

با سمه تعالی



گزارش تمرین کامپیوترا سری ۳

پردازش علائم بیولوژیک

دانشکده مهندسی برق

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گردآورنده: رادین خیام

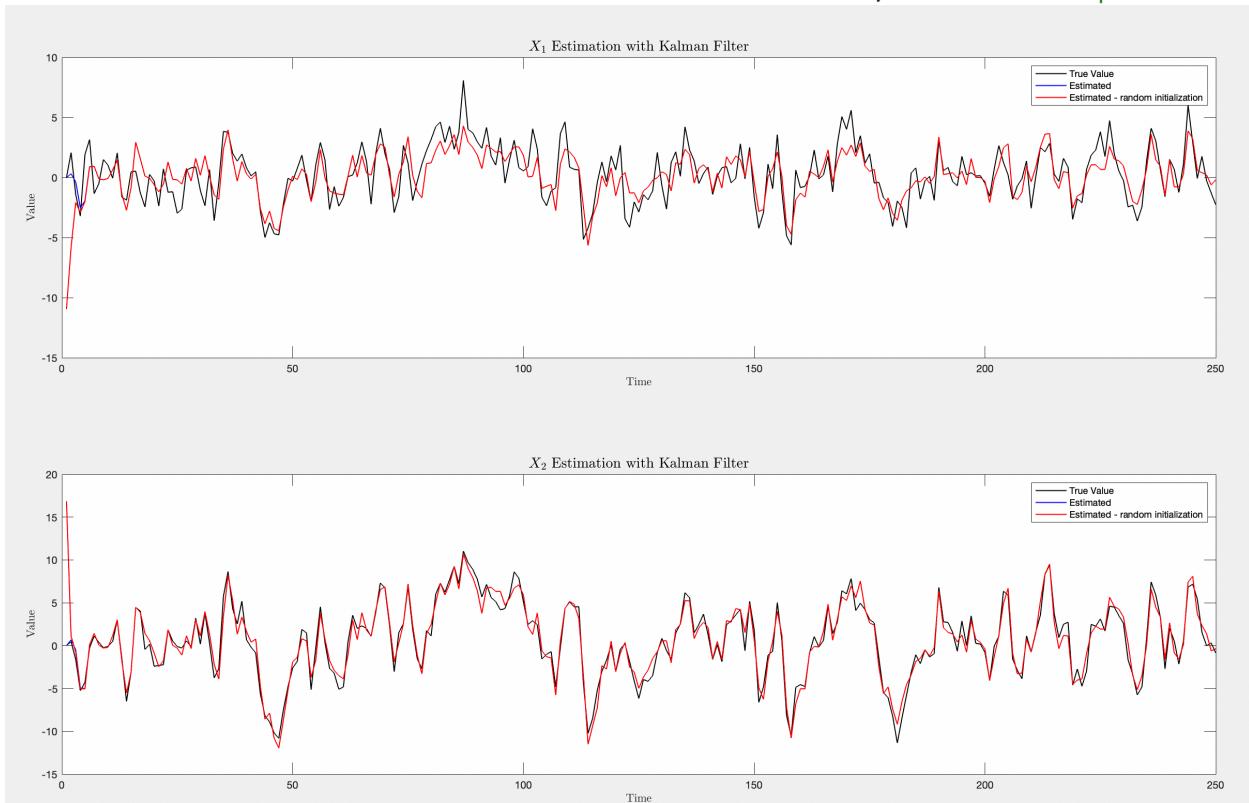
سؤال ۱-

(الف)

مقادیر اولیه در نظر گرفته شده:

```
sigma_v = 1;  
sigma_1 = 2;  
sigma_2 = 2;  
a = 0.5;  
b = 0.5;
```

N = 250; % Number of samples



Pearson Correlation Coeficient for X_1 = 0.79

Pearson Correlation Coeficient for X_2 = 0.97618

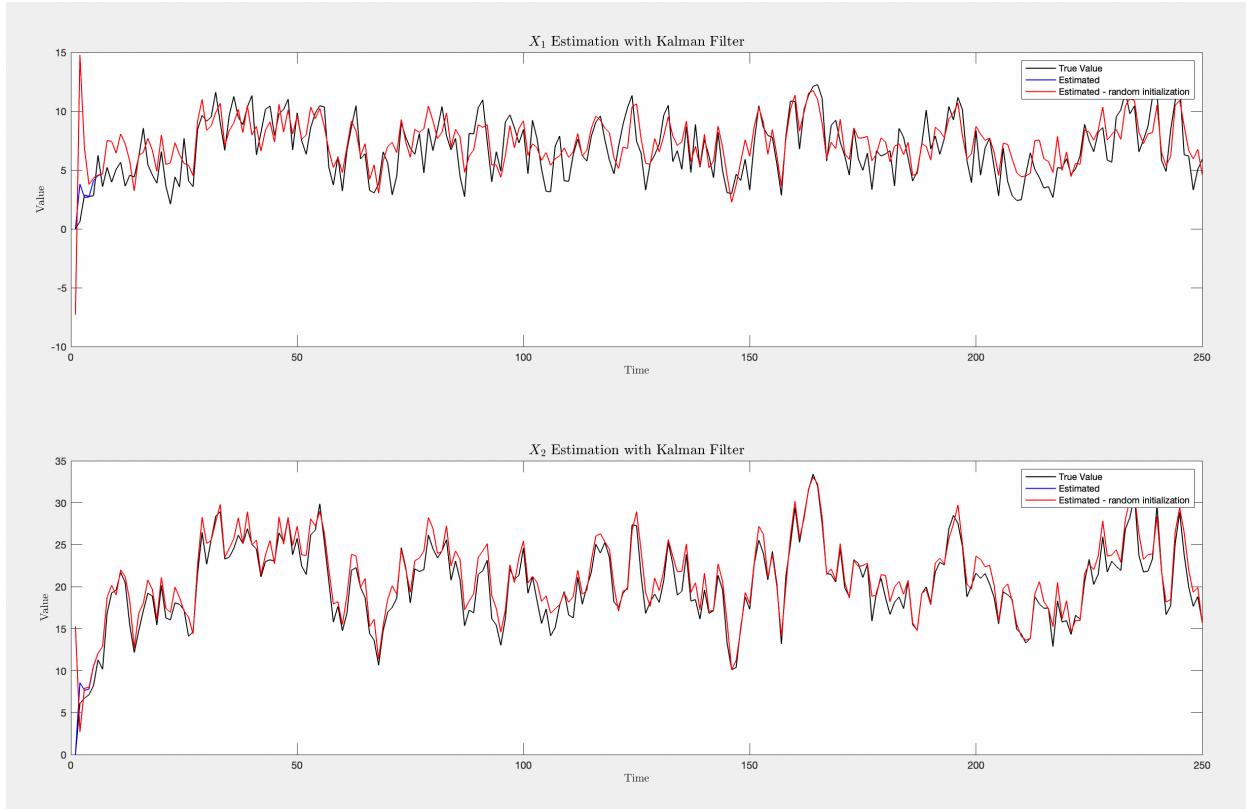
Pearson Correlation Coeficient for X_1 with random initialization = 0.71237

Pearson Correlation Coeficient for X_2 with random initialization = 0.94628

(ب)

به جای استفاده از توزیع گوسی برای تولید نویزها از توزیع یکنواخت استفاده کردیم.

```
V = sqrt(12) * sigma_v * rand(1, N); % generating noise with uniform distribution and variance sigma_v  
U_1 = sqrt(12) * sigma_1 * rand(1, N);  
U_2 = sqrt(12) * sigma_2 * rand(1, N);
```



Pearson Correlation Coeficient for X_1 = 0.77988
Pearson Correlation Coeficient for X_2 = 0.97882
Pearson Correlation Coeficient for X_1 with random initialization = 0.68609
Pearson Correlation Coeficient for X_2 with random initialization = 0.95948

نتایج خیلی متفاوت نشده است.

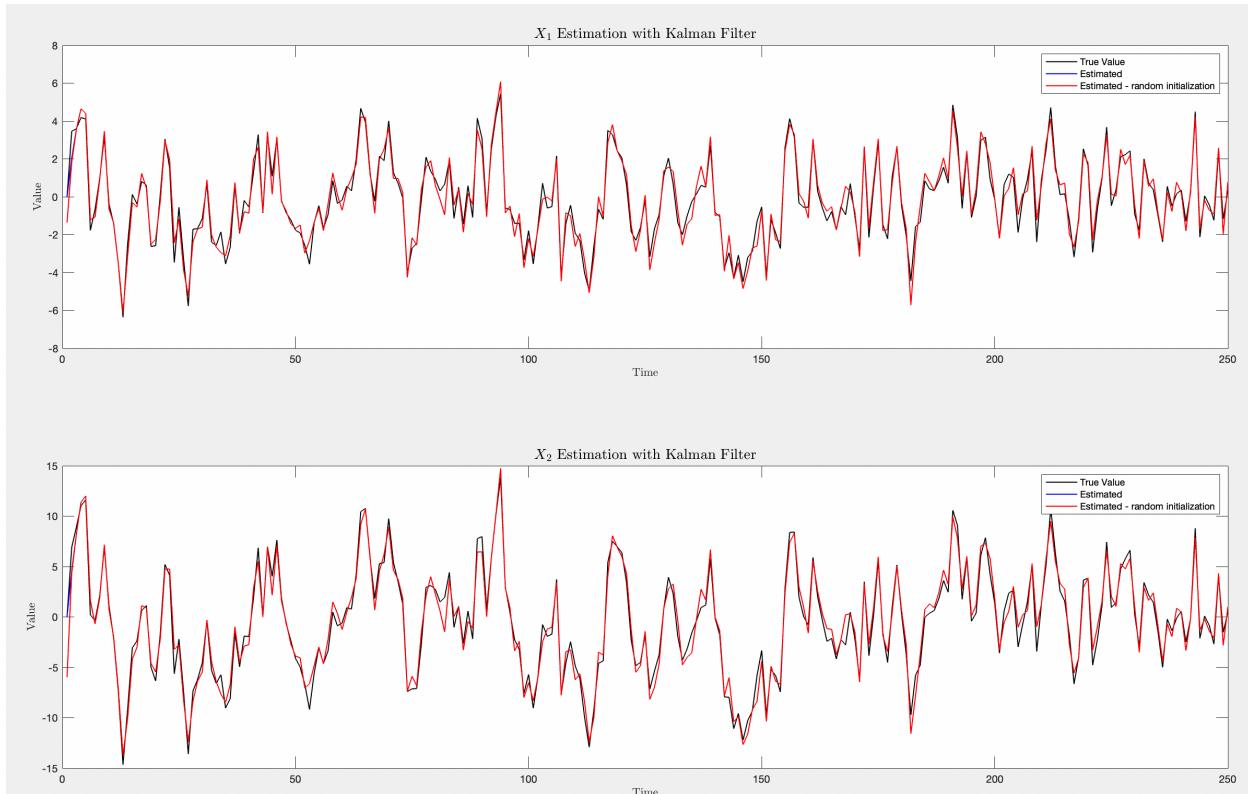
(پ)

نویز U_2 را ضریبی از نویز U_1 در نظر گرفتیم (به صورتی که واریانسش سیگما ۲ بشود)

```

V = sigma_v * randn(1, N);
U_1 = sigma_1 * randn(1, N);
U_2 = U_1 * sigma_2/sigma_1; % make correlation between U_1 and U_2

```



```

Pearson Correlation Coeficient for X_1 = 0.97524
Pearson Correlation Coeficient for X_2 = 0.98372
Pearson Correlation Coeficient for X_1 with random initialization = 0.9742
Pearson Correlation Coeficient for X_2 with random initialization = 0.98099

```

نتایج بهبود پیدا کرده است. در فیلتر کالمن نیازی نبود که حتماً نویزهای سیستم نسبت بهم مستقل باشند بلکه لازم بود که نویز مشاهدات و نویز سیستم از هم مستقل باشند.

(ت)

نتایج با هر دو حالت مقادیر اولیه مطابق با درس و با حالت اولیه رندوم در قسمت‌های الف، ب و پ گزارش شده است.

(ث)

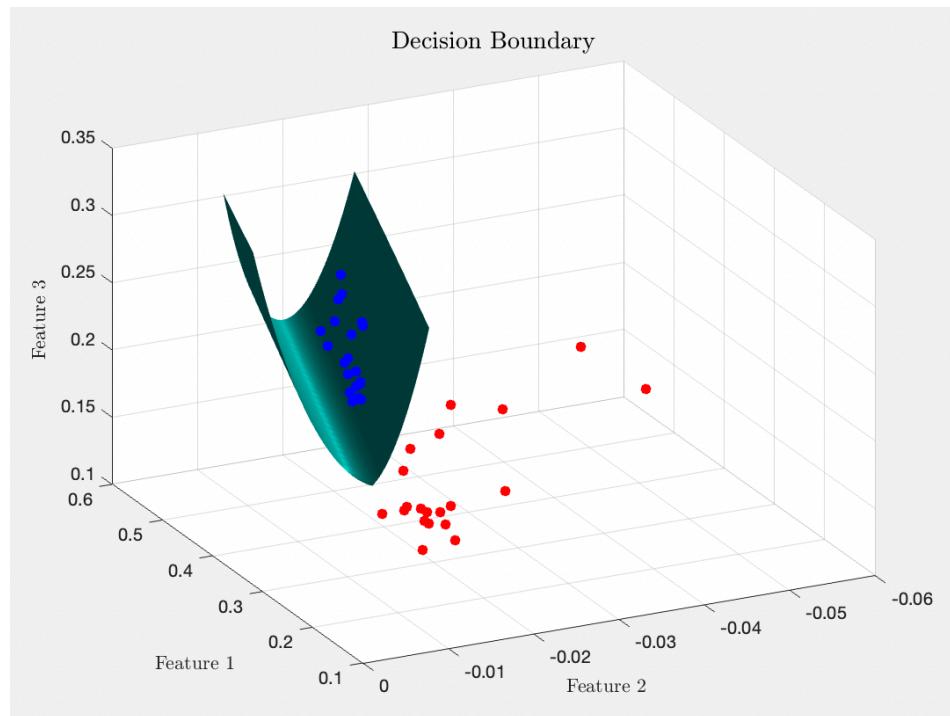
معیار ارزیابی ضریب همبستگی پیرسون در قسمت‌های الف، ب و پ گزارش شده است.

سؤال ۲-

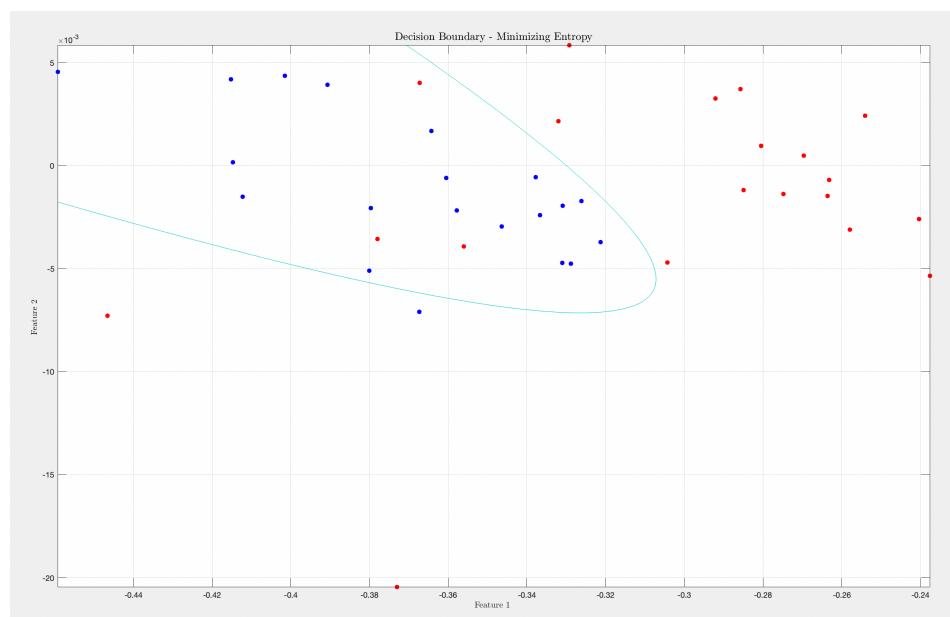
(الف)

ویژگی‌ها استخراج شدند.

(ب)

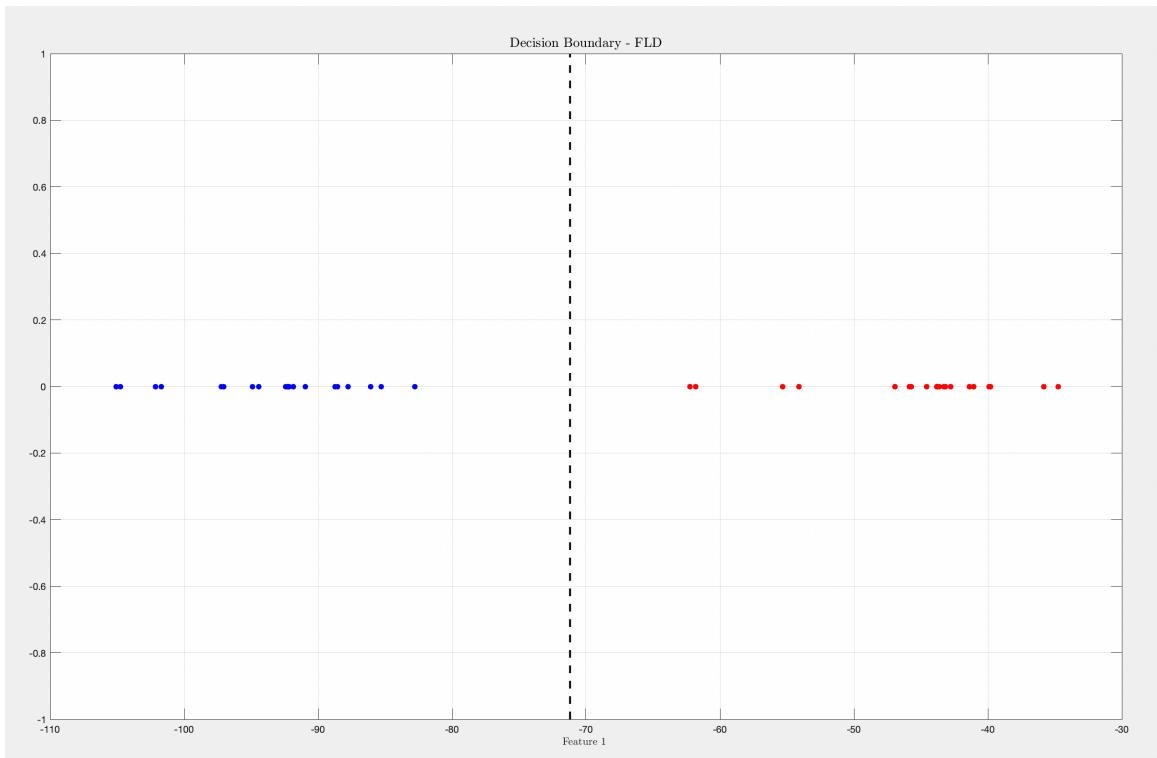


(پ)



جدایی پذیری کاہش پیدا کرده است اما اگر به آن مفهومی که در درس مطرح شد (یعنی اینکه اگر میانگین دو کلاس روی هم نیفتند یعنی جدایی پذیرند) بله جدایی پذیری حفظ شده است.

(ت)



بله جدابی پذیری کاملاً حفظ شده است.

(ث)

نتایج بدین صورت بدست آمده است.

Part b Results:

Accuracy of Mahalanobis Distance = 0.975

Accuracy of Euclidean Distance = 0.975

Accuracy of Bayes Classifier = 0.975

=====

Part c Results:

Accuracy of Mahalanobis Distance = 0.875

Accuracy of Euclidean Distance = 0.825

Accuracy of Bayes Classifier = 0.925

=====

Part d Results:

Accuracy of Mahalanobis Distance = 0.975

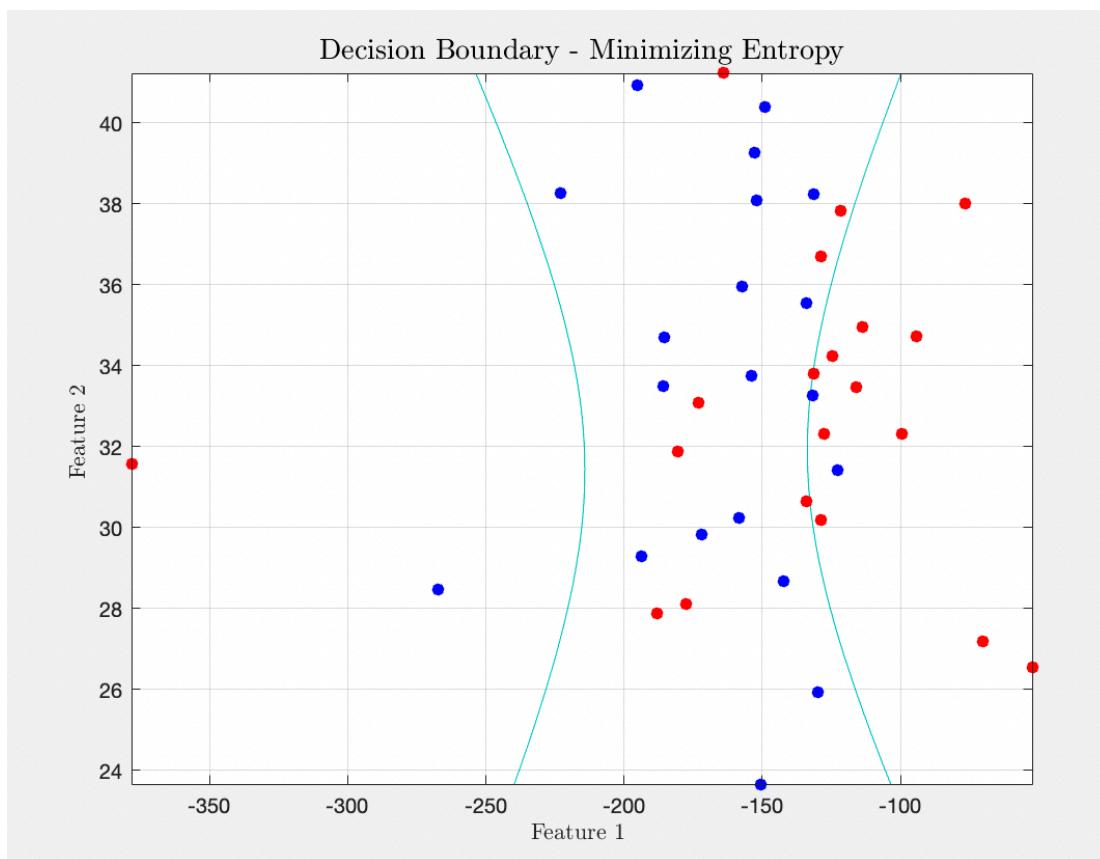
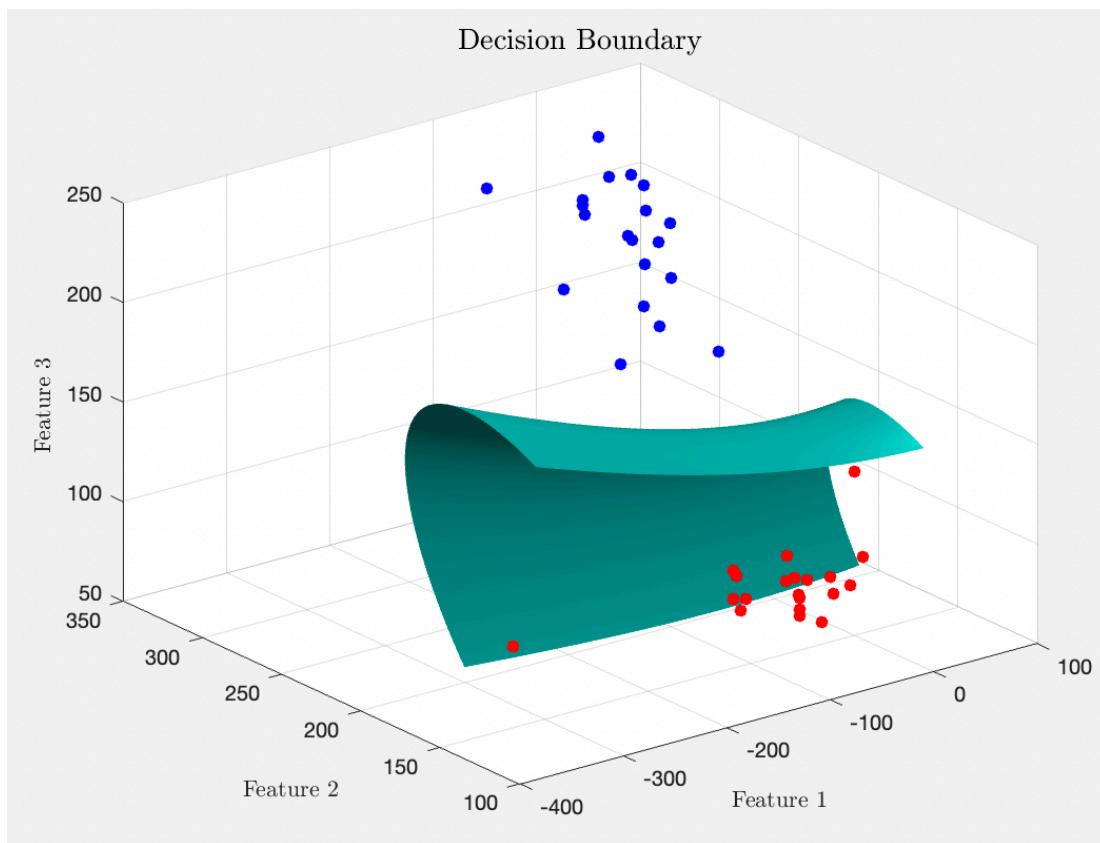
Accuracy of Euclidean Distance = 0.975

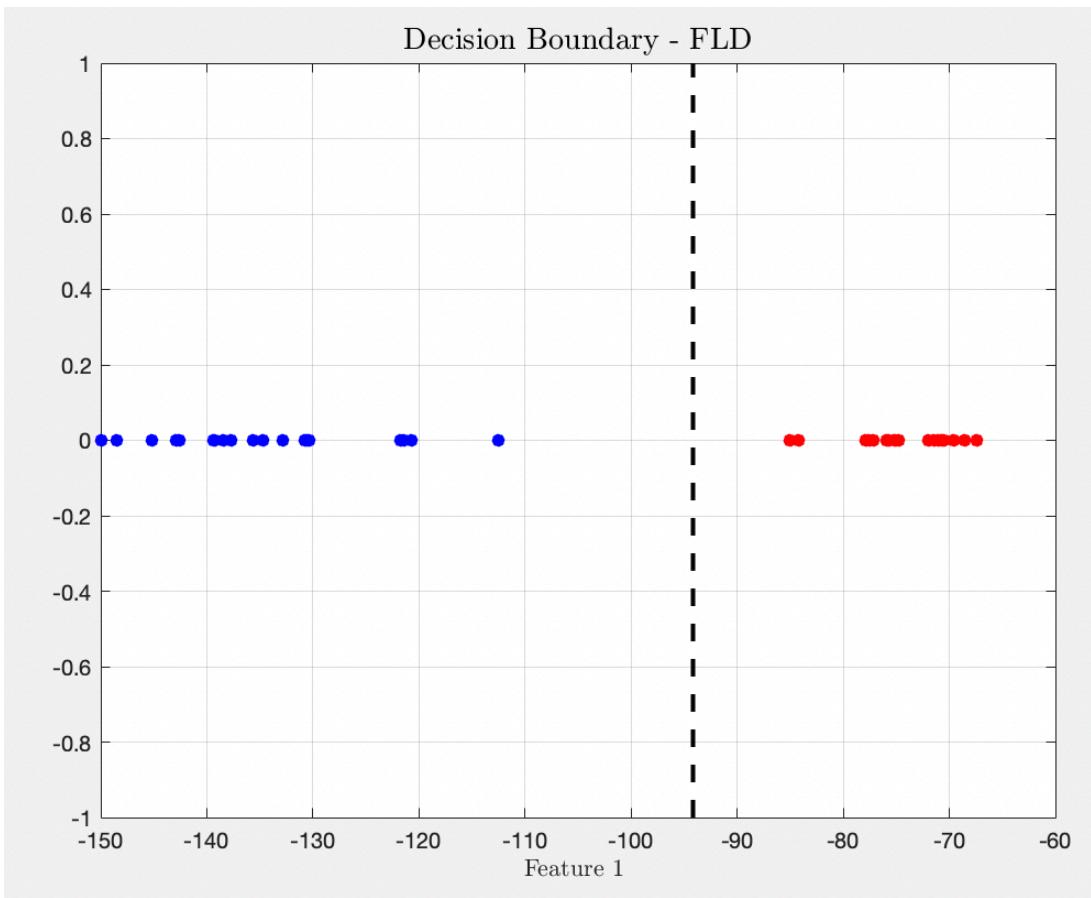
Accuracy of Bayes Classifier = 0.975

=====

(ج)

ویژگی‌های جدیدی که بدست آوریم، ویژگی‌های آماری میانگین سیگنال، انحراف معیار آن و Mean Absolute Deviation (MAD) بوده است.





Part b Results:

Accuracy of Mahalanobis Distance = 1

Accuracy of Euclidean Distance = 1

Accuracy of Bayes Classifier = 1

Part c Results:

Accuracy of Mahalanobis Distance = 0.575

Accuracy of Euclidean Distance = 0.675

Accuracy of Bayes Classifier = 0.7

Part d Results:

Accuracy of Mahalanobis Distance = 0.975

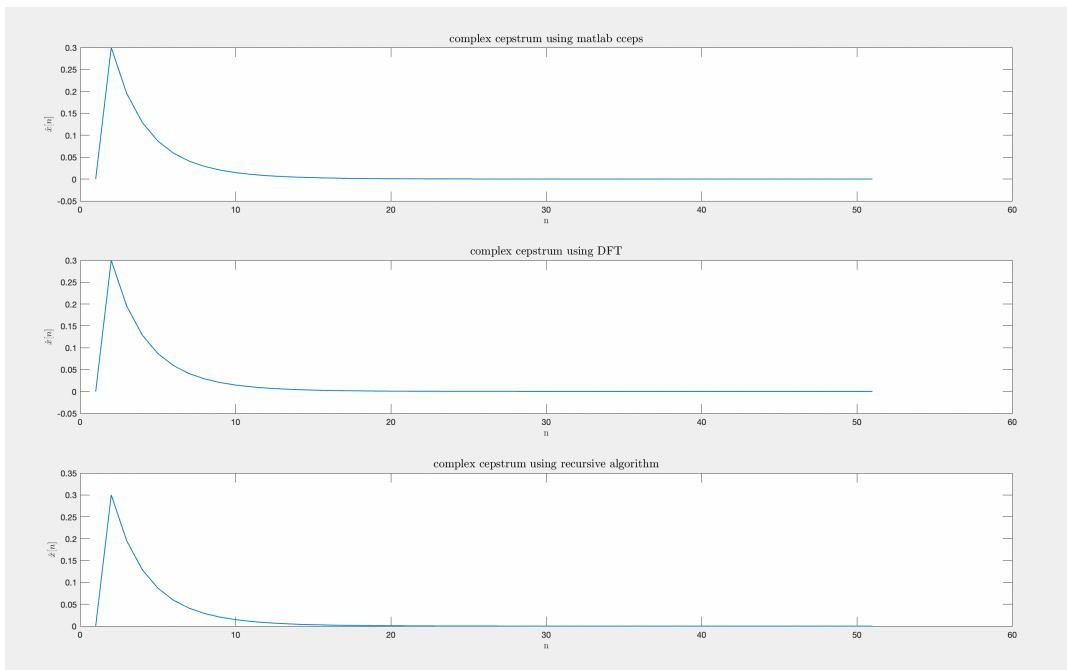
Accuracy of Euclidean Distance = 1

Accuracy of Bayes Classifier = 0.975

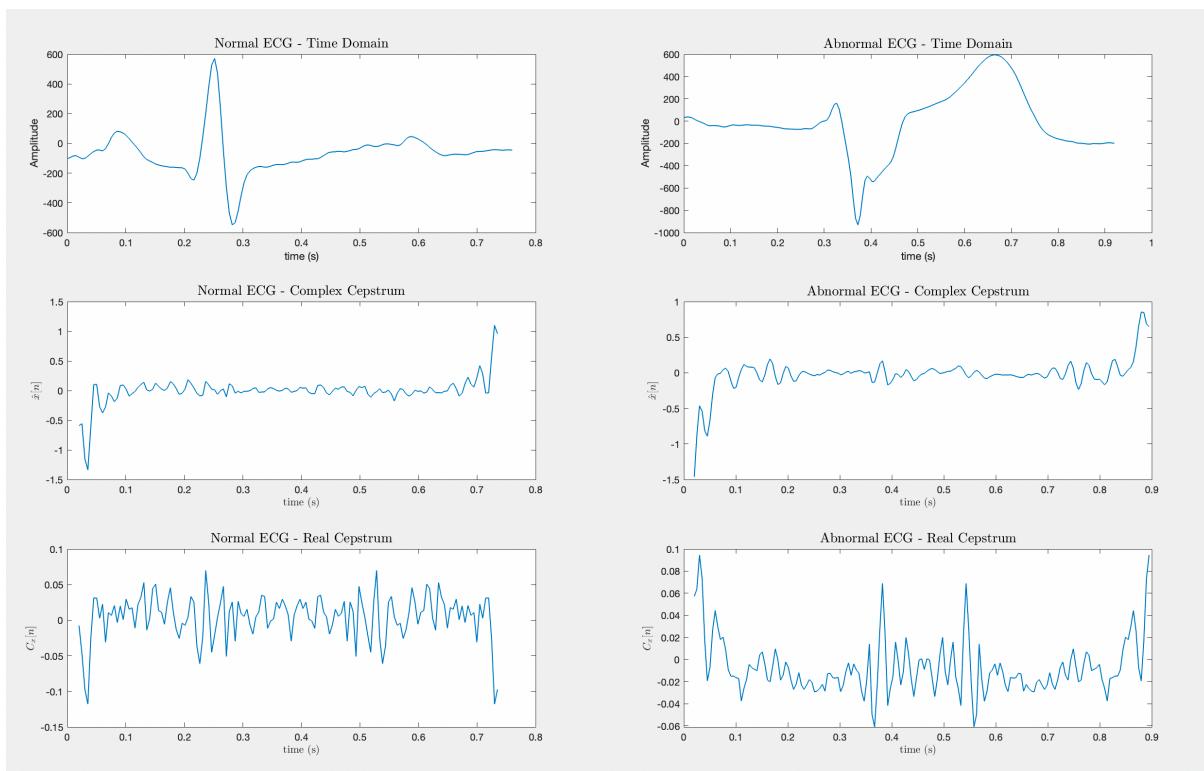
می‌بینیم که نتایج در حالت استفاده از سه ویژگی و همچنین حالت تک ویژگی کمی بهبود پیدا کرده است، اما در حالت کاهش بعد به دو ویژگی نتایج بدتر شده است.

سوال ۳-

(الف)



(ب)



از همان دیتاست موجود برای سوال ۲ استفاده کردیم.

فیلتر کالمن)

Driving fatigue detection based on brain source activity and ARMA model, Fahimeh Nadalizadeh, Mehdi Rajabioun, Amirreza Feyzi, Medical & Biological Engineering & Computing (2024) 62:1017–1030 , <https://doi.org/10.1007/s11517-023-02983-z>

مقاله‌ای که مورد بررسی قرار گرفت، روشی نوین برای تشخیص خستگی رانندگان با استفاده از سیگنال‌های الکتروانسفالوگرافی (EEG) ارائه داده است. این روش شامل استخراج فعالیت‌های مغزی از طریق روش‌های مکان‌یابی منبع و برآوردهای دینامیکی فعالیت منابع با استفاده از یک فیلتر دوگانه کالمن است. فیلتر کالمن دوگانه برای برآوردهای فعالیت منبع و روابط بین منابع به کار رفته و از این اطلاعات برای پیش‌بینی خستگی راننده استفاده می‌شود. مدل‌سازی زمانی فعالیت‌های مغزی و سیگنال‌های EEG نیز با استفاده از مدل‌های ARMA انجام شده است و از ویژگی‌های این مدل‌سازی برای تشخیص خستگی استفاده می‌کند. درنهایت، از روش‌های ترکیبی برای بهبود دقت تشخیص استفاده شده و نتایج نشان دهنده موفقیت این روش در مقایسه با روش‌های دیگر است.

طبقه بندی)

Analysis of ECG-based arrhythmia detection system using machine learning, Shikha Dhyani, Adesh Kumar, Sushabhan Choudhury, 2023, <https://doi.org/10.1016/j.mex.2023.102195>

مقاله مورد بررسی، به تجزیه و تحلیل سیستم تشخیص آریتمی قلبی با استفاده از سیگنال‌های الکتروکاردیوگرام (ECG) و روش‌های یادگیری ماشینی می‌پردازد. در این مطالعه، از تبدیل ویولت سه‌بعدی (3D Discrete Wavelet Transform) برای پیش‌پردازش و استخراج ویژگی‌های سیگنال ECG و همچنین استفاده از طبقه بند ماشین بردار پشتیبان (SVM) برای طبقه بندی ضربان‌های قلبی استفاده شده است. نتایج مقاله نشان می‌دهد که این روش‌ها قادر به تشخیص دقیق آریتمی‌های مختلف با دقت بالایی هستند. دقت کلی مدل SVM در طبقه بندی این داده‌ها بسیار بالا بوده و به بیش از 99٪ رسیده است. این نشان دهنده کارایی بالای ترکیب تبدیل موجک و SVM در تشخیص و طبقه بندی آریتمی‌ها می‌باشد. از این‌رو، این مدل می‌تواند به عنوان یک ابزار موثر در تشخیص زودهنگام و دقیق آریتمی‌های قلبی دریماران مورد استفاده قرار گیرد.

آنالیز کپستروم)

Cepstral Analysis-Based Artifact Detection, Recognition, and Removal in Prefrontal EEG. Issa, M. F., Tuboly, G., Kozmann, G., & Juhasz, Z. (2023). *IEEE Xplore*. DOI: 10.1109/MSR-2019-0016.

متاسفانه مقاله‌ای که از سال ۲۰۲۳ به بعد باشه و به صورت رایگان هم در دسترس باشه پیدا نکردم.

مدل مخفی مارکوف

N. J. Trujillo-Barreto, D. A. Galvez, A. Astudillo and W. El-Deredy, "Explicit Modeling of Brain State Duration Using Hidden Semi Markov Models in EEG Data," in *IEEE Access*, vol. 12, pp. 12335-12355, 2024, doi: 10.1109/ACCESS.2024.3354711.

در این مقاله، استفاده از مدل‌های نیمه‌مارکوف مخفی (HSMM) برای تجزیه و تحلیل داده‌های الکتروانسفالوگرافی (EEG) مورد بررسی قرار گرفته است. نویسنده‌گان بر محدودیت‌های مدل‌های مارکوف مخفی (HMM) که زمان‌های ماندگاری حالت را با توزیع هندسی مدل‌سازی می‌کنند تاکید و پیشنهاد داده‌اند که HSMM با امکان مدل‌سازی دقیق‌تر مدت زمان حالت مغزی می‌تواند این نقص را رفع کند. در این مطالعه، با استفاده از روش‌های بیزین و برآورد بیز متغیر، تعداد حالت‌های مغزی و توزیع‌های مدت زمان حالت تعیین و مدل‌ها بر روی داده‌های شبیه‌سازی شده و واقعی ارزیابی شده‌اند. نتایج نشان می‌دهند که HSMM در مقایسه با HMM عملکرد بهتری دارد و مدل‌سازی دقیق مدت زمان حالت برای دستیابی به پیش‌بینی‌های معتبر ضروری است.

کد:

```
%% CHW3 - Radin Khayyam - 99101579

%% Q1

clc; clear; close all;

% sigma_1 = input('Enter sigma_1: ');
% sigma_2 = input('Enter sigma_2: ');
% sigma_v = input('Enter sigma_v: ');
% a = input('Enter a: ');
% b = input('Enter b: ');

sigma_v = 1;
sigma_1 = 2;
sigma_2 = 2;
a = 0.5;
b = 0.5;

N = 250; % Number of samples

%% Q1 - Part a

clc;

v = sigma_v * randn(1, N);
U_1 = sigma_1 * randn(1, N);
```

```

U_2 = sigma_2 * randn(1, N);

% system real outputs

f1 = zeros(2,N);
f3 = zeros(1,N);

for k = 2:N
    f1(1,k) = a * f1(1,k-1) + U_1(k);
    f1(2,k) = b * f1(2,k-1) + f1(1,k) + U_2(k);
    f3(k) = f1(2,k) + V(k);
end

x_hat = myKalmanFilter(a,b,sigma_1,sigma_2,sigma_v,f3,N,0);
x_hat_rand_initial = myKalmanFilter(a,b,sigma_1,sigma_2,sigma_v,f3,N,1);

plotResults(f1, x_hat, x_hat_rand_initial);

PCC_X1 = corrcoef(f1(1,:),x_hat(1,:));
PCC_X2 = corrcoef(f1(2,:),x_hat(2,:));
PCC_X1_rand_initial = corrcoef(f1(1,:),x_hat_rand_initial(1,:));
PCC_X2_rand_initial = corrcoef(f1(2,:),x_hat_rand_initial(2,:));
disp(['Pearson Correlation Coeficient for X_1 = ',num2str(PCC_X1(1,2))]);
disp(['Pearson Correlation Coeficient for X_2 = ',num2str(PCC_X2(1,2))]);
disp(['Pearson Correlation Coeficient for X_1 with random initialization =
',num2str(PCC_X1_rand_initial(1,2))]);
disp(['Pearson Correlation Coeficient for X_2 with random initialization =
',num2str(PCC_X2_rand_initial(1,2))]);

%% Q1 - Part b

clc;

V = sqrt(12) * sigma_v * rand(1, N); % generating noise with uniform distribution and
variance sigma_v

U_1 = sqrt(12) * sigma_1 * rand(1, N);
U_2 = sqrt(12) * sigma_2 * rand(1, N);

```

```

% system real outputs

f1 = zeros(2,N);

f3 = zeros(1,N);

for k = 2:N

    f1(1,k) = a * f1(1,k-1) + U_1(k);

    f1(2,k) = b * f1(2,k-1) + f1(1,k) + U_2(k);

    f3(k) = f1(2,k) + V(k);

end

X_hat = myKalmanFilter(a,b,sigma_1,sigma_2,sigma_v,f3,N,0);

X_hat_rand_initial = myKalmanFilter(a,b,sigma_1,sigma_2,sigma_v,f3,N,1);

plotResults(f1, X_hat, X_hat_rand_initial);

PCC_X1 = corrcoef(f1(1,:),X_hat(1,:));

PCC_X2 = corrcoef(f1(2,:),X_hat(2,:));

PCC_X1_rand_initial = corrcoef(f1(1,:),X_hat_rand_initial(1,:));

PCC_X2_rand_initial = corrcoef(f1(2,:),X_hat_rand_initial(2,:));

disp(['Pearson Correlation Coeficient for X_1 = ',num2str(PCC_X1(1,2))]);

disp(['Pearson Correlation Coeficient for X_2 = ',num2str(PCC_X2(1,2))]);

disp(['Pearson Correlation Coeficient for X_1 with random initialization =
',num2str(PCC_X1_rand_initial(1,2))]);

disp(['Pearson Correlation Coeficient for X_2 with random initialization =
',num2str(PCC_X2_rand_initial(1,2))]);

%% Q1 - Part c

clc;

V = sigma_v * randn(1, N);

U_1 = sigma_1 * randn(1, N);

U_2 = U_1 * sigma_2/sigma_1; % make correlation between U_1 and U_2

% system real outputs

f1 = zeros(2,N);

```

```

f3 = zeros(1,N);

for k = 2:N

    f1(1,k) = a * f1(1,k-1) + U_1(k);
    f1(2,k) = b * f1(2,k-1) + f1(1,k) + U_2(k);
    f3(k) = f1(2,k) + V(k);

end

x_hat = myKalmanFilter(a,b,sigma_1,sigma_2,sigma_v,f3,N,0);
x_hat_rand_initial = myKalmanFilter(a,b,sigma_1,sigma_2,sigma_v,f3,N,1);

plotResults(f1, x_hat, x_hat_rand_initial);

PCC_X1 = corrcoef(f1(1,:),x_hat(1,:));
PCC_X2 = corrcoef(f1(2,:),x_hat(2,:));
PCC_X1_rand_initial = corrcoef(f1(1,:),x_hat_rand_initial(1,:));
PCC_X2_rand_initial = corrcoef(f1(2,:),x_hat_rand_initial(2,:));
disp(['Pearson Correlation Coeficient for X_1 = ',num2str(PCC_X1(1,2))]);
disp(['Pearson Correlation Coeficient for X_2 = ',num2str(PCC_X2(1,2))]);
disp(['Pearson Correlation Coeficient for X_1 with random initialization =
',num2str(PCC_X1_rand_initial(1,2))]);
disp(['Pearson Correlation Coeficient for X_2 with random initialization =
',num2str(PCC_X2_rand_initial(1,2))]);

%% Q2

clc; clear;

load('ECG.mat');

train_data = {ECG01, ECG02, ECG03, ECG04, ECG05, ECG06, ECG07, ECG08, ECG09, ECG10, ...
    ECG11, ECG12, ECG13, ECG14, ECG15, ECG16, ECG17, ECG18, ECG19, ECG20, ...
    ECG21, ECG22, ECG23, ECG24, ECG25, ECG26, ECG27, ECG28, ECG29, ECG30, ...
    ECG31, ECG32, ECG33, ECG34, ECG35, ECG36, ECG37, ECG38, ECG39, ECG40};

test_data = {ECG41, ECG42, ECG43, ECG44, ECG45, ECG46, ECG47, ECG48, ECG49, ECG50, ...};

```

```

    ECG51, ECG52, ECG53, ECG54, ECG55, ECG56, ECG57, ECG58, ECG59, ECG60, ...
    ECG61, ECG62, ECG63, ECG64, ECG65, ECG66, ECG67, ECG68, ECG69, ECG70, ...
    ECG71, ECG72, ECG73, ECG74, ECG75, ECG76, ECG77, ECG78, ECG79, ECG80};

%% Q2 - Part a

clc;

num_train = length(train_data);

features_train = zeros(3,num_train);

labels_train = zeros(1,num_train);

for i = 1:num_train

    signal = train_data{i};

    features_train (:, i) = featureExtractor(signal);

    if(i>20)

        labels_train(i) = 1;

    end

end

% % normalization - z score

% for i = 1:3

%     features_train(i,:) = (features_train(i,:) - mean(features_train(i,:)))/
var(features_train(i,:));

% end

%% Q2 - Part b

clc;

normal_features = features_train(:, labels_train == 0);

abnormal_features = features_train(:, labels_train == 1);

mu_normal = mean(normal_features,2);

```

```

mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

feature1_range = linspace(min(features_train(1,:)), max(features_train(1,:)), 100);
feature2_range = linspace(min(features_train(2,:)), max(features_train(2,:)), 100);
feature3_range = linspace(min(features_train(3,:)), max(features_train(3,:)), 100);

[f1, f2, f3] = meshgrid(feature1_range, feature2_range, feature3_range);

g_normal = zeros(size(f1));
g_abnormal = zeros(size(f1));

f1_flat = f1(:);
f2_flat = f2(:);
f3_flat = f3(:);

for i = 1:length(f1_flat)

    x = [f1_flat(i); f2_flat(i); f3_flat(i)];

    g_normal(i) = -0.5 * (x - mu_normal)' * inv(cov_normal) * (x-mu_normal) +
log(prior_normal) - 0.5 * log(det(cov_normal));

    g_abnormal(i) = -0.5 * (x - mu_abnormal)' * inv(cov_abnormal) * (x-mu_abnormal) +
log(prior_abnormal) - 0.5 * log(det(cov_abnormal));

end

g_normal = reshape(g_normal, size(f1));
g_abnormal = reshape(g_abnormal, size(f1));

```

```

g12 = g_normal - g_abnormal;

figure;
isosurface(f1, f2, f3, g12, 0,'k');
hold on;
scatter3(normal_features(1,:), normal_features(2,:), normal_features(3,:), 'r','filled');
hold on;
scatter3(abnormal_features(1,:), abnormal_features(2,:), abnormal_features(3,:),
'b','filled');
xlabel('Feature 1 ','Interpreter','latex');
ylabel('Feature 2 ','Interpreter','latex');
zlabel('Feature 3 ','Interpreter','latex');
title('Decision Boundary','Interpreter','latex','FontSize',14);
grid on;

%% Q2 - Part c
clc;

cov_all = cov(features_train');
[V, D] = eig(cov_all);
eigenvalues = diag(D);
[sortedEigenvalues, ind] = sort(eigenvalues, 'descend');
sortedEigenvectors = V(:, ind);

u1 = sortedEigenvectors(:,2:3);

features_train_reduced = u1' * features_train ;

normal_features = features_train_reduced(:, labels_train == 0);
abnormal_features = features_train_reduced(:, labels_train == 1);

```

```

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

feature1_range = linspace(min(features_train_reduced(1,:)),
max(features_train_reduced(1,:)), 100);
feature2_range = linspace(min(features_train_reduced(2,:)),
max(features_train_reduced(2,:)), 100);

[f1, f2] = meshgrid(feature1_range, feature2_range);

g_normal = zeros(size(f1));
g_abnormal = zeros(size(f1));

f1_flat = f1(:);
f2_flat = f2(:);

for i = 1:length(f1_flat)

x = [f1_flat(i); f2_flat(i)];

g_normal(i) = -0.5 * (x - mu_normal)' * inv(cov_normal) * (x-mu_normal) +
log(prior_normal) - 0.5 * log(det(cov_normal));

g_abnormal(i) = -0.5 * (x - mu_abnormal)' * inv(cov_abnormal) * (x-mu_abnormal) +
log(prior_abnormal) - 0.5 * log(det(cov_abnormal));

end

```

```

g_normal = reshape(g_normal, size(f1));
g_abnormal = reshape(g_abnormal, size(f1));

g12 = g_normal - g_abnormal;

figure;
contour(f1, f2, g12, [0 0]);
hold on;
scatter(normal_features(1,:), normal_features(2,:), 'r','filled');
hold on;
scatter(abnormal_features(1,:), abnormal_features(2,:), 'b','filled');
xlabel('Feature 1 ','Interpreter','latex');
ylabel('Feature 2 ','Interpreter','latex');
zlabel('Feature 3 ','Interpreter','latex');
title('Decision Boundary - Minimizing Entropy','Interpreter','latex','FontSize',14);
grid on;

%% Q2 - Part d

clc;

normal_features = features_train(:, labels_train == 0);
abnormal_features = features_train(:, labels_train == 1);

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

```

```

S_w = prior_normal * cov_normal + prior_abnormal * cov_abnormal;
w = inv(S_w)*(mu_normal - mu_abnormal);
features_train_reduced = w' * features_train;

normal_features = features_train_reduced(:, labels_train == 0);
abnormal_features = features_train_reduced(:, labels_train == 1);

mu_normal = mean(normal_features);
mu_abnormal = mean(abnormal_features);

var_normal = var(normal_features);
var_abnormal = var(abnormal_features);

feature1_range = linspace(min(features_train_reduced), max(features_train_reduced),
1000);

f1 = feature1_range;

g_normal = zeros(size(f1));
g_abnormal = zeros(size(f1));

f1_flat = f1(:);

for i = 1:length(f1_flat)

x = [f1_flat(i)];

g_normal(i) = -0.5 * (x - mu_normal)^2 / var_normal + log(prior_normal) - 0.5 *
log(var_normal);
g_abnormal(i) = -0.5 * (x - mu_abnormal)^2 / var_abnormal + log(prior_abnormal) - 0.5 *
log(var_abnormal);

end

```

```

g_normal = reshape(g_normal, size(f1));
g_abnormal = reshape(g_abnormal, size(f1));

g12 = g_normal - g_abnormal;

boundry = find(g12 == min(abs(g12))) / 1000 + feature1_range(1);
if isempty(boundry == 1)

    boundry = find(g12 == -1*min(abs(g12))) / 1000 * (feature1_range(end) -
feature1_range(1)) + feature1_range(1) ;

end

figure;
plot([boundry boundry], [-1 1], 'k--', 'LineWidth', 2);
hold on;
scatter(normal_features, 0, 'r','filled');
hold on;
scatter(abnormal_features, 0, 'b','filled');
xlabel('Feature 1 ','Interpreter','latex');
title('Decision Boundary - FLD','Interpreter','latex','FontSize',14);
grid on;

%% Q2 - Part e

clc;

% part a

num_test = length(test_data);
features_test = zeros(3,num_test);
labels_test = zeros(1,num_test);

for i = 1:num_test
    signal = test_data{i};

```

```

features_test (:, i) = featureExtractor(signal);

if(i>20)

labels_test(i) = 1;

end

end

% part b

normal_features_train = features_train(:, labels_train == 0);

abnormal_features_train = features_train(:, labels_train == 1);

mu_normal = mean(normal_features_train,2);

mu_abnormal = mean(abnormal_features_train,2);

cov_normal = cov(normal_features_train');

cov_abnormal = cov(abnormal_features_train');

predicted_labels_mah = zeros(1, num_test);

predicted_labels_euc = zeros(1, num_test);

predicted_labels_bayes = zeros(1, num_test);

for i=1:num_test

d_normal_mah = (features_test(:,i)-mu_normal)'*inv(cov_normal)*(features_test(:,i)-
mu_normal);

d_abnormal_mah = (features_test(:,i)-
mu_abnormal)'*inv(cov_abnormal)*(features_test(:,i)-mu_abnormal);

d_normal_euc = (features_test(:,i)-mu_normal)'*(features_test(:,i)-mu_normal);

d_abnormal_euc = (features_test(:,i)-mu_abnormal)'*(features_test(:,i)-mu_abnormal);

g_normal_bayes = -0.5 * (features_test(:,i) - mu_normal)' * inv(cov_normal) *
(features_test(:,i)-mu_normal) + log(prior_normal) - 0.5 * log(det(cov_normal));

g_abnormal_bayes = -0.5 * (features_test(:,i) - mu_abnormal)' * inv(cov_abnormal) *
(features_test(:,i)-mu_abnormal) + log(prior_abnormal) - 0.5 * log(det(cov_abnormal));

```

```

if (d_abnormal_mah < d_normal_mah)
    predicted_labels_mah(i) = 1;
elseif (d_abnormal_mah == d_normal_mah)
    if (randn > 0)
        predicted_labels_mah(i) = 1;
    end
end

if (d_abnormal_euc < d_normal_euc)
    predicted_labels_euc(i) = 1;
elseif (d_abnormal_euc == d_normal_euc)
    if (randn > 0)
        predicted_labels_euc(i) = 1;
    end
end

if (g_abnormal_bayes - g_normal_bayes > 0)
    predicted_labels_bayes(i) = 1;
elseif (g_abnormal_bayes - g_normal_bayes == 0)
    if (randn > 0)
        predicted_labels_bayes(i) = 1;
    end
end

end

acc_mah = sum(labels_test == predicted_labels_mah)/num_test;
acc_euc = sum(labels_test == predicted_labels_euc)/num_test;
acc_bayes = sum(labels_test == predicted_labels_bayes)/num_test;
disp('Part b Results:')
disp(['Accuracy of Mahalanobis Distance = ',num2str(acc_mah)]);
disp(['Accuracy of Euclidean Distance = ',num2str(acc_euc)]);

```

```

disp(['Accuracy of Bayes Classifier = ',num2str(acc_bayes )]);
disp('=====')
```

% part c

```

cov_all_train = cov(features_train');
[V, D] = eig(cov_all_train);
eigenvalues = diag(D);
[sortedEigenvalues, ind] = sort(eigenvalues, 'descend');
sortedEigenvectors = V(:, ind);

u1 = sortedEigenvectors(:,2:3);

features_train_reduced = u1' * features_train ;
features_test_reduced = u1' * features_test ;

normal_features = features_train_reduced(:, labels_train == 0);
abnormal_features = features_train_reduced(:, labels_train == 1);

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

predicted_labels_mah = zeros(1, num_test);
predicted_labels_euc = zeros(1, num_test);
predicted_labels_bayes = zeros(1, num_test);
for i=1:num_test

    d_normal_mah = (features_test_reduced(:,i)-
mu_normal)'*inv(cov_normal)*(features_test_reduced(:,i)-mu_normal);
    d_abnormal_mah = (features_test_reduced(:,i)-
mu_abnormal)'*inv(cov_abnormal)*(features_test_reduced(:,i)-mu_abnormal);
```

```

d_normal_euc = (features_test_reduced(:,i)-mu_normal)'*(features_test_reduced(:,i)-
mu_normal);

d_abnormal_euc = (features_test_reduced(:,i)-
mu_abnormal)*(features_test_reduced(:,i)-mu_abnormal);

g_normal_bayes = -0.5 * (features_test_reduced(:,i) - mu_normal)' * inv(cov_normal) *
(features_test_reduced(:,i)-mu_normal) + log(prior_normal) - 0.5 * log(det(cov_normal));

g_abnormal_bayes = -0.5 * (features_test_reduced(:,i) - mu_abnormal)' *
inv(cov_abnormal) * (features_test_reduced(:,i)-mu_abnormal) + log(prior_abnormal) - 0.5
* log(det(cov_abnormal));

if (d_abnormal_mah < d_normal_mah)
    predicted_labels_mah(i) = 1;

elseif (d_abnormal_mah == d_normal_mah)
    if (randn > 0)
        predicted_labels_mah(i) = 1;

    end
end

if (d_abnormal_euc < d_normal_euc)
    predicted_labels_euc(i) = 1;

elseif (d_abnormal_euc == d_normal_euc)
    if (randn > 0)
        predicted_labels_euc(i) = 1;

    end
end

if (g_abnormal_bayes - g_normal_bayes > 0)
    predicted_labels_bayes(i) = 1;

elseif (g_abnormal_bayes - g_normal_bayes == 0)
    if (randn > 0)
        predicted_labels_bayes(i) = 1;

    end
end

```

```

end

end

acc_mah = sum(labels_test == predicted_labels_mah)/num_test;
acc_euc = sum(labels_test == predicted_labels_euc)/num_test;
acc_bayes = sum(labels_test == predicted_labels_bayes)/num_test;
disp('Part c Results:')
disp(['Accuracy of Mahalanobis Distance = ',num2str(acc_mah)]);
disp(['Accuracy of Euclidean Distance = ',num2str(acc_euc)]);
disp(['Accuracy of Bayes Classifier = ',num2str(acc_bayes)]);
disp('=====')

% Part d

normal_features = features_train(:, labels_train == 0);
abnormal_features = features_train(:, labels_train == 1);

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

S_w = prior_normal * cov_normal + prior_abnormal * cov_abnormal;
w = inv(S_w)*(mu_normal - mu_abnormal);

features_train_reduced = w' * features_train;
features_test_reduced = w' * features_test;

```

```

normal_features = features_train_reduced(:, labels_train == 0);

abnormal_features = features_train_reduced(:, labels_train == 1);

mu_normal = mean(normal_features);

mu_abnormal = mean(abnormal_features);

var_normal = var(normal_features);

var_abnormal = var(abnormal_features);

predicted_labels_mah = zeros(1, num_test);

predicted_labels_euc = zeros(1, num_test);

predicted_labels_bayes = zeros(1, num_test);

for i=1:num_test

d_normal_mah = (features_test_reduced(i)-mu_normal)^2 / var_normal;

d_abnormal_mah = (features_test_reduced(i)-mu_abnormal)^2 / var_abnormal;

d_normal_euc = (features_test_reduced(:,i)-mu_normal)^2;

d_abnormal_euc = (features_test_reduced(:,i)-mu_abnormal)^2;

g_normal_bayes = -0.5 * (features_test_reduced(i) - mu_normal)^2 / var_normal + log(prior_normal) - 0.5 * log(var_normal);

g_abnormal_bayes = -0.5 * (features_test_reduced(i) - mu_abnormal)^2 / var_abnormal + log(prior_abnormal) - 0.5 * log(var_abnormal);

if (d_abnormal_mah < d_normal_mah)

predicted_labels_mah(i) = 1;

elseif (d_abnormal_mah == d_normal_mah)

if (randn > 0)

predicted_labels_mah(i) = 1;

end

end

```

```

if (d_abnormal_euc < d_normal_euc)

    predicted_labels_euc(i) = 1;

elseif (d_abnormal_euc == d_normal_euc)

    if (randn > 0)

        predicted_labels_euc(i) = 1;

    end

end


if (g_abnormal_bayes - g_normal_bayes > 0)

    predicted_labels_bayes(i) = 1;

elseif (g_abnormal_bayes - g_normal_bayes == 0)

    if (randn > 0)

        predicted_labels_bayes(i) = 1;

    end

end

end


acc_mah = sum(labels_test == predicted_labels_mah)/num_test;
acc_euc = sum(labels_test == predicted_labels_euc)/num_test;
acc_bayes = sum(labels_test == predicted_labels_bayes)/num_test;

disp('Part d Results:')

disp(['Accuracy of Mahalanobis Distance = ',num2str(acc_mah )]);
disp(['Accuracy of Euclidean Distance = ',num2str(acc_euc )]);
disp(['Accuracy of Bayes Classifier = ',num2str(acc_bayes )]);

disp('=====')  

%% Q2 - Part f - a  

clc; clear;  

load('ECG.mat');  

train_data = {ECG01, ECG02, ECG03, ECG04, ECG05, ECG06, ECG07, ECG08, ECG09, ECG10, ...
ECG11, ECG12, ECG13, ECG14, ECG15, ECG16, ECG17, ECG18, ECG19, ECG20, ...
ECG21, ECG22, ECG23, ECG24, ECG25, ECG26, ECG27, ECG28, ECG29, ECG30, ...
ECG31, ECG32, ECG33, ECG34, ECG35, ECG36, ECG37, ECG38, ECG39, ECG40};

```

```

test_data = {ECG41, ECG42, ECG43, ECG44, ECG45, ECG46, ECG47, ECG48, ECG49, ECG50, ...
             ECG51, ECG52, ECG53, ECG54, ECG55, ECG56, ECG57, ECG58, ECG59, ECG60, ...
             ECG61, ECG62, ECG63, ECG64, ECG65, ECG66, ECG67, ECG68, ECG69, ECG70, ...
             ECG71, ECG72, ECG73, ECG74, ECG75, ECG76, ECG77, ECG78, ECG79, ECG80};

num_train = length(train_data);
features_train = zeros(3,num_train);
labels_train = zeros(1,num_train);

for i = 1:num_train
    signal = train_data{i};
    features_train (:, i) = featureExtractor_new(signal);

    if(i>20)
        labels_train(i) = 1;
    end
end

% % normalization - z score
% for i = 1:3
%     features_train(i,:) = (features_train(i,:) - mean(features_train(i,:)))/
% var(features_train(i,:));
% end

%% Q2 - Part f - b
clc;

normal_features = features_train(:, labels_train == 0);
abnormal_features = features_train(:, labels_train == 1);

```

```

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

feature1_range = linspace(min(features_train(1,:)), max(features_train(1,:)), 100);
feature2_range = linspace(min(features_train(2,:)), max(features_train(2,:)), 100);
feature3_range = linspace(min(features_train(3,:)), max(features_train(3,:)), 100);

[f1, f2, f3] = meshgrid(feature1_range, feature2_range, feature3_range);

g_normal = zeros(size(f1));
g_abnormal = zeros(size(f1));

f1_flat = f1(:);
f2_flat = f2(:);
f3_flat = f3(:);

for i = 1:length(f1_flat)

    x = [f1_flat(i); f2_flat(i); f3_flat(i)];

    g_normal(i) = -0.5 * (x - mu_normal)' * inv(cov_normal) * (x-mu_normal) +
log(prior_normal) - 0.5 * log(det(cov_normal));

    g_abnormal(i) = -0.5 * (x - mu_abnormal)' * inv(cov_abnormal) * (x-mu_abnormal) +
log(prior_abnormal) - 0.5 * log(det(cov_abnormal));

end

g_normal = reshape(g_normal, size(f1));

```

```

g_abnormal = reshape(g_abnormal, size(f1));

g12 = g_normal - g_abnormal;

figure;

isosurface(f1, f2, f3, g12, 0,'k');

hold on;

scatter3(normal_features(1,:), normal_features(2,:), normal_features(3,:), 'r','filled');

hold on;

scatter3(abnormal_features(1,:), abnormal_features(2,:), abnormal_features(3,:),
'b','filled');

xlabel('Feature 1 ','Interpreter','latex');

ylabel('Feature 2 ','Interpreter','latex');

zlabel('Feature 3 ','Interpreter','latex');

title('Decision Boundary','Interpreter','latex','FontSize',14);

grid on;

%% Q2 - Part f - c

clc;

cov_all = cov(features_train');

[V, D] = eig(cov_all);

eigenvalues = diag(D);

[sortedEigenvalues, ind] = sort(eigenvalues, 'descend');

sortedEigenvectors = V(:, ind);

u1 = sortedEigenvectors(:,2:3);

features_train_reduced = u1' * features_train ;

normal_features = features_train_reduced(:, labels_train == 0);

abnormal_features = features_train_reduced(:, labels_train == 1);

```

```

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

feature1_range = linspace(min(features_train_reduced(1,:)),
max(features_train_reduced(1,:)), 100);
feature2_range = linspace(min(features_train_reduced(2,:)),
max(features_train_reduced(2,:)), 100);

[f1, f2] = meshgrid(feature1_range, feature2_range);

g_normal = zeros(size(f1));
g_abnormal = zeros(size(f1));

f1_flat = f1(:);
f2_flat = f2(:);

for i = 1:length(f1_flat)

x = [f1_flat(i); f2_flat(i)];

g_normal(i) = -0.5 * (x - mu_normal)' * inv(cov_normal) * (x-mu_normal) +
log(prior_normal) - 0.5 * log(det(cov_normal));

g_abnormal(i) = -0.5 * (x - mu_abnormal)' * inv(cov_abnormal) * (x-mu_abnormal) +
log(prior_abnormal) - 0.5 * log(det(cov_abnormal));

end

```

```

g_normal = reshape(g_normal, size(f1));
g_abnormal = reshape(g_abnormal, size(f1));

g12 = g_normal - g_abnormal;

figure;
contour(f1, f2, g12, [0 0]);
hold on;
scatter(normal_features(1,:), normal_features(2,:), 'r','filled');
hold on;
scatter(abnormal_features(1,:), abnormal_features(2,:), 'b','filled');
xlabel('Feature 1 ','Interpreter','latex');
ylabel('Feature 2 ','Interpreter','latex');
zlabel('Feature 3 ','Interpreter','latex');
title('Decision Boundary - Minimizing Entropy','Interpreter','latex','FontSize',14);
grid on;

%% Q2 - Part f - d
clc;

normal_features = features_train(:, labels_train == 0);
abnormal_features = features_train(:, labels_train == 1);

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

```

```

S_w = prior_normal * cov_normal + prior_abnormal * cov_abnormal;
w = inv(S_w)*(mu_normal - mu_abnormal);
features_train_reduced = w' * features_train;

normal_features = features_train_reduced(:, labels_train == 0);
abnormal_features = features_train_reduced(:, labels_train == 1);

mu_normal = mean(normal_features);
mu_abnormal = mean(abnormal_features);

var_normal = var(normal_features);
var_abnormal = var(abnormal_features);

feature1_range = linspace(min(features_train_reduced), max(features_train_reduced),
1000);

f1 = feature1_range;

g_normal = zeros(size(f1));
g_abnormal = zeros(size(f1));

f1_flat = f1(:);

for i = 1:length(f1_flat)

x = [f1_flat(i)];

g_normal(i) = -0.5 * (x - mu_normal)^2 / var_normal + log(prior_normal) - 0.5 *
log(var_normal);
g_abnormal(i) = -0.5 * (x - mu_abnormal)^2 / var_abnormal + log(prior_abnormal) - 0.5 *
log(var_abnormal);

```

```

end

g_normal = reshape(g_normal, size(f1));
g_abnormal = reshape(g_abnormal, size(f1));

g12 = g_normal - g_abnormal;

boundary = find(g12 == min(abs(g12))) / 1000 + feature1_range(1);
if isempty(boundary == 1)

    boundary = find(g12 == -1*min(abs(g12))) / 1000 * (feature1_range(end) -
feature1_range(1)) + feature1_range(1) ;

end

figure;

plot([boundary boundary], [-1 1], 'k--', 'LineWidth', 2);
hold on;
scatter(normal_features, 0, 'r','filled');
hold on;
scatter(abnormal_features, 0, 'b','filled');
xlabel('Feature 1 ','Interpreter','latex');
title('Decision Boundary - FLD','Interpreter','latex','FontSize',14);
grid on;

%% Q2 - Part f - e

clc;

% part a

num_test = length(test_data);
features_test = zeros(3,num_test);
labels_test = zeros(1,num_test);

for i = 1:num_test

```

```

signal = test_data{i};

features_test (:, i) = featureExtractor_new(signal);

if(i>20)

    labels_test(i) = 1;

end

end

% part b

normal_features_train = features_train(:, labels_train == 0);

abnormal_features_train = features_train(:, labels_train == 1);

mu_normal = mean(normal_features_train,2);

mu_abnormal = mean(abnormal_features_train,2);

cov_normal = cov(normal_features_train');

cov_abnormal = cov(abnormal_features_train');

predicted_labels_mah = zeros(1, num_test);

predicted_labels_euc = zeros(1, num_test);

predicted_labels_bayes = zeros(1, num_test);

for i=1:num_test

    d_normal_mah = (features_test(:,i)-mu_normal)'*inv(cov_normal)*(features_test(:,i)-mu_normal);

    d_abnormal_mah = (features_test(:,i)-mu_abnormal)'*inv(cov_abnormal)*(features_test(:,i)-mu_abnormal);

    d_normal_euc = (features_test(:,i)-mu_normal)'*(features_test(:,i)-mu_normal);

    d_abnormal_euc = (features_test(:,i)-mu_abnormal)'*(features_test(:,i)-mu_abnormal);

    g_normal_bayes = -0.5 * (features_test(:,i) - mu_normal)' * inv(cov_normal) *
(features_test(:,i)-mu_normal) + log(prior_normal) - 0.5 * log(det(cov_normal));

    g_abnormal_bayes = -0.5 * (features_test(:,i) - mu_abnormal)' * inv(cov_abnormal) *
(features_test(:,i)-mu_abnormal) + log(prior_abnormal) - 0.5 * log(det(cov_abnormal));

```

```

if (d_abnormal_mah < d_normal_mah)
    predicted_labels_mah(i) = 1;
elseif (d_abnormal_mah == d_normal_mah)
    if (randn > 0)
        predicted_labels_mah(i) = 1;
    end
end

if (d_abnormal_euc < d_normal_euc)
    predicted_labels_euc(i) = 1;
elseif (d_abnormal_euc == d_normal_euc)
    if (randn > 0)
        predicted_labels_euc(i) = 1;
    end
end

if (g_abnormal_bayes - g_normal_bayes > 0)
    predicted_labels_bayes(i) = 1;
elseif (g_abnormal_bayes - g_normal_bayes == 0)
    if (randn > 0)
        predicted_labels_bayes(i) = 1;
    end
end

end

acc_mah = sum(labels_test == predicted_labels_mah)/num_test;
acc_euc = sum(labels_test == predicted_labels_euc)/num_test;
acc_bayes = sum(labels_test == predicted_labels_bayes)/num_test;
disp('Part b Results:')
disp(['Accuracy of Mahalanobis Distance = ',num2str(acc_mah)]);

```

```

disp(['Accuracy of Euclidean Distance = ',num2str(acc_euc)]);
disp(['Accuracy of Bayes Classifier = ',num2str(acc_bayes)]);
disp('=====')
```

% part c

```

cov_all_train = cov(features_train');
[V, D] = eig(cov_all_train);
eigenvalues = diag(D);
[sortedEigenvalues, ind] = sort(eigenvalues, 'descend');
sortedEigenvectors = V(:, ind);

u1 = sortedEigenvectors(:,2:3);

features_train_reduced = u1' * features_train ;
features_test_reduced = u1' * features_test ;

normal_features = features_train_reduced(:, labels_train == 0);
abnormal_features = features_train_reduced(:, labels_train == 1);

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

predicted_labels_mah = zeros(1, num_test);
predicted_labels_euc = zeros(1, num_test);
predicted_labels_bayes = zeros(1, num_test);

for i=1:num_test

    d_normal_mah = (features_test_reduced(:,i)-
mu_normal)'*inv(cov_normal)*(features_test_reduced(:,i)-mu_normal);

```

```

d_abnormal_mah = (features_test_reduced(:,i)-
mu_abnormal)'*inv(cov_abnormal)*(features_test_reduced(:,i)-mu_abnormal);

d_normal_euc = (features_test_reduced(:,i)-mu_normal)'*(features_test_reduced(:,i)-
mu_normal);

d_abnormal_euc = (features_test_reduced(:,i)-
mu_abnormal)'*(features_test_reduced(:,i)-mu_abnormal);

g_normal_bayes = -0.5 * (features_test_reduced(:,i) - mu_normal)' * inv(cov_normal) *
(features_test_reduced(:,i)-mu_normal) + log(prior_normal) - 0.5 * log(det(cov_normal));

g_abnormal_bayes = -0.5 * (features_test_reduced(:,i) - mu_abnormal)' *
inv(cov_abnormal) * (features_test_reduced(:,i)-mu_abnormal) + log(prior_abnormal) - 0.5
* log(det(cov_abnormal));

if (d_abnormal_mah < d_normal_mah)
    predicted_labels_mah(i) = 1;
elseif (d_abnormal_mah == d_normal_mah)
    if (randn > 0)
        predicted_labels_mah(i) = 1;
    end
end

if (d_abnormal_euc < d_normal_euc)
    predicted_labels_euc(i) = 1;
elseif (d_abnormal_euc == d_normal_euc)
    if (randn > 0)
        predicted_labels_euc(i) = 1;
    end
end

if (g_abnormal_bayes - g_normal_bayes > 0)
    predicted_labels_bayes(i) = 1;
elseif (g_abnormal_bayes - g_normal_bayes == 0)
    if (randn > 0)

```

```

predicted_labels_bayes(i) = 1;

end

end

acc_mah = sum(labels_test == predicted_labels_mah)/num_test;
acc_euc = sum(labels_test == predicted_labels_euc)/num_test;
acc_bayes = sum(labels_test == predicted_labels_bayes)/num_test;
disp('Part c Results:')
disp(['Accuracy of Mahalanobis Distance = ',num2str(acc_mah)]);
disp(['Accuracy of Euclidean Distance = ',num2str(acc_euc)]);
disp(['Accuracy of Bayes Classifier = ',num2str(acc_bayes)]);
disp('=====')

% Part d

normal_features = features_train(:, labels_train == 0);
abnormal_features = features_train(:, labels_train == 1);

mu_normal = mean(normal_features,2);
mu_abnormal = mean(abnormal_features,2);

cov_normal = cov(normal_features');
cov_abnormal = cov(abnormal_features');

prior_normal = 0.5;
prior_abnormal = 0.5;

S_w = prior_normal * cov_normal + prior_abnormal * cov_abnormal;
w = inv(S_w)*(mu_normal - mu_abnormal);

features_train_reduced = w' * features_train;

```

```

features_test_reduced = w' * features_test;

normal_features = features_train_reduced(:, labels_train == 0);
abnormal_features = features_train_reduced(:, labels_train == 1);

mu_normal = mean(normal_features);
mu_abnormal = mean(abnormal_features);

var_normal = var(normal_features);
var_abnormal = var(abnormal_features);

predicted_labels_mah = zeros(1, num_test);
predicted_labels_euc = zeros(1, num_test);
predicted_labels_bayes = zeros(1, num_test);

for i=1:num_test

d_normal_mah = (features_test_reduced(i)-mu_normal)^2 / var_normal;
d_abnormal_mah = (features_test_reduced(i)-mu_abnormal)^2 / var_abnormal;

d_normal_euc = (features_test_reduced(:,i)-mu_normal)^2;
d_abnormal_euc = (features_test_reduced(:,i)-mu_abnormal)^2;

g_normal_bayes = -0.5 * (features_test_reduced(i) - mu_normal)^2 / var_normal +
log(prior_normal) - 0.5 * log(var_normal);
g_abnormal_bayes = -0.5 * (features_test_reduced(i) - mu_abnormal)^2 / var_abnormal +
log(prior_abnormal) - 0.5 * log(var_abnormal);

if (d_abnormal_mah < d_normal_mah)
predicted_labels_mah(i) = 1;
elseif (d_abnormal_mah == d_normal_mah)
if (randn > 0)
predicted_labels_mah(i) = 1;
end

```

```

end

if (d_abnormal_euc < d_normal_euc)
    predicted_labels_euc(i) = 1;
elseif (d_abnormal_euc == d_normal_euc)
    if (randn > 0)
        predicted_labels_euc(i) = 1;
    end
end

if (g_abnormal_bayes - g_normal_bayes > 0)
    predicted_labels_bayes(i) = 1;
elseif (g_abnormal_bayes - g_normal_bayes == 0)
    if (randn > 0)
        predicted_labels_bayes(i) = 1;
    end
end

end

acc_mah = sum(labels_test == predicted_labels_mah)/num_test;
acc_euc = sum(labels_test == predicted_labels_euc)/num_test;
acc_bayes = sum(labels_test == predicted_labels_bayes)/num_test;
disp('Part d Results:')
disp(['Accuracy of Mahalanobis Distance = ',num2str(acc_mah)]);
disp(['Accuracy of Euclidean Distance = ',num2str(acc_euc)]);
disp(['Accuracy of Bayes Classifier = ',num2str(acc_bayes)]);
disp('=====')

%% Q3 - Part a

clc; clear;

n = 0:50;

```

```

x = (0.8.^n) - 0.5 * (0.8.^(n-1)) .* (n > 0);

c_cep_matlab = cceps(x);

fft_x = fft(x);

complex_cepstrum_definition = ifft(log(fft_x));

c_cep_rec = [];
c_cep_rec(1) = log(x(1));
for i=2:length(n)
    a = 0;
    for j=1:i-1
        a = a + (j-1)/(i-1)*c_cep_rec(j)*x(i-(j-1))/x(1);
    end
    c_cep_rec(i) = x(i)/x(1) - a;
end

figure;
subplot(3,1,1)
plot(c_cep_matlab,'LineWidth',1)
title('complex cepstrum using matlab cceps','Interpreter','latex','FontSize',14)
ylabel('$\hat{x}[n]$', 'Interpreter', 'latex')
xlabel('n', 'Interpreter', 'latex')
subplot(3,1,2)
plot(complex_cepstrum_definition,'LineWidth',1)
title('complex cepstrum using DFT','Interpreter','latex','FontSize',14)
ylabel('$\hat{x}[n]$', 'Interpreter', 'latex')
xlabel('n', 'Interpreter', 'latex')
subplot(3,1,3)
plot(c_cep_rec,'LineWidth',1)
title('complex cepstrum using recursive algorithm','Interpreter','latex','FontSize',14)
ylabel('$\hat{x}[n]$', 'Interpreter', 'latex')

```

```

xlabel('n','Interpreter','latex')

%% part b

clc; clear;

data = load('ECG.mat');

fs = 200;

normal_ECG = data.ECG05;

abnormal_ECG = data.ECG34;

t_normal = linspace(0,length(normal_ECG)/fs,length(normal_ECG));

t_abnormal = linspace(0,length(abnormal_ECG)/fs,length(abnormal_ECG));

normal_ECG_cceps = cceps(normal_ECG);

abnormal_ECG_cceps = cceps(abnormal_ECG);

normal_ECG_rceps = rceps(normal_ECG);

abnormal_ECG_rceps = rceps(abnormal_ECG);

figure;

subplot(3,2,1);

plot(t_normal,normal_ECG,'LineWidth',1);

title('Normal ECG - Time Domain','Interpreter','latex','FontSize',14);

xlabel('time (s)');

ylabel('Amplitude');

subplot(3,2,2);

plot(t_abnormal,abnormal_ECG,'LineWidth',1);

title('Abnormal ECG - Time Domain','Interpreter','latex','FontSize',14);

xlabel('time (s)');

ylabel('Amplitude');

subplot(3,2,3);

plot(t_normal(5:end-5),normal_ECG_cceps(5:end-5),'LineWidth',1);

title('Normal ECG - Complex Cepstrum','Interpreter','latex','FontSize',14);

xlabel('time (s)','Interpreter','latex');

```

```

ylabel('$\hat{x}[n]$', 'Interpreter', 'latex');

subplot(3,2,4);

plot(t_abnormal(5:end-5),abnormal_ECG_cceps(5:end-5), 'LineWidth',1);
title('Abnormal ECG - Complex Cepstrum', 'Interpreter', 'latex', 'FontSize',14);
xlabel('time (s)', 'Interpreter', 'latex');
ylabel('$\hat{x}[n]$', 'Interpreter', 'latex');

subplot(3,2,5);

plot(t_normal(5:end-5),normal_ECG_rceps(5:end-5), 'LineWidth',1);
title('Normal ECG - Real Cepstrum', 'Interpreter', 'latex', 'FontSize',14);
xlabel('time (s)', 'Interpreter', 'latex');
ylabel('$C_x[n]$', 'Interpreter', 'latex');

subplot(3,2,6);

plot(t_abnormal(5:end-5),abnormal_ECG_rceps(5:end-5), 'LineWidth',1);
title('Abnormal ECG - Real Cepstrum', 'Interpreter', 'latex', 'FontSize',14);
xlabel('time (s)', 'Interpreter', 'latex');
ylabel('$C_x[n]$', 'Interpreter', 'latex');

%% Function

function [X_hat] = myKalmanFilter(a,b,sigma_1,sigma_2,sigma_v,z,N,initialization)

F_k = [a 0; a b];
H_k = [0 1];
Q_k = [sigma_1^2 sigma_1^2; sigma_1^2 sigma_1^2+sigma_2^2];
R_k = sigma_v^2;

% Initial condition

if initialization == 0
    X_hat = zeros(2,N);

```

```

P = [sigma_1^2/(1-a^2) sigma_2^2/((1-a^2)*(1-a*b));
      sigma_2^2/((1-a^2)*(1-a*b)) (sigma_1^2 +(1-a^2)*sigma_2^2)/((1-a^2)*(1-
b^2))];
else
    X_hat = 10*randn(2,N);
    P = 10*randn(2,2);
end

for k = 2:N
    % Prediction
    X_hat(:,k) = F_k * X_hat(:,k-1);
    P = F_k*P*F_k' + Q_k;

    % Update
    G = P * H_k'* inv(H_k*P*H_k'+R_k);
    X_hat(:,k) = X_hat(:,k)+G*(Z(k)-H_k*X_hat(:,k));
    P = P - G*H_k*P;
end

end

function plotResults(X, X_hat, X_hat_rand_initial)
    figure;
    subplot(2,1,1);
    plot(X(1, :), 'k', 'LineWidth', 1);
    hold on;
    plot(X_hat(1, :), 'b', 'LineWidth', 1);
    hold on;
    plot(X_hat_rand_initial(1, :), 'r', 'LineWidth', 1);
    title('$X_1$ Estimation with Kalman Filter', 'Interpreter', 'latex', 'FontSize', 14);
    xlabel('Time', 'Interpreter', 'latex');
    ylabel('Value', 'Interpreter', 'latex');

```

```

legend('True Value','Estimated','Estimated - random initialization')

subplot(2,1,2);
plot(X(2, :), 'k', 'LineWidth',1);
hold on;
plot(X_hat(2, :), 'b', 'LineWidth',1);
hold on;
plot(X_hat_rand_initial(2, :), 'r', 'LineWidth',1);
title('$X_2$ Estimation with Kalman Filter', 'Interpreter', 'latex', 'FontSize', 14);
xlabel('Time', 'Interpreter', 'latex');
ylabel('Value', 'Interpreter', 'latex');
legend('True Value','Estimated','Estimated - random initialization')

end

function [features] = featureExtractor(signal)

m = mean(signal); % Mean of the signal

smallerThanMean = signal(signal < m);
largerThanMean = signal(signal > m);

% Feature 1: Average ratio of samples smaller than m to the minimum value
minVal = min(signal);

if minVal == 0
    features(1) = mean(smallerThanMean) / minVal;
else
    features(1) = NaN; % Handle division by zero
end

% Feature 2: Ratio of mean to variance of samples smaller than m
varSmallerThanMean = var(smallerThanMean);

if varSmallerThanMean ~= 0
    features(2) = mean(smallerThanMean) / varSmallerThanMean;
end

```

```

else

    features(2) = NaN; % Handle division by zero

end


% Feature 3: d/D ratio where d is the average difference between samples
% larger than m and smaller than m, and D is the range of the signal

maxVal = max(signal);

D = maxVal - minValue;

d = mean(largerThanMean) - mean(smallerThanMean);

if D ~= 0

    features(3) = d / D;

else

    features(3) = NaN; % Handle division by zero

end

end


function [features] = featureExtractor_new(ecgSignal)

meanECG = mean(ecgSignal);

stdECG = std(ecgSignal);

madECG = mean(abs(ecgSignal - meanECG));



features = [meanECG, stdECG, madECG];

end


function [meanECG, stdECG, madECG] = extractSimpleECGFeatures(ecgSignal)

% Simple ECG Feature Extraction Function

% Inputs:

%   ecgSignal - the raw ECG signal

% Outputs:

%   meanECG - Mean value of the ECG signal
%   stdECG - Standard deviation of the ECG signal

```

```
% madECG - Mean absolute deviation of the ECG signal

% Calculate Mean
meanECG = mean(ecgSignal);

% Calculate Standard Deviation
stdECG = std(ecgSignal);

% Calculate Mean Absolute Deviation
madECG = mean(abs(ecgSignal - meanECG));

% Display Results
disp(['Mean of ECG: ', num2str(meanECG)]);
disp(['Standard Deviation of ECG: ', num2str(stdECG)]);
disp(['Mean Absolute Deviation of ECG: ', num2str(madECG)]);

end
```