

Random forests

STAT 4710

November 3, 2022

Where we are

- ✓ **Unit 1:** R for data mining
- ✓ **Unit 2:** Prediction fundamentals
- ✓ **Unit 3:** Regression-based methods
- Unit 4:** Tree-based methods
- Unit 5:** Deep learning

Lecture 1: Growing decision trees

Lecture 2: Tree pruning and bagging

Lecture 3: Random forests

Lecture 4: Boosting

Lecture 5: Unit review and quiz in class

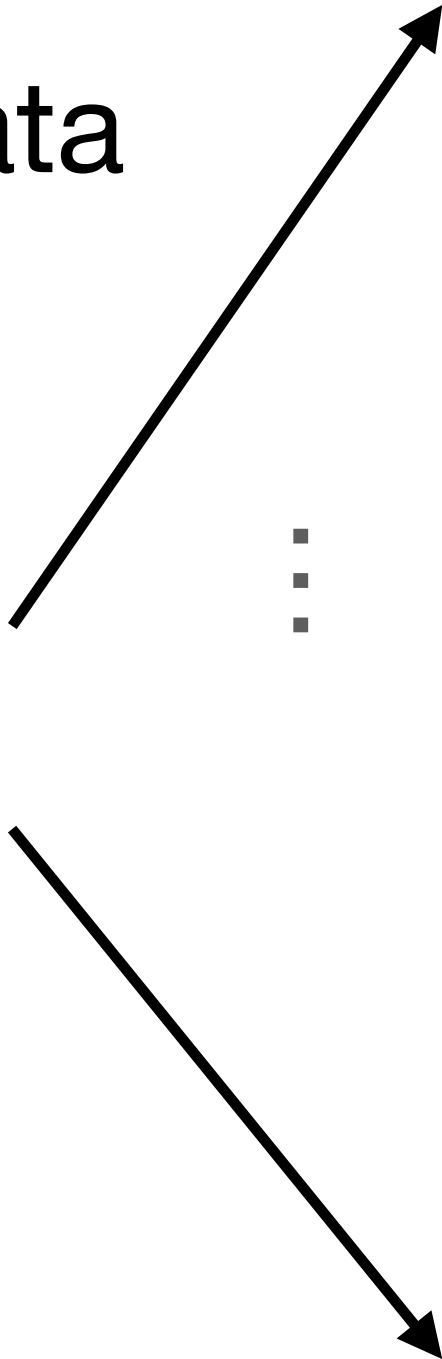
Recall: Bagging

Bootstrap sample 1

Obs ID	X	Y
5	X_5	Y_5
3	X_3	Y_3
2	X_2	Y_2
3	X_3	Y_3
1	X_1	Y_1

Original training data

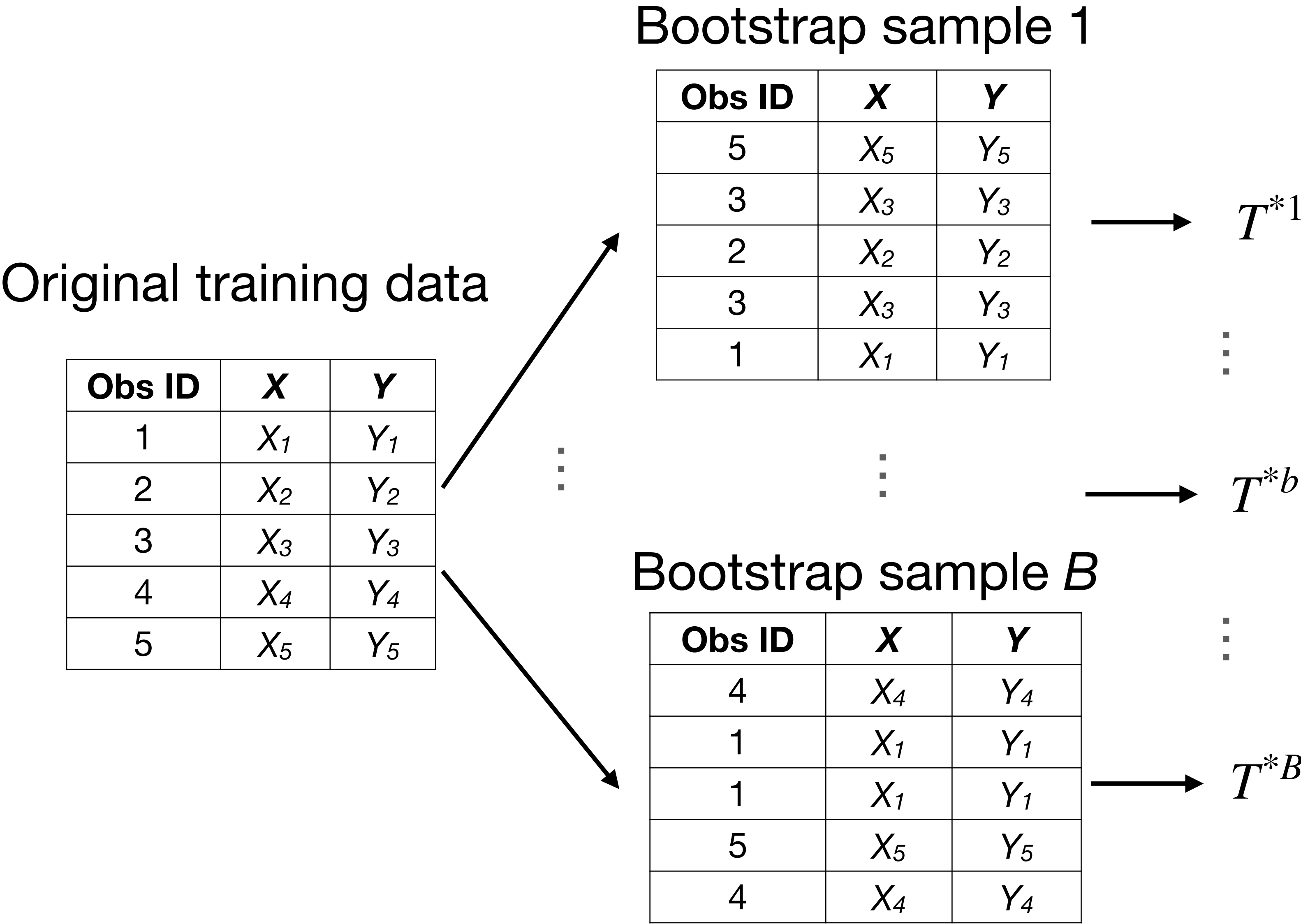
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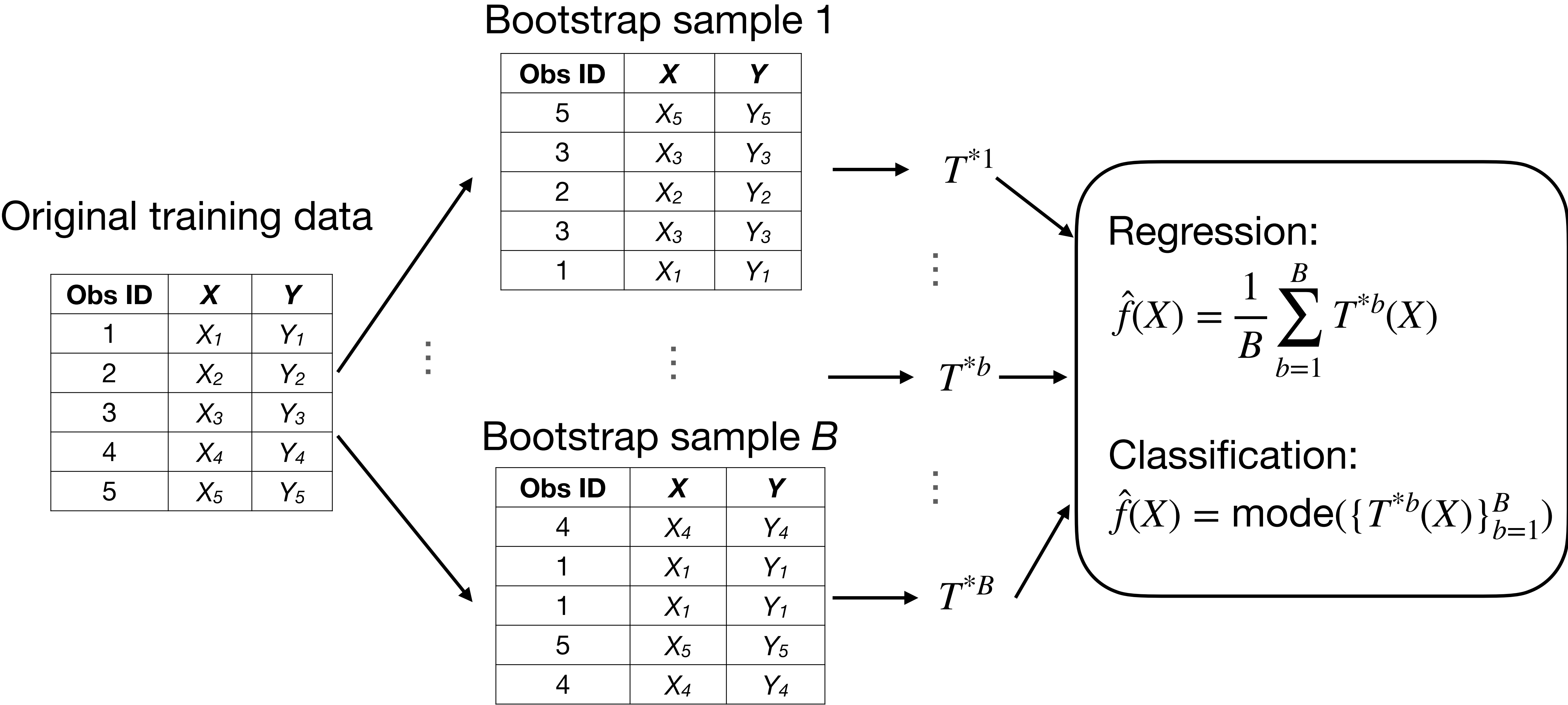
Bootstrap sample B

Obs ID	X	Y
4	X_4	Y_4
1	X_1	Y_1
1	X_1	Y_1
5	X_5	Y_5
4	X_4	Y_4

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$$\text{Var}[\hat{f}(X)] \approx \left(\frac{1}{B} + \frac{B-1}{B} \rho \right) \text{Var}[T(X)] \approx \rho \cdot \text{Var}[T(X)],$$

where $T(X)$ is a single decision tree.

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- The variance is reduced by a factor of $\rho = \text{Corr}[T^{*b_1}(X), T^{*b_2}(X)]$, so the less correlated the bootstrapped trees prediction are, the better.
- As long as B is large enough, the variance reduction is about the same.

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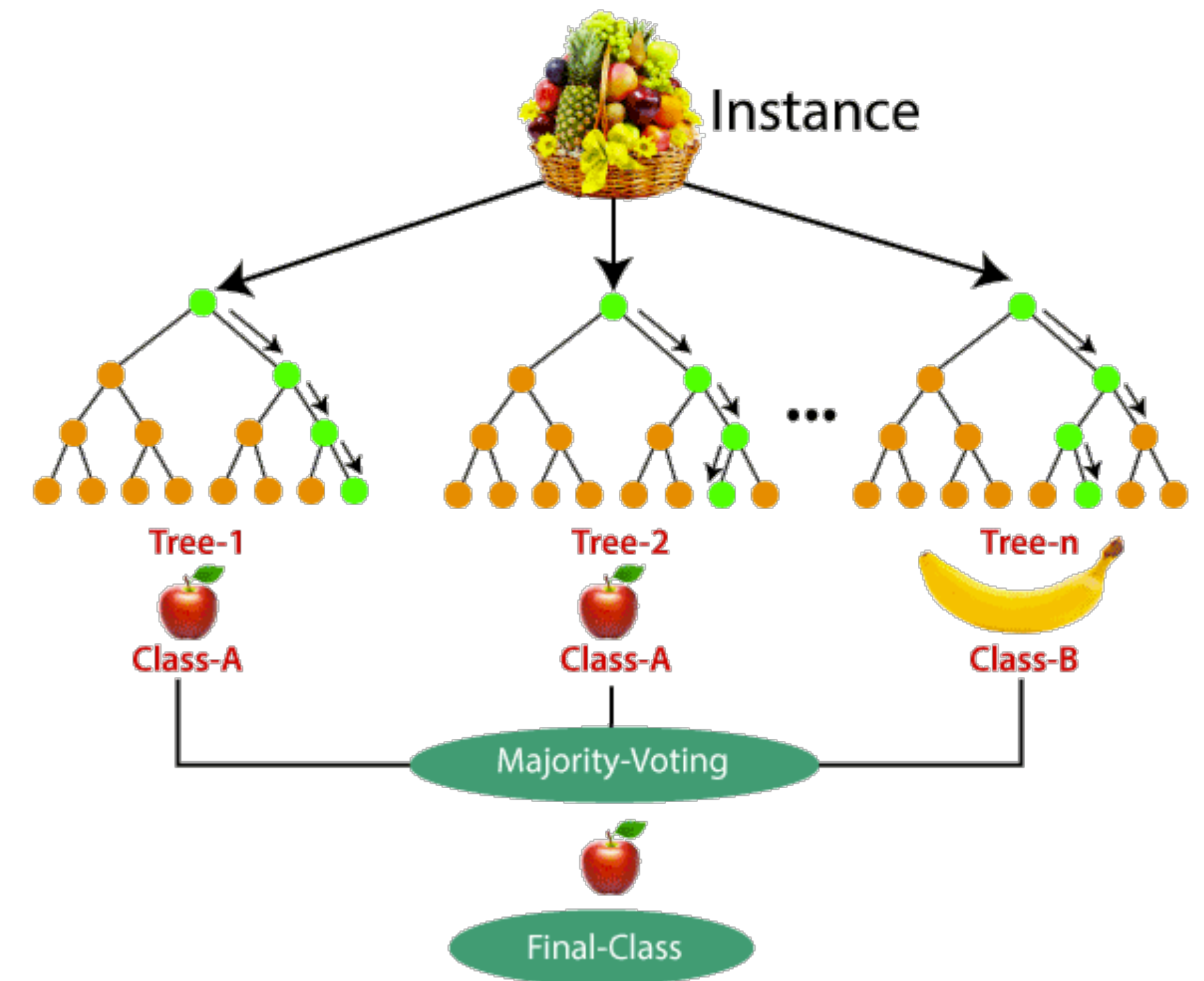


Image source: <https://www.section.io/engineering-education/introduction-to-random-forest-in-machine-learning/>

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Intuition: Sampling features at each split decorrelates the trees, reducing variance and therefore boosting prediction performance.

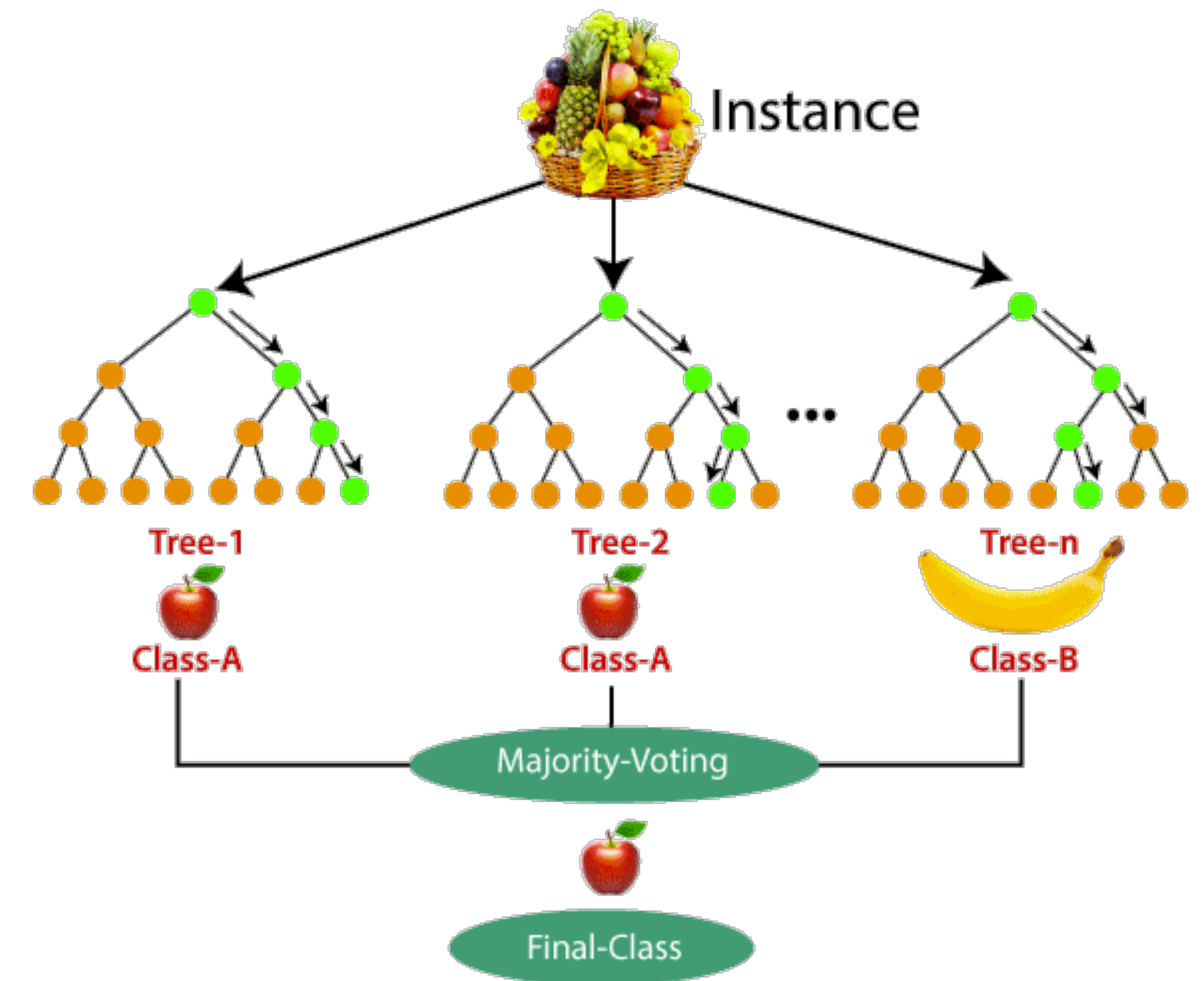


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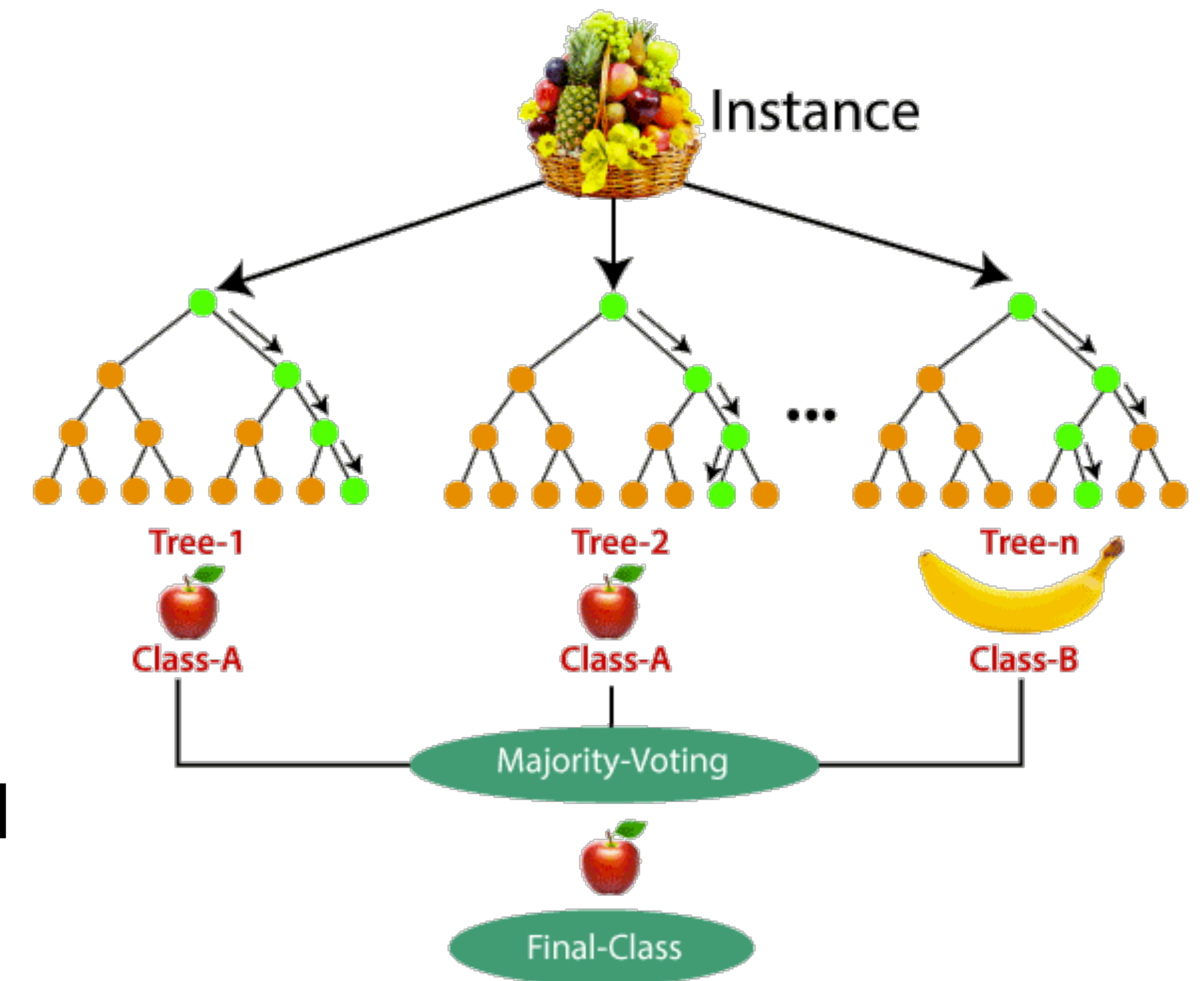


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Note that setting $m = p$ recovers bagging.

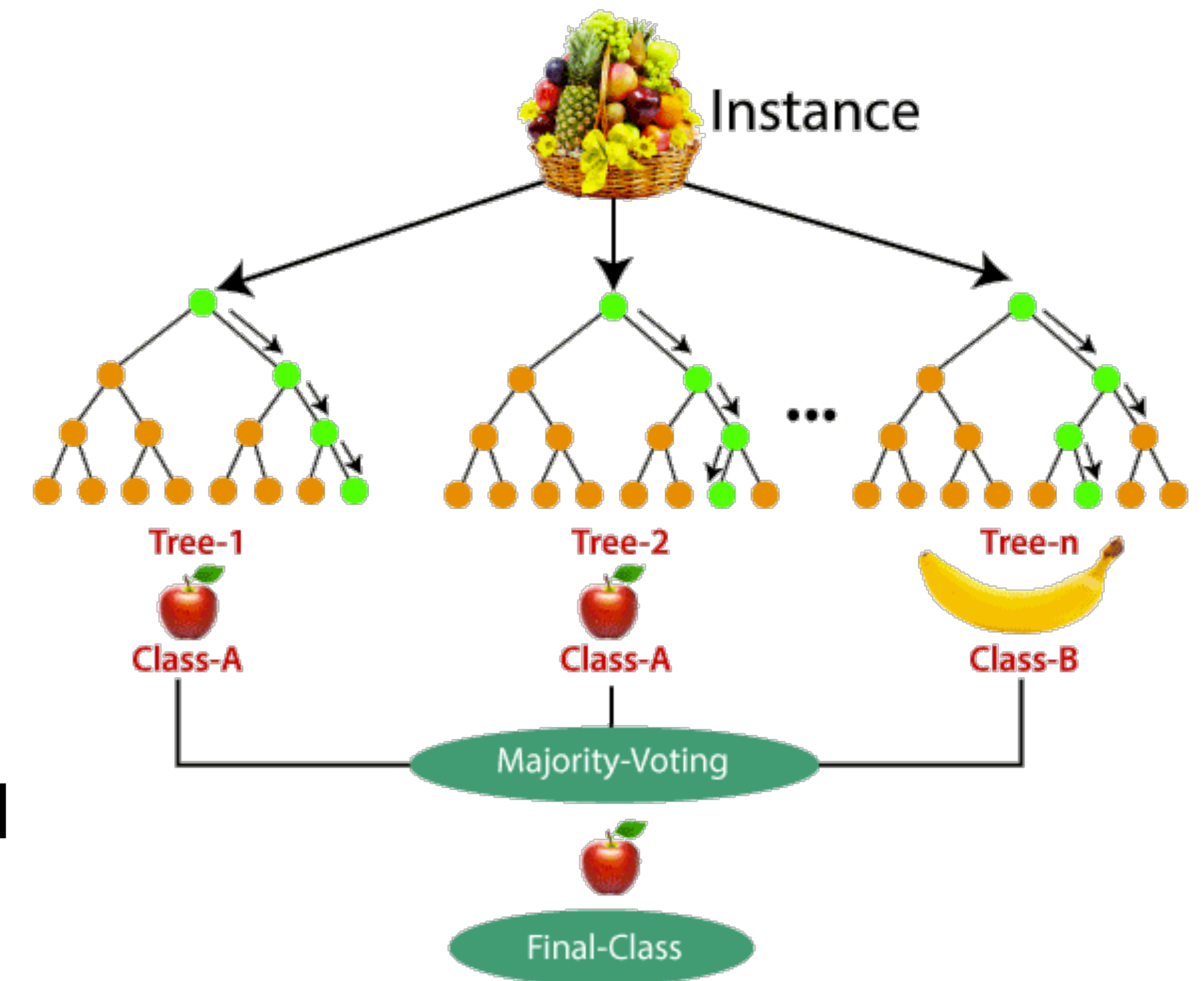


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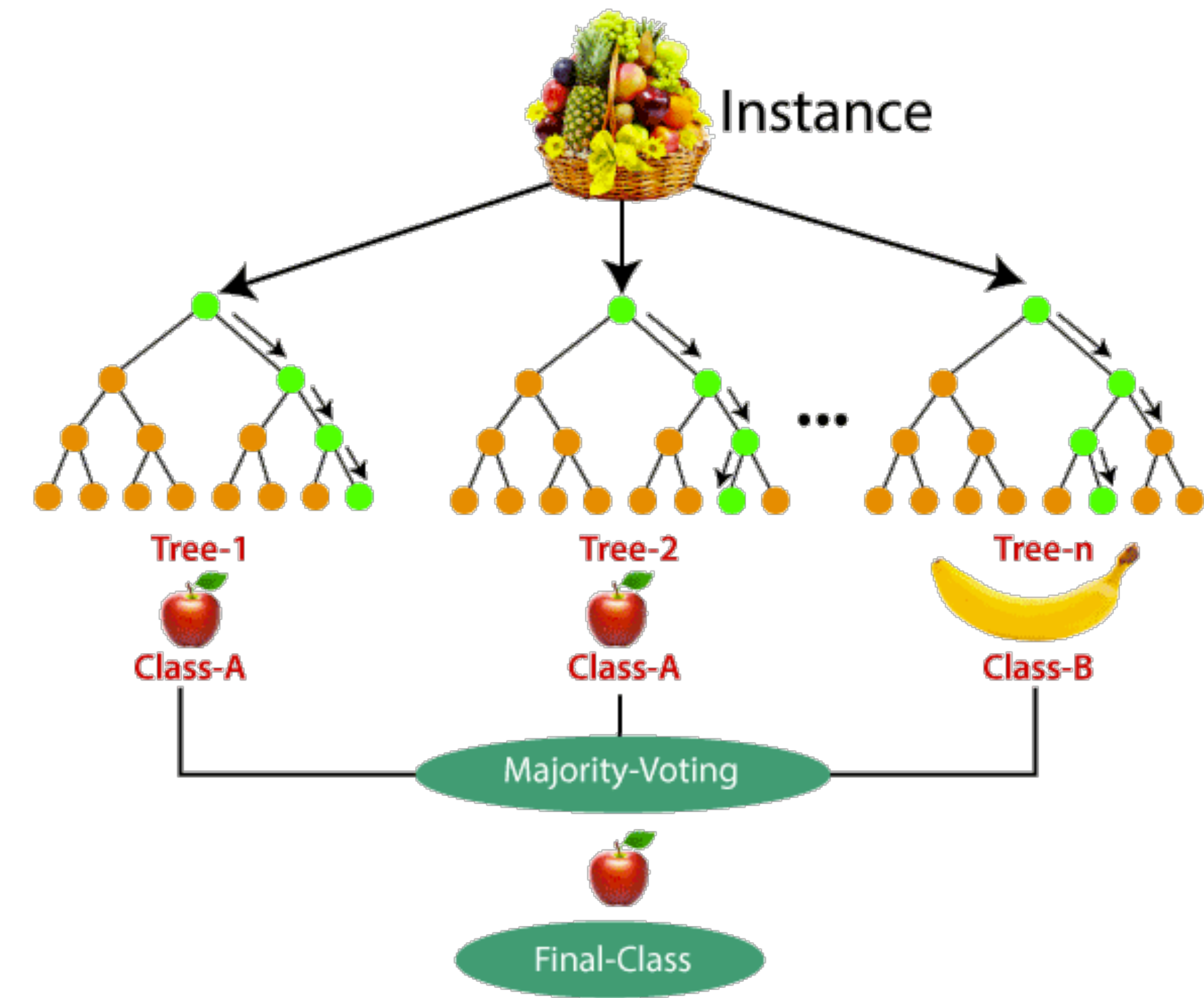


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Random forests

Parameters:

- B : number of bootstrap samples
- m : number of variables to sample at each split
- criterion to stop splitting, like max number of nodes and/or min samples per node

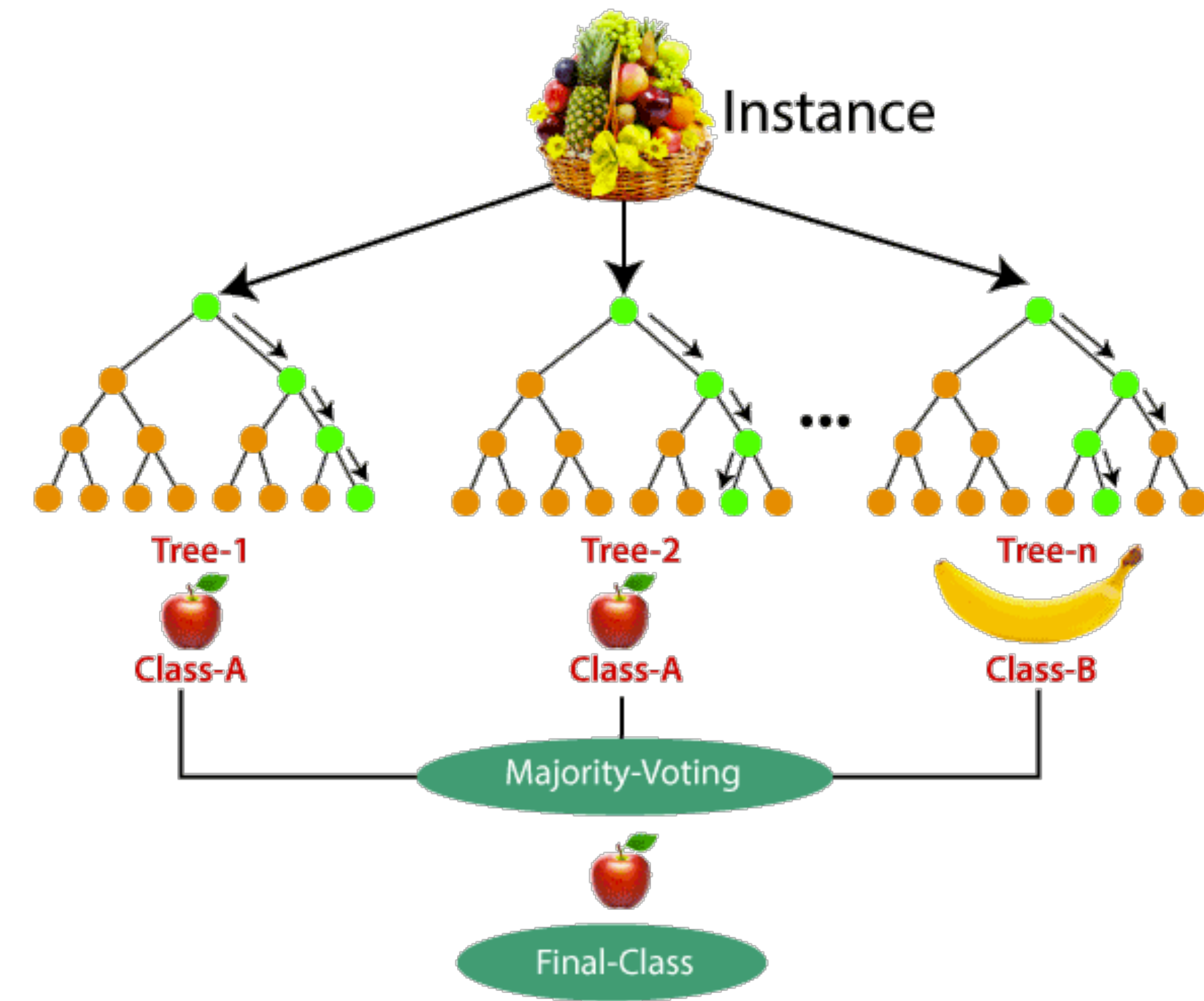


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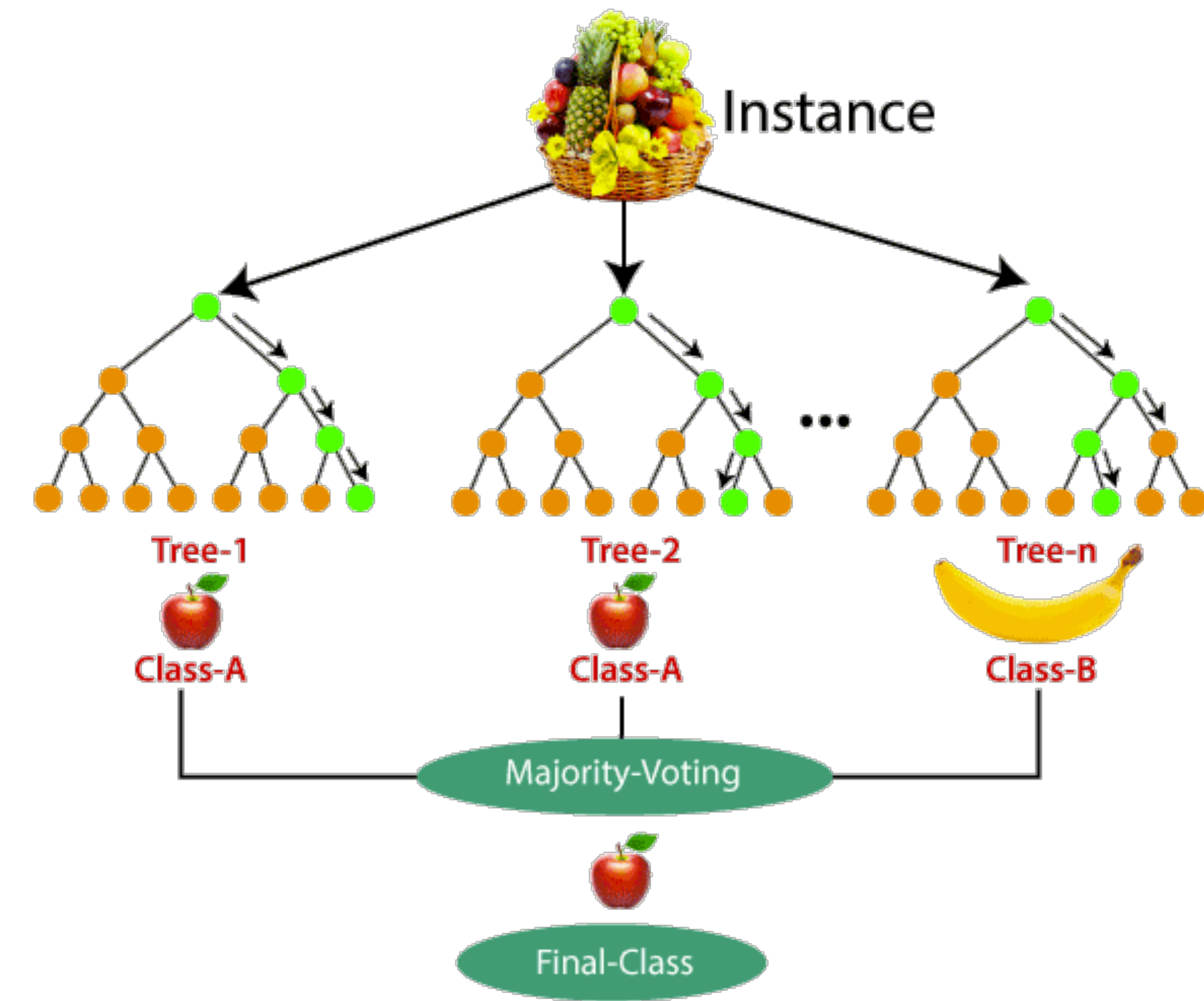


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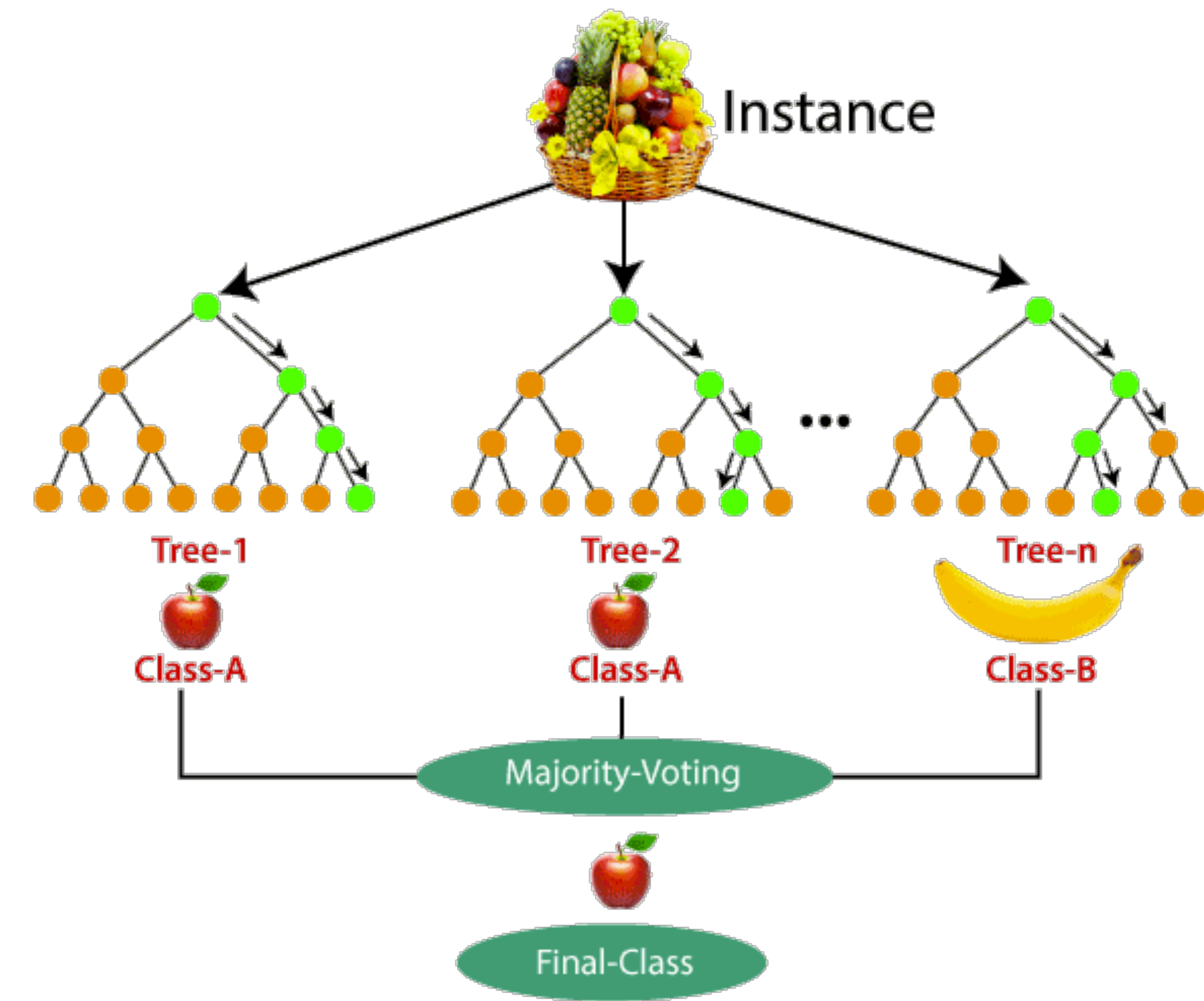


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Prediction:

- aggregate the decision trees using the mean (for regression) or mode (for classification)

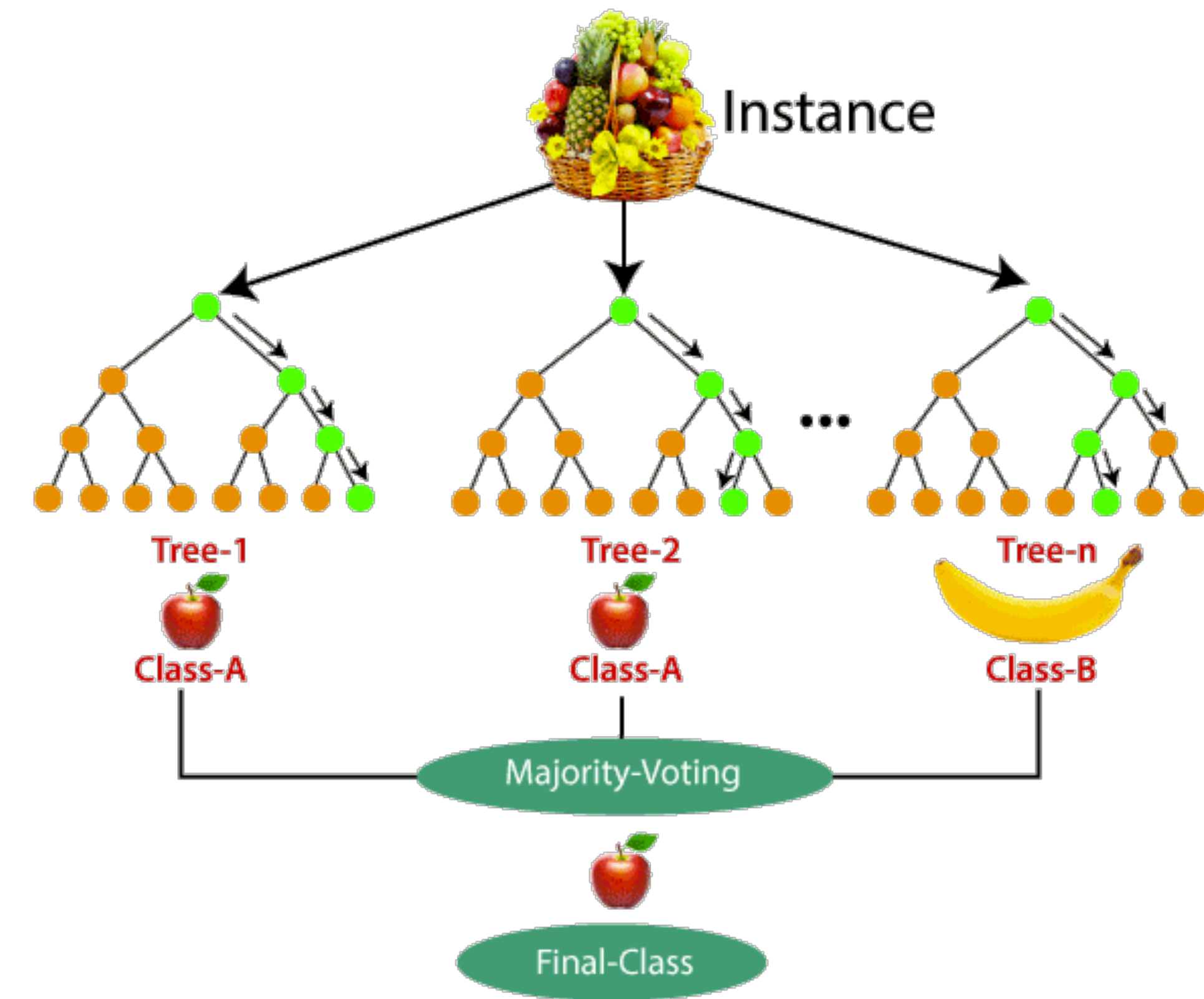


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Default choices: $m = p/3$ for regression and $m = \sqrt{p}$ for classification.

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For best predictive performance, m should be tuned.

Tuning random forests via out-of-bag error

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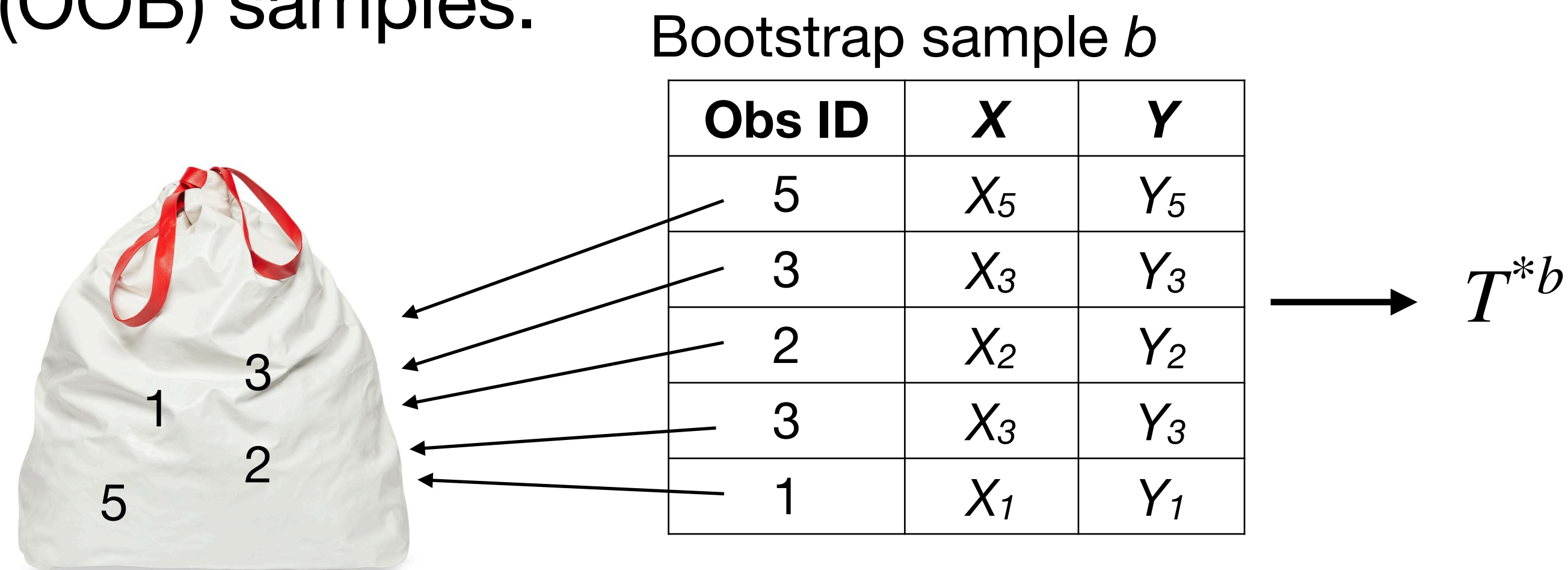
→ T^{*b}

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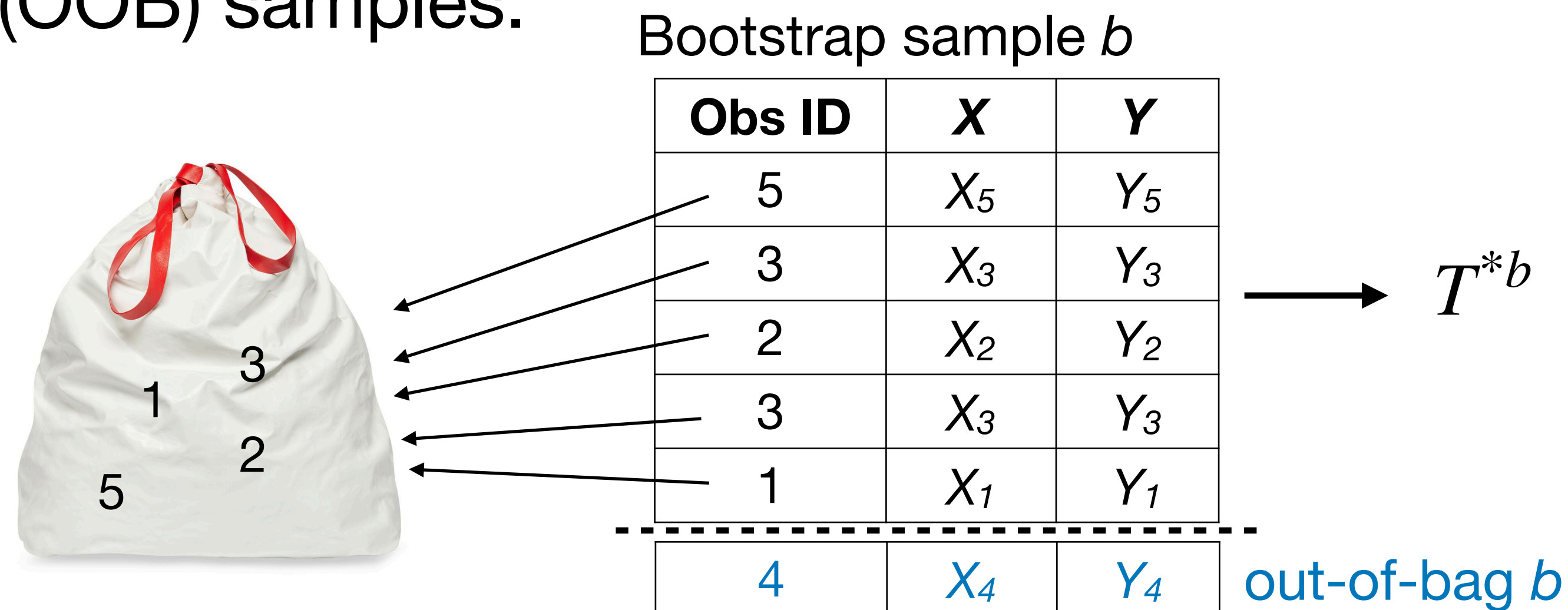


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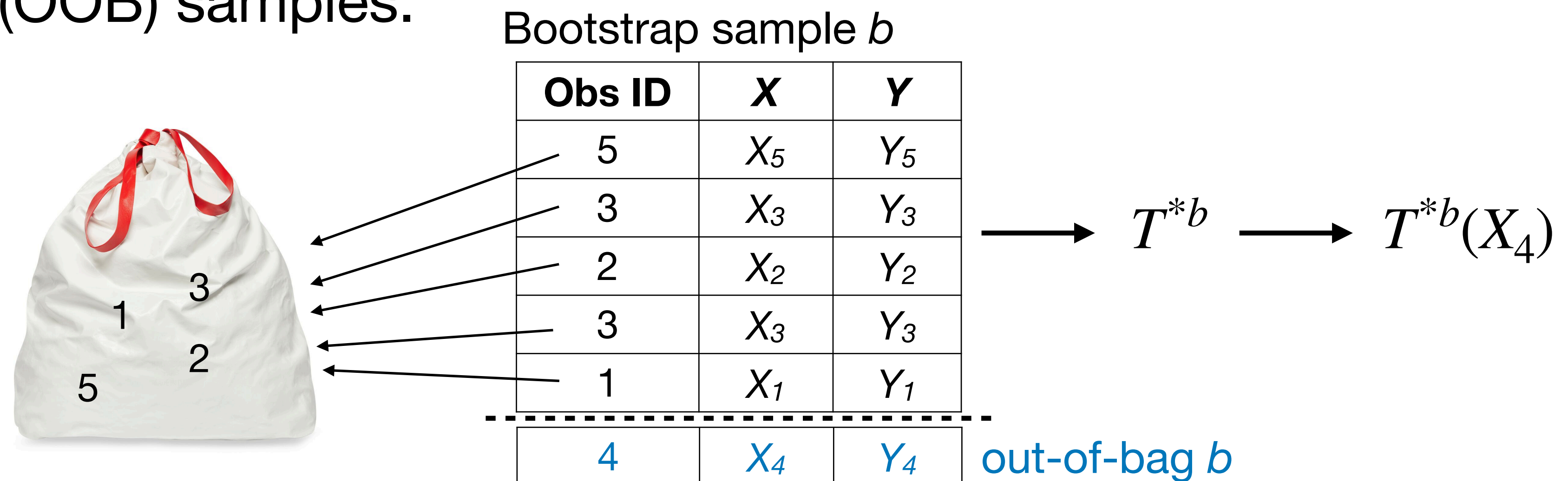


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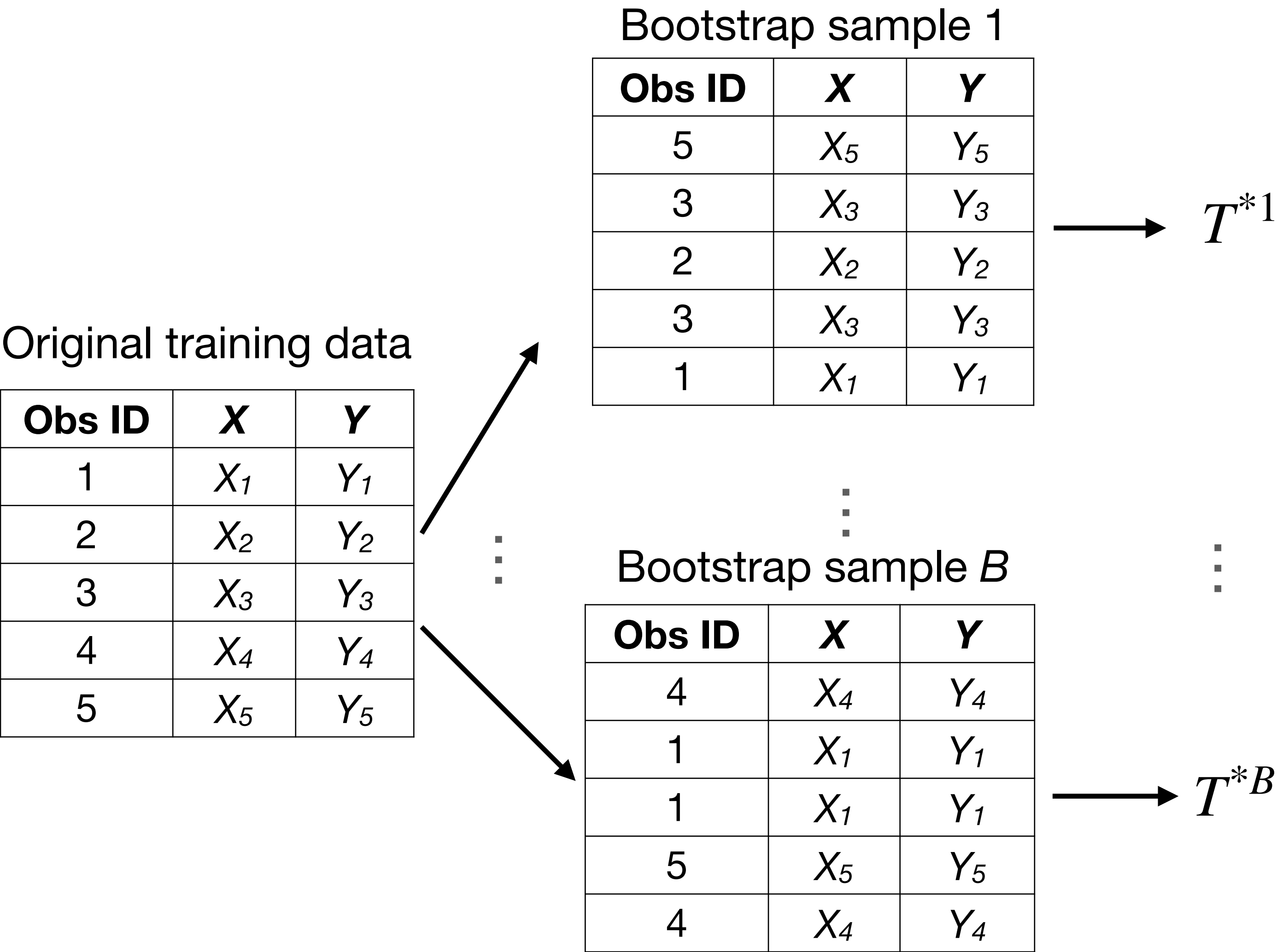
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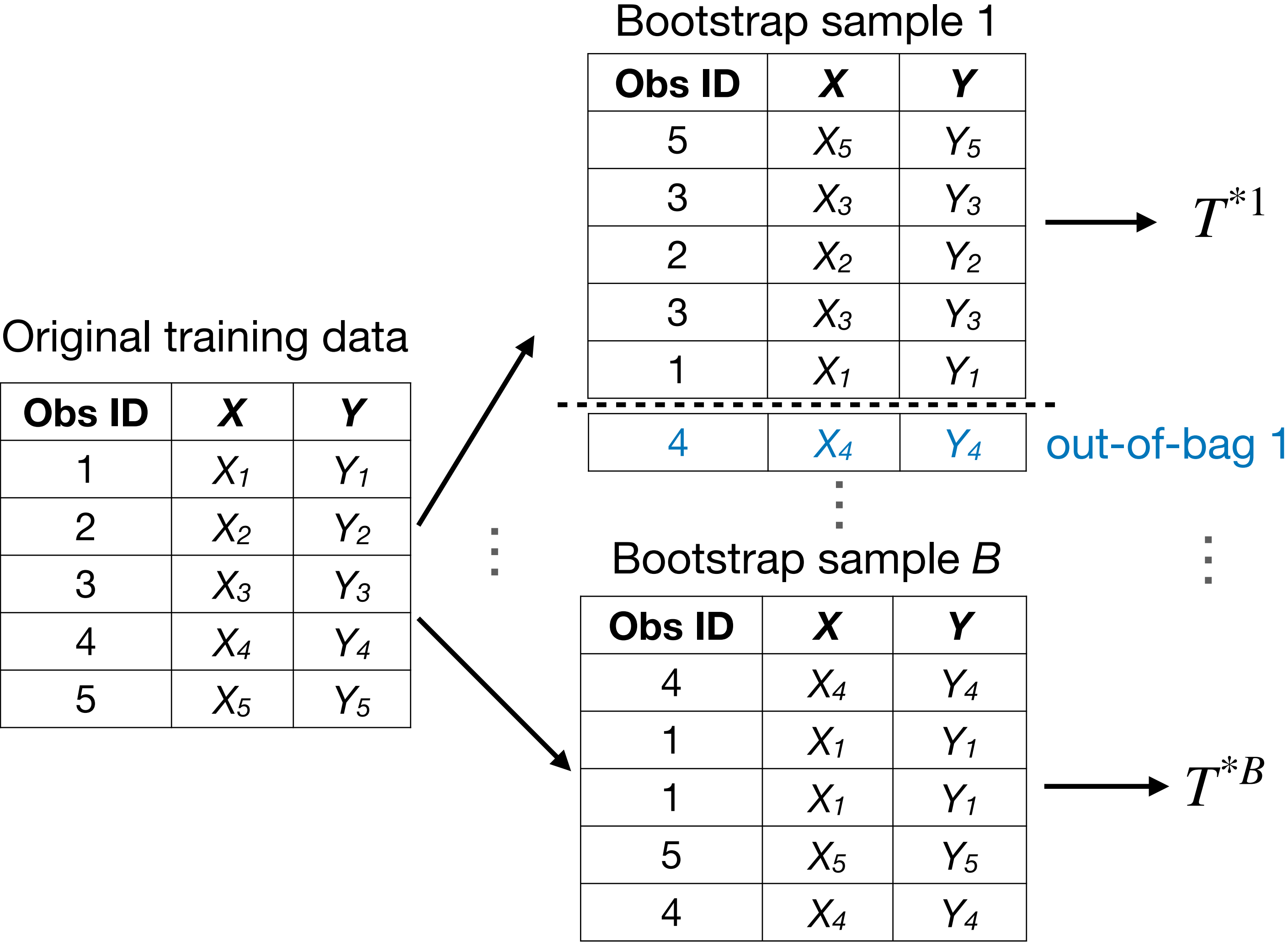
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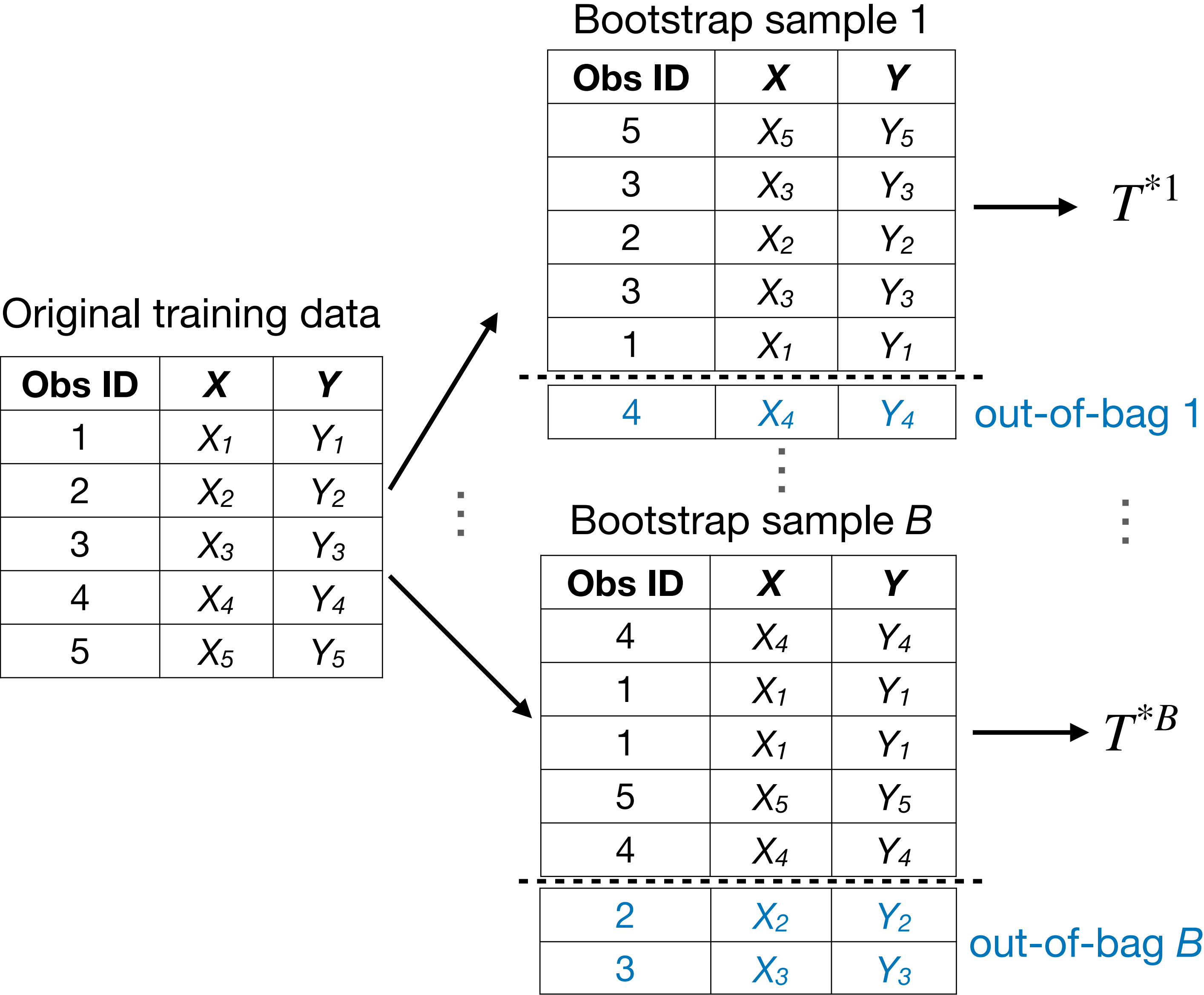
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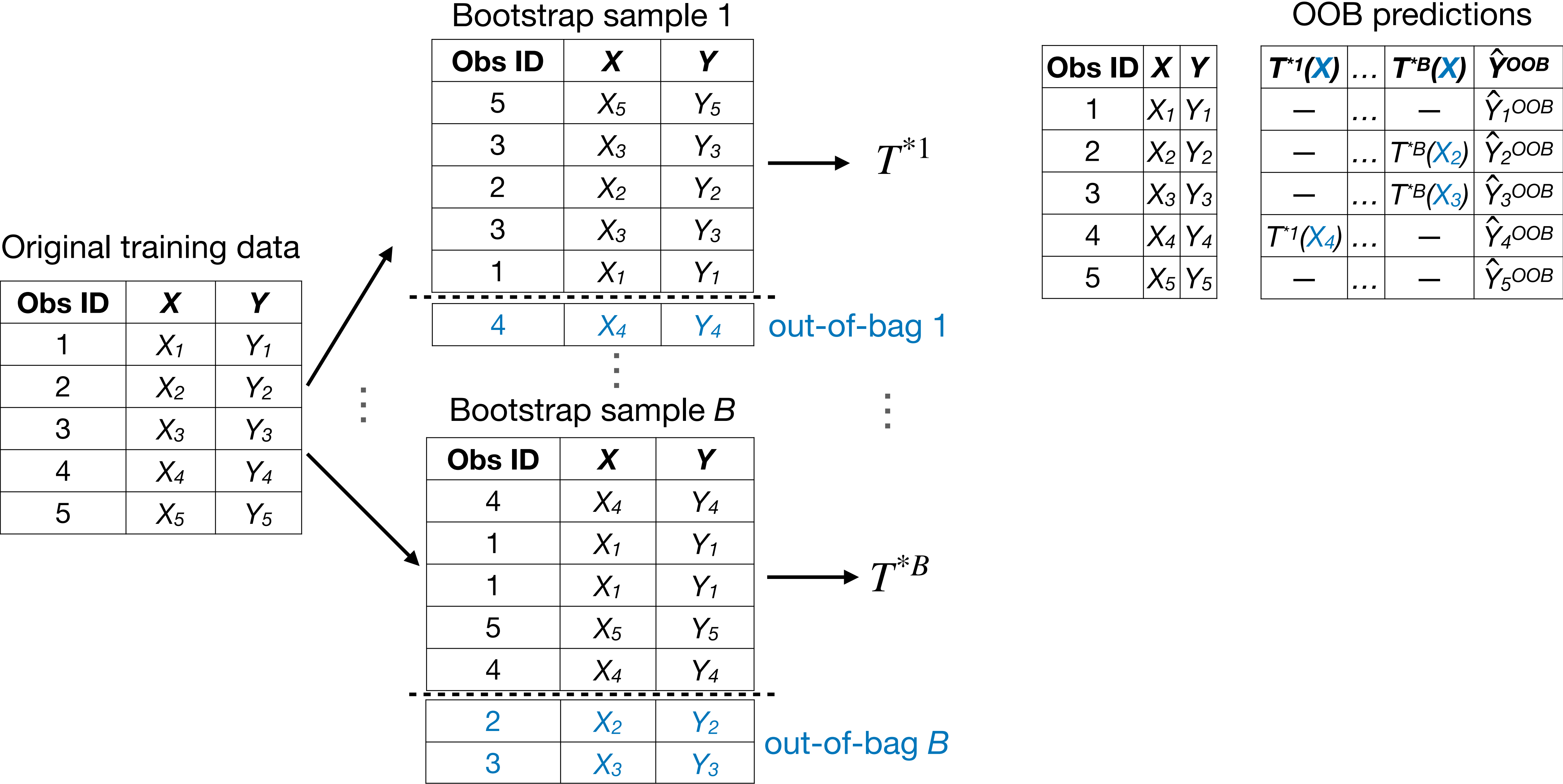
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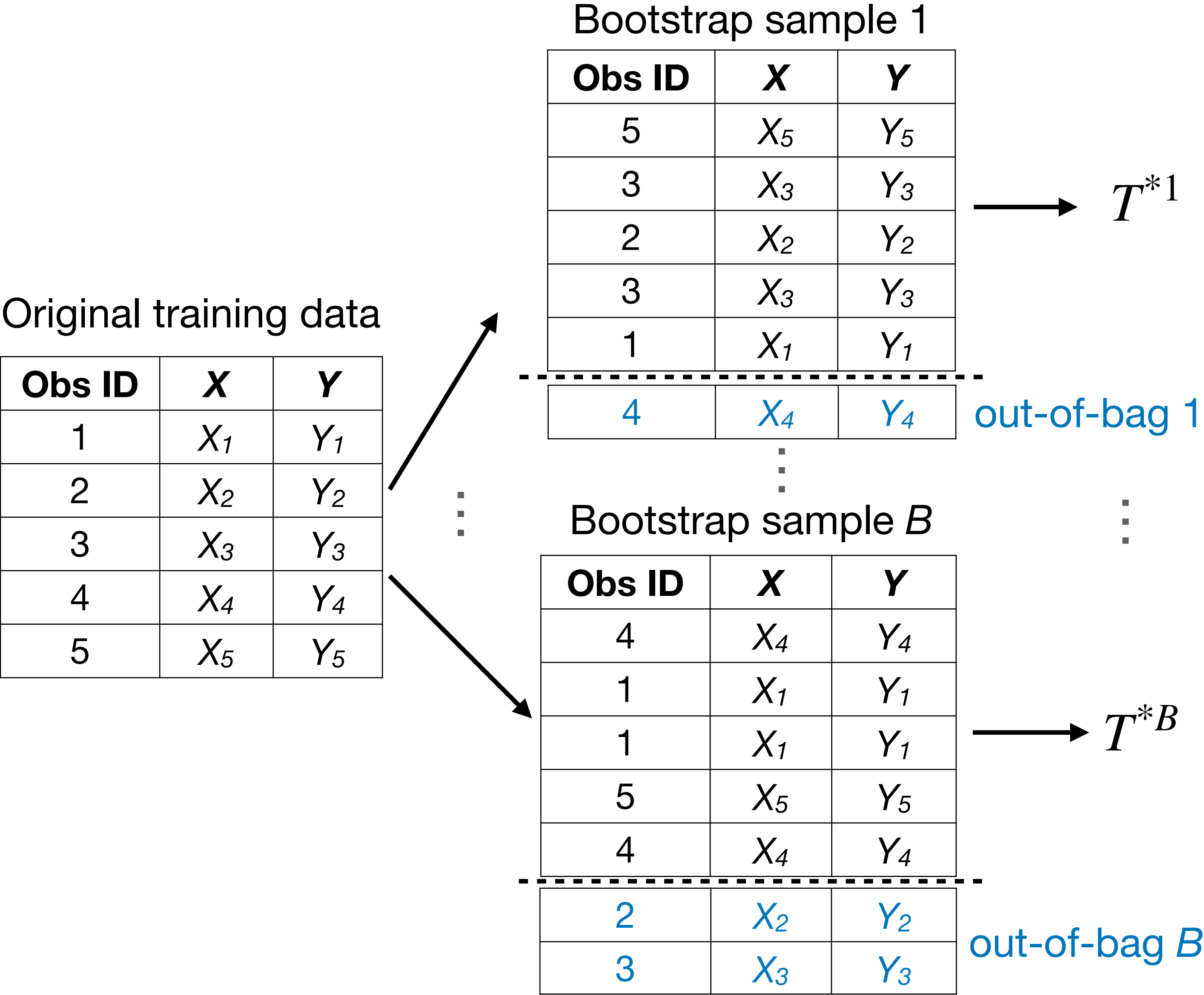
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OOB predictions

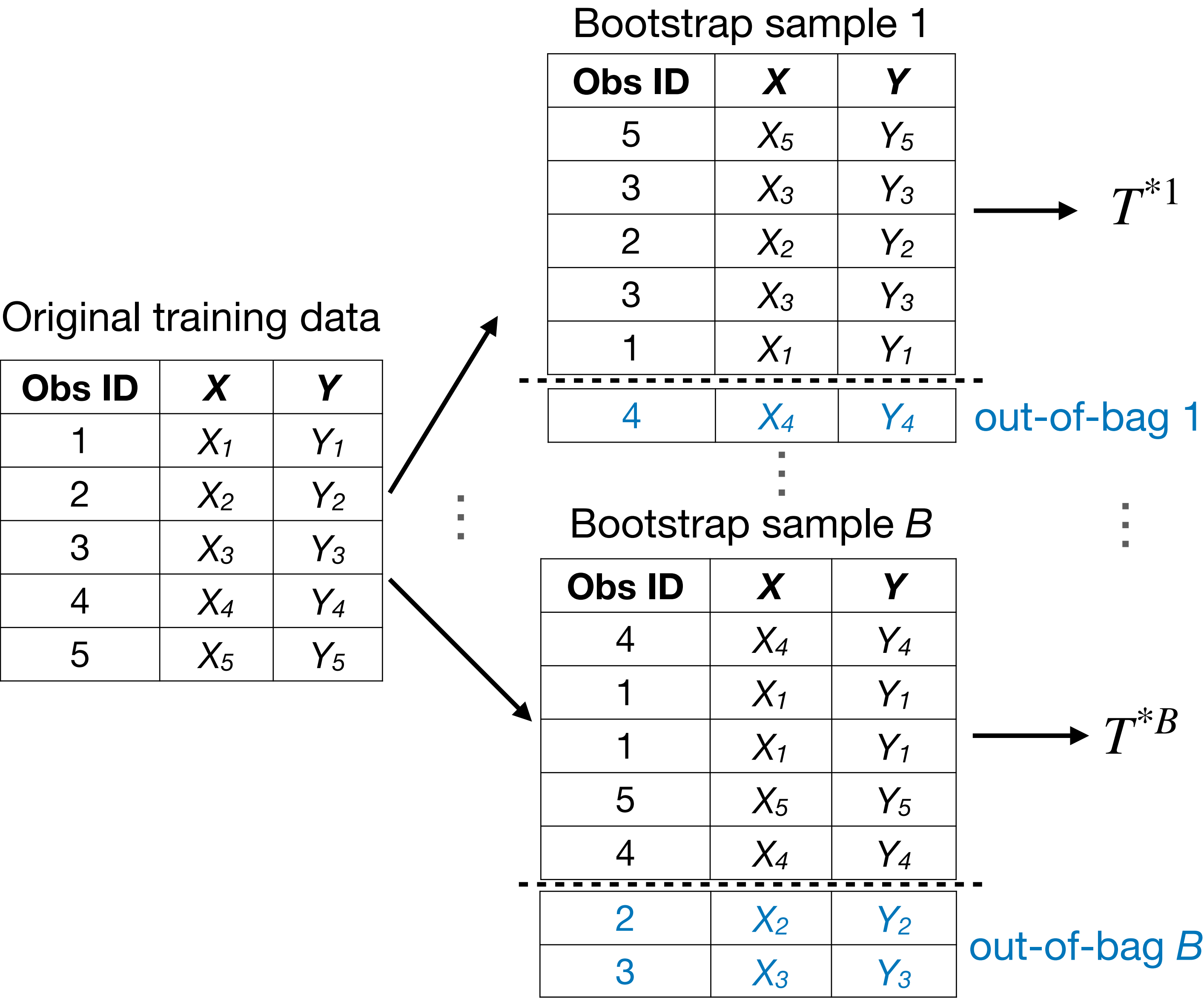
Obs ID	X	Y	$T^{*1}(X)$...	$T^{*B}(X)$	\hat{Y}^{OOB}
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4	X_4	Y_4	$T^{*1}(X_4)$...	—	\hat{Y}_4^{OOB}
5	X_5	Y_5	—	...	—	\hat{Y}_5^{OOB}

Regression:

$$\hat{Y}_i^{OOB} = \text{mean}\{T^{*b}(X_i)\}_{i \in \text{OOB}_b}$$

$$\text{OOB err} = \frac{1}{n} \sum_{i=1}^n (Y_i - \hat{Y}_i^{OOB})^2$$

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Classification:

$$\hat{Y}_i^{OOB} = \text{mode}\{T^{*b}(X_i)\}_{i \in \text{OOB}_b}$$

$$\text{OOB err} = \frac{1}{n} \sum_{i=1}^n I(Y_i \neq \hat{Y}_i^{OOB})$$

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- *m*: most important tuning parameter
- *criteria to stop splitting*: can be tuned but growing trees about as deep as possible generally works pretty well
- *B*: least necessary to tune; just choose a large value like 100-1000.

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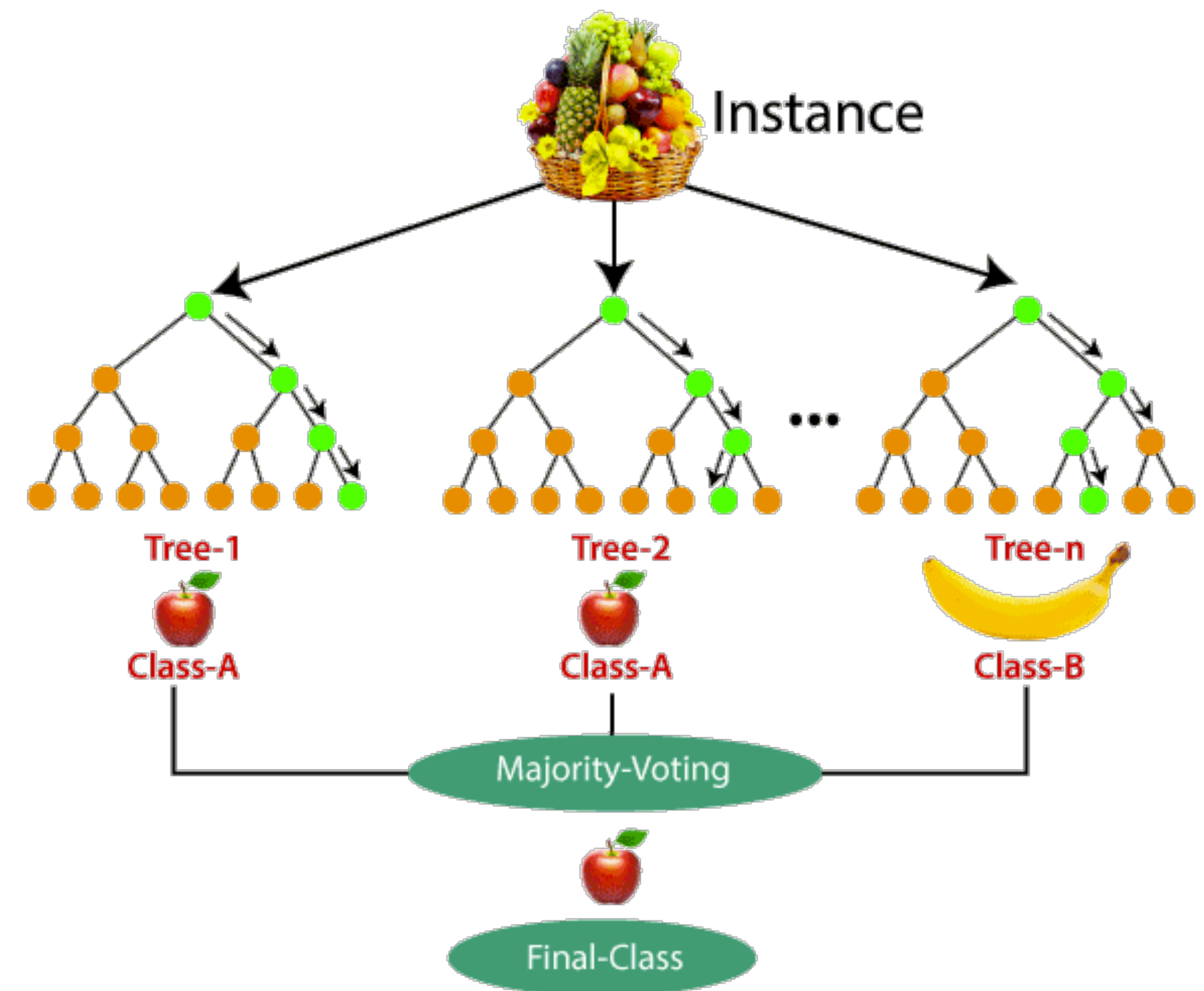
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Two types of variable importance measures are used for random forests:

- purity based importance: how much improvement in node purity results from splitting on a feature
- OOB prediction based importance: how much deterioration in prediction accuracy results from scrambling a feature out of bag

Purity-based variable importance

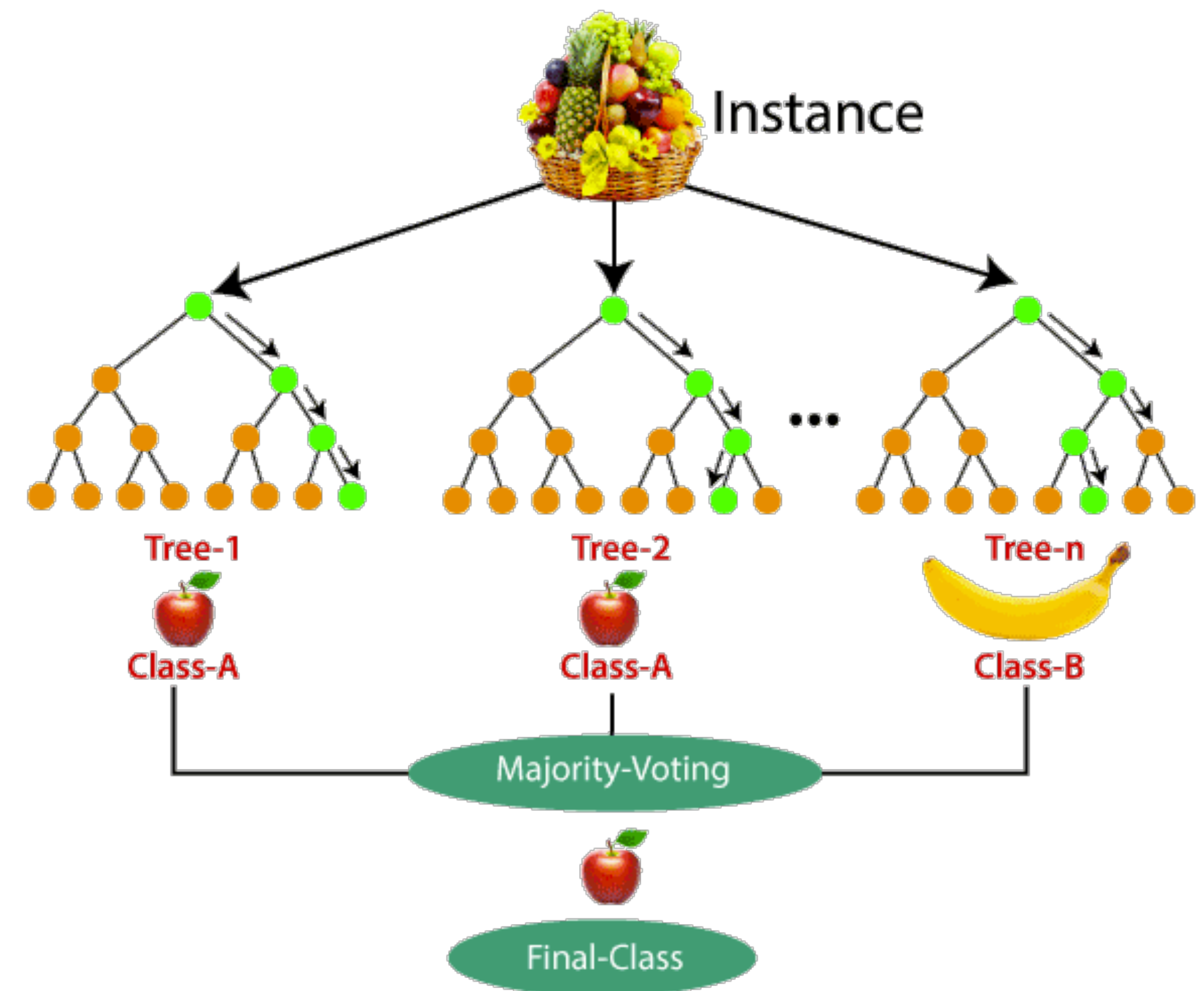
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Purity-based variable importance

Consider the construction of one tree. For each split, note the feature that was split on and resulting reduction in RSS or Gini index (i.e. improvement in purity).

Define the importance of each feature in this single tree by summing up the improvement in purity for all splits based on this feature.

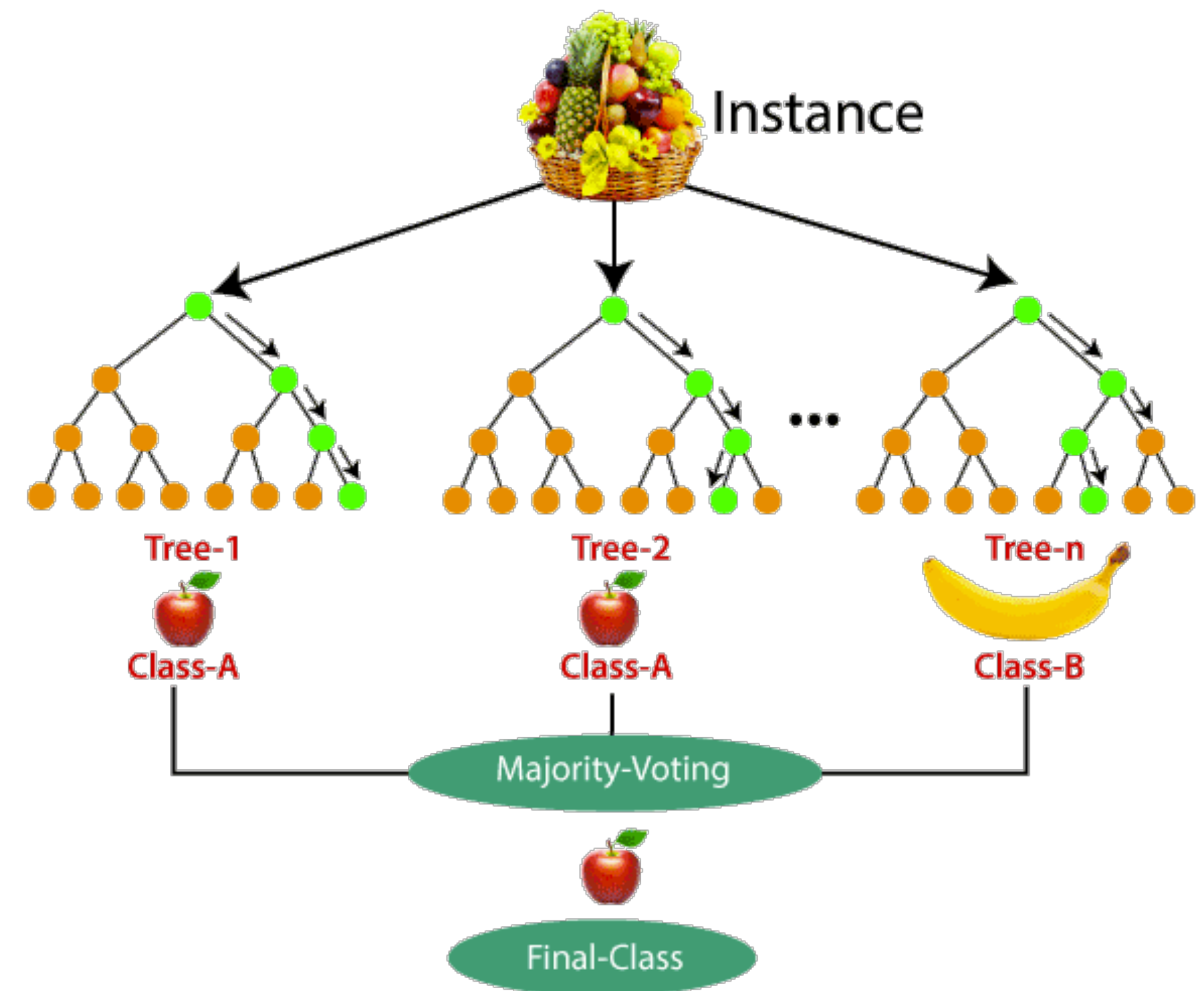


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For random forests, we can average this quantity over all of the trees to get a purity-based variable importance metric.



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X =

X_0	X_1	...	X_j	...	X_{p-1}
12	0		a		1.5
-3	1		b		-0.7
5	0		c		0.2
16	0		d		-3.5
-7	1		e		0.9

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Regular OOB predictions

$T^{*1}(\mathbf{x})$...	$T^{*B}(\mathbf{x})$	\hat{y}^{OOB}
—	...	—	\hat{y}_1^{OOB}
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—	...	$T^{*B}(x_3)$	\hat{y}_3^{OOB}
$T^{*1}(x_4)$...	—	\hat{y}_4^{OOB}
—	...	—	\hat{y}_5^{OOB}

→ Regular OOB error

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OOB prediction based variable importance

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For each feature j and each tree, consider making predictions on the OOB data after first scrambling feature j . We can therefore get a scrambled OOB error.

Regular OOB predictions

$T^{*1}(\mathbf{X})$...	$T^{*B}(\mathbf{X})$	\hat{Y}^{OOB}
—	...	—	\hat{Y}_1^{OOB}
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$T^{*1}(X_4)$...	—	\hat{Y}_4^{OOB}
—	...	—	\hat{Y}_5^{OOB}

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→ scramble

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-3	1		a		-0.7
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$T^{*1}(\mathbf{X})$...	$T^{*B}(\mathbf{X})$	\hat{Y}^{OOB}
—	...	—	\hat{Y}_1^{OOB}
—	...	$T^{*B}(X_2)$	\hat{Y}_2^{OOB}
—	...	$T^{*B}(X_3)$	\hat{Y}_3^{OOB}
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Regular
OOB error

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Scrambled
OOB error

\mathbf{X} =

X_0	X_1	...	X_j	...	X_{p-1}
12	0		a		1.5
-3	1		b		-0.7
5	0		c		0.2
16	0		d		-3.5
-7	1		e		0.9

scramble

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OOB prediction based variable importance

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For each feature j , we can define an OOB-prediction-based variable importance by the difference in OOB error when this feature is scrambled and when it is not.

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scramble →

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Scrambled
OOB error

Var. Imp. = scrambled OOB err - regular OOB err

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Quiz Practice

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