

# Classification models

MODELING WITH TIDYMODELS IN R



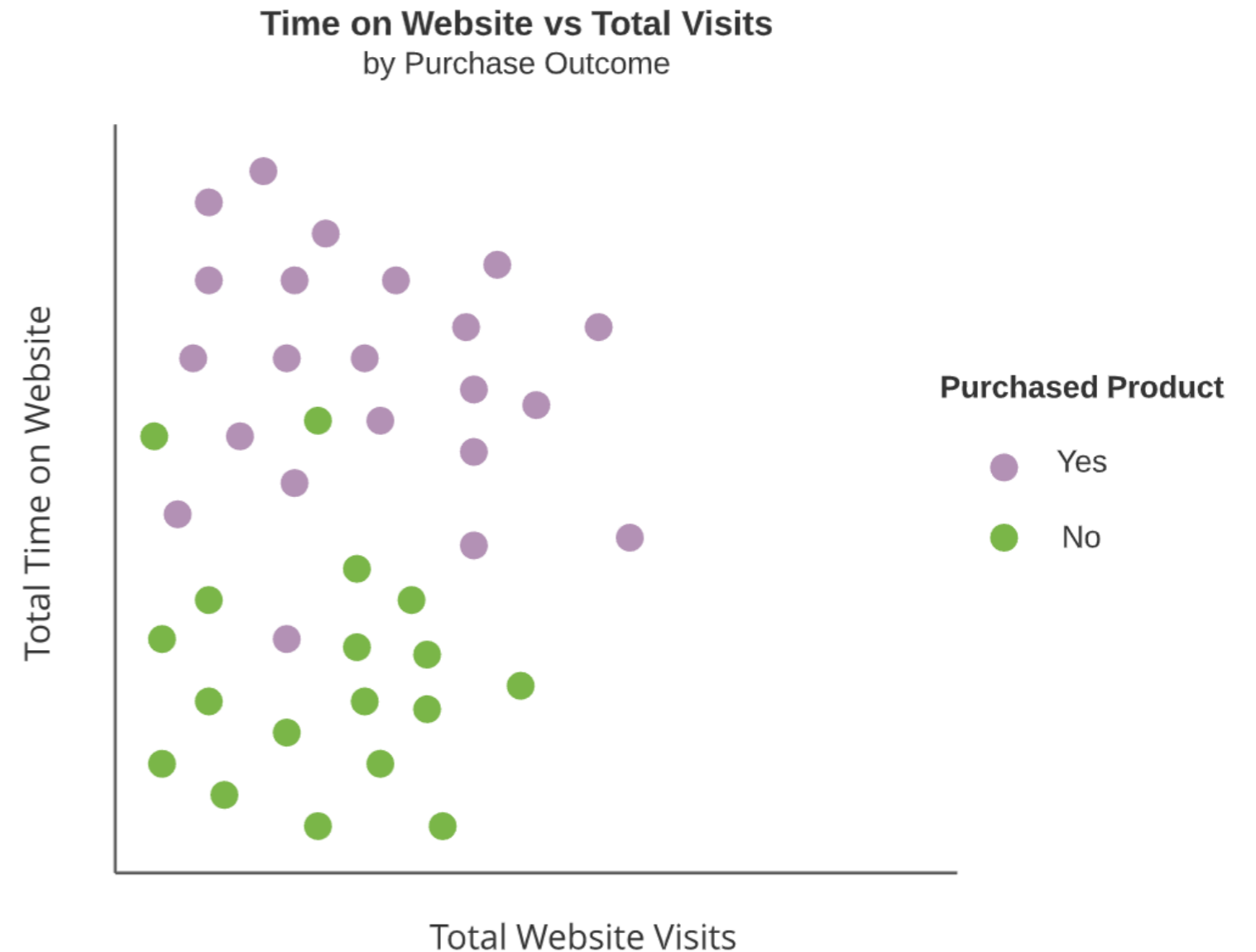
**David Svancer**  
Data Scientist

# Predicting product purchases

Classification models predict categorical outcome variables

- Predicting product purchases

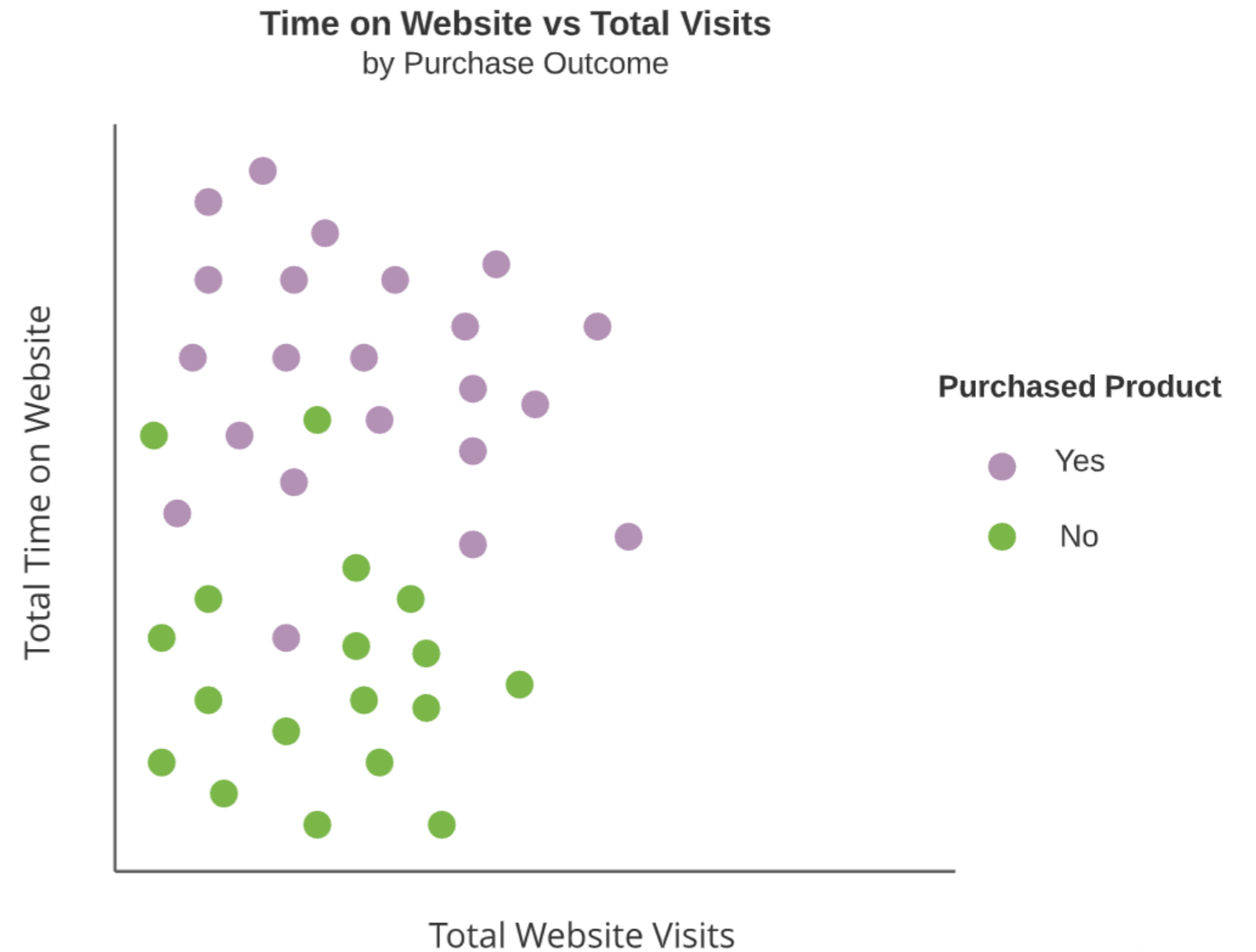
purchased	total_time	total_visits
yes	800	3
yes	978	7
no	220	4
no	124	5
yes	641	4



# Classification algorithms

**Goal:** Create distinct, non-overlapping regions along set of predictor variable values

- Predict the same categorical outcome in each region



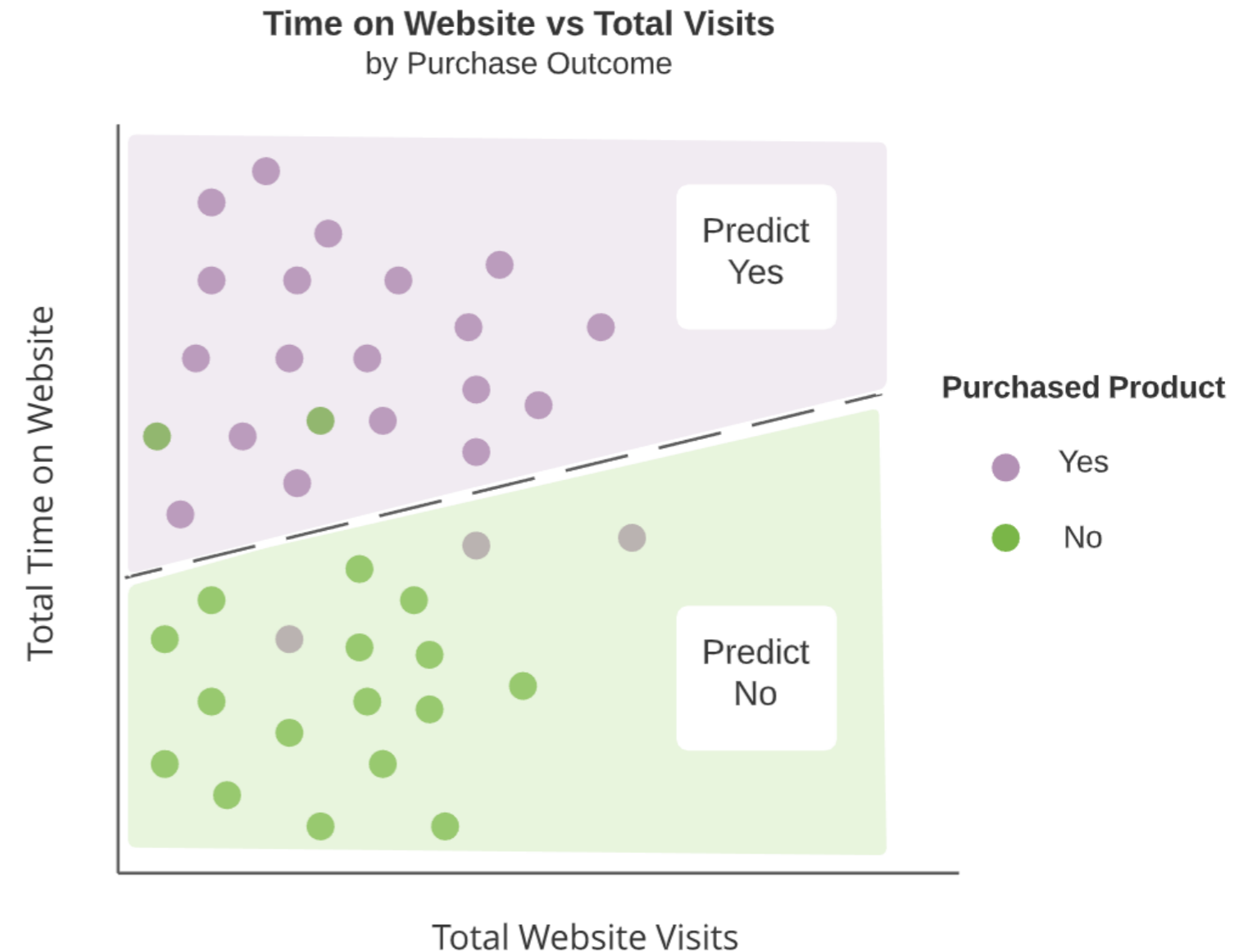
# Classification algorithms

**Goal:** Create distinct, non-overlapping regions along set of predictor variable values

- Predict the same categorical outcome in each region

## Logistic Regression

- Popular classification algorithm which creates a *linear* separation between outcome categories



# Lead scoring data

leads\_df

```
# A tibble: 1,328 x 7
  purchased total_visits total_time pages_per_visit total_clicks lead_source us_location
  <fct>      <dbl>      <dbl>      <dbl>          <dbl>      <dbl> <fct>      <fct>
1 yes        7        1148         7            59 direct_traffic west
2 no         8         100         2.67          24 direct_traffic west
3 no         5         228         2.5           25 email        southeast
4 no         7         481         2.33          21 organic_search west
5 no         4         177         4             37 direct_traffic west
6 no         2        1273         2             26 email        midwest
7 no         3         711         3             28 organic_search west
8 no         3         166         3             32 direct_traffic southeast
9 no         3          7         3             23 organic_search west
10 no        6         562         6            48 organic_search southeast
# ... with 1,318 more rows
```

# Data resampling

First step in fitting a model

- Create data split object with `initial_split()`
- Create training and test datasets with `training()` and `testing()`

```
leads_split <- initial_split(leads_df,  
                             prop = 0.75,  
                             strata = purchased)  
  
leads_training <- leads_split %>%  
  training()  
  
leads_test <- leads_split %>%  
  testing()
```

# Logistic regression model specification

Model specification in `parsnip`

- `logistic_reg()`
  - General interface to logistic regression models in `parsnip`
  - Common engine is 'glm'
  - Mode is 'classification'

```
logistic_model <- logistic_reg() %>%  
  set_engine('glm') %>%  
  set_mode('classification')
```

# Model fitting

Once model is specified, the `fit()` function is used for model training

- Pass model object to `fit()`
- Specify model formula
- Provide training data, `data`

```
logistic_fit <- logistic_model %>%  
  fit(purchased ~ total_visits + total_time,  
      data = leads_training)
```



# Predicting outcome categories

The `predict()` function

- `new_data` specifies dataset on which to predict new values
- `type`
  - `'class'` provides categorical predictions

Standardized output from `predict()`

1. Returns a tibble
2. When `type` is `'class'`, returns a factor column named `.pred_class`

```
class_preds <- logistic_fit %>%  
  predict(new_data = leads_test,  
         type = 'class')
```

```
class_preds
```

```
# A tibble: 332 x 1  
  .pred_class  
  <fct>  
1 no  
2 yes  
3 no  
4 no  
5 yes  
# ... with 327 more rows
```

# Estimated probabilities

Setting `type` to `'prob'` provides estimated probabilities for each outcome category

The `predict()` function will return a tibble with multiple columns

- One for each category of the outcome variable
- Naming convention is `.pred_{outcome_category}`

```
prob_preds <- logistic_fit %>%  
  predict(new_data = leads_test,  
         type = 'prob')
```

```
prob_preds
```

```
# A tibble: 332 x 2  
  .pred_yes .pred_no  
    <dbl>    <dbl>  
1  0.134    0.866  
2  0.729    0.271  
3  0.133    0.867  
4  0.0916   0.908  
5  0.598    0.402  
# ... with 327 more rows
```

# Combining results

For model evaluation with the `yardstick` package, a results tibble will be needed

The outcome variable from the test dataset and prediction tibbles can be combined with `bind_cols()`

```
leads_results <- leads_test %>%  
  select(purchased) %>%  
  bind_cols(class_preds, prob_preds)
```

```
leads_results
```

```
# A tibble: 332 x 4  
  purchased .pred_class .pred_yes .pred_no  
  <fct>     <fct>         <dbl>   <dbl>  
1 no       no             0.134   0.866  
2 yes     yes            0.729   0.271  
3 no       no             0.133   0.867  
4 no       no             0.0916  0.908  
5 yes     yes            0.598   0.402  
# ... with 327 more rows
```

# Telecommunications data

telecom\_df

```
# A tibble: 975 x 9
  canceled_service cellular_service avg_data_gb avg_call_mins avg_intl_mins internet_service contract months_with_company monthly_charges
  <fct>             <fct>             <dbl>         <dbl>         <dbl>         <fct>         <fct>         <dbl>         <dbl>
1 yes              single_line        7.78          497          127          fiber_optic    month_to_month 7           76.4
2 yes              single_line        9.04          336           88          fiber_optic    month_to_month 10          94.9
3 no               single_line       10.3          262           55          fiber_optic    one_year      50          103.
4 yes              multiple_lines     5.08          250          107          digital        one_year      53          60.0
5 no               multiple_lines     8.05          328          122          digital        two_year      50          75.2
6 no               single_line        9.3           326          114          fiber_optic    month_to_month 25          95.7
7 yes              multiple_lines     8.01          525           97          fiber_optic    month_to_month 19          83.6
8 no               multiple_lines     9.4           312          147          fiber_optic    one_year      50          99.4
9 yes              single_line        5.29          417           96          digital        month_to_month 8           49.8
10 no              multiple_lines     9.96          340          136          fiber_optic    month_to_month 61          106.
# ... with 965 more rows
```

# Let's practice!

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# Assessing model fit

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# Binary classification

Outcome variable with two levels

- **Positive class**
  - Event of interest to predict
  - "yes" in `purchased` variable
- **Negative class**
  - "no"
- In `tidymodels` outcome variable needs to be a factor
  - First level is positive class
  - Check order with `levels()`

```
leads_df
```

```
# A tibble: 1,328 x 7
  purchased total_visits ... us_location
  <fct>      <dbl>   ... <fct>
1 yes         7     ... west
2 no          8     ... west
3 no          5     ... southeast
# ... with 1,325 more rows
```

```
levels(leads_df[['purchased']])
```

```
[1] "yes" "no"
```

# Confusion matrix

Matrix with counts of all combinations of actual and predicted outcome values

## Correct Predictions

- True Positive (TP)
- True Negative (TN)

## Classification Errors

- False Positive (FP)
- False Negative (FN)

		Truth	
		Positive (+)	Negative (-)
Predicted	Positive (+)	TP	FP
	Negative (-)	FN	TN



# Classification metrics with yardstick

Creating confusion matrices and other model fit metrics with yardstick

- Requires a tibble of model results which contain:
  - True outcome values
    - `purchased`
  - Predicted outcome categories
    - `.pred_class`
  - Estimated probabilities of each category
    - `.pred_yes`
    - `.pred_no`

```
leads_results
```

```
# A tibble: 332 x 4
  purchased .pred_class .pred_yes .pred_no
  <fct>     <fct>         <dbl>  <dbl>
1 no       no             0.134   0.866
2 yes      yes            0.729   0.271
3 no       no             0.133   0.867
4 no       no             0.0916  0.908
5 yes      yes            0.598   0.402
6 no       no             0.128   0.872
7 yes      no             0.112   0.888
8 no       no             0.169   0.831
9 no       no             0.158   0.842
10 yes     yes            0.520   0.480
# ... with 322 more rows
```

# Confusion matrix with yardstick

The `conf_mat()` function

- Tibble of model results
- `truth` - column with true outcomes
- `estimate` - column with predicted outcomes

Logistic regression on `leads_df`

- Correctly classified 252 out of 332 customers (76%)
- 46 false negatives
- 34 false positives

```
conf_mat(leads_results,  
         truth = purchased,  
         estimate = .pred_class)
```

	Truth	
Prediction	yes	no
yes	74	34
no	46	178

# Classification accuracy

The `accuracy()` function

- Takes same arguments as `conf_mat()`
- Calculates classification accuracy

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

- `yardstick` functions always return a tibble
  - `.metric` - type of metric
  - `.estimate` - calculated value

```
accuracy(leads_results,  
         truth = purchased,  
         estimate = .pred_class)
```

```
# A tibble: 1 x 3  
  .metric .estimator .estimate  
  <chr>   <chr>         <dbl>  
1 accuracy binary      0.759
```

# Sensitivity

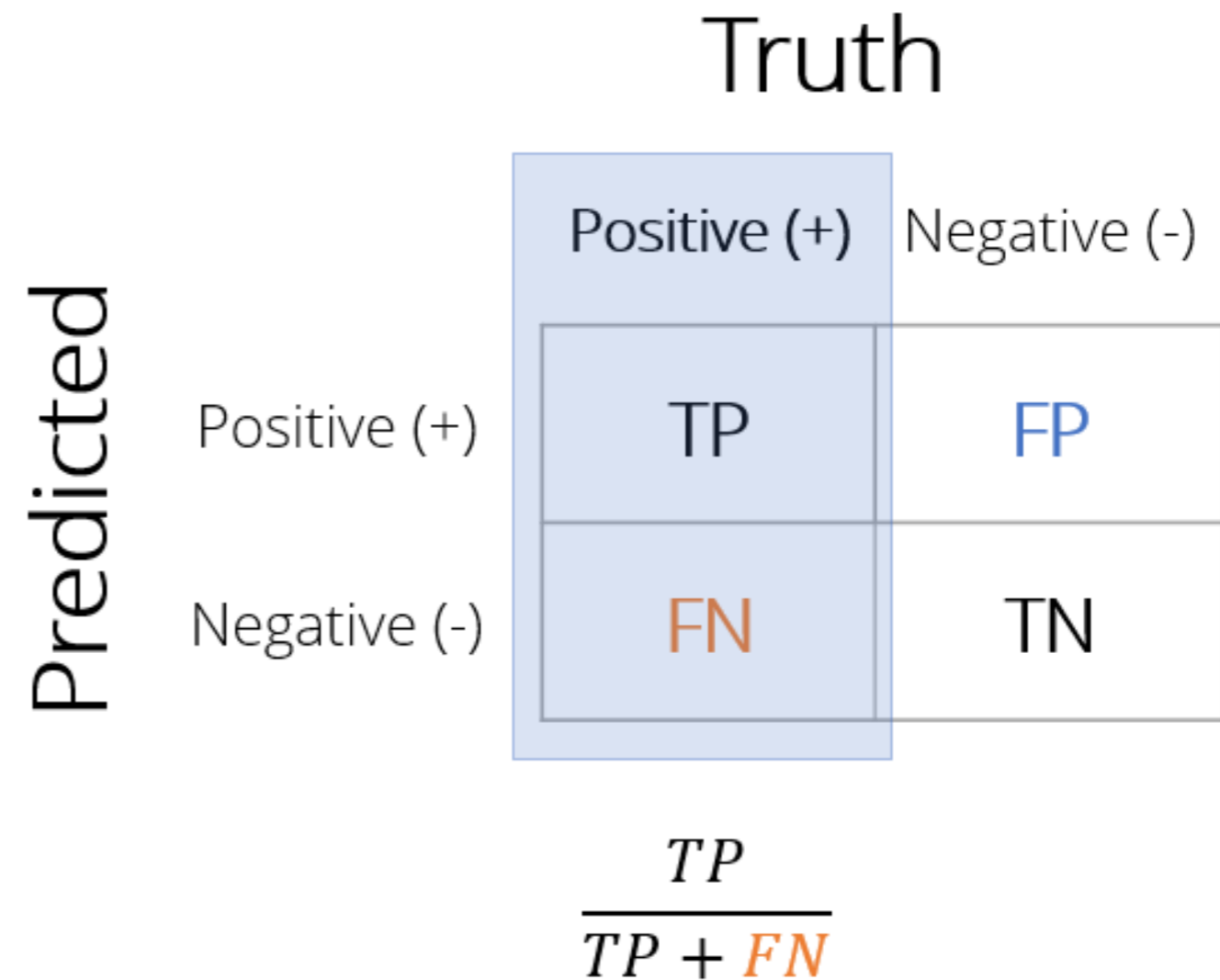
In many cases *accuracy* is not the best metric

- `leads_df` data
  - Classifying all as 'no' gives 64% accuracy

## Sensitivity

Proportion of all positive cases that were correctly classified

- Of customers *who did purchase*, what proportion did our model predict correctly?
  - Lower false negatives increase sensitivity



# Calculating sensitivity

The `sens()` function

- Takes same arguments as `conf_mat()` and `accuracy()`
- Returns sensitivity calculation in `.estimate` column

```
sens(leads_results,  
     truth = purchased,  
     estimate = .pred_class)
```

```
# A tibble: 1 x 3  
  .metric .estimator .estimate  
  <chr>   <chr>         <dbl>  
1 sens   binary         0.617
```

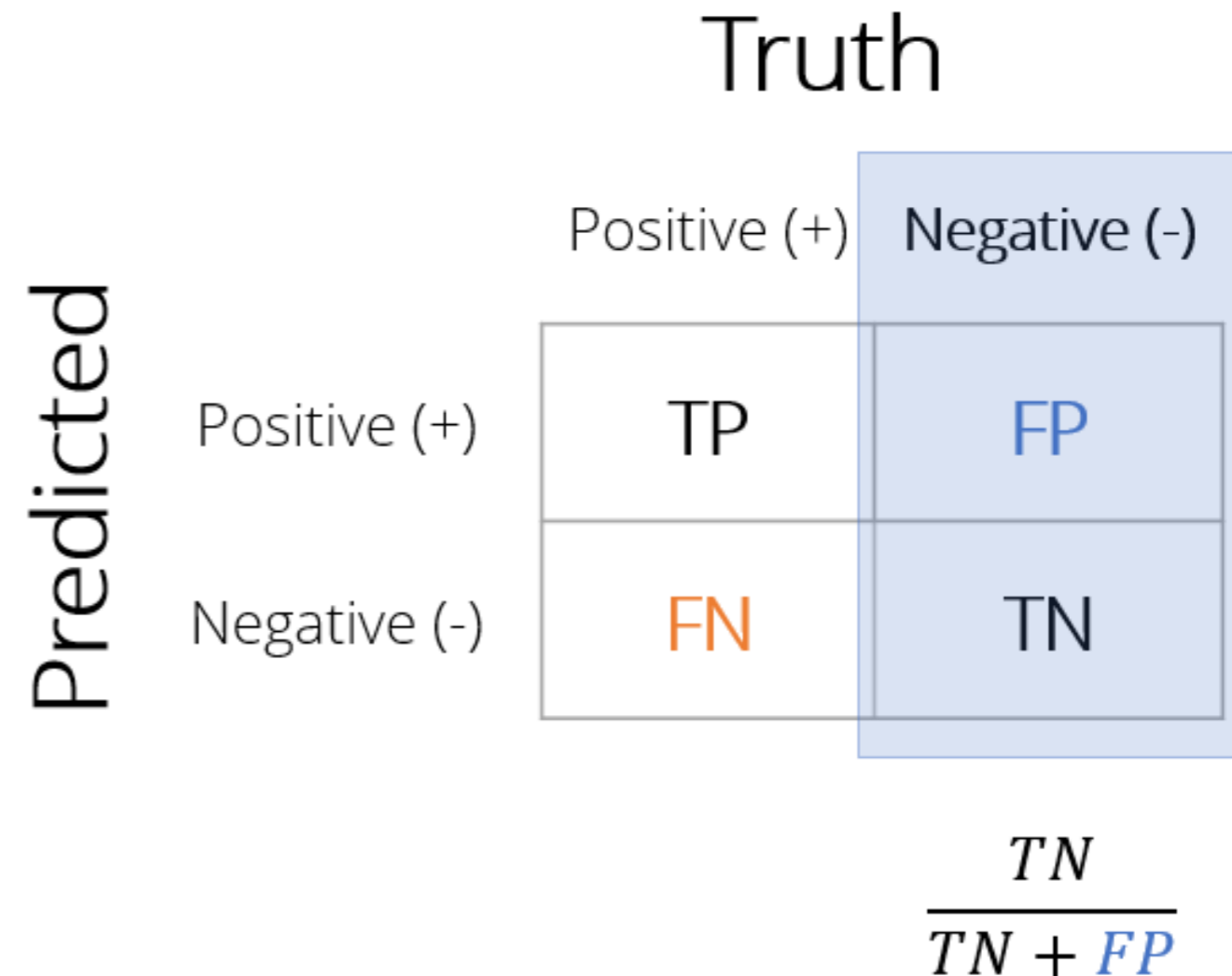
# Specificity

**Specificity** is the proportion of all negative cases that were correctly classified

- Of customers who *did not purchase*, what proportion did our model predict correctly?
  - Lower false positives increase specificity

## 1 - Specificity

- Also called the *false positive rate* (FPR)
- Proportion of false positives among true negatives



# Calculating specificity

The `spec()` function

- Takes same arguments as `sens()`
- Returns specificity calculation in `.estimate` column

```
spec(leads_results,  
     truth = purchased,  
     estimate = .pred_class)
```

```
# A tibble: 1 x 3  
  .metric .estimator .estimate  
  <chr>   <chr>         <dbl>  
1 spec   binary         0.840
```

# Creating a metric set

## User-defined metric sets

- `metric_set()` function
  - Creates user-defined metric function with selected `yardstick` metrics
  - Pass `yardstick` metric function names into `metric_set()`
  - Use custom function to calculate metrics

```
custom_metrics <-  
  metric_set(accuracy, sens, spec)
```

```
custom_metrics(leads_results,  
               truth = purchased,  
               estimate = .pred_class)
```

```
# A tibble: 3 x 3  
  .metric .estimator .estimate  
  <chr>   <chr>         <dbl>  
1 accuracy binary         0.759  
2 sens    binary         0.617  
3 spec    binary         0.840
```



# Many metrics

## Binary classification metrics

- Wide variety of binary classification metrics
  - `accuracy()`, `kap()`, `sens()`, `spec()`,  
`ppv()`, `npv()`, `mcc()`, `j_index()`,  
`bal_accuracy()`,  
`detection_prevalence()`, `precision()`,  
`recall()`, `f_meas()`
- Pass results of `conf_mat()` to `summary()` to calculate all

<https://yardstick.tidymodels.org/reference>

```
conf_mat(leads_results, truth = purchased,  
         estimate = .pred_class) %>%  
summary()
```

```
# A tibble: 13 x 3  
  .metric      .estimator .estimate  
  <chr>        <chr>      <dbl>  
1 accuracy    binary     0.759  
2 kap         binary     0.466  
3 sens        binary     0.617  
4 spec        binary     0.840  
5 ppv         binary     0.685  
6 npv         binary     0.795  
7 mcc         binary     0.468  
8 j_index     binary     0.456  
9 bal_accuracy binary     0.728  
10 detection_prevalence binary     0.325  
11 precision   binary     0.685  
12 recall      binary     0.617  
13 f_meas      binary     0.649
```

# Let's practice!

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# Visualizing model performance

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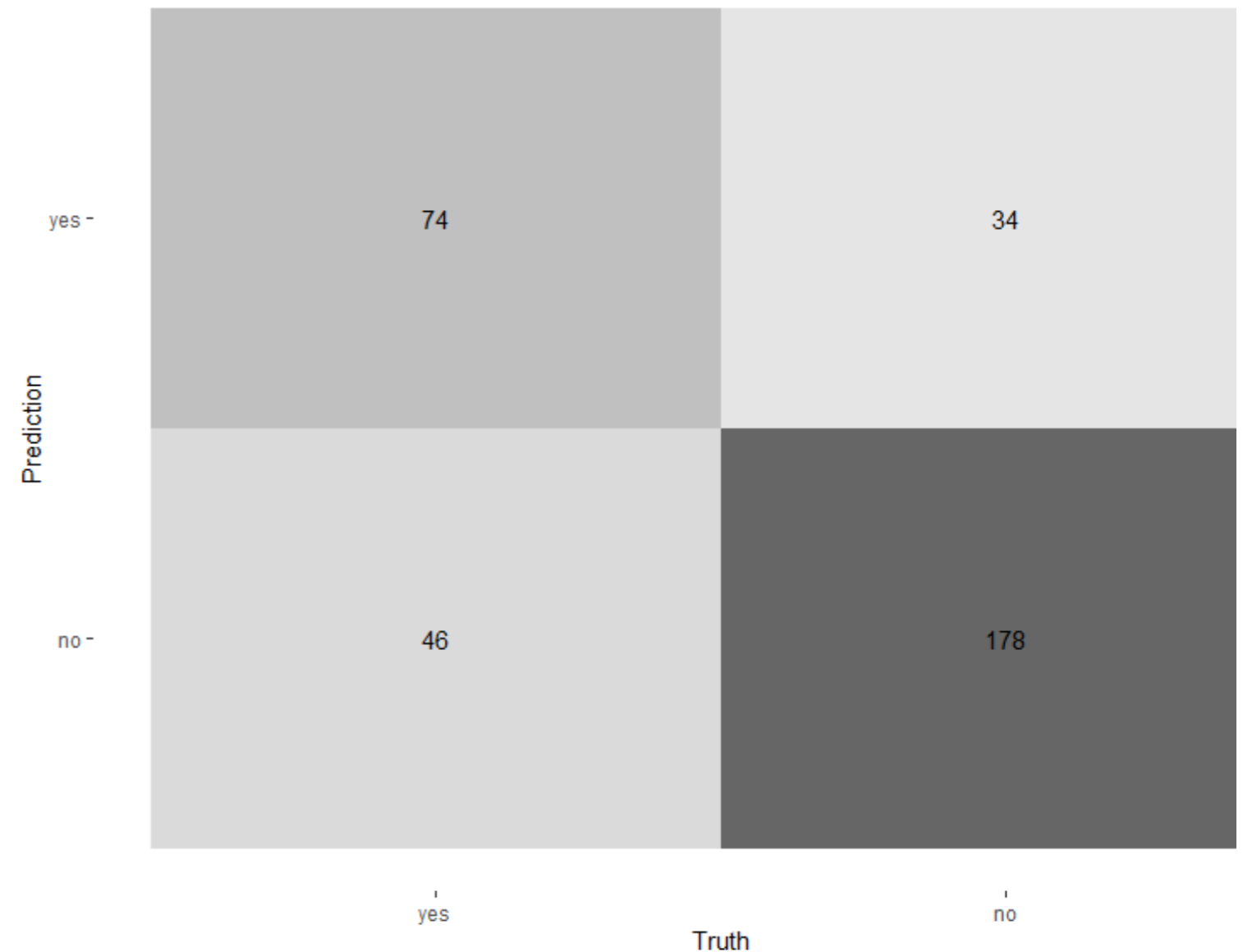
**David Svancer**  
Data Scientist

# Plotting the confusion matrix

Heatmap with `autoplot()`

- Pass confusion matrix object into `autoplot()`
- Set `type` to `'heatmap'`
- Visualize the most prevalent counts

```
conf_mat(leads_results,  
         truth = purchased,  
         estimate = .pred_class) %>%  
  autoplot(type = 'heatmap')
```

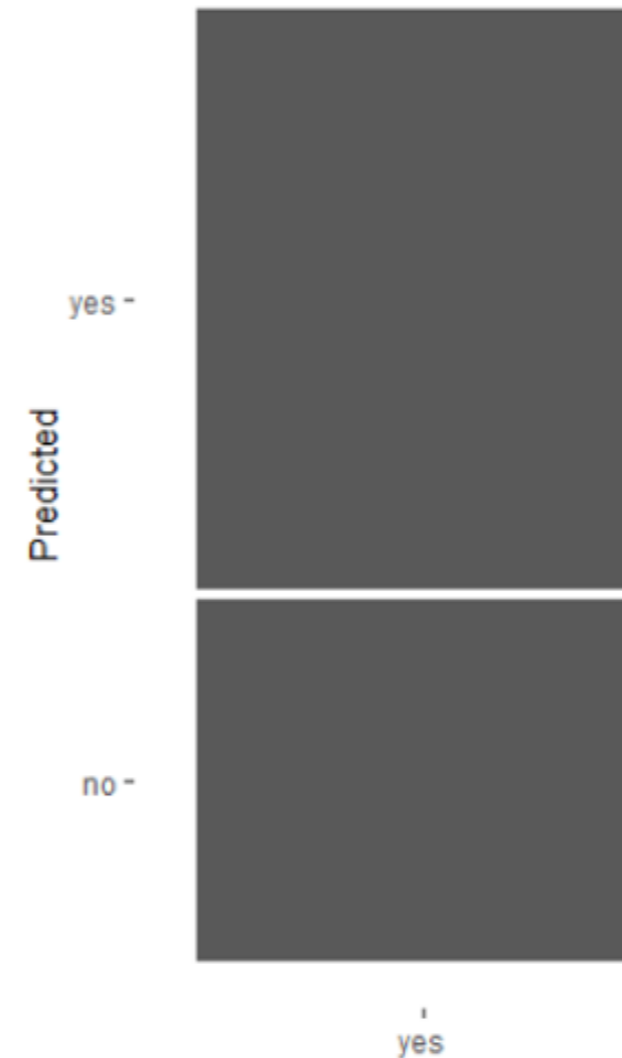


# Mosaic plot

Mosaic with `autoplot()`

- Set `type` to `'mosaic'`
- Each vertical bar represents 100% of actual outcome value in column
- Visually displays
  - sensitivity

```
conf_mat(leads_results,  
         truth = purchased,  
         estimate = .pred_class) %>%  
  autoplot(type = 'mosaic')
```

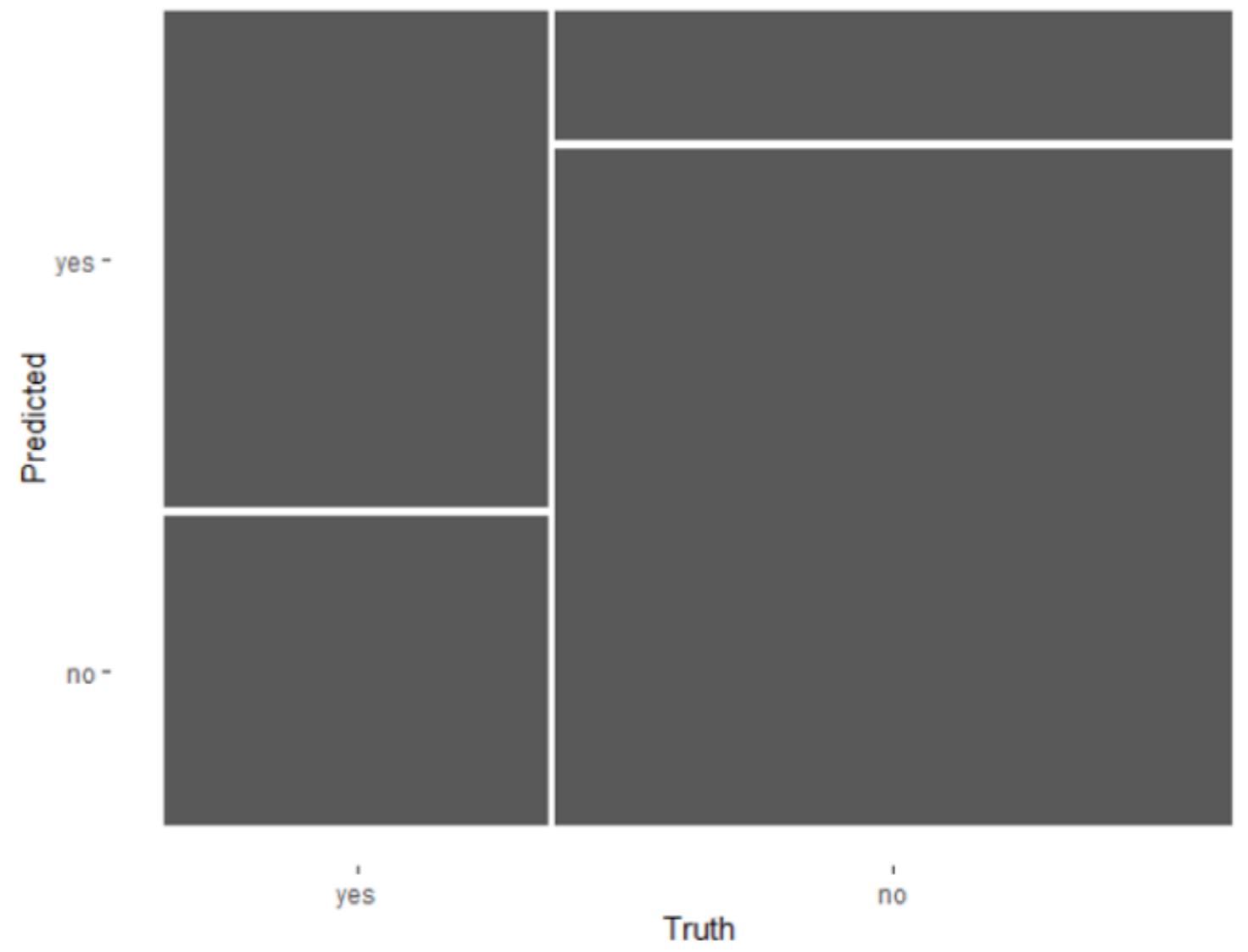


# Mosaic plot

Mosaic with `autoplot()`

- Set `type` to `'mosaic'`
- Each vertical bar represents 100% of actual outcome value in column
- Visually displays
  - sensitivity
  - specificity

```
conf_mat(leads_results,  
         truth = purchased,  
         estimate = .pred_class) %>%  
  autoplot(type = 'mosaic')
```



# Probability thresholds

Default probability threshold in binary classification is 0.5

- If the estimated probability of the positive class is greater than or equal to 0.5, the positive class is predicted

leads\_results

- If `.pred_yes` is greater than or equal to 0.5 then `.pred_class` is set to 'yes' by the `predict()` function in `tidymodels`

leads\_results

```
# A tibble: 332 x 4
  purchased .pred_class .pred_yes .pred_no
  <fct>     <fct>         <dbl>  <dbl>
1 no       no             0.134   0.866
2 yes      yes            0.729   0.271
3 no       no             0.133   0.867
4 no       no             0.0916  0.908
5 yes      yes            0.598   0.402
6 no       no             0.128   0.872
7 yes      no             0.112   0.888
8 no       no             0.169   0.831
9 no       no             0.158   0.842
10 yes     yes            0.520   0.480
# ... with 322 more rows
```

# Exploring performance across thresholds

How does a classification model perform across a range of thresholds?

- Unique probability thresholds in the `.pred_yes` column of the test dataset results
  - Calculate specificity and sensitivity for each

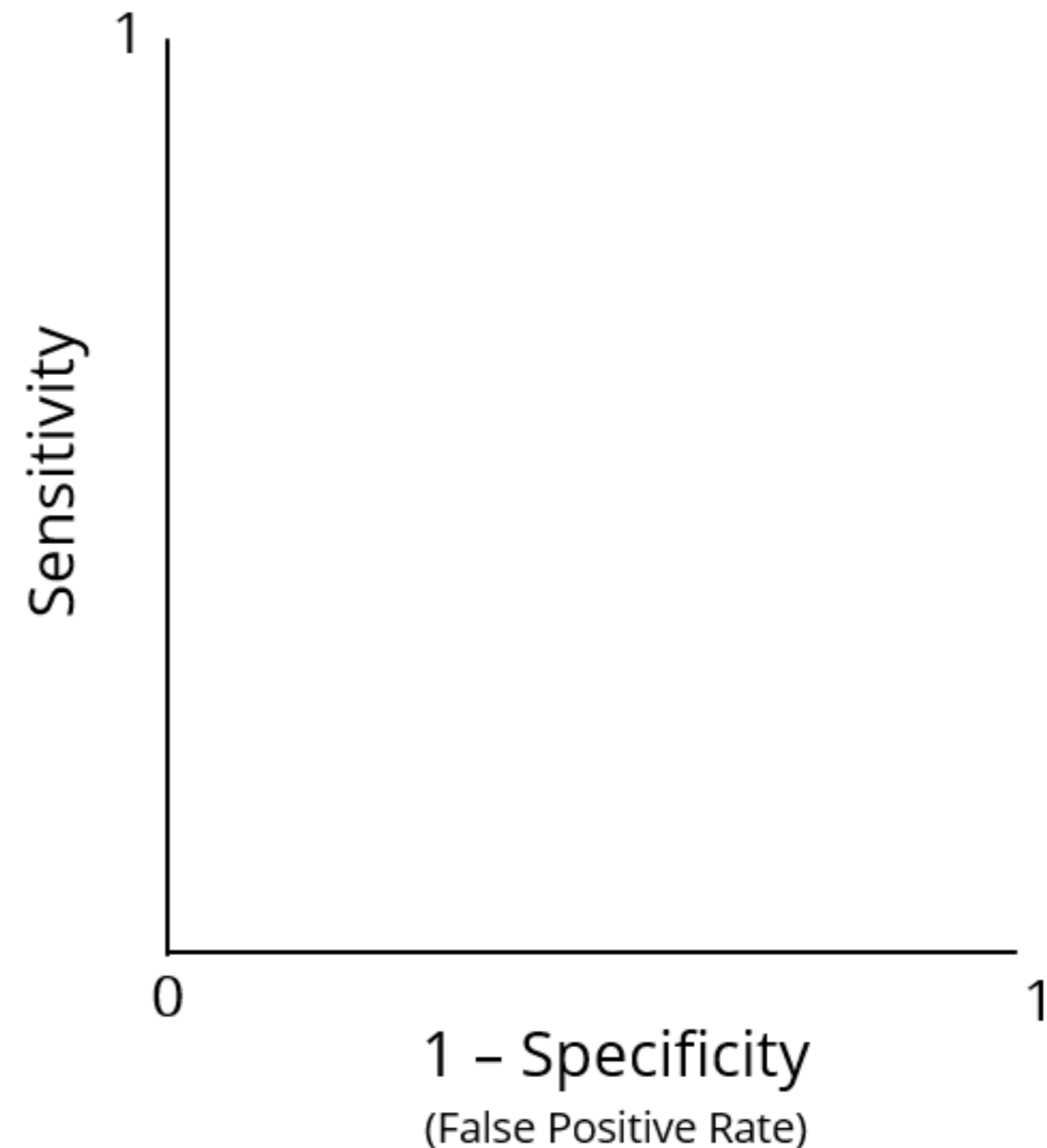
threshold	specificity	sensitivity
0	0	1
0.11	0.01	0.98
0.15	0.05	0.97
...	...	...
0.84	0.89	0.08
0.87	0.94	0.02
0.91	0.99	0
1	1	0



# Visualizing performance across thresholds

Receiver operating characteristic (ROC) curve

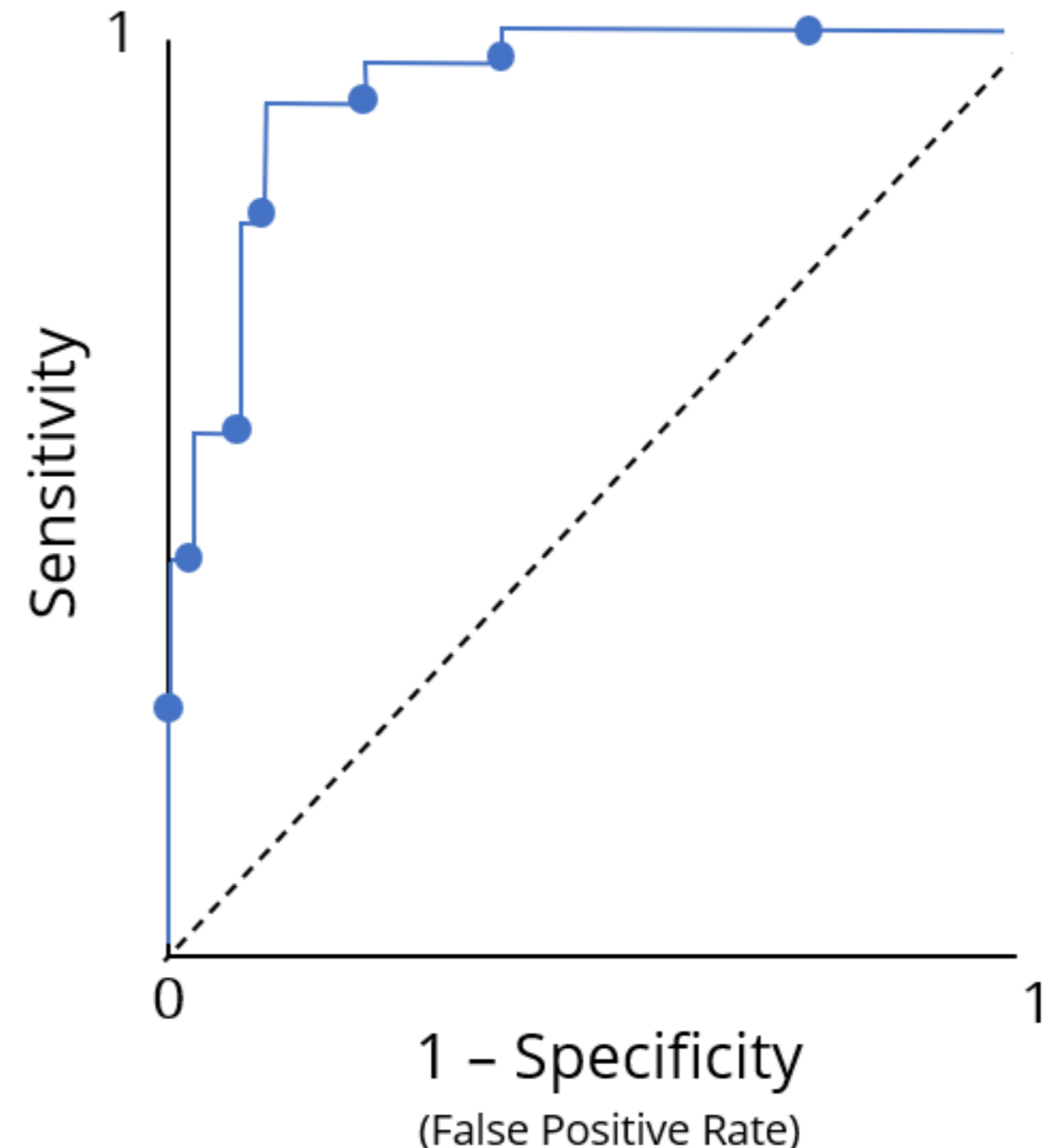
- Used to visualize performance across probability thresholds
- Sensitivity vs. (1 - specificity) across unique thresholds in test set results



# Visualizing performance across thresholds

Receiver operating characteristic (ROC) curve

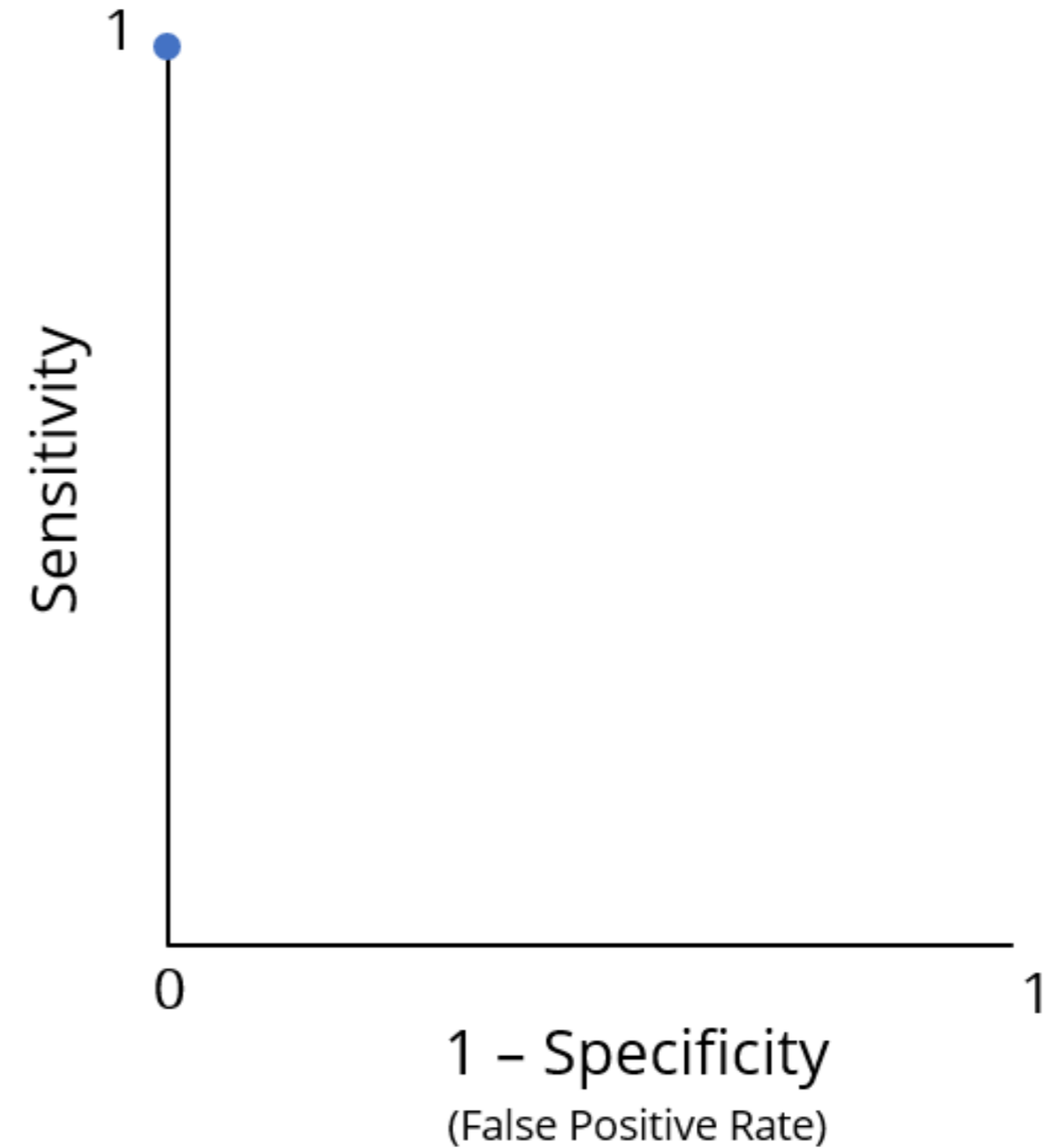
- Used to visualize performance across probability thresholds
- Sensitivity vs (1 - specificity) across unique thresholds in test set results
  - Proportion correct among actual positives vs. proportion **incorrect** among actual negatives



# ROC curves

**Optimal performance** is at the point (0, 1)

- Ideally, a classification model produces points close to left upper edge across all thresholds



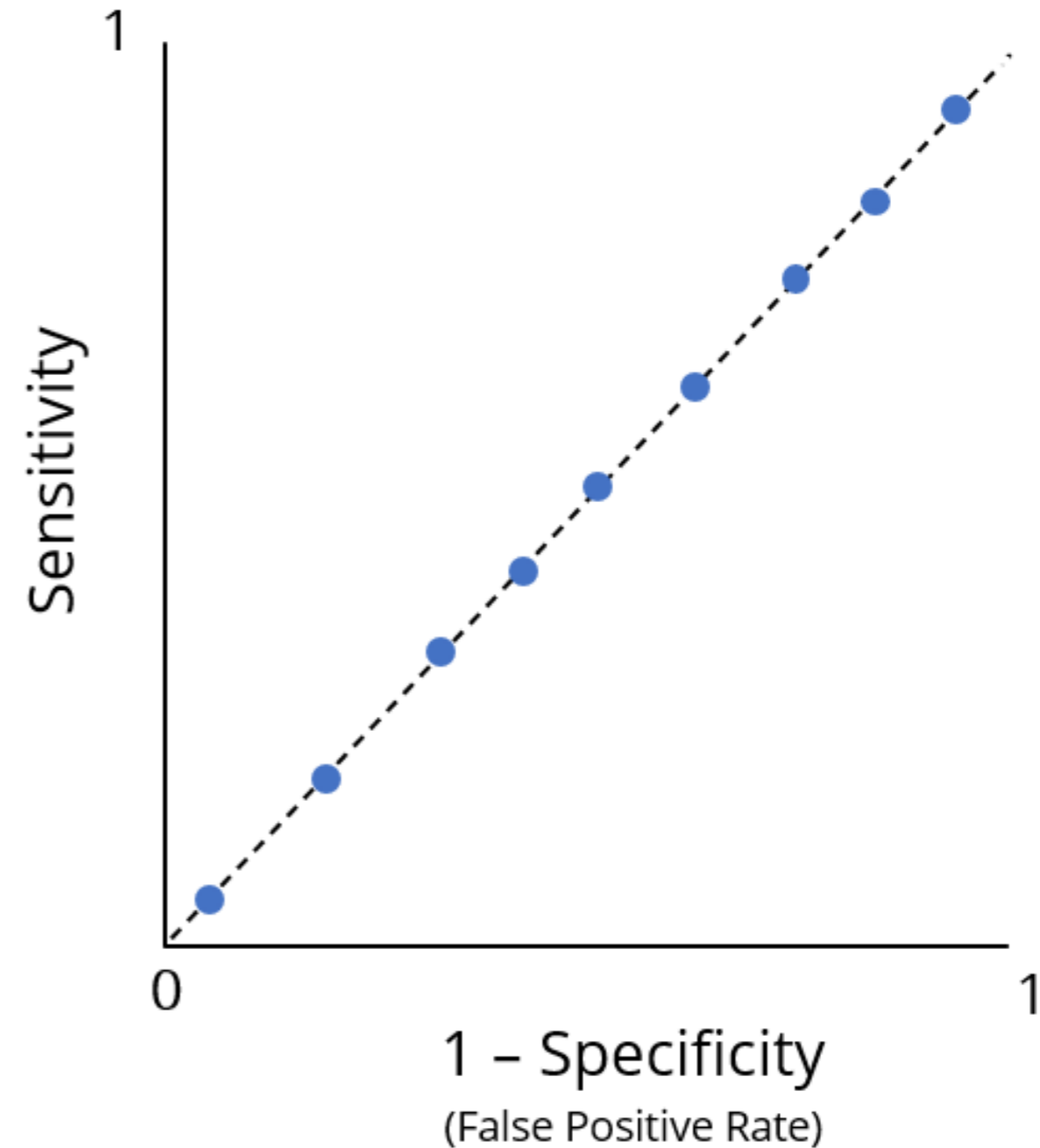
# ROC curves

**Optimal performance** is at the point (0, 1)

- Ideally, a classification model produces points close to left upper edge across all thresholds

## Poor performance

- Sensitivity and (1 - specificity) are equal across all thresholds
  - Corresponds to a classification model that predicts outcomes based on the result of randomly flipping a fair coin

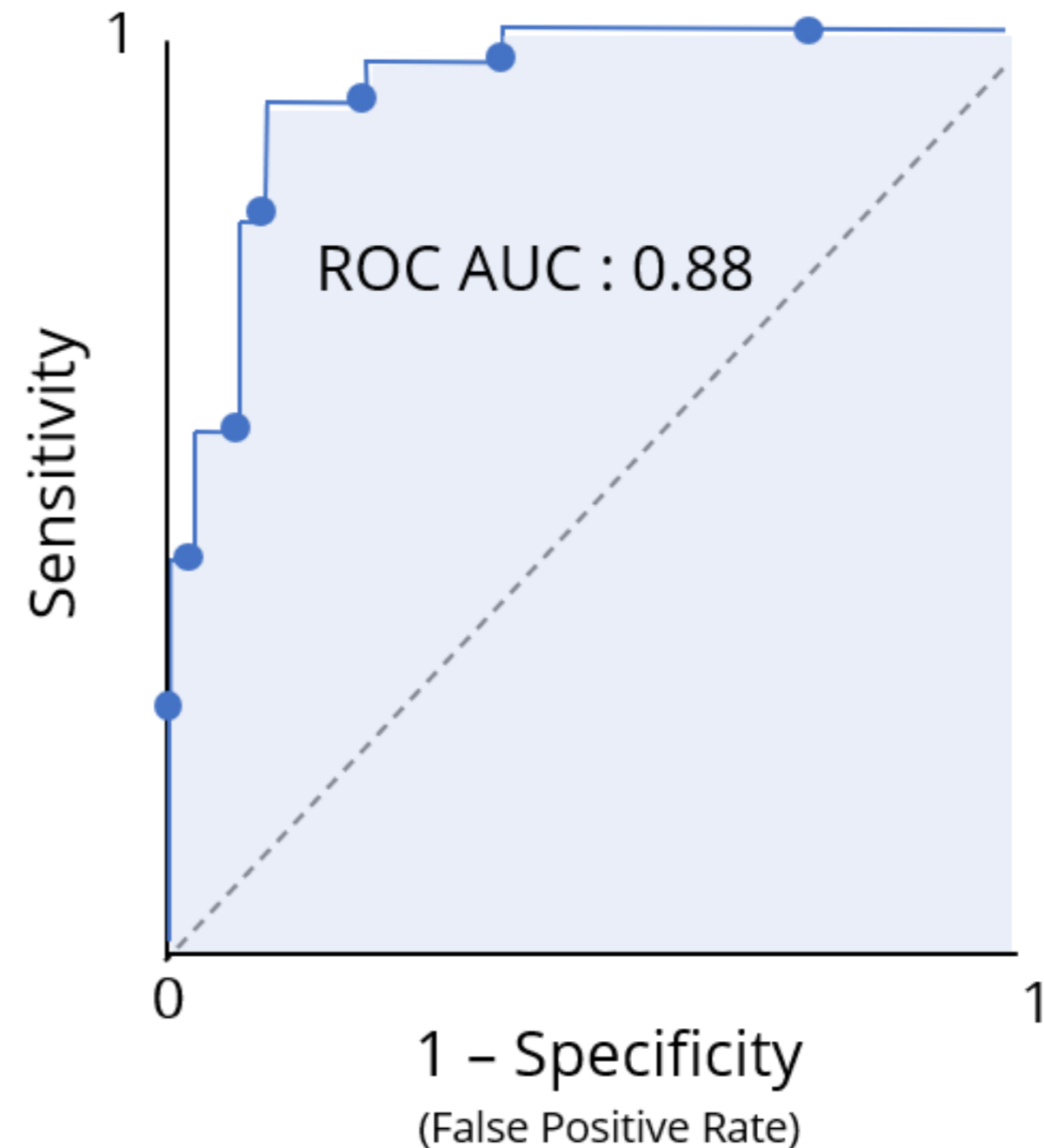


# Summarizing the ROC curve

The area under the ROC curve (ROC AUC) captures the ROC curve information of a classification model in a single number

Useful interpretation as a letter grade of classification performance

- A - [0.9, 1]
- B - [0.8, 0.9)
- C - [0.7, 0.8)
- D - [0.6, 0.7)
- F - [0.5, 0.6)



# Calculating performance across thresholds

The `roc_curve()` function

- Takes a results tibble as the first argument
- `truth` column with true outcome categories
- Column with estimated probabilities for the positive class
  - `.pred_yes` in `leads_results` tibble
- Returns a tibble with specificity and sensitivity for all unique thresholds in `.pred_yes`

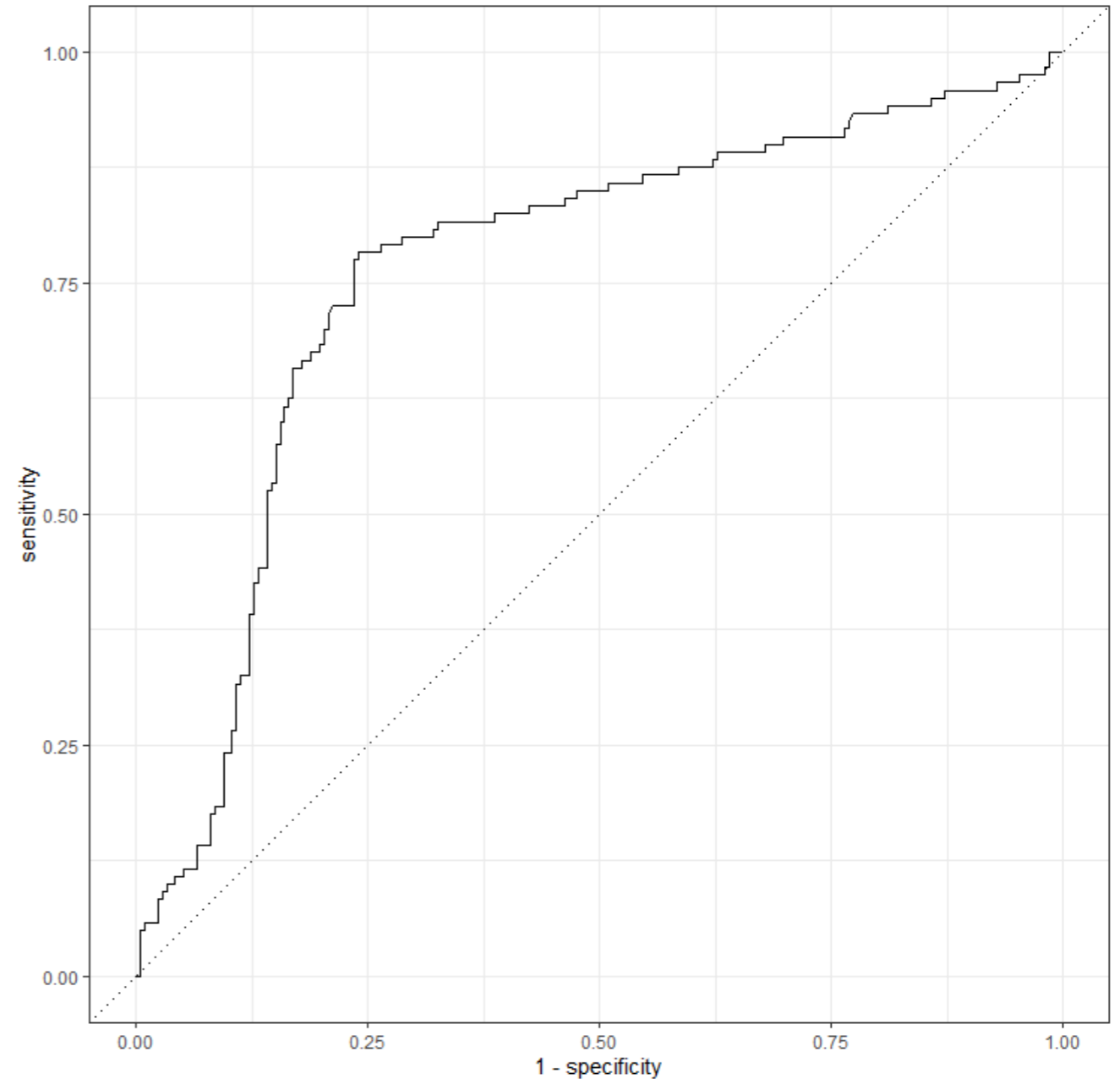
```
leads_results %>%  
  roc_curve(truth = purchased, .pred_yes)
```

```
# A tibble: 331 x 3  
  .threshold specificity sensitivity  
    <dbl>         <dbl>         <dbl>  
1    -Inf             0             1  
2    0.0871            0             1  
3    0.0888            0.00472       1  
4    0.0893            0.00943       1  
5    0.0896            0.0142        1  
6    0.0902            0.0142       0.992  
7    0.0916            0.0142       0.983  
8    0.0944            0.0189       0.983  
# ... with 323 more rows
```

# Plotting the ROC curve

Passing the results of `roc_curve()` to the `autoplot()` function returns an ROC curve plot

```
leads_results %>%  
  roc_curve(truth = purchased, .pred_yes) %>%  
  autoplot()
```



# Calculating ROC AUC

The `roc_auc()` function from `yardstick` will calculate the ROC AUC

- Tibble of model results
- `truth` column
- Column with estimated probabilities for the positive class

```
roc_auc(leads_results,  
        truth = purchased,  
        .pred_yes)
```

```
# A tibble: 1 x 3  
  .metric .estimator .estimate  
  <chr>   <chr>         <dbl>  
1 roc_auc binary       0.763
```



# Let's practice!

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# Automating the modeling workflow

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# Streamlining the workflow

The `last_fit()` function

- Also accepts classification models
- Speeds up the modeling process
- Fits the model to the training data and produces predictions on the test dataset

Similar to using `fit()`, the first steps include:

- Creating a data split object with `rsample`
- Specifying a model with `parsnip`

```
leads_split <- initial_split(leads_df,  
                             strata = purchased)  
  
logistic_model <- logistic_reg() %>%  
  set_engine('glm') %>%  
  set_mode('classification')
```

# Fitting the model and collecting metrics

The `last_fit()` function

- `parsnip` model object
- Model formula
- Data split object

The `collect_metrics()` function calculates metrics using the test dataset

- Accuracy and ROC AUC by default

```
logistic_last_fit <- logistic_model %>%  
  last_fit(purchased ~ total_visits + total_time,  
          split = leads_split)
```

```
logistic_last_fit %>%  
  collect_metrics()
```

```
# A tibble: 2 x 3  
  .metric .estimator .estimate  
  <chr>   <chr>         <dbl>  
1 accuracy binary         0.759  
2 roc_auc binary         0.763
```

# Collecting predictions

```
collect_predictions()
```

- Creates a tibble with all necessary columns for `yardstick` functions
- Actual and predicted outcomes with the test data
- Estimated probability columns for all outcome categories

```
last_fit_results <- logistic_last_fit %>%  
  collect_predictions()
```

```
last_fit_results
```

```
# A tibble: 332 x 6  
  id          .pred_yes .pred_no .row .pred_class purchased  
  <chr>      <dbl>    <dbl> <int> <fct>      <fct>  
1 train/test split 0.134    0.866     2    no        no  
2 train/test split 0.729    0.271    17    yes       yes  
3 train/test split 0.133    0.867    21    no        no  
4 train/test split 0.0916   0.908    22    no        no  
5 train/test split 0.598    0.402    24    yes       yes  
# ... with 327 more rows
```

# Custom metric sets

The `metric_set()` function

- `accuracy()`, `sens()`, and `spec()`
  - Require `truth` and `estimate` arguments
- `roc_auc()`
  - Requires `truth` and column of estimated probabilities

The `custom_metrics()` function will need all three, with `.pred_yes` as the last argument

```
custom_metrics <- metric_set(accuracy, sens,  
                             spec, roc_auc)
```

```
custom_metrics(last_fit_results,  
               truth = purchased,  
               estimate = .pred_class,  
               .pred_yes)
```

```
# A tibble: 4 x 3  
  .metric .estimator .estimate  
  <chr>   <chr>         <dbl>  
1 accuracy binary       0.759  
2 sens    binary       0.617  
3 spec    binary       0.840  
4 roc_auc binary       0.763
```

# Let's practice!

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