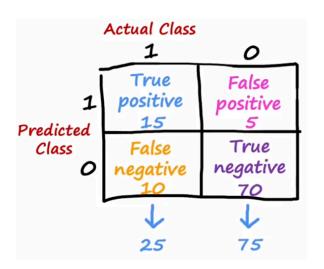
## **Skewed datasets**

## **Error metrics for skewed datasets**

- Traditional accuracy metrics can be misleading when dealing with skewed datasets, as they may not reflect the true performance of the model.
- An example illustrates that a simple algorithm predicting all negatives can achieve high accuracy, but is not useful for diagnosis. Here's a paraphrased version of your sentence:

A dataset is considered **skewed** when there is a significant imbalance between the number of positive and negative examples, meaning one class heavily outnumbers the other.

## Precision/recall



y=1 in presence of rare class we want to detect

**Precision:**(of all patients where we predicted y=1, what fraction actually have the rare disease?)

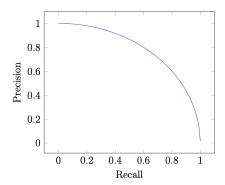
$$\frac{\mathit{TruePositives}}{\mathit{PredictedPositive}} = \frac{\mathit{TruePositives}}{\mathit{TruePos+FalsePos}} = \frac{15}{15+5} = 0.75$$

Recall: (of all patients that actually have the rare disease, what fraction did we correctly detect as having it?)

$$\frac{TruePositives}{ActualPositive} = \frac{TruePositives}{TruePos+FalseNeg} = \frac{15}{15+10} = 0.6$$

## **Trading off precision and recall**

The trade-off between Precision and Recall follows this curve:



When selecting a model, it's important to consider both precision P and recall R. To simplify this process, we introduce the  $F_1-score$ , a metric that combines P and R, defined as:

$$F_1=rac{1}{rac{1}{2}\left(rac{1}{P}+rac{1}{R}
ight)}=rac{2PR}{P+R}.$$

With this measure, we can select the model that achieves the highest  ${\it F_1-score}.$