

# Gradient boosting Hyper-parameters

Math behind Gradient Boosting: <https://explained.ai/gradient-boosting/>

	Parameter	Default Values	Description	Impact
Boosting Parameters	learning_rate	0.1	the effect of each tree on the outcome is shrunk by this factor.	> lower always preferred > inversely proportional to n_estimators > use high value for tuning and lower for final submissions
	n_estimators	100	the number of trees to fit sequentially	> tune using CV for a given learning rate > higher value for low learning rate but computationally expensive
	subsample	1	the fraction of observations to be used in individual tree	> typical value 0.8 > reduces variance in model > tune using CV
Tree-Specific Parameters	min_samples_split	2	the minimum number of observations required to split an internal node	> higher values prevent overfitting > risk of underfitting with too high values > tune using CV > ~0.5-2% of total observations
	min_samples_leaf	1	the minimum number of observations required in a terminal node for a split to be valid	> higher values prevent overfitting > risk of underfitting with too high values > tune using CV

	min_weight_fraction_leaf	0	same as min_samples_leaf but defined as a ratio of total number of observations	<ul style="list-style-type: none"> <li>&gt; higher values prevent overfitting</li> <li>&gt; risk of underfitting with too high values</li> <li>&gt; tune using CV</li> </ul>
	max_depth	None	the maximum depth of each tree. None specified no limit on depth.	<ul style="list-style-type: none"> <li>&gt; lower values prevent overfitting</li> <li>&gt; risk of underfitting with too low values</li> <li>&gt; tune using CV</li> <li>&gt; typical 5-20</li> </ul>
	max_leaf_nodes	None	the maximum number of terminal nodes in each tree. If None then no limit	<ul style="list-style-type: none"> <li>&gt; lower values prevent overfitting</li> <li>&gt; risk of underfitting with too low values</li> <li>&gt; tune using CV</li> </ul>
	max_features	None	the number of features to consider for each split	<ul style="list-style-type: none"> <li>&gt; typical sqrt to 30-40% of total features</li> <li>&gt; possible values: <ul style="list-style-type: none"> <li>- Int: actual number of features</li> <li>- Float: the fraction of total number of features</li> <li>- 'auto' or 'sqrt': square root of total features</li> <li>- 'log2': log to base 2 of total features</li> <li>- None: equal to total number of features</li> </ul> </li> </ul>
Miscellaneous Parameters	loss	deviance (class) ls (regression)	the cost function to be minimized by optimization	<ul style="list-style-type: none"> <li>&gt; use default value if not sure</li> <li>&gt; options: <ul style="list-style-type: none"> <li>- class: deviance / exponential</li> <li>- regr: ls / lad / huber / quantile</li> </ul> </li> </ul>
	init	None	estimator for getting the initial predictions	<ul style="list-style-type: none"> <li>&gt; BaseEstimator can be any class object with fit and predict functions</li> </ul>
	random_state	None	set the seed of random number generator	<ul style="list-style-type: none"> <li>&gt; use fixed value for parameter tuning</li> <li>&gt; try different values only if possible computationally</li> </ul>

	verbose	0	select the type of output to be generated by model fit	>options: - 0: No output - 1: output for few trees (selected automatically) - >1: output all trees
	warm_start	False	if True, it will start from the previous call to fit and add more trees to it	> should be used judiciously for advanced applications
	presort	'auto'	whether to presort data to speed up the finding of best splits	> mostly 'auto' works fine



# XGBoost Hyper-parameters

	Parameter	Default Values	Decription & Impact
<b>Boosting Parameters</b>	eta/learning_rate	0.3	<ul style="list-style-type: none"> <li>&gt; Makes the model more robust by shrinking the weights on each step</li> <li>&gt; Typical final values to be used: 0.01-0.2 (0.015,0.025,0.01,0.05,0.1..)</li> </ul>
	min_child_weight	1	<ul style="list-style-type: none"> <li>&gt; Used to control over-fitting. Higher values prevent a model from learning relations which might be highly specific to the particular sample selected for a tree.</li> <li>&gt; Too high values can lead to under-fitting hence, it should be tuned using CV.</li> <li>&gt; Typical Value Range - (1, 3, 5, 7....15)</li> </ul>
	max_depth	6	<ul style="list-style-type: none"> <li>&gt; The maximum depth of a tree, same as GBM.</li> <li>&gt; Used to control over-fitting as higher depth will allow model to learn relations very specific to a particular sample.</li> <li>&gt; Should be tuned using CV. Typical values: 3-10.</li> </ul>
	gamma/min_split_loss	0	<ul style="list-style-type: none"> <li>&gt; Minimum loss reduction required to make a further partition on a leaf node of the tree.</li> <li>&gt; The larger gamma is, the more conservative the algorithm will be.</li> <li>&gt; Typical range: 0.05,0.1,0.3,0.5,0.7,0.9.1</li> </ul>
	subsample	1	<ul style="list-style-type: none"> <li>&gt; Lower values make the algorithm more conservative and prevents overfitting but too small values might lead to under-fitting.</li> <li>&gt; Typical values: 0.5-1</li> </ul>
	colsample_bytree	1	<ul style="list-style-type: none"> <li>&gt; Similar to max_features in GBM. Denotes the fraction of columns to be randomly samples for each tree.</li> <li>&gt; Typical values: 0.5-1</li> </ul>
	lambda/reg_lambda	1	<ul style="list-style-type: none"> <li>&gt; L2 regularization term on weights. Increasing this value will make model more conservative.</li> <li>&gt; Typical range: 0.01-0.1/ 1.0</li> </ul>
	alpha/reg_alpha	0	<ul style="list-style-type: none"> <li>&gt; L1 regularization term on weights. Increasing this value will make model more conservative.</li> <li>&gt; Typical range: 0.01-0.1/ 1.0</li> </ul>

	scale_pos_weight	1	<ul style="list-style-type: none"><li>&gt; Control the balance of positive and negative weights, useful for unbalanced classes.</li><li>&gt; A typical value to consider: <math>\text{sum}(\text{negative instances}) / \text{sum}(\text{positive instances})</math>.</li></ul>
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