmnet_python-0.2.0-py3.9.egg/glmnet_python/cvgl mnetPlot.py:113: MatplotlibDeprecationWarning: Case-insensitive properties were deprecated in 3.3 and support will be removed two minor releases later

ax2.set(XLim=xlim1, XTicks = atdf, XTickLabels = prettydf)





In [33]: lam_min = fit1['lambda_min']
 print(lam_min)

[0.01077568]

上がlogloss, 下がaccuracy基準での λ ごとの値 今回はlogloss基準でモデル選択を行う CVの結果lambda=0.01077568が最適である

In [34]: pred = cvgImnetPredict(fit1, X_test, s = lam_min, ptype='class')

In [35]: from sklearn.metrics import accuracy_score accuracy_score(pred,y_test)

Out[35]: 1.0

従って正答率100%で完璧に予測することができた...