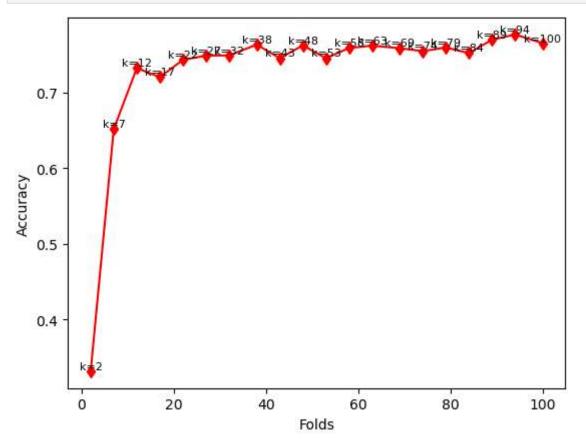
```
import numpy as np
In [11]:
         import pandas as pd
          import itertools
          import matplotlib.pyplot as plt
In [12]: headers = [
            "Label", "Alchohol", "Malic Acid", "Ash",
            "Alcalinity of Ash", "Magnesium", "Total Phenols",
            "Flavanoids", "Nonflavanoid Phenols", "Proanthocyanins",
            "Color Intensity", "Hue", "OD280/OD315 of Diluted Wines", "Proline"
         df = pd.read_csv("../data/HW2/wine.data", names=headers)
In [13]: y = np.array(df["Label"], dtype=np.float32)
         X = np.array(df.iloc[:,1:], dtype=np.float32)
         print(X.shape, y.shape)
         (178, 13) (178,)
In [14]: def Euclid(x, y):
           return np.sum(np.square(y - x))
         def NN(dist_fn, X, y, query):
            idx = np.argmin([dist_fn(r, query) for r in X])
            return y[idx]
         def K_Fold_CV(X, y, k):
           n = len(y)
           errs = []
            conf_data = []
            for i in range(k):
             lo, hi = (i * n) // k, ((i + 1) * n) // k
             hold_X, hold_y = X[lo:hi], y[lo:hi]
             tr_X, tr_y = np.vstack((X[:lo,:], X[hi:,:])), np.hstack((y[:lo], y[hi:]))
             preds = [NN(Euclid, tr_X, tr_y, query) for query in hold_X]
             if k == n:
                conf_data.append((preds[0], hold_y.item()))
             mismatch = np.not_equal(preds, hold_y)
             err = np.sum(mismatch) / len(mismatch)
             errs.append(err)
           if k == n:
             return np.mean(errs), conf_data
             return np.mean(errs)
```

## 11a,b

```
In [15]: def Do_LOOCV(X, y):
    err, conf_data = K_Fold_CV(X, y, len(y))
    conf_mat = np.zeros((3, 3))
```

```
for pred, lab in conf_data:
              conf_mat[int(pred)-1,int(lab)-1] += 1
            return conf_mat, err
         conf mat, err = Do LOOCV(X, y)
         print(f"=== Accuracy")
         print(1 - err)
         print("=== Confusion Matrix")
         print(conf_mat)
         === Accuracy
         0.7696629213483146
         === Confusion Matrix
         [[52. 5. 3.]
          [ 3. 54. 14.]
          [ 4. 12. 31.]]
In [16]: folds = np.linspace(2, 100, 20).astype(np.int32)
         err = [1 - K_Fold_CV(X, y, fold) for fold in folds]
         plt.plot(folds, err, "-dr")
         plt.ylabel("Accuracy")
         plt.xlabel("Folds")
         for f, e in zip(folds, err):
            plt.annotate(f"k={f}", (f, e), ha="center", va="bottom", fontsize=8)
         plt.show()
         best = np.argmax(err)
         print("=== Best")
         print(f"k={folds[best]}, acc={err[best]}")
```



=== Best k=94, acc=0.7765957446808511

## 11c

```
In [17]: ranges = np.max(X, axis=0, keepdims=True) - np.min(X, axis=0, keepdims=True)
    X2 = (X - np.min(X, axis=0, keepdims=True)) / ranges

In [18]: conf_mat, err = Do_LOOCV(X2, y)
    print(f"=== Accuracy")
    print(1 - err)
    print("=== Confusion Matrix")
    print(conf_mat)

    print("The Normaliation helped a significant amount, increasing accuracy from 0.770 to
    === Accuracy
    0.949438202247191
    === Confusion Matrix
    [[59.     5.     0.]
        [          0.     62.     0.]
        [          0.     4.     48.]]
        The Normaliation helped a significant amount, increasing accuracy from 0.770 to 0.949
```

## 12

```
In []:
```