## Mudcard

- For quiz 1, if we display the output of the confusion matrix, I am curious about how to interpret this output [[5 2 0] [1 3 1] [1 1 2]]
  - it is a 3x3 confusion matrix because there are three different classes
  - the meaning of the confusi9on matrix doesn't change, C\_ij is equal to the number of observations known to be in group but predicted to be in group
  - the only difference is that i and j can be 0, 1, or 2.
- why not use multiple imputation?
  - imputation generally can be very misleading
  - imputation can reduce the variance of the feature, it can distort the relationship between the imputed feature and the target variable, it can hide data quality issues, and it can reduce model performance
- still confused on how to choose a metric should we use fscore by default?
  - no, there is no default metric
  - it depends on what your model will do once it is deployed
  - in classification models, the question usually comes down to the 'cost' of model's mistakes
  - which is worse? false positives or false negatives?
- What is the meaning of f-beta score? What does the score indicate about the dataset?
  - evaluation metrics don't usually indicate anything about your dataset
  - they measure the quality of the predictions of a machine learning model
  - let's discuss first the f1 score
    - precision and recall are often in tension with each other
    - o a model that has good recall, often has poor precision and vice versa
    - the f1 score tries to force the model to be good at both at the same time
    - the way this is achieved by the geometric mean of precision and recall
    - if either precision or recall are low, the overall f1 score is low as well
    - that's not necessarily true for other ways of averaging the two scores
  - the f-beta score is a generalization of the f1 score where you can choose how much you care about precision vs recall
    - o if beta > 1, recall gets more weight, it is considered more important
    - if beta < 1, precision gets more weight, it is considered more important</li>

# Evaluation metrics in supervised ML, part 2, predicted probabilities and regression metrics

By the end of this lecture, you will be able to

- Summarize the ROC and precision-recall curves, and the logloss metric
- Describe the most commonly used regression metrics

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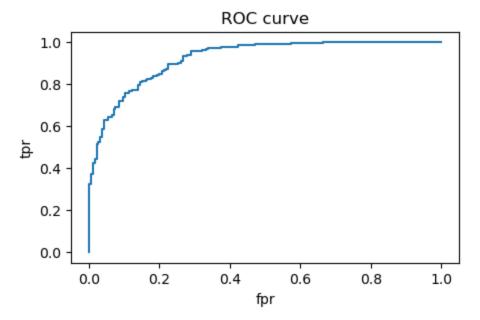
#### The ROC curve

- Receiver Operating Characteristic
  - x axis: false positive rate (fpr = FP / (FP + TN))
  - y axis: true positive rate (R = TP / (TP + FN))
  - the curve shows fpr and R value pairs for various class 1 critical probabilities
- upper left corner: perfect predictor
- · diagonal point: chance level predictions
- lower right corner: worst predictor

```
In [1]: import pandas as pd
        import matplotlib.pyplot as plt
        import numpy as np
        from sklearn.metrics import confusion matrix
        df = pd.read_csv('../data/true_labels_pred_probs.csv')
        y_true = df['y_true']
        pred_prob_class1 = df['pred_prob_class1']
        pred_prob_class0 = df['pred_prob_class0']
        fpr = np.zeros(len(y_true))
        tpr = np.zeros(len(y true))
        p_crits = np.sort(pred_prob_class1) # the sorted predicted probabilities ser
        for i in range(len(p crits)):
            p_crit = p_crits[i]
            y pred = np.zeros(len(y true))
            y_pred[pred_prob_class1 < p_crit] = 0</pre>
            y_pred[pred_prob_class1 >= p_crit] = 1
            C = confusion_matrix(y_true,y_pred)
            tpr[i] = C[1,1]/(C[1,0]+C[1,1])
            fpr[i] = C[0,1]/(C[0,0]+C[0,1])
        from sklearn.metrics import roc curve
```

```
# # the roc_curve function performs the same calculation
fpr,tpr,p_crits = roc_curve(y_true,pred_prob_class1)
```

```
In [2]: plt.figure(figsize=(5,3))
   plt.plot(fpr,tpr)
   plt.xlabel('fpr')
   plt.ylabel('tpr')
   plt.title('ROC curve')
   plt.show()
```



# Quiz 1

What's the (fpr,tpr) coordinate on the ROC curve if p\_crit = 1?

## **ROC AUC**

- ROC is useful but it is not a single number metric
  - it cannot be directly used to compare various classification models
- summary statistics based on the ROC curve (for a complete list, see here)
- most commonly used metric is ROC AUC ROC Area Under the Curve
  - AUC = 1 is a perfect classifier
  - AUC > 0.5 is above chance-level predictor
  - AUC = 0.5 is a chance-level classifier
  - AUC < 0.5 is a bad predictor
  - AUC = 0 classifies all points incorrectly

```
In [3]: from sklearn.metrics import roc_auc_score
  print(roc_auc_score(y_true,pred_prob_class1))
```

0.9236524315231854

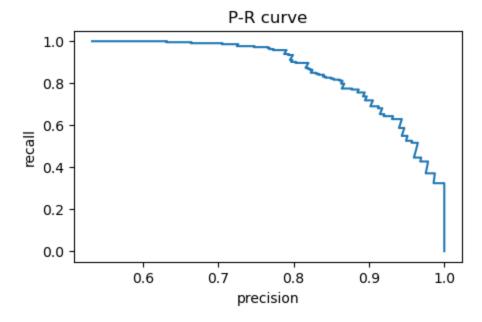
### Precision-recall curve

- the drawback of ROC is that it uses TN, not good for imbalanced problems.
- the precision-recall curve doesn't use TN, ideal for imbalanced problems.

```
In [4]: from sklearn.metrics import precision_recall_curve
    from sklearn.metrics import average_precision_score # the AUC of the P-R cur
    p,r,p_crits = precision_recall_curve(y_true,pred_prob_class1)
    print(average_precision_score(y_true,pred_prob_class1))
```

#### 0.9315588971251672

```
In [5]: plt.figure(figsize=(5,3))
   plt.plot(p,r)
   plt.xlabel('precision')
   plt.ylabel('recall')
   plt.title('P-R curve')
   plt.show()
```



## Quiz 2

What's the (p,r) coordinate on the curve if  $p_{crit} = 0$ ?

## The logloss metric

```
\log s = -\frac{1}{m}\sum_{i=0}^{m-1} (y_{i, true}\ln(p_{i, pred}) + (1-y_{i, true})\ln(1-p_{i, pred}))
```

\$p\_{pred}\$ is the predicted probability of the positive class

- the predicted probabilities are not converted into predicted classes
- excellent choice if you need accurate probabilities (e.g., when it is expensive/costly
  to act due to limited resources so you need to rank your points based on
  probabilities)
- two scenarios:
  - y\_true = 0 left term disappears
  - y\_true = 1 right term disappears
- log(0) is undefined
  - \$p\_{pred}\$ is replaced with \$\max(\min(p,1-10^{-15}),10^{-15})\$ to avoid this issue

#### The extreme cases

- the classifier is confidently wrong
  - \$p\_{pred} = 10^{-15}\$ for points in class 1
  - $p_{pred} = 1 10^{-15}$  for points in class 0  $\log \log = -\frac{1}{m}\sum_{n=0}^{-15} = -\ln(10^{-15})$   $\log \log \sum \sin 34.5$
- the classifier is correct
  - \$p\_{pred} = 10^{-15}\$ for points in class 0
  - \$p\_{pred} = 1 10^{-15}\$ for points in class 1
    \$logloss = -\frac{1}{m}\sum (1-0)(1-\ln(1-10^{-15})) = 10^{-15}\$ for class 0
    \$logloss = -\frac{1}{m}\sum 1\*\ln(1-10^{-15}) = 10^{-15}\$ for class 1
    \$logloss \sim 0\$

```
In [6]: from sklearn.metrics import log_loss
    print(log_loss(y_true,pred_prob_class1))
    help(log_loss)
```

```
lecture13 - evaluation metrics part 2
0.3501519054532857
Help on function log_loss in module sklearn.metrics._classification:
log_loss(y_true, y_pred, *, normalize=True, sample_weight=None, labels=None)
    Log loss, aka logistic loss or cross-entropy loss.
    This is the loss function used in (multinomial) logistic regression
    and extensions of it such as neural networks, defined as the negative
    log-likelihood of a logistic model that returns ``y pred`` probabilities
    for its training data ``y_true``.
    The log loss is only defined for two or more labels.
    For a single sample with true label :math: 'y \in \{0,1\}' and
    a probability estimate :math:p = \operatorname{probability}(y = 1), the log
    loss is:
    .. math::
        L_{\log}(y, p) = -(y \log (p) + (1 - y) \log (1 - p))
    Read more in the :ref:`User Guide <log loss>`.
    Parameters
    y_true : array-like or label indicator matrix
        Ground truth (correct) labels for n_samples samples.
    y_pred : array-like of float, shape = (n_samples, n_classes) or (n_sampl
es,)
        Predicted probabilities, as returned by a classifier's
        predict_proba method. If ``y_pred.shape = (n_samples,)``
        the probabilities provided are assumed to be that of the
        positive class. The labels in ``y_pred`` are assumed to be
        ordered alphabetically, as done by
        :class:`~sklearn.preprocessing.LabelBinarizer`.
        `y_pred` values are clipped to `[eps, 1-eps]` where `eps` is the mac
hine
        precision for `y_pred`'s dtype.
    normalize : bool, default=True
        If true, return the mean loss per sample.
        Otherwise, return the sum of the per-sample losses.
    sample_weight : array-like of shape (n_samples,), default=None
        Sample weights.
    labels : array-like, default=None
        If not provided, labels will be inferred from y_true. If ``labels``
        is ``None`` and ``y pred`` has shape (n samples,) the labels are
        assumed to be binary and are inferred from ``y_true``.
        .. versionadded:: 0.18
    Returns
```

\_\_\_\_\_

loss : float

Log loss, aka logistic loss or cross-entropy loss.

#### Notes

\_\_\_\_

The logarithm used is the natural logarithm (base-e).

#### References

C.M. Bishop (2006). Pattern Recognition and Machine Learning. Springer, p. 209.

#### Examples

```
>>> from sklearn.metrics import log_loss
>>> log_loss(["spam", "ham", "ham", "spam"],
... [[.1, .9], [.9, .1], [.8, .2], [.35, .65]])
0.21616...
```

# Evaluation metrics in supervised ML, part 2, predicted probabilities and regression

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- Describe the most commonly used regression metrics

## Regression metrics

- the target variable is continuous
- the predicted values are also continuous
- regression metrics measure some type of difference between y (true values) and y' (predicted values)

## Mean Squared Error

$$MSE(y,y') = \frac{1}{m}\sum_{i=1}^{m}(y_i-y_i')^2$$

No description has been provided for this image

The unit of MSE is not the same as the target variable.

## **Root Mean Square Error**

$$RMSE(y,y') = \sqrt{\frac{1}{m}\sum_{i=1}^{m}(y_i-y_i')^2}$$

### Mean Absolute Error

$$MAE(y,y') = \frac{1}{m}\sum_{i=1}^{m}|y_i-y_i'|$$

Both RMSE and MAE have the same unit as the target variable.

## R2 score - coefficient of determination

$$R^2(y,y') = 1 - \frac{(i=1)^{m}(y_i-y_i')^2}{\sum_{i=1}^{m}(y_i-bar{y})^2},$$

where \$\bar{y}\$ is the mean of \$y\$.

- R2 = 1 is the perfect regression model (\$y == y'\$)
- R2 = 0 is as good as a constant model that always predicts the expected value of y (\$\bar{y}\$)
- R2 < 0 is a bad regression model

#### R2 is dimensionless.

```
In [7]: from sklearn.metrics import mean_squared_error
    from sklearn.metrics import mean_absolute_error
    from sklearn.metrics import r2_score
    help(mean_squared_error)
```

```
Help on function mean_squared_error in module sklearn.metrics._regression:
mean squared error(y true, y pred, *, sample weight=None, multioutput='unifo
rm average')
    Mean squared error regression loss.
    Read more in the :ref:`User Guide <mean squared error>`.
    Parameters
    y_true : array-like of shape (n_samples,) or (n_samples, n_outputs)
        Ground truth (correct) target values.
    y_pred : array-like of shape (n_samples,) or (n_samples, n_outputs)
        Estimated target values.
    sample_weight : array-like of shape (n_samples,), default=None
        Sample weights.
    multioutput : {'raw_values', 'uniform_average'} or array-like of shape
(n_outputs,), default='uniform_average'
        Defines aggregating of multiple output values.
        Array-like value defines weights used to average errors.
        'raw values':
            Returns a full set of errors in case of multioutput input.
        'uniform average':
            Errors of all outputs are averaged with uniform weight.
    Returns
    loss: float or array of floats
        A non-negative floating point value (the best value is 0.0), or an
        array of floating point values, one for each individual target.
    Examples
    >>> from sklearn.metrics import mean_squared_error
    >>> y_true = [3, -0.5, 2, 7]
    >>> y_pred = [2.5, 0.0, 2, 8]
    >>> mean_squared_error(y_true, y_pred)
    0.375
    >>> y true = [[0.5, 1], [-1, 1], [7, -6]]
    >>> y_pred = [[0, 2], [-1, 2], [8, -5]]
    >>> mean_squared_error(y_true, y_pred)
    0.708...
    >>> mean_squared_error(y_true, y_pred, multioutput='raw_values')
    array([0.41666667, 1.
                                 1)
    >>> mean squared error(y true, y pred, multioutput=[0.3, 0.7])
    0.825...
```

 RMSE is not implemented in sklearn, but you can calculate it as np.sqrt(mean\_squared\_error(y\_true,y\_pred)) • you can find more on regression metrics here

# Quiz 3

Read in data/reg\_preds.csv . It contains two columns:

- y\_true: value of owner-occupied homes in \$1000's in Boston
- y\_pred: predictions of a regression model

What's the ratio between the MSE and the variance of the home values? How does this ratio relate to the R2 score?

In [ ]:

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