Ressourcen •

fitcdiscr

Fit discriminant analysis classifier

Syntax

```
Mdl = fitcdiscr(Tbl,ResponseVarName)
Mdl = fitcdiscr(Tbl,formula)
Mdl = fitcdiscr(Tbl,Y)

Mdl = fitcdiscr(X,Y)

Mdl = fitcdiscr(__,Name=Value)
[Mdl,AggregateOptimizationResults] = fitcdiscr(__)
```

Description

Mdl = fitcdiscr(Tbl, ResponseVarName) returns a fitted discriminant analysis model based on the input variables (also known as predictors, features, or attributes) contained in the table Tbl and output (response or labels) contained in ResponseVarName.

Mdl = fitcdiscr(Tbl, formula) returns a fitted discriminant analysis model based on the input variables contained in the table Tbl. formula is an explanatory model of the response and a subset of predictor variables in Tbl used to fit Mdl.

Mdl = fitcdiscr(Tbl,Y) returns a fitted discriminant analysis model based on the input variables contained in the table Tbl and response Y.

Mdl = fitcdiscr(X,Y) returns a discriminant analysis classifier based on the input variables X and response Y.

example

Md1 = fitcdiscr(___, Name=Value) fits a classifier with additional options specified by one or more name-value arguments, using any of the previous syntaxes. For example, you can optimize hyperparameters to minimize the model's cross-validation loss, or specify the cost of misclassification, the prior probabilities for each class, or the observation weights.

[Mdl,AggregateOptimizationResults] = fitcdiscr(___) also returns AggregateOptimizationResults, which contains hyperparameter optimization results when you specify the OptimizeHyperparameters and HyperparameterOptimizationOptions name-value arguments. You must also specify the ConstraintType and ConstraintBounds options of HyperparameterOptimizationOptions. You can use this syntax to optimize on compact model size instead of cross-validation loss, and to perform a set of multiple optimization problems that have the same options but different constraint bounds.



Note

For a list of supported syntaxes when the input variables are tall arrays, see Tall Arrays.

Examples collapse all

Train Discriminant Analysis Model

Load Fisher's iris data set.



```
rng(1)
Mdl = fitcdiscr(meas, species, 'OptimizeHyperparameters', 'auto',...
    'HyperparameterOptimizationOptions',...
    struct('AcquisitionFunctionName', 'expected-improvement-plus'))
```

Iter	 Eval	 Objective	Objective	BestSoFar	BestSoFar	 Delta	
	result	l 	runtime	(observed)	(estim.)		
1	Best	 0.66667	 0.5599	 0.66667	 0.66667	13.261	
2	Best	0.02	0.20361	0.02	0.064227	2.7404e-05	
3	Accept	0.04	0.1408	0.02	0.020084	3.2455e-06	
4	Accept	0.66667	0.05779	0.02	0.020118	14.879	
5	Accept	0.046667	0.041831	0.02	0.019907	0.00031449	
6	Accept	0.04	0.037865	0.02	0.028438	4.5092e-05	
7	Accept	0.046667	0.063158	0.02	0.031424	2.0973e-05	
8	Accept	0.02	0.035232	0.02	0.022424	1.0554e-06	0
9	Accept	0.02	0.037108	0.02	0.021105	1.1232e-06	0.
10	Accept	0.02	0.042567	0.02	0.020948	0.00011837	0
11	Accept	0.02	0.03459	0.02	0.020172	1.0292e-06	
12	Accept	0.02	0.055514	0.02	0.020105	9.7792e-05	0
13	Accept	0.02	0.045267	0.02	0.020038	0.00036014	0
14	Accept	0.02	0.033937	0.02	0.019597	0.00021059	0
15	Accept	0.02	0.070328	0.02	0.019461	1.1911e-05	0
16	Accept	0.02	0.081124	0.02	0.01993	0.0017896	0.
17	Accept	0.02	0.039471	0.02	0.019551	0.00073745	0
18	Accept	0.02	0.078447	0.02	0.019776	0.00079304	0.
19	Accept	0.02	0.049007	0.02	0.019678	0.007292	0
20	Accept	0.046667	0.041023	0.02	0.019785	0.0074408	
===== Iter	======= Eval	======================================	======================================	======================================	======================================	======================================	:====:
	result		runtime	(observed)	(estim.)		
21	======= Accept	======================================	======================================	======================================	======================================	 0.0036004	 0
22	Accept	0.02	0.043938	0.02	0.019755	2.5238e-05	0
23	Accept	0.02	0.09227	0.02	0.0191	1.5478e-05	0
24	Accept	0.02	0.034113	0.02	0.019081	0.0040557	0.
25	Accept		0.032803	0.02	0.019333	2.959e-05	0
	Accept	_					
	Accept				:	:	
	Accept						0
	Accept				:	:	
	Accept						

Optimization completed.

MaxObjectiveEvaluations of 30 reached.

Total function evaluations: 30 Total elapsed time: 21.161 seconds

Total objective function evaluation time: 2.2371

Best observed feasible point:

Delta Gamma

2.7404e-05 0.073264

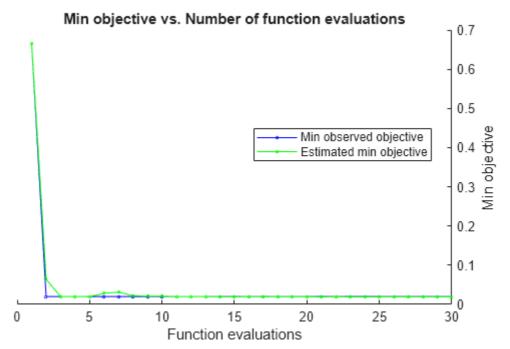
Observed objective function value = 0.02Estimated objective function value = 0.022693Function evaluation time = 0.20361

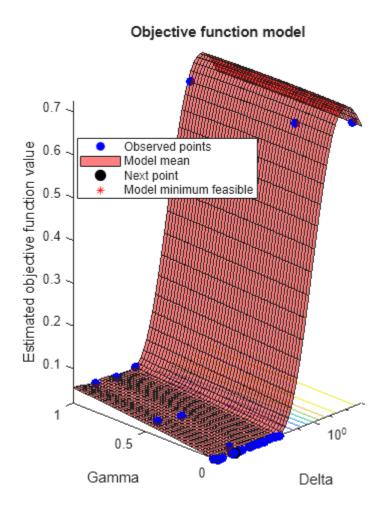
Best estimated feasible point (according to models):

Delta Gamma

2.5238e-05 0.0015542

Estimated objective function value = 0.01947 Estimated function evaluation time = 0.05596





Md1 = ClassificationDiscriminant

ResponseName: 'Y'

CategoricalPredictors: []

ClassNames: {'setosa' 'versicolor' 'virginica'}

ScoreTransform: 'none'

NumObservations: 150

HyperparameterOptimizationResults: [1x1 BayesianOptimization]

DiscrimType: 'linear'

Mu: [3×4 double]
Coeffs: [3×3 struct]

Properties, Methods

The fit achieves about 2% loss for the default 5-fold cross validation.

Optimize Discriminant Analysis Model on Tall Array

This example shows how to optimize hyperparameters of a discriminant analysis model automatically using a tall array. The sample data set airlinesmall.csv is a large data set that contains a tabular file of airline flight data. This example creates a tall table containing the data and uses it to run the optimization procedure.

When you perform calculations on tall arrays, MATLAB® uses either a parallel pool (default if you have Parallel Computing Toolbox $^{\text{\tiny{M}}}$) or the local MATLAB

This example uses: Statistics and Machine Learning Toolbox Parallel Computing Toolbox session. If you want to run the example using the local MATLAB session when you have Parallel Computing Toolbox, you can change the global execution environment by using the mapreducer function.

Open in MATLAB
Online

Copy Command

Create a datastore that references the folder location with the data. Select a subset of the variables to work with, and treat NA values as missing data so that datastore replaces them with NaN values. Create a tall table that contains the data in the datastore.

Starting parallel pool (parpool) using the 'Processes' profile ... 07-Dec-2023 09:05:49: Job Queued. Waiting for parallel pool job with ID 1 to start ... 07-Dec-2023 09:06:50: Job Queued. Waiting for parallel pool job with ID 1 to start ... Connected to parallel pool with 6 workers.

tt =

M×7 tall table

Month	DayofMonth	DayOfWeek	DepTime	ArrDelay	Distance	DepDelay
10	21	3	642	8	308	12
10	26	1	1021	8	296	1
10	23	5	2055	21	480	20
10	23	5	1332	13	296	12
10	22	4	629	4	373	-1
10	28	3	1446	59	308	63
10	8	4	928	3	447	-2
10	10	6	859	11	954	-1
:	:	:	:	:	:	:
:	:	:	:	:	:	:

Determine the flights that are late by 10 minutes or more by defining a logical variable that is true for a late flight. This variable contains the class labels. A preview of this variable includes the first few rows.

```
Y = tt.DepDelay > 10 % Class labels □ Get ▼
```

Y =

M×1 tall logical array

1 0

1

0

1

0

:

Create a tall array for the predictor data.

```
I□ Get ¬
X = tt{:,1:end-1} % Predictor data
X =
  M×6 tall double matrix
           10
                         21
                                        3
                                                   642
                                                                   8
                                                                              308
           10
                         26
                                                  1021
                                                                   8
                                                                              296
           10
                         23
                                        5
                                                  2055
                                                                  21
                                                                              480
                                        5
           10
                         23
                                                  1332
                                                                  13
                                                                              296
                                        4
           10
                         22
                                                   629
                                                                   4
                                                                              373
                         28
                                        3
           10
                                                  1446
                                                                  59
                                                                              308
                          8
                                                   928
                                                                   3
           10
                                        4
                                                                              447
           10
                         10
                                        6
                                                   859
                                                                  11
                                                                              954
                                                    :
                                                                               :
```

Remove rows in X and Y that contain missing data.

```
R = rmmissing([X Y]); % Data with missing entries removed
X = R(:,1:end-1);
Y = R(:,end);
```

Standardize the predictor variables.

```
Z = zscore(X); ☐ Get ▼
```

Optimize hyperparameters automatically using the OptimizeHyperparameters name-value argument. Note that when you use tall arrays, DiscrimType is the only hyperparameter you can optimize, regardless of whether you specify "auto" or "all". Find the optimal DiscrimType value that minimizes holdout cross-validation loss. For reproducibility, use the "expected-improvement-plus" acquisition function and set the seeds of the random number generators using rng and tallrng. The results can vary depending on the number of workers and the execution environment for the tall arrays. For details, see Control Where Your Code Runs.

```
rng("default")
                                                                           ı□ Get 🕶
tallrng("default")
[Mdl,FitInfo,HyperparameterOptimizationResults] = fitcdiscr(Z,Y, ...
    "OptimizeHyperparameters", "auto", ...
   "HyperparameterOptimizationOptions", struct("Holdout", 0.3, ...
    "AcquisitionFunctionName", "expected-improvement-plus"))
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 2: Completed in 7.3 sec
- Pass 2 of 2: Completed in 3.8 sec
Evaluation completed in 19 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 4 sec
Evaluation completed in 4.3 sec
|-----|
| Iter | Eval
              | Objective
                            | Objective
                                         BestSoFar
                                                      BestSoFar
      | result |
                            | runtime
                                         (observed)
                                                     (estim.)
                    0.11354
                                  27.449
                                              0.11354
                                                           0.11354
                                                                       quadratic |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1.3 sec
Evaluation completed in 2.5 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
```

```
Evaluation completed in 1.2 sec
    2 | Accept | 0.11354 | 5.4566 | 0.11354 | 0.11354 | pseudoQuadra |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.96 sec
Evaluation completed in 2.2 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.89 sec
Evaluation completed in 1.1 sec
    3 | Accept | 0.12869 | 4.8549 | 0.11354 | 0.11859 | pseudoLinear |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.99 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.89 sec
Evaluation completed in 1.1 sec
                                 4.2867 | 0.11354 | 0.1208 | diagLinear |
    4 | Accept | 0.12745 |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.98 sec
Evaluation completed in 2 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.84 sec
Evaluation completed in 1 sec
    5 | Accept | 0.12869 | 4.6497 | 0.11354 | 0.12238 | linear |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.7 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.89 sec
Evaluation completed in 1.1 sec
                                                           0.12082 | diagQuadrati |
    6 | Best | 0.11301 |
                                4.0594
                                             0.11301
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.96 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.82 sec
Evaluation completed in 1 sec
                                4.0419 | 0.11301 | 0.11301 | diagQuadrati |
    7 | Accept |
                  0.11301
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.85 sec
Evaluation completed in 1 sec
    8 | Accept | 0.11301 | 4.0382 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.97 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.81 sec
Evaluation completed in 1 sec
    9 | Accept | 0.11301 |
                                3.9186 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.84 sec
```

```
Evaluation completed in 1 sec
| 10 | Accept | 0.11301 | 4.0947 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.88 sec
Evaluation completed in 1.1 sec
   11 | Accept | 0.11301 | 4.3088 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.94 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.8 sec
Evaluation completed in 1 sec
                                3.9644 | 0.11301 | 0.11301 | diagQuadrati |
   12 | Accept | 0.11301 |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.86 sec
Evaluation completed in 1.1 sec
| 13 | Accept | 0.11301 | 4.0673 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.93 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.94 sec
Evaluation completed in 1.2 sec
                                                           0.11301 | diagQuadrati |
  14 | Accept | 0.11301 | 4.1285 | 0.11301 |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1.2 sec
Evaluation completed in 2 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.93 sec
Evaluation completed in 1.1 sec
                                4.4217 | 0.11301 | 0.11301 | diagQuadrati |
   15 | Accept | 0.11301 |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.98 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.89 sec
Evaluation completed in 1.1 sec
| 16 | Accept | 0.11301 |
                                4.0631 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.99 sec
Evaluation completed in 1.7 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.86 sec
Evaluation completed in 1.1 sec
| 17 | Accept | 0.11301 |
                                 4.0227 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.88 sec
```

```
Evaluation completed in 1.1 sec
| 18 | Accept | 0.11354 | 4.3391 | 0.11301 | 0.11301 | pseudoQuadra |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.98 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.85 sec
Evaluation completed in 1 sec
   19 | Accept | 0.11301 | 4.139 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.97 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.91 sec
Evaluation completed in 1.1 sec
                             4.2078 | 0.11301 | 0.11301 | diagQuadrati |
   20 | Accept | 0.11301 |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.86 sec
Evaluation completed in 1.1 sec
|-----|
| Iter | Eval | Objective | Objective | BestSoFar | BestSoFar | DiscrimType |
result runtime (observed) (estim.)
|-----|
| 21 | Accept | 0.11301 | 4.1129 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1.1 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.86 sec
Evaluation completed in 1.1 sec
   22 | Accept | 0.11354 |
                              4.2473 | 0.11301 | 0.11301 | quadratic |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.98 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.87 sec
Evaluation completed in 1.1 sec
   23 | Accept | 0.11301 | 4.0342 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1.1 sec
Evaluation completed in 1.9 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.88 sec
Evaluation completed in 1.1 sec
   24 | Accept | 0.11354 |
                             4.173 | 0.11301 | 0.11301 | pseudoQuadra |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.99 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.82 sec
Evaluation completed in 1.1 sec
                             3.9707 | 0.11301 | 0.11301 | diagQuadrati |
   25 | Accept | 0.11301 |
Evaluating tall expression using the Parallel Pool 'Processes':
```

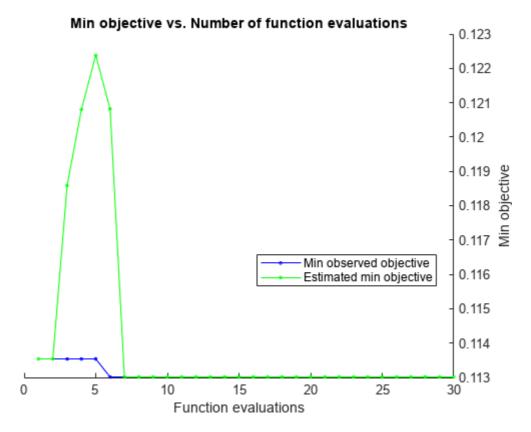
```
- Pass 1 of 1: Completed in 1 sec
Evaluation completed in 1.7 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.98 sec
Evaluation completed in 1.2 sec
   26 | Accept | 0.11354 |
                                 4.1135 | 0.11301 | 0.11301 | quadratic |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 1.1 sec
Evaluation completed in 2 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.92 sec
Evaluation completed in 1.1 sec
   27 | Accept |
                   0.11301
                                  4.2567
                                               0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.94 sec
Evaluation completed in 1.5 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.88 sec
Evaluation completed in 1.1 sec
                     0.11301 | 3.7988 | 0.11301 | 0.11301 | diagQuadrati |
   28 | Accept |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.97 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.88 sec
Evaluation completed in 1.1 sec
   29 | Accept |
                   0.11301
                                   3.9926 | 0.11301 | 0.11301 | diagQuadrati |
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.95 sec
Evaluation completed in 1.8 sec
Evaluating tall expression using the Parallel Pool 'Processes':
- Pass 1 of 1: Completed in 0.85 sec
Evaluation completed in 1 sec
  30 | Accept | 0.11301 | 3.9793 | 0.11301 | 0.11301 | diagQuadrati |
Optimization completed.
MaxObjectiveEvaluations of 30 reached.
Total function evaluations: 30
Total elapsed time: 180.1662 seconds
Total objective function evaluation time: 149.1907
Best observed feasible point:
    DiscrimType
   diagQuadratic
Observed objective function value = 0.11301
Estimated objective function value = 0.11301
Function evaluation time = 4.0594
Best estimated feasible point (according to models):
    DiscrimType
```

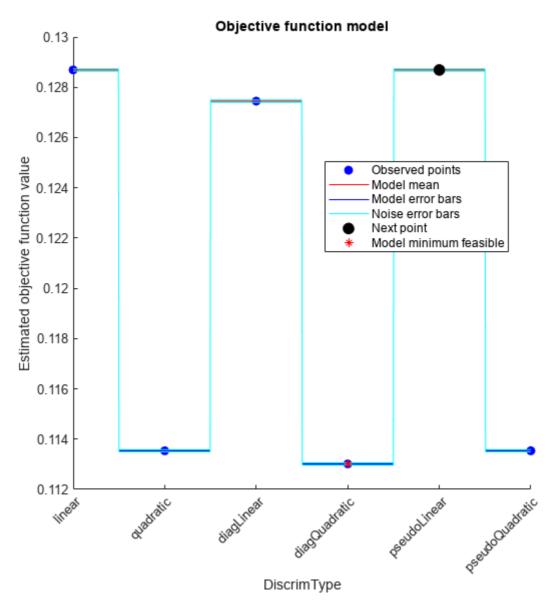
diagQuadratic

Estimated objective function value = 0.11301 Estimated function evaluation time = 4.1702

Evaluating tall expression using the Parallel Pool 'Processes':

- Pass 1 of 1: Completed in 0.8 sec Evaluation completed in 1.6 sec





Mdl =

 ${\tt CompactClassificationDiscriminant}$

PredictorNames: {'x1' 'x2' 'x3' 'x4' 'x5' 'x6'}

ResponseName: 'Y'

CategoricalPredictors: []

ClassNames: [0 1]

ScoreTransform: 'none'

DiscrimType: 'diagQuadratic'

Mu: [2×6 double]

Coeffs: [2×2 struct]

Properties, Methods

FitInfo = struct with no fields.

HyperparameterOptimizationResults =

BayesianOptimization with properties:

ObjectiveFcn: @createObjFcn/tallObjFcn

VariableDescriptions: [1x1 optimizableVariable]

Options: [1×1 struct]

```
MinObjective: 0.1130
               XAtMinObjective: [1x1 table]
         MinEstimatedObjective: 0.1130
      XAtMinEstimatedObjective: [1x1 table]
       NumObjectiveEvaluations: 30
              TotalElapsedTime: 180.1662
                     NextPoint: [1x1 table]
                        XTrace: [30×1 table]
                ObjectiveTrace: [30×1 double]
              ConstraintsTrace: []
                 UserDataTrace: {30×1 cell}
 ObjectiveEvaluationTimeTrace: [30×1 double]
            IterationTimeTrace: [30x1 double]
                    ErrorTrace: [30×1 double]
              FeasibilityTrace: [30×1 logical]
  FeasibilityProbabilityTrace: [30×1 double]
           IndexOfMinimumTrace: [30×1 double]
         ObjectiveMinimumTrace: [30×1 double]
EstimatedObjectiveMinimumTrace: [30x1 double]
```

Input Arguments collapse all

```
Tb1 — Sample data table
```

Sample data used to train the model, specified as a table. Each row of Tb1 corresponds to one observation, and each column corresponds to one predictor variable. Categorical predictor variables are not supported. Optionally, Tb1 can contain one additional column for the response variable, which can be categorical. Multicolumn variables and cell arrays other than cell arrays of character vectors are not allowed.

- If Tbl contains the response variable, and you want to use all remaining variables in Tbl as predictors, then specify the response variable by using ResponseVarName.
- If Tbl contains the response variable, and you want to use only a subset of the remaining variables in Tbl as predictors, then specify a formula by using formula.
- If Tb1 does not contain the response variable, then specify a response variable by using Y. The length of the response variable and the number of rows in Tb1 must be equal.

```
ResponseVarName — Response variable name name of variable in Tb1
```

Response variable name, specified as the name of a variable in Tb1.

You must specify ResponseVarName as a character vector or string scalar. For example, if the response variable Y is stored as Tbl.Y, then specify it as "Y". Otherwise, the software treats all columns of Tbl, including Y, as predictors when training the model.

The response variable must be a categorical, character, or string array; a logical or numeric vector; or a cell array of character vectors. If Y is a character array, then each element of the response variable must correspond to one row of the array.

A good practice is to specify the order of the classes by using the ClassNames name-value argument.

Data Types: char | string

formula — Explanatory model of response variable and subset of predictor variables character vector | string scalar

Explanatory model of the response variable and a subset of the predictor variables, specified as a character vector or string scalar in the form "Y~x1+x2+x3". In this form, Y represents the response variable, and x1, x2, and x3 represent the predictor variables.

To specify a subset of variables in Tbl as predictors for training the model, use a formula. If you specify a formula, then the software does not use any variables in Tbl that do not appear in formula.

The variable names in the formula must be both variable names in Tb1 (Tb1.Properties.VariableNames) and valid MATLAB[®] identifiers. You can verify the variable names in Tb1 by using the isvarname function. If the variable names are not valid, then you can convert them by using the matlab.lang.makeValidName function.

Data Types: char | string

Y - Class labels

categorical array | character array | string array | logical vector | numeric vector | cell array of character vectors

Class labels, specified as a categorical, character, or string array, a logical or numeric vector, or a cell array of character vectors. Each row of Y represents the classification of the corresponding row of X.

The software considers NaN, '' (empty character vector), "" (empty string), <missing>, and <undefined> values in Y to be missing values. Consequently, the software does not train using observations with a missing response.

Data Types: categorical | char | string | logical | single | double | cell

X - Predictor data

numeric matrix

Predictor values, specified as a numeric matrix. Each column of X represents one variable, and each row represents one observation. Categorical predictor variables are not supported.

fitcdiscr considers NaN values in X as missing values. fitcdiscr does not use observations with missing values for X in the fit.

Data Types: single | double

Name-Value Arguments

expand all

Specify optional pairs of arguments as Name1=Value1,..., NameN=ValueN, where Name is the argument name and Value is the corresponding value. Name-value arguments must appear after other arguments, but the order of the pairs does not matter.

Before R2021a, use commas to separate each name and value, and enclose Name in quotes.

Example: 'DiscrimType', 'quadratic', 'SaveMemory', 'on' specifies a quadratic discriminant classifier and does not store the covariance matrix in the output object.



Note

You cannot use any cross-validation name-value argument together with the OptimizeHyperparameters name-value argument. You can modify the cross-validation for OptimizeHyperparameters only by using the HyperparameterOptimizationOptions name-value argument.

Model Parameters collapse all

ClassNames — Names of classes to use for training

categorical array | character array | string array | logical vector | numeric vector | cell array of character vectors

Names of classes to use for training, specified as a categorical, character, or string array; a logical or numeric vector; or a cell array of character vectors. ClassNames must have the same data type as the response variable in Tbl or Y.

If ClassNames is a character array, then each element must correspond to one row of the array.

Use ClassNames to:

- · Specify the order of the classes during training.
- Specify the order of any input or output argument dimension that corresponds to the class order. For example, use ClassNames to specify the order of the dimensions of Cost or the column order of classification scores returned by predict.
- Select a subset of classes for training. For example, suppose that the set of all distinct class names in Y is ["a", "b", "c"]. To train the model using observations from classes "a" and "c" only, specify ClassNames=["a", "c"].

The default value for ClassNames is the set of all distinct class names in the response variable in Tb1 or Y.

Example: ClassNames=["b", "g"]

Data Types: categorical | char | string | logical | single | double | cell

Cost - Cost of misclassification

square matrix | structure

Cost of misclassification of a point, specified as one of the following:

- Square matrix, where Cost(i,j) is the cost of classifying a point into class j if its true class is i (that is, the rows correspond to the true class and the columns correspond to the predicted class). To specify the class order for the corresponding rows and columns of Cost, additionally specify the ClassNames name-value pair argument.
- Structure S having two fields: S.ClassNames containing the group names as a variable of the same type as Y, and S.ClassificationCosts containing the cost matrix.

The default is Cost(i,j)=1 if $i\sim j$, and Cost(i,j)=0 if i=j.

Data Types: single | double | struct

Delta - Linear coefficient threshold

0 (default) | nonnegative scalar value

Linear coefficient threshold, specified as the comma-separated pair consisting of 'Delta' and a nonnegative scalar value. If a coefficient of Mdl has magnitude smaller than Delta, Mdl sets this coefficient to 0, and you can

eliminate the corresponding predictor from the model. Set Delta to a higher value to eliminate more predictors.

Delta must be 0 for quadratic discriminant models.

Data Types: single | double

DiscrimType — Discriminant type

'linear' (default) | 'quadratic' | 'diaglinear' | 'diagquadratic' | 'pseudolinear' |

Discriminant type, specified as the comma-separated pair consisting of 'DiscrimType' and a character vector or string scalar in this table.

Value	Description	Predictor Covariance Treatment
'linear'	Regularized linear discriminant analysis (LDA)	• All classes have the same covariance matrix. $ \hat{\Sigma}_{\gamma} = (1-\gamma)\hat{\Sigma} + \gamma \mathrm{diag}(\hat{\Sigma}). $ $ \hat{\Sigma} \text{ is the empirical, pooled covariance matrix and } \gamma \text{ is the amount of regularization.} $
'diaglinear'	LDA	All classes have the same, diagonal covariance matrix.
'pseudolinear'	LDA	All classes have the same covariance matrix. The software inverts the covariance matrix using the pseudo inverse.
'quadratic'	Quadratic discriminant analysis (QDA)	The covariance matrices can vary among classes.
'diagquadratic'	QDA	The covariance matrices are diagonal and can vary among classes.
'pseudoquadratic'	QDA	The covariance matrices can vary among classes. The software inverts the covariance matrix using the pseudo inverse.



Note

To use regularization, you must specify 'linear'. To specify the amount of regularization, use the Gamma name-value pair argument.

Example: 'DiscrimType', 'quadratic'

FillCoeffs — Coeffs property flag

'on'|'off'

^{&#}x27;pseudoquadratic'

Coeffs property flag, specified as the comma-separated pair consisting of 'FillCoeffs' and 'on' or 'off'. Setting the flag to 'on' populates the Coeffs property in the classifier object. This can be computationally intensive, especially when cross-validating. The default is 'on', unless you specify a cross-validation name-value pair, in which case the flag is set to 'off' by default.

Example: 'FillCoeffs','off'

${\bf Gamma-Amount\ of\ regularization}$

scalar value in the interval [0,1]

Amount of regularization to apply when estimating the covariance matrix of the predictors, specified as the comma-separated pair consisting of 'Gamma' and a scalar value in the interval [0,1]. Gamma provides finer control over the covariance matrix structure than <code>DiscrimType</code>.

- If you specify 0, then the software does not use regularization to adjust the covariance matrix. That is, the software estimates and uses the unrestricted, empirical covariance matrix.
 - For linear discriminant analysis, if the empirical covariance matrix is singular, then the software automatically applies the minimal regularization required to invert the covariance matrix. You can display the chosen regularization amount by entering Mdl.Gamma at the command line.
 - For quadratic discriminant analysis, if at least one class has an empirical covariance matrix that is singular, then the software throws an error.
- If you specify a value in the interval (0,1), then you must implement linear discriminant analysis, otherwise the software throws an error. Consequently, the software sets DiscrimType to 'linear'.
- If you specify 1, then the software uses maximum regularization for covariance matrix estimation. That is, the software restricts the covariance matrix to be diagonal. Alternatively, you can set DiscrimType to 'diagLinear' or 'diagQuadratic' for diagonal covariance matrices.

Example: 'Gamma',1

Data Types: single | double

PredictorNames - Predictor variable names

string array of unique names | cell array of unique character vectors

Predictor variable names, specified as a string array of unique names or cell array of unique character vectors. The functionality of PredictorNames depends on the way you supply the training data.

- If you supply X and Y, then you can use PredictorNames to assign names to the predictor variables in X.
 - The order of the names in PredictorNames must correspond to the column order of X. That is, $PredictorNames\{1\} is the name of X(:,1), PredictorNames\{2\} is the name of X(:,2), and so on. \\ Also, size(X,2) and numel(PredictorNames) must be equal.$
 - By default, PredictorNames is {'x1', 'x2',...}.
- If you supply Tb1, then you can use PredictorNames to choose which predictor variables to use in training. That is, fitcdiscr uses only the predictor variables in PredictorNames and the response variable during training.
 - PredictorNames must be a subset of Tbl.Properties.VariableNames and cannot include the name of the response variable.
 - By default, PredictorNames contains the names of all predictor variables.
 - A good practice is to specify the predictors for training using either PredictorNames or formula, but not both.

Example: "PredictorNames",["SepalLength","SepalWidth","PetalLength","PetalWidth"]
Data Types: string | cell

Prior - Prior probabilities

"empirical" (default) | "uniform" | vector of scalar values | structure

Prior probabilities for each class, specified as a value in this table.

Value	Description
"empirical"	The class prior probabilities are the class relative frequencies in Y.
"uniform"	All class prior probabilities are equal to $1/K$, where K is the number of classes.
numeric vector	Each element is a class prior probability. Order the elements according to Mdl.ClassNames or specify the order using the ClassNames namevalue pair argument. The software normalizes the elements such that they sum to 1.
structure	A structure S with two fields: • S.ClassNames contains the class names as a variable of the same type as Y. • S.ClassProbs contains a vector of corresponding prior probabilities. The software normalizes the elements such that they sum to 1.

If you set values for both Weights and Prior, the weights are renormalized to add up to the value of the prior probability in the respective class.

Example: Prior="uniform"

Data Types: char | string | single | double | struct

ResponseName — Response variable name

"Y" (default) | character vector | string scalar

Response variable name, specified as a character vector or string scalar.

- If you supply Y, then you can use ResponseName to specify a name for the response variable.
- If you supply ResponseVarName or formula, then you cannot use ResponseName.

Example: ResponseName="response"

Data Types: char | string

SaveMemory — Flag to save covariance matrix

'off' (default) | 'on'

Flag to save covariance matrix, specified as the comma-separated pair consisting of 'SaveMemory' and either 'on' or 'off'. If you specify 'on', then fitcdiscr does not store the full covariance matrix, but instead stores enough information to compute the matrix. The predict method computes the full covariance matrix for prediction, and does not store the matrix. If you specify 'off', then fitcdiscr computes and stores the full covariance matrix in Mdl.

Specify SaveMemory as 'on' when the input matrix contains thousands of predictors.

Example: 'SaveMemory', 'on'

ScoreTransform — Score transformation

"none" (default) | "doublelogit" | "invlogit" | "ismax" | "logit" | function handle | ...

Score transformation, specified as a character vector, string scalar, or function handle.

This table summarizes the available character vectors and string scalars.

Value	Description
"doublelogit"	$1/(1 + e^{-2x})$
"invlogit"	$\log(x / (1 - x))$
"ismax"	Sets the score for the class with the largest score to 1, and sets the scores for all other classes to 0
"logit"	$1/(1 + e^{-x})$
"none" or "identity"	x (no transformation)
"sign"	-1 for x < 0 0 for $x = 0$ 1 for $x > 0$
"symmetric"	2 <i>x</i> – 1
"symmetricismax"	Sets the score for the class with the largest score to 1, and sets the scores for all other classes to −1
"symmetriclogit"	$2/(1+e^{-x})-1$

For a MATLAB function or a function you define, use its function handle for the score transform. The function handle must accept a matrix (the original scores) and return a matrix of the same size (the transformed scores).

Example: ScoreTransform="logit"

Data Types: char | string | function_handle

Weights — Observation weights

numeric vector of positive values | name of variable in Tbl

Observation weights, specified as a numeric vector of positive values or name of a variable in Tbl. The software weighs the observations in each row of X or Tbl with the corresponding value in Weights. The size of Weights must equal the number of rows of X or Tbl.

If you specify the input data as a table Tb1, then Weights can be the name of a variable in Tb1 that contains a numeric vector. In this case, you must specify Weights as a character vector or string scalar. For example, if the weights vector W is stored as Tb1.W, then specify it as "W". Otherwise, the software treats all columns of Tb1, including W, as predictors or the response when training the model.

By default, Weights is ones (n, 1), where n is the number of observations in X or Tb1.

The software normalizes Weights to sum up to the value of the prior probability in the respective class. Inf weights are not supported.

Data Types: double | single | char | string

Cross-Validation Options

collapse all

CrossVal - Cross-validation flag 'off' (default) | 'on'

 ${\it Cross-validation flag, specified as the comma-separated pair consisting of "Crossval" and "on" or "off".}$

If you specify 'on', then the software implements 10-fold cross-validation.

To override this cross-validation setting, use one of these name-value pair arguments: CVPartition, Holdout, KFold, or Leaveout. To create a cross-validated model, you can use one cross-validation name-value pair argument at a time only.

Alternatively, cross-validate later by passing Mdl to crossval.

Example: 'CrossVal', 'on'

CVPartition — Cross-validation partition

[] (default) | cvpartition object

Cross-validation partition, specified as a cvpartition object that specifies the type of cross-validation and the indexing for the training and validation sets.

To create a cross-validated model, you can specify only one of these four name-value arguments: CVPartition, Holdout, KFold, or Leaveout.

Example: Suppose you create a random partition for 5-fold cross-validation on 500 observations by using cvp = cvpartition(500, KFold=5). Then, you can specify the cross-validation partition by setting CVPartition=cvp.

Holdout - Fraction of data for holdout validation

scalar value in the range (0,1)

Fraction of the data used for holdout validation, specified as a scalar value in the range (0,1). If you specify Holdout=p, then the software completes these steps:

- 1. Randomly select and reserve p*100% of the data as validation data, and train the model using the rest of the data.
- 2. Store the compact trained model in the Trained property of the cross-validated model.

To create a cross-validated model, you can specify only one of these four name-value arguments: CVPartition, Holdout, KFold, or Leaveout.

Example: Holdout=0.1

Data Types: double | single

KFold — **Number of folds**

10 (default) | positive integer value greater than 1

Number of folds to use in the cross-validated model, specified as a positive integer value greater than 1. If you specify KFold=k, then the software completes these steps:

- 1. Randomly partition the data into k sets.
- 2. For each set, reserve the set as validation data, and train the model using the other k-1 sets.
- 3. Store the k compact trained models in a k-by-1 cell vector in the Trained property of the cross-validated model.

To create a cross-validated model, you can specify only one of these four name-value arguments: CVPartition, Holdout, KFold, or Leaveout.

Example: KFold=5

Data Types: single | double

Leaveout — Leave-one-out cross-validation flag

"off" (default) | "on"

Leave-one-out cross-validation flag, specified as "on" or "off". If you specify Leaveout="on", then for each of the n observations (where n is the number of observations, excluding missing observations, specified in the NumObservations property of the model), the software completes these steps:

- 1. Reserve the one observation as validation data, and train the model using the other n-1 observations.
- 2. Store the n compact trained models in an n-by-1 cell vector in the Trained property of the cross-validated model.

To create a cross-validated model, you can specify only one of these four name-value arguments: CVPartition, Holdout, KFold, or Leaveout.

Example: Leaveout="on"

Data Types: char | string

Hyperparameter Optimization Options

expand all

OptimizeHyperparameters — Parameters to optimize

'none' (default) | 'auto' | 'all' | string array or cell array of eligible parameter names | vector of optimizable Variable objects

Parameters to optimize, specified as the comma-separated pair consisting of 'OptimizeHyperparameters' and one of the following:

- 'none' Do not optimize.
- 'auto' Use {'Delta', 'Gamma'}.
- 'all' Optimize all eligible parameters.
- String array or cell array of eligible parameter names.
- Vector of optimizableVariable objects, typically the output of hyperparameters.

The optimization attempts to minimize the cross-validation loss (error) for fitcdiscr by varying the parameters. To control the cross-validation type and other aspects of the optimization, use the

HyperparameterOptimizationOptions name-value argument. When you use

HyperparameterOptimizationOptions, you can use the (compact) model size instead of the cross-validation loss as the optimization objective by setting the ConstraintType and ConstraintBounds options.



Note

The values of OptimizeHyperparameters override any values you specify using other name-value arguments. For example, setting OptimizeHyperparameters to "auto" causes fitcdiscr to optimize hyperparameters corresponding to the "auto" option and to ignore any specified values for the hyperparameters.

The eligible parameters for fitcdiscr are:

- Delta fitcdiscr searches among positive values, by default log-scaled in the range [1e-6,1e3].
- DiscrimType fitcdiscr searches among 'linear', 'quadratic', 'diagLinear', 'diagQuadratic', 'pseudoLinear', and 'pseudoQuadratic'.
- Gamma fitcdiscr searches among real values in the range [0,1].

Set nondefault parameters by passing a vector of optimizable Variable objects that have nondefault values. For example,

```
load fisheriris
params = hyperparameters('fitcdiscr', meas, species);
params(1).Range = [1e-4,1e6];
```

Pass params as the value of OptimizeHyperparameters.

By default, the iterative display appears at the command line, and plots appear according to the number of hyperparameters in the optimization. For the optimization and plots, the objective function is the misclassification rate. To control the iterative display, set the Verbose option of the HyperparameterOptimizationOptions name-value argument. To control the plots, set the ShowPlots field of the HyperparameterOptimizationOptions name-value argument.

For an example, see Optimize Discriminant Analysis Model.

Example: 'auto'

HyperparameterOptimizationOptions — Options for optimization

HyperparameterOptimizationOptions object | structure

Output Arguments

collapse all

Md1 - Trained discriminant analysis classification model

 ${\tt ClassificationDiscriminant\ model\ object\ |\ ClassificationPartitioned Model\ cross-validated\ model\ object\ |\ Classificationed\ cross-validated\ model\ object\ |\ Classificationed\ cross-validated\ model\ object\ |\ Classificationed\ cross-validated\ cross-validated\ nodel\ object\ |\ Classificationed\ cross-validated\ cross-validated\ nodel\ object\ |\ Classificationed\ cross-validated\ cross-validate$

Trained discriminant analysis classification model, returned as a ClassificationDiscriminant model object or a ClassificationPartitionedModel cross-validated model object.

If you set any of the name-value pair arguments KFold, Holdout, CrossVal, or CVPartition, then Mdl is a ClassificationPartitionedModel cross-validated model object. Otherwise, Mdl is a ClassificationDiscriminant model object.

To reference properties of Md1, use dot notation. For example, to display the estimated component means at the Command Window, enter Md1.Mu.

If you specify OptimizeHyperparameters and set the ConstraintType and ConstraintBounds options of HyperparameterOptimizationOptions, then Mdl is an N-by-1 cell array of model objects, where N is equal to the number of rows in ConstraintBounds. If none of the optimization problems yields a feasible model, then each cell array value is [].

AggregateOptimizationResults - Aggregate optimization results

AggregateBayesianOptimization object

Aggregate optimization results for multiple optimization problems, returned as an AggregateBayesianOptimization object. To return AggregateOptimizationResults, you must specify OptimizeHyperparameters and HyperparameterOptimizationOptions. You must also specify the ConstraintType and ConstraintBounds options of HyperparameterOptimizationOptions. For an example that shows how to produce this output, see Hyperparameter Optimization with Multiple Constraint Bounds.

More About collapse all

Discriminant Classification

The model for discriminant analysis is:

- Each class (Y) generates data (X) using a multivariate normal distribution. That is, the model assumes X has a Gaussian mixture distribution (gmdistribution).
 - For linear discriminant analysis, the model has the same covariance matrix for each class, only the means vary.
 - For quadratic discriminant analysis, both means and covariances of each class vary.

predict classifies so as to minimize the expected classification cost:

$$\widehat{y} = \underset{y=1,\dots,K}{\operatorname{argmin}} \sum_{k=1}^{K} \widehat{P}(k|x)C(y|k),$$

where

- \hat{y} is the predicted classification.
- K is the number of classes.
- $\widehat{P}(k|x)$ is the posterior probability of class k for observation x.
- C(y|k) is the cost of classifying an observation as y when its true class is k.

For details, see Prediction Using Discriminant Analysis Models.

Tips

After training a model, you can generate C/C++ code that predicts labels for new data. Generating C/C++ code requires MATLAB Coder™. For details, see Introduction to Code Generation.

Algorithms

• If you specify the Cost, Prior, and Weights name-value arguments, the output model object stores the specified values in the Cost, Prior, and W properties, respectively. The Cost property stores the user-specified cost matrix

as is. The Prior and W properties store the prior probabilities and observation weights, respectively, after normalization. For details, see Misclassification Cost Matrix, Prior Probabilities, and Observation Weights.

• The software uses the Cost property for prediction, but not training. Therefore, Cost is not read-only; you can change the property value by using dot notation after creating the trained model.

Alternative Functionality

Functions

The classify function also performs discriminant analysis. classify is usually more awkward to use.

- classify requires you to fit the classifier every time you make a new prediction.
- classify does not perform cross-validation or hyperparameter optimization.
- classify requires you to fit the classifier when changing prior probabilities.

Extended Capabilities

expand all

Tall Arrays

Calculate with arrays that have more rows than fit in memory.

Automatic Parallel Support

Accelerate code by automatically running computation in parallel using Parallel Computing Toolbox™.

Version History

Introduced in R2014a expand all

R2025a: Compute serially when parallel hyperparameter optimization is not available ▲

See Also

ClassificationDiscriminant | ClassificationPartitionedModel | predict | crossval | classify

Topics

Discriminant Analysis Classification
Improving Discriminant Analysis Models

Regularize Discriminant Analysis Classifier