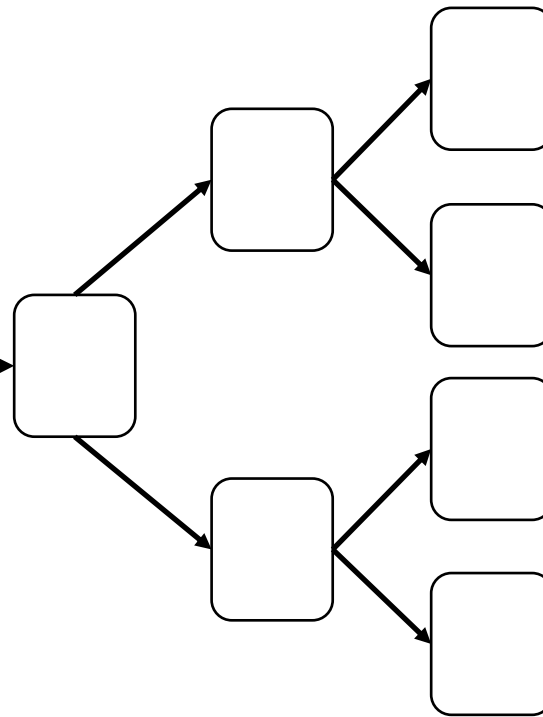


X, Y

Single Cell RNA-Seq Dataset
 X = Gene expression
 Y = Tissue types



`Sklearn.tree.DecisionTreeClassifier`

$M(X)$

$M(X)$ = Tree model for X