

Intelligent Systems

MECHANICAL ENGINEERING

Class Assignment 1

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Contents

1	Dataset 1: Diabetes Dataset (Regression)	2
2	Dataset 2: Pima Indians Diabetes Dataset (Classification)	3
3	GitHub Repo	5
Δη	noves	6

1 Dataset 1: Diabetes Dataset (Regression)

Fuzzy C-Means Clustering was applied with a range of values for the number of clusters of 2 to 10 with an increment of 1 and the fuzziness coefficient was varied between 1.5 and 2.5 with an increment of 0.1. The results were evaluated using the Fuzzy Partition Coefficient (FPC) and culminated in a number of 2 clusters and a fuzziness coefficient of 1.5 achieving a FPC of 0.939724280208228.

The resulting TSK model achieved a MSE of 2534.134033203125.

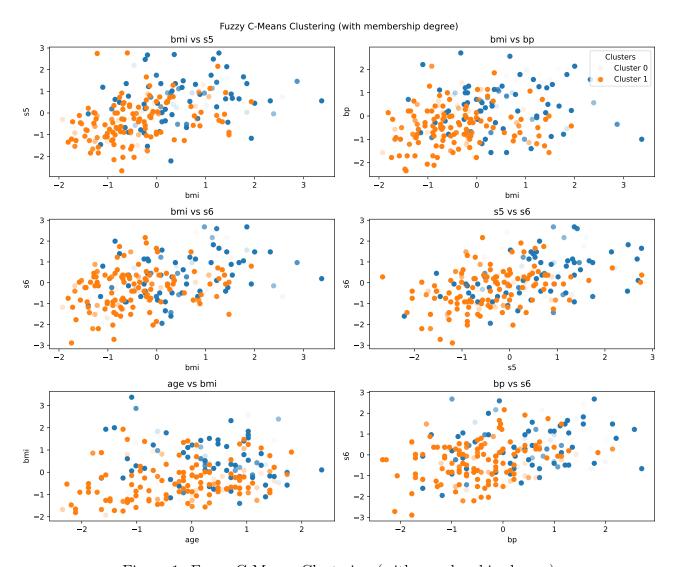


Figure 1: Fuzzy C-Means Clustering (with membership degree)

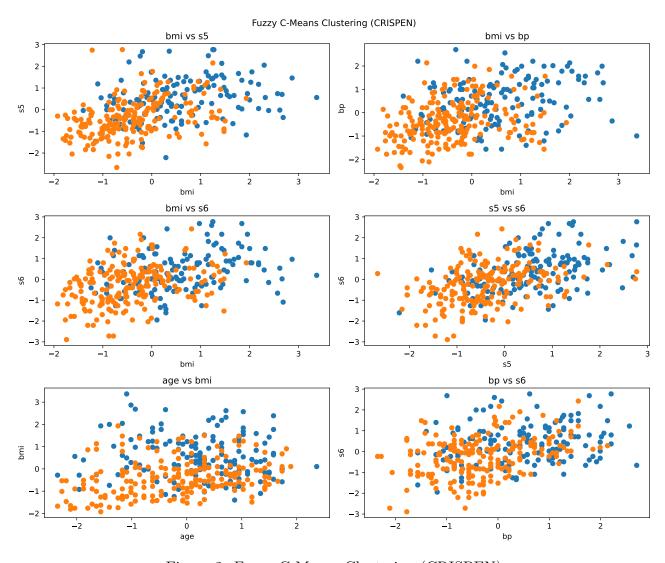


Figure 2: Fuzzy C-Means Clustering (CRISPEN)

2 Dataset 2: Pima Indians Diabetes Dataset (Classification)

Regarding data processing, several things were done to improve the dataset. Firstly, the target variable was converted from text strings to binary. The *Skin Thickness* and *Insulin* features lacked many rows of measurements; as such, these were discarded. Despite this, some measurements had 0 values that were equivalent to impossible; the rows containing these were discarded. The remaining data processing tasks were already implemented. The performance metrics improved after these measures were implemented.

Fuzzy C-Means Clustering was applied with a range of values for the number of clusters of 2 to 10 with an increment of 1 and the fuzziness coefficient was varied between 1.5 and 2.5 with an increment of 0.1. The results were evaluated using the Fuzzy Partition Coefficient (FPC) and culminated in a number of 2 clusters and a fuzziness coefficient of 1.5 achieving a FPC of 0.7231894414858948.

The resulting TSK model achieved an ACC of 0.8.

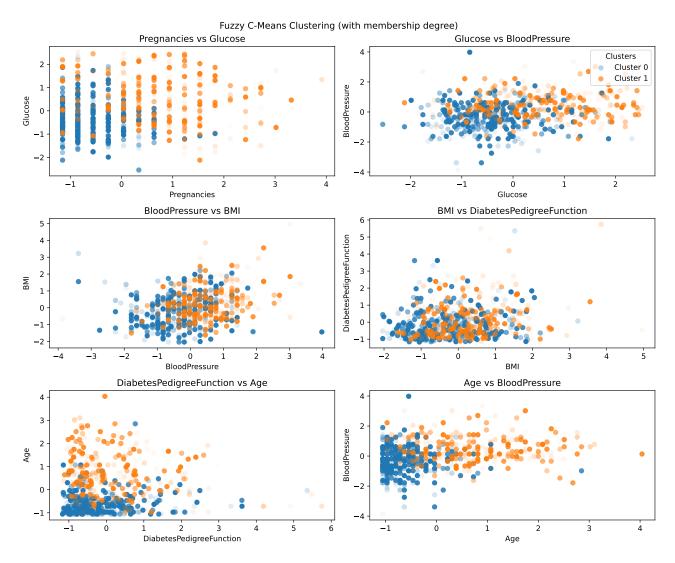


Figure 3: Fuzzy C-Means Clustering (with membership degree)

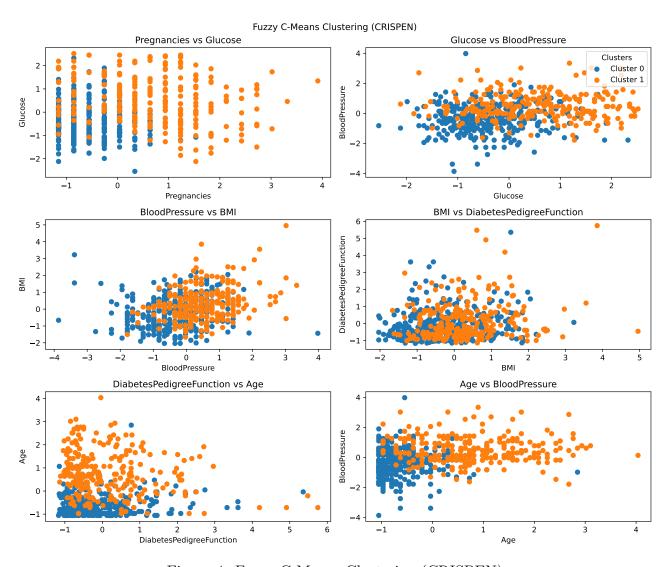


Figure 4: Fuzzy C-Means Clustering (CRISPEN)

3 GitHub Repo

 $github.com/Pereira 98/SI_Individual_87172.git$

TSK_pytorch_REG

September 26, 2025

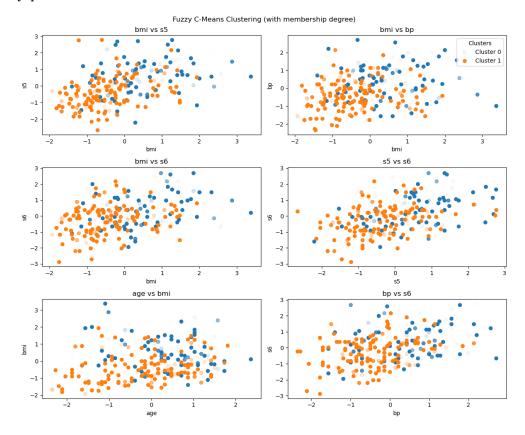
```
[1]: import numpy as np
     from sklearn import datasets
     from sklearn.preprocessing import StandardScaler
     from sklearn.model_selection import train_test_split
     from sklearn.metrics import
     -mean_squared_error,accuracy_score,classification_report
     import skfuzzy as fuzz
     import matplotlib.pyplot as plt
     import torch
     import torch.nn as nn
     import torch.optim as optim
     import pandas
[2]: # CHOOSE DATASET
     # Regression dataset
     data = datasets.load_diabetes(as_frame=True)
     X = data.data.values
     y = data.target.values
     X.shape
[2]: (442, 10)
[3]: #train test spliting
     test_size=0.2
     Xtr, Xte, ytr, yte = train_test_split(X, y, test_size=test_size,_
      →random_state=42)
[4]: # Standardize features
     scaler=StandardScaler()
     Xtr= scaler.fit_transform(Xtr)
     Xte= scaler.transform(Xte)
[5]: # Number of clusters
     n_{clusters} = 2
     m=2
```

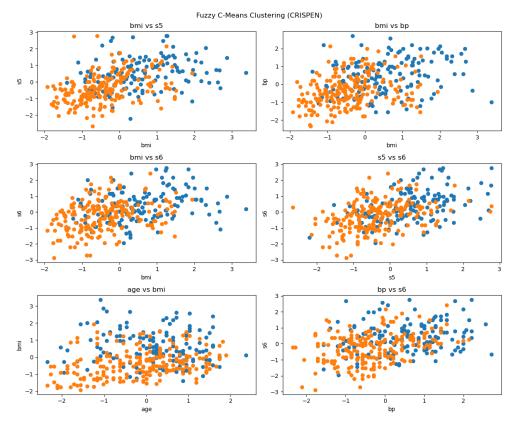
```
# Concatenate target for clustering
     Xexp=np.concatenate([Xtr, ytr.reshape(-1, 1)], axis=1)
     #Xexp=Xtr
     # Transpose data for skfuzzy (expects features x samples)
     Xexp_T = Xexp.T
     best_fpc = -1
     best_params = None
     for n_clusters in np.arange(2, 10, 1):
                                              # number of clusters to test
         for m in np.arange(1.5, 2.5, 0.1):
         # Fuzzy C-means clustering
             centers, u, u0, d, jm, p, fpc = fuzz.cluster.cmeans(
                 Xexp_T, n_clusters, m, error=0.005, maxiter=1000, init=None,
             if fpc > best_fpc:
                 best_fpc = fpc
                 best_params = (n_clusters, m)
                 print(best_params)
     print(best_fpc)
     print(best_params[0])
     print(best_params[1])
     n_clusters=best_params[0]
     m=best_params[1]
     centers, u, u0, d, jm, p, fpc = fuzz.cluster.cmeans(
         Xexp_T, n_clusters, m, error=0.005, maxiter=10000, init=None,
    (np.int64(2), np.float64(1.5))
    0.93972435685851
    2
    1.5
[6]: centers.shape
[6]: (2, 11)
[7]: # Compute sigma (spread) for each cluster
     sigmas = []
     for j in range(n_clusters):
         # membership weights for cluster j, raised to m
        u_{j} = u[j, :] ** m
         # weighted variance for each feature
         var_j = np.average((Xexp - centers[j])**2, axis=0, weights=u_j)
```

```
sigma_j = np.sqrt(var_j)
sigmas.append(sigma_j)
sigmas=np.array(sigmas)
```

```
[8]: # Hard clustering from fuzzy membership
    cluster_labels = np.argmax(u, axis=0)
    print("Fuzzy partition coefficient (FPC):", fpc)
     # Diabetes feature names (sklearn's load_diabetes ordering)
    feature_names = [
        "age", # 0
        "sex", # 1
        "bmi", # 2
        "bp",
                # 3
        "s1",
              # 4 (tc)
        "s2",
              # 5 (ldl)
        "s3",
              # 6 (hdl)
        "s4", # 7 (tch)
        "s5", # 8 (ltg)
        "s6"
               # 9 (qlu)
    # Choose 6 feature pairs (indices in Xexp)
    pairs = [
        (2, 8), # bmi vs s5 (both strong predictors)
        (2, 3), # bmi vs bp
        (2, 9), # bmi vs glu
        (8, 9), # s5 vs glu
         (0, 2), # age vs bmi
         (3, 9) # bp vs qlu
    ]
    fig, axes = plt.subplots(3, 2, figsize=(12, 10))
    for ax, (i, j) in zip(axes.ravel(), pairs):
        # Plot 2 features with fuzzy membership
        for k in range(n_clusters):
            ax.scatter(
                Xexp[cluster_labels == k, i],
                                              # Feature i
                Xexp[cluster_labels == k, j],  # Feature j
                                                # transparency ~ membership degree
                alpha=u[k, :],
                label=f'Cluster {k}'
            )
        ax.set_xlabel(feature_names[i])
        ax.set_ylabel(feature_names[j])
        ax.set_title(f"{feature_names[i]} vs {feature_names[j]}")
```

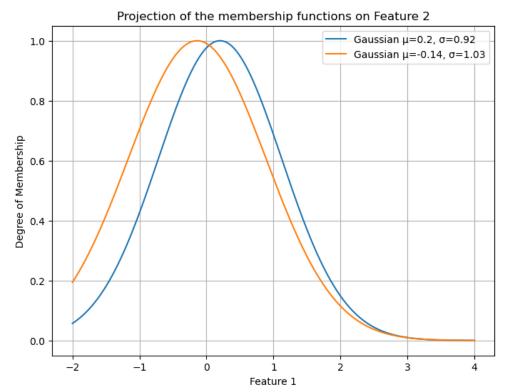
Fuzzy partition coefficient (FPC): 0.9397242309334268





```
[10]: # Gaussian formula
def gaussian(x, mu, sigma):
    return np.exp(-0.5 * ((x - mu)/sigma)**2)
```

```
lin=np.linspace(-2, 4, 500)
plt.figure(figsize=(8,6))
y_aux=[]
feature=0
for j in range(n_clusters):
# Compute curves
    y_aux.append(gaussian(lin, centers[j,feature], sigmas[j,feature]))
# Plot
    {\tt plt.plot(lin, y\_aux[j], label=f"Gaussian = \{np.}\\
 →round(centers[j,feature],2)}, ={np.round(sigmas[j,feature],2)}")
plt.title("Projection of the membership functions on Feature 2")
plt.xlabel("Feature 1")
plt.ylabel("Degree of Membership")
plt.legend()
plt.grid(True)
plt.show()
```



```
[11]: # -----
      # Gaussian Membership Function
      # -----
      class GaussianMF(nn.Module):
          def __init__(self, centers, sigmas, agg_prob):
              super().__init__()
              self.centers = nn.Parameter(torch.tensor(centers, dtype=torch.float32))
              self.sigmas = nn.Parameter(torch.tensor(sigmas, dtype=torch.float32))
              self.agg_prob=agg_prob
          def forward(self, x):
              # Expand for broadcasting
              # x: (batch, 1, n_dims), centers: (1, n_rules, n_dims), sigmas: (1,u
       \hookrightarrow n\_rules, n\_dims)
              diff = abs((x.unsqueeze(1) - self.centers.unsqueeze(0))/self.sigmas.

unsqueeze(0)) #(batch, n_rules, n_dims)

              # Aggregation
              if self.agg_prob:
                  dist = torch.norm(diff, dim=-1) # (batch, n_rules) # probablistic_
       \hookrightarrow intersection
                  dist = torch.max(diff, dim=-1).values # (batch, n_rules) # min_
       \hookrightarrow intersection (min instersection of normal funtion is the same as the max on \sqcup
       \hookrightarrow dist)
              return torch.exp(-0.5 * dist ** 2)
      # TSK Model
      class TSK(nn.Module):
          def __init__(self, n_inputs, n_rules, centers, sigmas,agg_prob=False):
              super().__init__()
              self.n_inputs = n_inputs
              self.n rules = n rules
              # Antecedents (Gaussian MFs)
              self.mfs=GaussianMF(centers, sigmas,agg_prob)
              # Consequents (linear functions of inputs)
              # Each rule has coeffs for each input + bias
              self.consequents = nn.Parameter(
                  torch.randn(n_inputs + 1,n_rules)
```

```
def forward(self, x):
       # x: (batch, n_inputs)
       batch_size = x.shape[0]
        # Compute membership values for each input feature
        # firing_strengths: (batch, n_rules)
        firing_strengths = self.mfs(x)
        # Normalize memberships
        # norm_fs: (batch, n_rules)
        norm\_fs = firing\_strengths / (firing\_strengths.sum(dim=1, keepdim=True)_{\sqcup}
 →+ 1e-9)
        # Consequent output (linear model per rule)
        x_aug = torch.cat([x, torch.ones(batch_size, 1)], dim=1) # add bias
       rule_outputs = torch.einsum("br,rk->bk", x_aug, self.consequents) #__
 \hookrightarrow (batch, rules)
        # Weighted sum
        output = torch.sum(norm_fs * rule_outputs, dim=1, keepdim=True)
        return output, norm_fs, rule_outputs
# Least Squares Solver for Consequents (TSK)
# -----
def train_ls(model, X, y):
   with torch.no_grad():
        _, norm_fs, _ = model(X)
        # Design matrix for LS: combine normalized firing strengths with input
        X_aug = torch.cat([X, torch.ones(X.shape[0], 1)], dim=1)
        Phi = torch.einsum("br,bi->bri", X_aug, norm_fs).reshape(X.shape[0], -1)
        # Solve LS: consequents = (Phi^T Phi)^-1 Phi^T y
        theta= torch.linalg.lstsq(Phi, y).solution
```

```
[13]: # ------
# Gradient Descent Training
```

model.consequents.data = theta.reshape(model.consequents.shape)

```
def train_gd(model, X, y, epochs=100, lr=1e-3):
         optimizer = optim.Adam(model.parameters(), lr=lr)
         criterion = nn.MSELoss()
         for _ in range(epochs):
             optimizer.zero_grad()
             y_pred, _, _ = model(X)
             loss = criterion(y_pred, y)
             print(loss)
             loss.backward()
             optimizer.step()
[14]: # -----
     # Hybrid Training (Classic ANFIS)
     # -----
     def train_hybrid_anfis(model, X, y, max_iters=10, gd_epochs=20, lr=1e-3):
         train_ls(model, X, y)
         for _ in range(max_iters):
             # Step A: GD on antecedents (freeze consequents)
             model.consequents.requires_grad = False
             train_gd(model, X, y, epochs=gd_epochs, lr=lr)
             # Step B: LS on consequents (freeze antecedents)
             model.consequents.requires_grad = True
             model.mfs.requires_grad = False
             train_ls(model, X, y)
             # Re-enable antecedents
             model.mfs.requires_grad = True
     # Alternative Hybrid Training (LS+ gradient descent on all)
     # -----
     def train_hybrid(model, X, y, epochs=100, lr=1e-4):
         # Step 1: LS for consequents
         train_ls(model, X, y)
         # Step 2: GD fine-tuning
         train_gd(model, X, y, epochs=epochs, lr=lr)
[16]: # Build model
     model = TSK(n_inputs=Xtr.shape[1], n_rules=n_clusters, centers=centers[:,:-1],__
      ⇔sigmas=sigmas[:,:-1])
     Xtr = torch.tensor(Xtr, dtype=torch.float32)
     ytr = torch.tensor(ytr, dtype=torch.float32)
     Xte = torch.tensor(Xte, dtype=torch.float32)
     yte = torch.tensor(yte, dtype=torch.float32)
```

MSE:2534.13330078125

TSK_pytorch-CLF

September 26, 2025

```
[1]: import numpy as np
    from sklearn import datasets
    from sklearn.preprocessing import StandardScaler
    from sklearn.model_selection import train_test_split
    from sklearn.metrics import
     -mean_squared_error,accuracy_score,classification_report
    import skfuzzy as fuzz
    import matplotlib.pyplot as plt
    import torch
     import torch.nn as nn
     import torch.optim as optim
     import pandas
    import copy
[2]: # CHOOSE DATASET
    # Binary classification dataset
    data = datasets.fetch_openml(name="diabetes",version=1, as_frame=True)
    X = data.data.values
```

```
# Binary classification dataset
data = datasets.fetch_openml(name="diabetes",version=1, as_frame=True)

X = data.data.values
y = data.target.values
X.shape

# Keep as DataFrame for named-column ops
df = data.data.copy()
y = np.where(data.target.values == "tested_positive", 1, 0).astype(np.float32)

# indices of features with invalid zeros
invalid_idx = [1, 2, 3, 4, 5, 7]

# count zeros per feature
zero_counts = (X[:, invalid_idx] == 0).sum(axis=0)
rows_with_zero = (X[:, invalid_idx] == 0).any(axis=1).sum()

print("Zeros per feature:\n", zero_counts)
print(f"Rows with 1 zero: {rows_with_zero} / {len(df)}")
```

```
(len(X), zero_counts, rows_with_zero)
     # Drop columns 3 and 4 (0-based indexing)
    X = np.delete(X, [3, 4], axis=1)
    # Keep only rows where Glucose, BloodPressure, BMI are non-zero
    mask = (X[:, [1, 2, 3, 4]] != 0).all(axis=1)
    X = X[mask]
    y = y[mask]
    Zeros per feature:

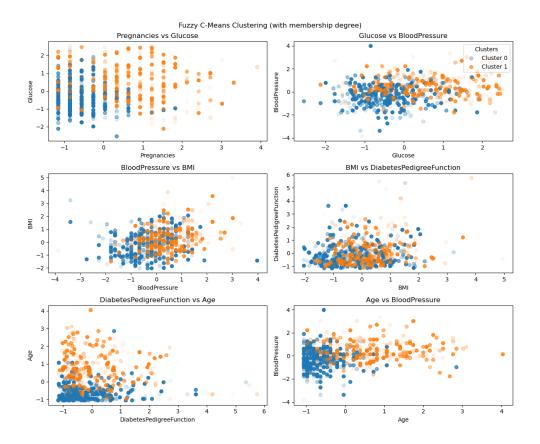
    5 35 227 374 11

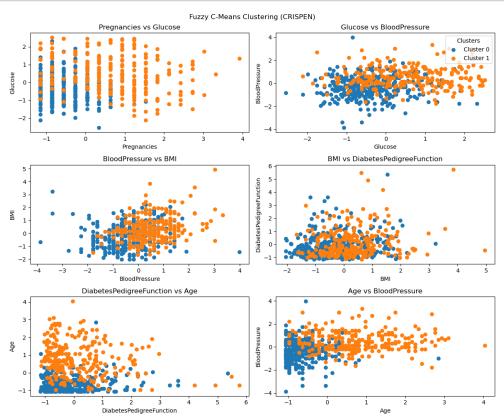
    Rows with 1 zero: 376 / 768
[3]: #train test spliting
    test_size=0.2
    Xtr, Xte, ytr, yte = train_test_split(X, y, test_size=test_size,__
      ⇔random_state=42)
[4]: # Standardize features
    scaler=StandardScaler()
    Xtr= scaler.fit_transform(Xtr)
    Xte= scaler.transform(Xte)
[5]: # Concatenate target for clustering
    Xexp=np.concatenate([Xtr, ytr.reshape(-1, 1)], axis=1)
     #Xexp=Xtr
     # Transpose data for skfuzzy (expects features x samples)
    Xexp_T = Xexp.T
    best_fpc = -1
    best_params = None
    for n_clusters in np.arange(2, 10, 1): # number of clusters to test
        for m in np.arange(1.5, 2.5, 0.01):
        # Fuzzy C-means clustering
             centers, u, u0, d, jm, p, fpc = fuzz.cluster.cmeans(
                 Xexp_T, n_clusters, m, error=0.005, maxiter=1000, init=None,
             if fpc > best_fpc:
                best_fpc = fpc
                best_params = (n_clusters, m)
    print(best_fpc)
    print(best_params[0])
```

```
print(best_params[1])
     n_clusters=best_params[0]
     m=best_params[1]
     centers, u, u0, d, jm, p, fpc = fuzz.cluster.cmeans(
         Xexp_T, n_clusters, m, error=0.005, maxiter=10000, init=None,
    0.7231811657391913
    2
    1.5
[6]: centers.shape
[6]: (2, 7)
[7]: # Compute sigma (spread) for each cluster
     sigmas = []
     for j in range(n_clusters):
         # membership weights for cluster j, raised to m
         u_j = u[j, :] ** m
         # weighted variance for each feature
         var_j = np.average((Xexp - centers[j])**2, axis=0, weights=u_j)
         sigma_j = np.sqrt(var_j)
         sigmas.append(sigma_j)
     sigmas=np.array(sigmas)
[8]: # Hard clustering from fuzzy membership
     cluster_labels = np.argmax(u, axis=0)
     print("Fuzzy partition coefficient (FPC):", fpc)
     feature_names = [
         "Pregnancies", "Glucose", "BloodPressure",
         "BMI", "DiabetesPedigreeFunction", "Age"
     ]
     # Choose 4 feature pairs (indices in Xexp)
     pairs = [(0, 1), # Pregnancies vs Glucose
              (1, 2), # Glucose vs BloodPressure
              (2, 3), # BloodPressure vs BMI
              (3, 4), # BMI vs Pedigree
              (4, 5), # Pedigree vs Age
              (5, 2)] # Age vs Glucose
     fig, axes = plt.subplots(3, 2, figsize=(12, 10))
     for ax, (i, j) in zip(axes.ravel(), pairs):
```

```
# Plot 2 features with fuzzy membership
   for k in range(n_clusters):
       ax.scatter(
           Xexp[cluster_labels == k, i],
                                                    # Feature 1
           Xexp[cluster_labels == k, j],
                                                   # Feature 2
           alpha=u[k, :],
                                 # transparency ~ membership
           label=f'Cluster {k}'
       )
   ax.set_xlabel(feature_names[i])
   ax.set_ylabel(feature_names[j])
   ax.set_title(f"{feature_names[i]} vs {feature_names[j]}")
fig.suptitle("Fuzzy C-Means Clustering (with membership degree)")
axes[0,1].legend(title="Clusters", loc="upper right")
plt.tight_layout()
plt.show()
fig.savefig("Fuzzy C-Means Clustering (with membership degree) CLF.pdf", _
 oformat="pdf", bbox_inches="tight")
```

Fuzzy partition coefficient (FPC): 0.723188876981547

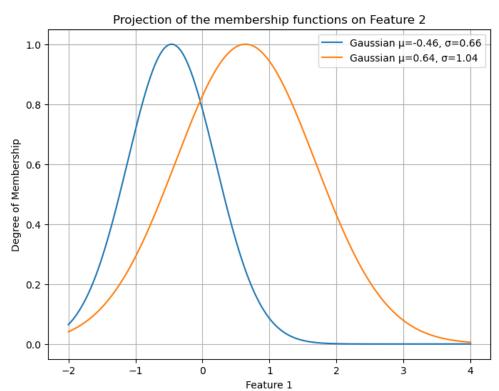




```
[10]: # Gaussian formula
def gaussian(x, mu, sigma):
    return np.exp(-0.5 * ((x - mu)/sigma)**2)

lin=np.linspace(-2, 4, 500)
plt.figure(figsize=(8,6))

y_aux=[]
feature=0
for j in range(n_clusters):
# Compute curves
    y_aux.append(gaussian(lin, centers[j,feature], sigmas[j,feature]))
```



```
# ------
# Gaussian Membership Function
# ------
class GaussianMF(nn.Module):
    def __init__(self, centers, sigmas, agg_prob):
        super().__init__()
        self.centers = nn.Parameter(torch.tensor(centers, dtype=torch.float32))
        self.sigmas = nn.Parameter(torch.tensor(sigmas, dtype=torch.float32))
```

```
self.agg_prob=agg_prob
    def forward(self, x):
        # Expand for broadcasting
        # x: (batch, 1, n_dims), centers: (1, n_dims), n_dims), n_dims), n_dims: (1,n_dims)
 \hookrightarrow n_rules, n_dims)
        diff = abs((x.unsqueeze(1) - self.centers.unsqueeze(0))/self.sigmas.
 →unsqueeze(0)) #(batch, n_rules, n_dims)
        # Aggregation
        if self.agg_prob:
            dist = torch.norm(diff, dim=-1) # (batch, n_rules) # probablistic_
 \hookrightarrow intersection
        else:
            dist = torch.max(diff, dim=-1).values # (batch, n_rules) # min_
 →intersection (min instersection of normal funtion is the same as the max on
 \rightarrow dist)
        return torch.exp(-0.5 * dist ** 2)
# TSK Model
class TSK(nn.Module):
    def __init__(self, n_inputs, n_rules, centers, sigmas,agg_prob=False):
        super().__init__()
        self.n_inputs = n_inputs
        self.n_rules = n_rules
        # Antecedents (Gaussian MFs)
        self.mfs=GaussianMF(centers, sigmas,agg_prob)
        # Consequents (linear functions of inputs)
        # Each rule has coeffs for each input + bias
        self.consequents = nn.Parameter(
            torch.randn(n_inputs + 1,n_rules)
        )
    def forward(self, x):
        # x: (batch, n_inputs)
        batch_size = x.shape[0]
        # Compute membership values for each input feature
        # firing_strengths: (batch, n_rules)
        firing_strengths = self.mfs(x)
```

```
# Normalize memberships
             # norm_fs: (batch, n_rules)
             norm_fs = firing_strengths / (firing_strengths.sum(dim=1, keepdim=True)_
       →+ 1e-9)
             # Consequent output (linear model per rule)
             x_aug = torch.cat([x, torch.ones(batch_size, 1)], dim=1) # add bias
             rule_outputs = torch.einsum("br,rk->bk", x_aug, self.consequents) #__
       \hookrightarrow (batch, rules)
             # Weighted sum
             output = torch.sum(norm_fs * rule_outputs, dim=1, keepdim=True)
             return output, norm_fs, rule_outputs
[12]: # -----
      # Least Squares Solver for Consequents (TSK)
     def train_ls(model, X, y):
         with torch.no_grad():
             _, norm_fs, _ = model(X)
             # Design matrix for LS: combine normalized firing strengths with input
             X_aug = torch.cat([X, torch.ones(X.shape[0], 1)], dim=1)
             Phi = torch.einsum("br,bi->bri", X_aug, norm_fs).reshape(X.shape[0], -1)
             # Solve LS: consequents = (Phi^T Phi)^-1 Phi^T y
             theta= torch.linalg.lstsq(Phi, y).solution
             model.consequents.data = theta.reshape(model.consequents.shape)
[13]: # -----
      # Gradient Descent Training
     def train_gd(model, X, y, epochs=100, lr=1e-3):
         optimizer = optim.Adam(model.parameters(), lr=lr)
         criterion = nn.MSELoss()
         for _ in range(epochs):
             optimizer.zero_grad()
             y_pred, _, _ = model(X)
             loss = criterion(y_pred, y)
```

```
print(loss)
             loss.backward()
             optimizer.step()
[14]: # -----
      # Hybrid Training (Classic ANFIS)
      # -----
     def train_hybrid_anfis(model, X, y, max_iters=10, gd_epochs=20, lr=1e-3):
         train_ls(model, X, y)
         for _ in range(max_iters):
             # Step A: GD on antecedents (freeze consequents)
             model.consequents.requires_grad = False
             train_gd(model, X, y, epochs=gd_epochs, lr=lr)
             # Step B: LS on consequents (freeze antecedents)
             model.consequents.requires_grad = True
             model.mfs.requires_grad = False
             train_ls(model, X, y)
             # Re-enable antecedents
             model.mfs.requires_grad = True
[15]: # -----
     # Alternative Hybrid Training (LS+ gradient descent on all)
     def train_hybrid(model, X, y, epochs=100, lr=1e-4):
         # Step 1: LS for consequents
         train_ls(model, X, y)
         # Step 2: GD fine-tuning
         train_gd(model, X, y, epochs=epochs, lr=lr)
[16]: # Build model
     model = TSK(n_inputs=Xtr.shape[1], n_rules=n_clusters, centers=centers[:,:-1],__
      ⇔sigmas=sigmas[:,:-1])
     Xtr = torch.tensor(Xtr, dtype=torch.float32)
     ytr = torch.tensor(ytr, dtype=torch.float32)
     Xte = torch.tensor(Xte, dtype=torch.float32)
     yte = torch.tensor(yte, dtype=torch.float32)
     # Training with LS:
     train_ls(model, Xtr, ytr.reshape(-1,1))
[17]: y_pred, _, _=model(Xte)
     #performance metric for classification
     print(f'ACC:{accuracy_score(yte.detach().numpy(),y_pred.detach().numpy()>0.5)}')
```

ACC:0.8