```
def decisiontree( X, Y, method = 'CART'):
  if method == 'CART':
method = 'CART'를 입력하면 CART로 one-level tree가 되고, 그 외 아무거나 입력 시
C4.5가 되도록 함수를 정의했습니다.
```

□ cart

```
if method == 'CART':
      A=[]
      B = np.unique(Y)
      for i in range(0,len(X.iloc[0])):
   if type(X.iloc[0,i]) == str:
             options = []
for L in range(1,len(np.unique(X.iloc[:,i]))):
                for subset in itertools.combinations(np.unique(X,iloc[:,i]),L):
    subset = list(subset)
             options.append(subset)
A.append(lén(options))
         else:
             A.append(len(X.iloc[:,i].sort_values().unique()))
      c = np.max(A)

K = pd.DataFrame(index = range(0,len(X,iloc[0])), columns = range(0,c))

I = pd.DataFrame(index = range(0,len(X,iloc[0])), columns = range(0,c))

number =pd.DataFrame(index = range(0,len(X,iloc[0])), columns = range(0,c))
      for i in range(0,len(X,iloc[0])):
Z=[]
         if type(X.iloc[0,i]) == str:
    options = []
    for L in range(1,len(np.unique(X.iloc[:,i]))):
        for subset in itertools.combinations(np.unique(X.iloc[:,i]),L):
        subset = list(subset)
                    options.append(subset)
             options.append(subset)
for j in range(0,len(options)):
  condition = X.iloc[:,i].isin(options[j])
| = len(Y[condition==True][Y[condition==True]==np.unique(Y)[1]])
g = len(Y[condition==True])/len(Y)*gini(Y[condition==True]) + len(Y[condition==False])/len(Y)*gini(Y[condition==False])
                  s = gini(Y) - g
                s = gin(t) - g

K[j][i] = s

[[j][i] = np.array([options[j], |ist(np.delete(np.unique(X.iloc[:,i]).np.where(np.unique(X.iloc[:,i])==options[j])))]).flatten().tolist()

number[j][i] = [[l,len(Y[condition == True])-l ], [len(Y[Y==np.unique(Y)[i]])-l, len(Y[Y==np.unique(Y)[0]])-(len(Y[condition == True])-l)]]
             for j in range(0,len(X.iloc[:,i].sort_values().unique())-1):
z1 = X.iloc[:,i].sort_values().unique()[i]
z2 = X.iloc[:,i].sort_values().unique()[i+1]
Z.append((z1+z2)/2)
                 for k in range(0,len(Z)):
                    condition = X.iloc[:,i]<=Z[k]
l = len(Y[condition==True][Y[condition==True]==np.unique(Y)[1]])</pre>
                     g = len(Y[condition==True])/len(Y)*gini(Y[condition==True]) + len(Y[condition==False])/len(Y)*gini(Y[condition==False])
                      = aini(Y) - a
                    K[k][i] = s
|[k][i] = Z[k]
                     number[k][i] = [[l,len(Y[condition == True]-l)], [len(Y[Y==np.unique(Y)[i]])-l, len(Y[Y==np.unique(Y)[0]])-(len(Y[condition == True])-l)]]
```

cart로 진행시 위와 같은 코드로 결과가 도출되고,

```
else:
            A=[]
            B = np.unique(Y)
             for i in range(0,len(X,iloc[0])):
                  if type(X.iloc[0,i]) == str:
                        A.append(1)
                   else:
                         A.append(len(X.iloc[:,i].sort_values().unique()))
            c = np.max(A)
            K = pd.DataFrame(index = range(0, len(X.iloc[0])), columns = range(0,c))
             I = pd.DataFrame(index = range(0,len(X.iloc[0])), columns = range(0,c))
            number =pd.DataFrame(index = range(0,len(X.iloc[0])), columns = range(0,c))
             e = len(Y[Y==np.unique(Y)[1]])/len(Y)
             entropyY = -e*np.log2(e) -(1-e)*np.log2(1-e)
             for i in range(0,len(X.iloc[0])):
                  if type(X.iloc[0,i]) == str:
                        p=[]
                         n=\Gamma 1
                         num=[]
                         for j in range(0, len(np.unique(X.iloc[:,i])));
                              condition = (X.iloc[:,i]==np.unique(X.iloc[:,i])[j])
                                I = len(Y[condition==True][Y[condition==True]==np.unique(Y)[1]])
                               p1 = I/len(Y[condition==True])
                               p.append((-p1*np.log2(p1) - (1-p1)*np.log2(1-p1))*(len(Y[condition==True])/(len(Y))))
                               \label{eq:nappend} n.append(-(len(Y[condition==True])/len(Y))*np.log2(len(Y[condition==True])/len(Y)))
                                num.append([I,len(Y[condition == True])-I ])
                         K[0][i] = (entropyY-np.sum(p))/np.sum(n)
                         [[0][i] = np.unique(X.iloc[:,i])
                         number[0][i] = num
                   else:
                         Z=[]
                         for j in range(0,len(X.iloc[:,i].sort_values().unique())-1):
                              z1 = X.iloc[:,i].sort_values().unique()[j]
                               z2 = X.iloc[:,i].sort_values().unique()[j+1]
                               Z.append((z1+z2)/2)
                               for k in range(0, len(Z)):
                                      condition = X.iloc[:,i]<=Z[k]
                                       I = len(Y[condition==True][Y[condition==True]==np.unique(Y)[1]])
                                      p1 = len(Y[condition==True][Y[condition==True]==np.unique(Y)[1]])/len(Y[condition==True])
                                      g = len(Y[condition==True])/len(Y)*(-p1*np.log2(p1)) + len(Y[condition==False])/len(Y)*(-(1-p1)*np.log2(1-p1)) + len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/len(Y[condition==False])/l
                                      n = -len(Y[condition == True])/len(Y) * np. log2(len(Y[condition == True])/len(Y)) - (1-len(Y[condition == True])/len(Y)) * np. log2(len(Y[condition == True])/len(Y[condition ==
                                      K[k][i] = (entropyY - g)/n
                                      I[k][i] = Z[k]
                                      number[k][i] = [[I,len(Y[condition == True]-I)], [len(Y[Y==np.unique(Y)[1]])-I, len(Y[Y==np.unique(Y)[0]])-(len(Y[Y=np.unique(Y)[0]])-I)]
```

C4.5는 위와 같은 코드를 따라 결과를 도출합니다.

```
print('tree structure (', method, ')')
 if len(Y[Y==np.unique(Y)[1]]) >= len(Y)/2:
  print( 'node 1 : yes (', len(Y[Y==np.unique(Y)[1]]), ', ', len(Y) - len(Y[Y==np.unique(Y)[1]]), ')')
else:
  print( 'node 1 : no (', len(Y[Y==np.unique(Y)[1]]),',', len(Y) - len(Y[Y==np.unique(Y)[1]]), ')')
 for i in range(0,len(|[idx_c][idx_r])):
  if number[idx_c][idx_r][i][0]>= number[idx_c][idx_r][i][1];
    print(f'node {i+2} :' 'class = ', I[idx_c][idx_r][i], ' yes (',number[idx_c][idx_r][i][0], ',' , number[idx_c
   else!
    print(f'node {i+2} :' 'class = ', ![idx_c][idx_r][i], ' no (',number[idx_c][idx_r][i][0], ',' , number[idx_c]
 if number[idx_c][idx_r][0][0]>= number[idx_c][idx_r][0][1]:
  for i in range(0, len(Y)):
    if X.iloc[:,idx_r][i] == I[idx_c][idx_r][0]:
      Y2.append(np.unique(Y)[1])
    else:
      Y2.append(np.unique(Y)[0])
else:
    if X.iloc[:,idx_r][i] == I[idx_c][idx_r][1]:
      Y2.append(np.unique(Y)[1])
    else:
      Y2.append(np.unique(Y)[0])
confusion_tst = confusion_matrix(Y,Y2)
accu_tst = 0
for i in range(len(np.unique(Y))):
  accu_tst = accu_tst + confusion_tst[i][i]
accuracy_tst = accu_tst / X.shape[0]
print('\m\nconfusion matrix (train)')
print('---
print('
                predicted class \mathfrak{Wn} Actual 1 ' ,confusion_tst[0], '\mathfrak{Wn} class 2 ', confusion_tst[1])
for i in range(2,len(np.unique(Y)) ):
 print(f' {i+1} ', confusion_tst[i])
print('model summary')
print('-----
print('Overall accuracy = ' ,accuracy_tst)
위의 코드를 통해 결과도출 까지 이루어지도록 작성했습니다.
```

출력 결과는 다음과 같습니다.

```
□cart
tree structure ( CART )
node 1: no (711, 1490)
node 2 :class = Female yes ( 344 , 126 )
node 3 :class = Male no ( 367 , 1364 )
confusion matrix (train)
      predicted class
 Actual 1 [1364 126]
 class 2 [367 344]
model summary
Overall accuracy = 0.7760109041344844
□C4.5
tree structure ( C4_5 )
node 1: no (711, 1490)
node 2 :class = Female yes ( 344 , 126 )
node 3 :class = Male no ( 367 , 1364 )
confusion matrix (train)
        predicted class
 Actual 1 [1364 126]
 class 2 [367 344]
model summary
Overall accuracy = 0.7760109041344844
둘 다 동일하게, 성별을 기준으로 분류가 되는 결과가 나왔습니다.
```