

# Lab2 实验报告

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## part\_1 传统机器学习

### 1. 决策树

① 根据训练数据构建决策树并在测试集上测试。构建决策树的过程主体是一个递归的过程：

○ 读入数据 → 生成节点 → 判断节点是否可以划分属性 → 选择最优划分 → 生成分支  
递归进行。

① 判断部分（出口）：若为空则赋值 `Null`；若样本都属于类别C则标记为C类叶节点。判断结束本次递归结束。

② 选择最优划分部分的依据为信息增益，依据实验文档给出的公式计算。

③ 连续值处理采用二分法离散化。

② 具体过程：

① 计算信息熵：

```
def entropy(y):
    classes, counts = np.unique(y, return_counts=True)
    probabilities = counts / len(y)
    return -np.sum(probabilities * np.log2(probabilities))
```

② 计算信息增益（二分法处理连续值）：

```
# 信息增益
def info_gain(self, X, y, feature, threshold):
    left_idx = X[:, feature] < threshold
    right_idx = X[:, feature] ≥ threshold
    left_entropy = self.entropy(y[left_idx])
    right_entropy = self.entropy(y[right_idx])
    return self.entropy(y) - np.mean(left_idx) * left_entropy -
    np.mean(right_idx) * right_entropy
```

③ 遍历分割点选择最佳分割：

#### # 选择最佳分割点

```
def best_split(self, X, y):
    best_feature, best_threshold, best_info_gain = None, None, -
np.inf
    n_features = X.shape[1]
    for feature in range(n_features):
        thresholds = np.unique(X[:, feature])
        for threshold in thresholds:
            ig = self.info_gain(X, y, feature, threshold)
            if ig > best_info_gain:
                best_feature, best_threshold, best_info_gain =
feature, threshold, ig
    return best_feature, best_threshold, best_info_gain
```

#### 4 构建决策树

#### # 递归构建决策树

```
def build_tree(self, X, y, max_depth, min_samples_leaf):
    if len(y) == 0:
        return {'threshold': None}
    if max_depth == 0 or len(y) < min_samples_leaf:
        return {'threshold': np.argmax(np.bincount(y))}
    feature, threshold, info_gain = self.best_split(X, y)
    # 信息增益为0
    if info_gain == 0:
        return {'threshold': np.argmax(np.bincount(y))}
    left_idx = X[:, feature] < threshold
    right_idx = X[:, feature] ≥ threshold
    left_tree = self.build_tree(X[left_idx], y[left_idx],
max_depth-1, min_samples_leaf)
    right_tree = self.build_tree(X[right_idx], y[right_idx],
max_depth-1, min_samples_leaf)
    return {'feature': feature,
            'threshold': threshold,
            'left_tree': left_tree,
            'right_tree': right_tree}
```

#### 5 做出预测

```
def predict(self, X):
    # X: [n_samples_test, n_features],
    # return: y: [n_samples_test, ]
```

```

X = np.array(X)
y = np.zeros(X.shape[0])

for i in range(X.shape[0]):
    node = self.tree
    while 'feature' in node:
        if X[i][node['feature']] < node['threshold']:
            node = node['left_tree']
        else:
            node = node['right_tree']
    y[i] = node['threshold']
return y

```

### 3 调试参数

```

accuracy_init = 0
for i in range(5, 12):
    max_depth = i
    for j in range(2, 10):
        min_samples_leaf = j
        clf.fit(X_train, y_train, max_depth=max_depth,
min_samples_leaf=min_samples_leaf)
        y_pred = clf.predict(X_test)
        if accuracy(y_test, y_pred) > accuracy_init:
            accuracy_init = accuracy(y_test, y_pred)
            max_depth_best = max_depth
            min_samples_leaf_best = min_samples_leaf

print(f"准确度: {accuracy_init}, 最佳最大深度: {max_depth_best}, 最佳最小样本
量: {min_samples_leaf_best}")

```

最大深度从5到11，最小样本量从2到9。

最大深度为最佳准确率和最佳参数如下：

**准确度：0.9527186761229315，最佳最大深度：10，最佳最小样本量：7**

## 2. PCAKMeans

### 1 PCA

#### 1 核函数：

```

def get_kernel_function(kernel):
    if kernel == "rbf":
        def rbf_kernel(x, y, gamma=1.0):
            return np.exp(-gamma * np.linalg.norm(x - y) ** 2)
        return rbf_kernel
    # 这里使用线性核
    elif kernel == "linear":
        def linear_kernel(x, y):
            return np.dot(x, y)
        return linear_kernel
    else:
        raise ValueError("Unsupported kernel")

```

2 主成分分析:

```

def fit(self, X: np.ndarray):
    # X: [n_samples, n_features]
    n_samples = X.shape[0]
    K = np.zeros((n_samples, n_samples))

    for i in range(n_samples):
        for j in range(n_samples):
            K[i, j] = self.kernel_f(X[i], X[j])

    # 中心化
    one_n = np.ones((n_samples, n_samples)) / n_samples
    K_centered = K - one_n @ K - K @ one_n + one_n @ K @ one_n
    self.K = K_centered

    # Eigenvalue decomposition
    eigenvalues, eigenvectors = np.linalg.eig(K_centered)
    idx = eigenvalues.argsort()[::-1]

    self.eigenvectors = eigenvectors[:, idx[:self.n_components]]
    self.eigenvalues = eigenvalues[idx[:self.n_components]]

```

3 投影:  $x'_j = \frac{1}{\sqrt{\lambda}} [K(x_1, x_j), \dots, K(x_n, x_j)] \cdot u$

```
def transform(self, X: np.ndarray):
    # X: [n_samples, n_features]
    self.fit(X)
    X_reduced = (self.K @ self.eigenvectors) /
np.sqrt(self.eigenvalues[:self.n_components])

    return X_reduced
```

## 2 KMeans

- 1 初始化。
- 2 计算每个点到各个中心的距离，并分配到最近的聚类。

```
def assign_points(self, points):
    n_samples = points.shape
    self.labels = np.zeros(n_samples, dtype = int)
    for i in range(n_samples):
        distances = np.linalg.norm(points[i] - self.centers,
axis=1)
        self.labels[i] = np.argmin(distances)
    return self.labels
```

- 3 更新聚类中心。

```
def update_centers(self, points):
    for i in range(self.k):
        cluster_points = points[self.labels == i]
        if len(cluster_points) > 0:
            self.centers[i] = cluster_points.mean(axis=0)
        else:
            self.centers[i] =
points[np.random.choice(len(points))]
```

- 4 KMeans执行。（使用 `allclose` 判断两个数组的元素是否逐一接近）

```
def fit(self, points):
    self.centers = self.initialize_centers(points)
    for _ in range(self.max_iter):
        previous_centers = self.centers.copy()
        self.labels = self.assign_points(points)
        self.update_centers(points)
        if np.allclose(previous_centers, self.centers):
            break
```

3 测试：分解到2维，使用 **linear** 核。7个聚类。

