Chapter 5

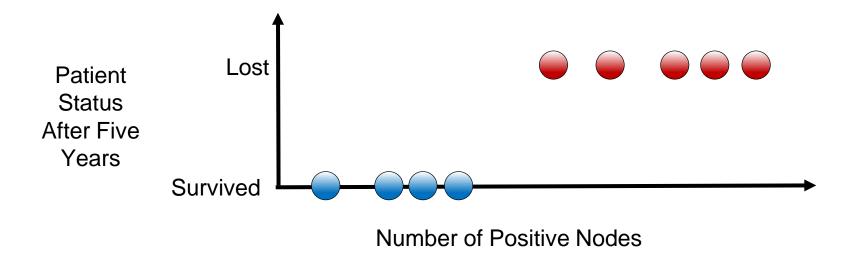
Logistic Regression Classification Error Metrics



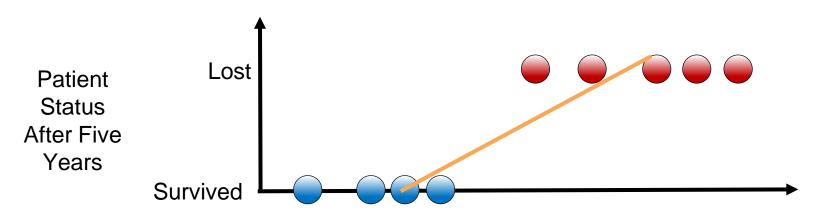
Logistic Regression



Introduction to Logistic Regression

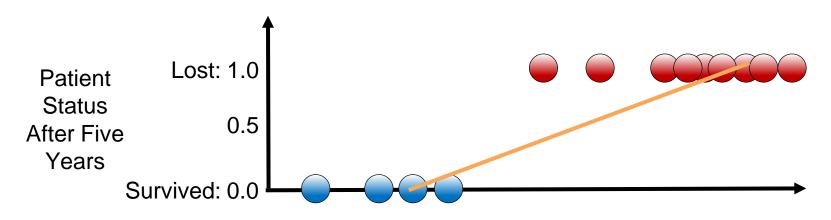






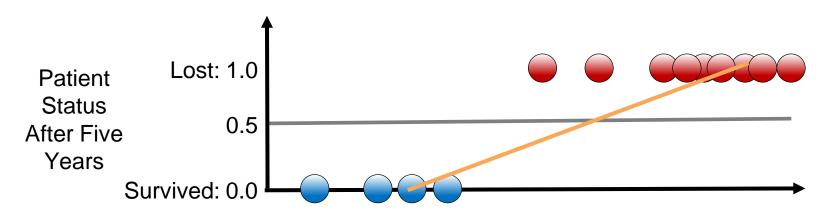
$$y_{\beta}(x) = \beta_0 + \beta_1 x + \varepsilon$$





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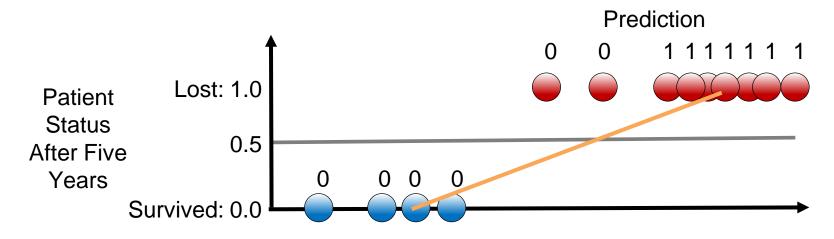




Number of Positive Nodes

If model result > 0.5: predict lost
If model result < 0.5: predict survived



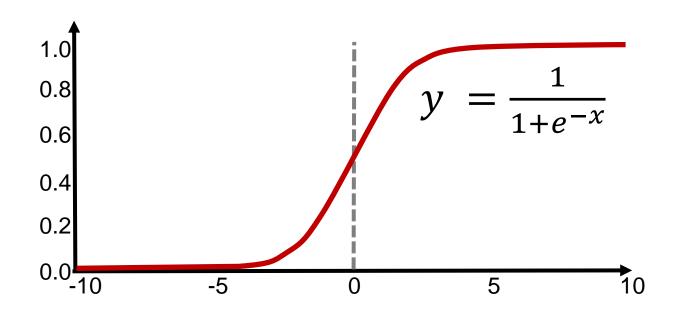


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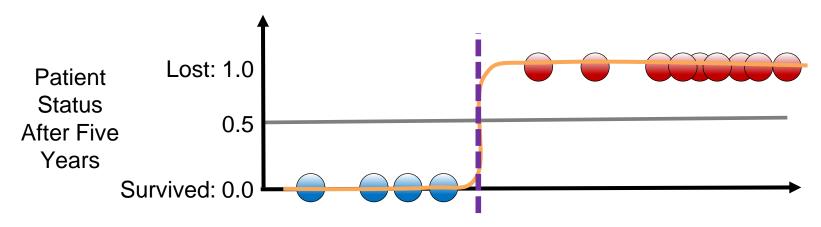


What is this Function?





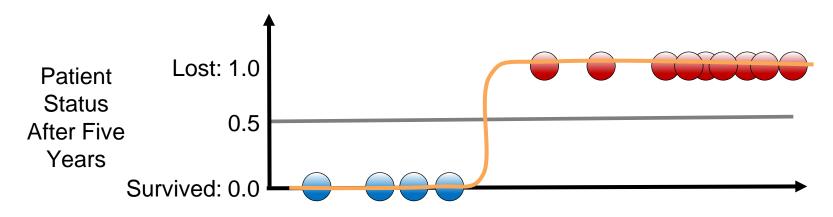
The Decision Boundary



$$y_{\beta}(x) = \frac{1}{1 + e^{-(\beta_0 + \beta_1 x + \varepsilon)}}$$



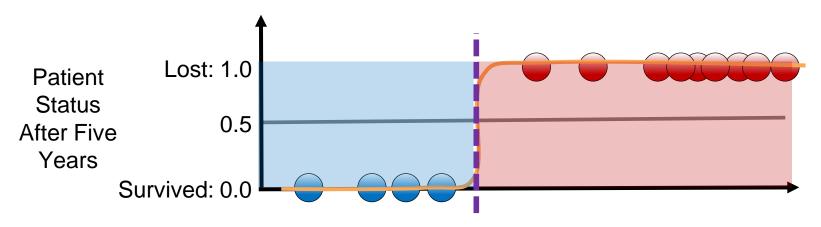
Logistic Regression



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The Decision Boundary



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$$P(x) = \frac{e^{(\beta_0 + \beta_1 x)}}{1 + e^{(\beta_0 + \beta_1 x)}}$$



$$\frac{\text{Log}}{\text{Odds}} \quad log \left| \frac{P(x)}{1 - P(x)} \right| = \beta_0 + \beta_1 x$$



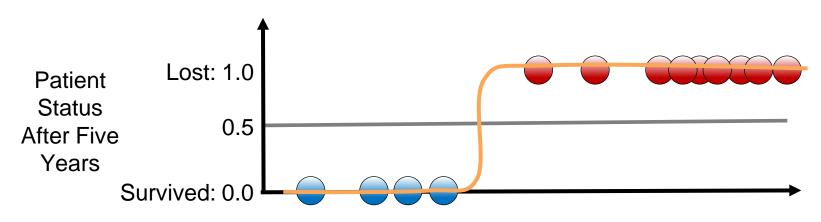
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$$\frac{\text{Log}}{\text{Odds}} \quad log \left[\frac{P(x)}{1 - P(x)} \right] = \left[\beta_0 + \beta_1 x \right]$$



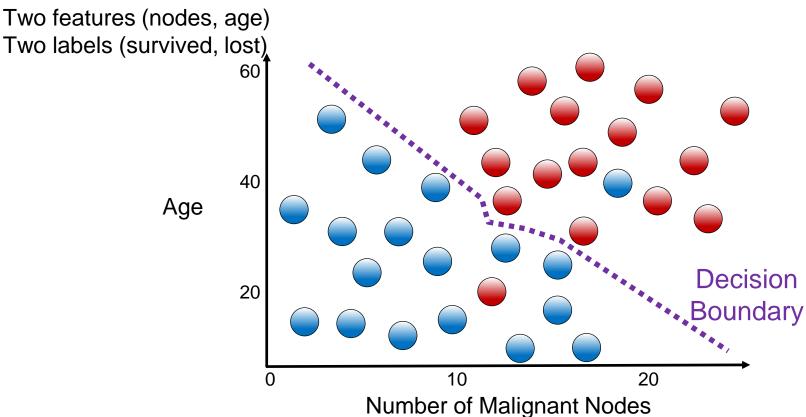
One feature (nodes)
Two labels (survived, lost)



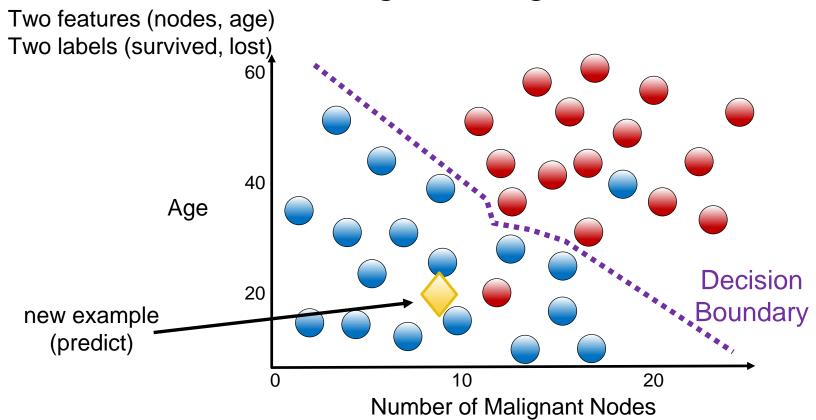


Two features (nodes, age) Two labels (survived, lost) 60 40 Age 20 0 10 20 Number of Malignant Nodes







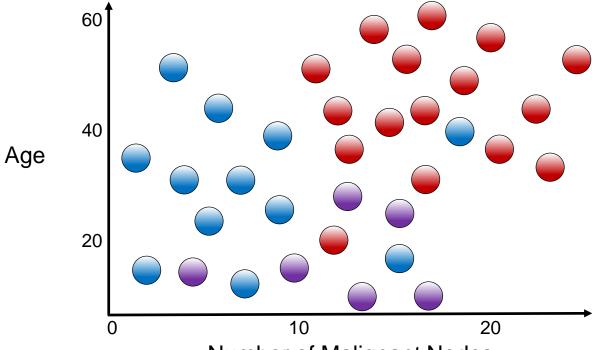




Multiclass Classification with Logistic Regression Two features (nodes, age)

Three labels (survived, complications,

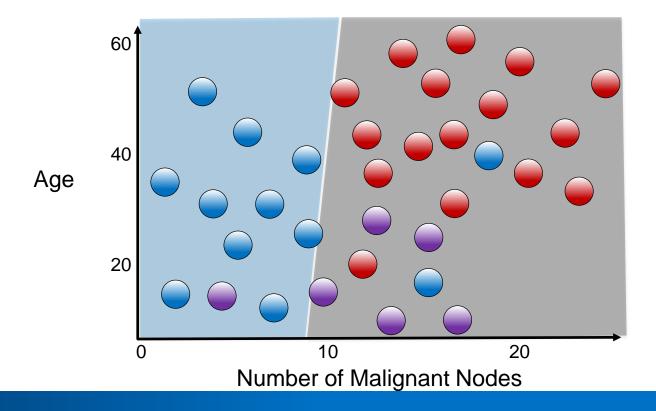
lost)



Number of Malignant Nodes

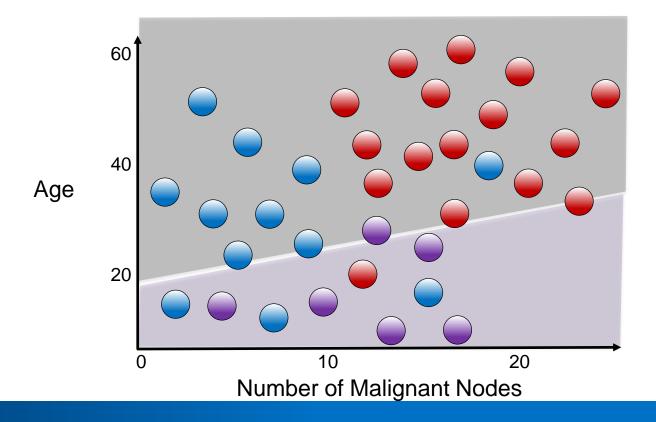


One vs All: Survived vs All



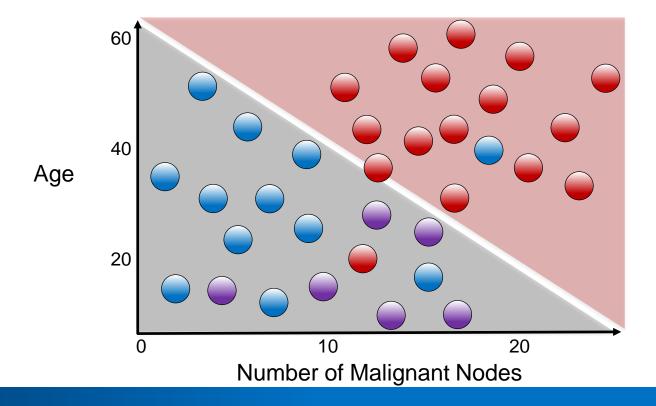


One vs All: Complications vs All





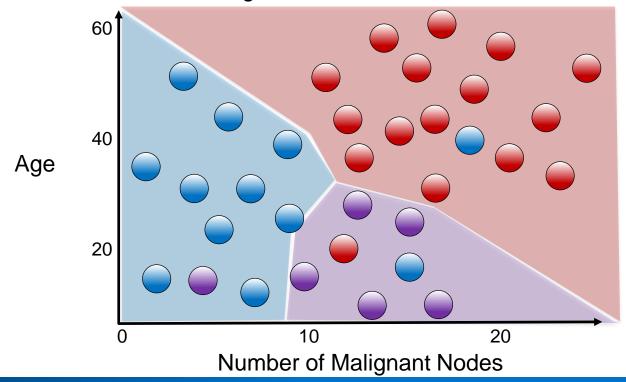
One vs All: Loss vs All





Multiclass Decision Boundary

Assign most probable class to each region





Import the class containing the classification method

from sklearn.linear_model import LogisticRegression



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Create an instance of the class

LR = LogisticRegression(penalty='l2', c=10.0)

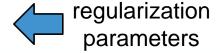


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LR = LR.fit(X_train, y_train)
y_predict = LR.predict(X_test)
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LR = LR.fit(X_train, y_train)
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```

Tune regularization parameters with cross-validation: LogisticRegressionCV.



Classification Error Metrics



Choosing the Right Error Measurement

- You are asked to build a classifier for leukemia
- Training data: 1% patients with leukemia, 99% healthy
- Measure accuracy: total % of predictions that are correct



Choosing the Right Error Measurement

- You are asked to build a classifier for leukemia
- Training data: 1% patients with leukemia, 99% healthy
- Measure accuracy: total % of predictions that are correct
- Build a simple model that always predicts "healthy"
- Accuracy will be 99%...

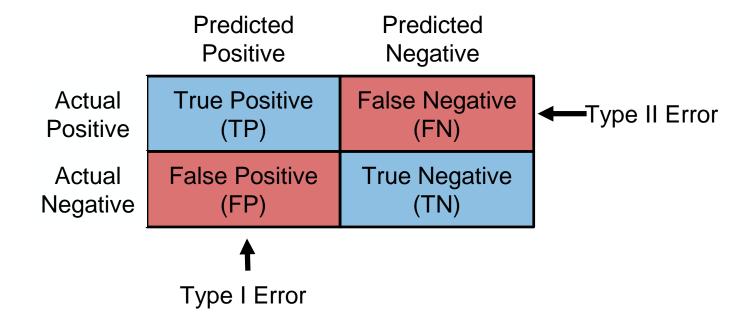


Confusion Matrix

	Predicted Positive	Predicted Negative
Actual	True Positive	False Negative
Positive	(TP)	(FN)
Actual	False Positive	True Negative
Negative	(FP)	(TN)



Confusion Matrix





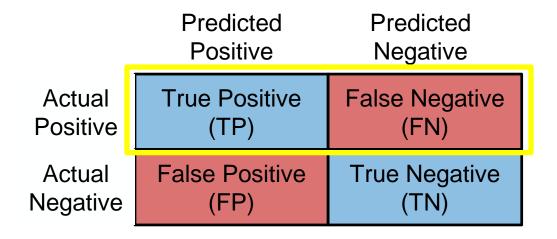
Accuracy: Predicting Correctly

	Predicted Positive	Predicted Negative
Actual	True Positive	False Negative
Positive	(TP)	(FN)
Actual	False Positive	True Negative
Negative	(FP)	(TN)

Accuracy =
$$\frac{TP + TN}{TP + FN + FP + TN}$$



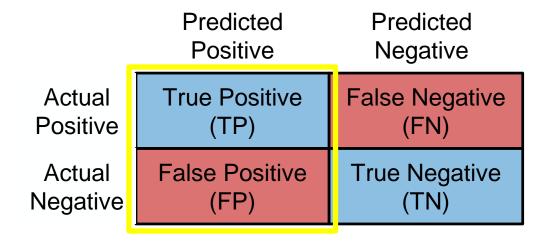
Recall: Identifying All Positive Instances



Recall or
$$=$$
 $\frac{TP}{Sensitivity}$ $=$ $\frac{TP}{TP + FN}$



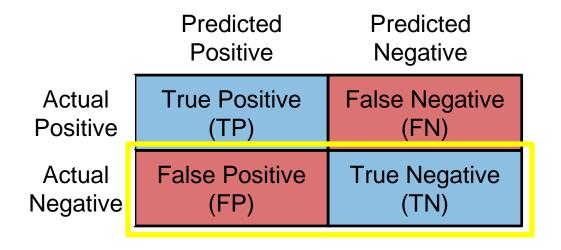
Precision: Identifying Only Positive Instances



Precision =
$$\frac{TP}{TP + FP}$$



Specificity: Avoiding False Alarms



Specificity =
$$\frac{TN}{FP + TN}$$



Error Measurements

	Predicted Positive	Predicted Negative	
Actual	True Positive	False Negative	
Positive	(TP)	(FN)	
Actual	False Positive	True Negative	
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Accuracy =
$$\frac{TP + TN}{TP + FN + FP + TN}$$
Precision =
$$\frac{TP}{TP + FP}$$



Error Measurements

	Predicted Positive	Predicted Negative	
Actual	True Positive	False Negative	
Positive	(TP)	(FN)	
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Negative	(FP)	(TN)	

Accuracy =
$$\frac{TP + TN}{TP + FN + FP + TN}$$
 Recall or Sensitivity = $\frac{TP}{TP + FN}$

Precision = $\frac{TP}{TP + FP}$ Specificity = $\frac{TN}{FP + TN}$



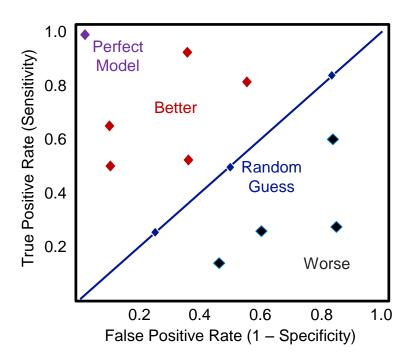
Error Measurements

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Actual	True Positive	False Negative	
Positive	(TP)	(FN)	
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Accuracy =
$$\frac{TP + TN}{TP + FN + FP + TN}$$
 Recall or $= \frac{TP}{Sensitivity}$ $= \frac{TP}{TP + FN}$ F1 = 2 $= \frac{Precision * Recall}{Precision + Recall}$ Precision = $= \frac{TP}{TP + FN}$ Specificity = $= \frac{TP}{TN}$ F1 = 2 $= \frac{Precision * Recall}{Precision + Recall}$



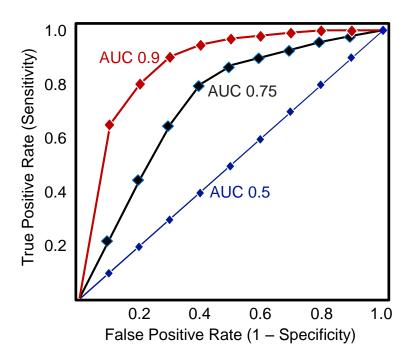
Receiver Operating Characteristic (ROC)



Evaluation of model at all possible thresholds



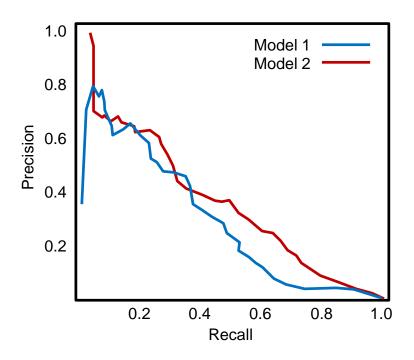
Area Under Curve (AUC)



Measures total area under ROC curve



Precision Recall Curve (PR Curve)



Measures trade-off between precision and recall



Multiple Class Error Metrics

	Predicted Class 1	Predicted Class 2	Predicted Class 3
Actual Class 1	TP1		
Actual Class 2		TP2	
Actual Class 3			TP3



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	Predicted Class 1	Predicted Class 2	Predicted Class 3
Actual Class 1	TP1		
Actual Class 2		TP2	
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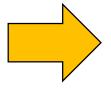
Accuracy =
$$\frac{TP1 + TP2 + TP3}{Total}$$



Multiple Class Error Metrics

	Predicted Class 1	Predicted Class 2	Predicted Class 3
Actual Class 1	TP1		
Actual Class 2		TP2	
Actual Class 3			TP3

Accuracy =
$$\frac{TP1 + TP2 + TP3}{Total}$$



Most multi-class error metrics are similar to binary versions— just expand elements as a sum



Classification Error Metrics: The Syntax

Import the desired error function

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accuracy_value = accuracy_score(y_test, y_pred)



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```

Lots of other error metrics and diagnostic tools:

```
from sklearn.metrics import precision_score, recall_score,
f1_score, roc_auc_score,
confusion_matrix, roc_curve,
precision_recall_curve
```



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