

# Title: AutoML with Monte Carlo Tree Search and Neural Network

Author: Shaohang Chen

Supervisor : Yiran Huang

CHAIR FOR PERVASIVE COMPUTER SYSTEMS, INSTITUTE OF TELEMATICS, DEPARTMENT OF COMPUTER SCIENCE



# 0. Design

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- AutoML
- Monte Carlo Tree Search
- MCTML Structure
- MCTML Details
- Evaluation
- Future Work

# 1. AutoML

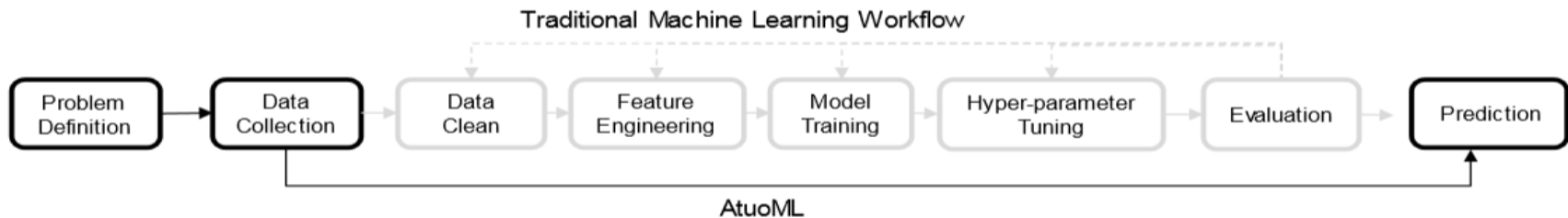


Figure 1.1: Traditional machine learning workflow and AutoML

Task: Algorithm selection and hyperparameter selection --- black box optimization[1]

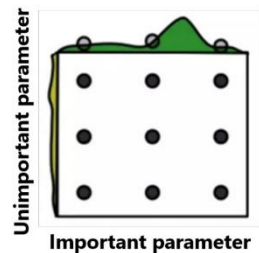


Figure 1.2: Grid Search

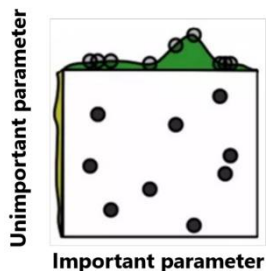


Figure 1.3: Random Search

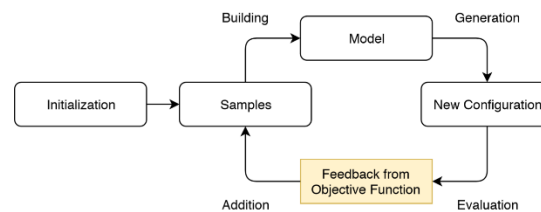


Figure 1.4: Bayesian Optimization

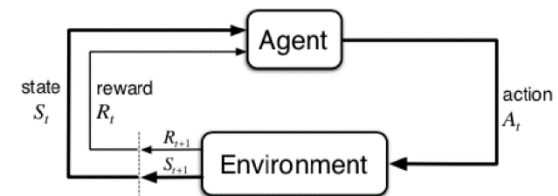


Figure 1.5: Reinforcement Learning

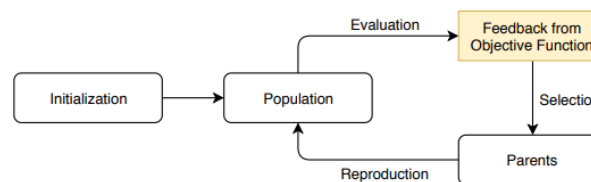


Figure 1.5: Genetic Algorithm

Method	Tool
Bayesian Optimization	Autosklearn[2]
Genetic Algorithm	TPOT[3]
Reinforcement Learning	—

Table 1: AutoML methods and tools

[1] Automated Machine Learning: Methods, Systems, Challenges, Hutter, Frank, 2018

[2] Efficient and Robust Automated Machine Learning, Matthias Feurer, Aaron Klein, Katharina Eggensperger, 2015

[3] TPOT: A Tree-Based Pipeline Optimization Tool for Automating Machine Learning, Randal S. Olson, et al, 2019

## 2. Monte Carlo Tree Search

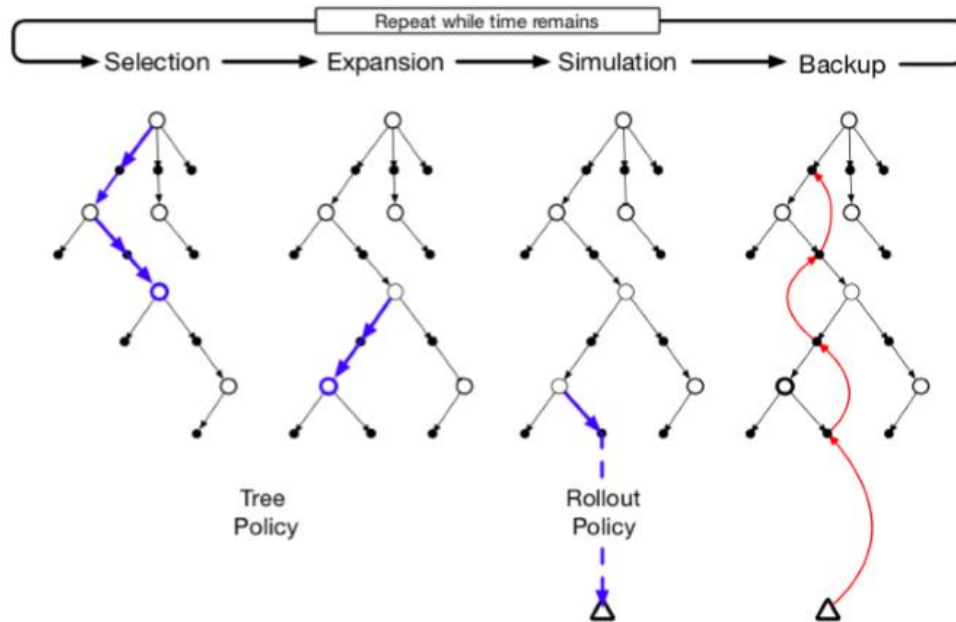


Figure 2.1: Outline of Monte Carlo Tree Search[1]

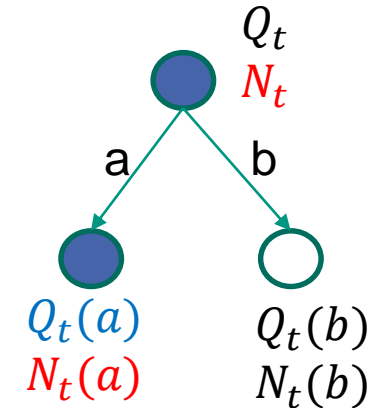


Figure 2.2: Illustration of parent node and children nodes

$$A_t \doteq \operatorname{argmax}_a [Q_t(a) + c \sqrt{\frac{N_t}{N_t(a)}}]$$

Equation 1: Upper Confidence Bound Algorithm.

**Selection:** traverses and selects the best-scored child node(UCB algorithm)

**Expansion:** a new child node is added to the tree

**Simulation:** choosing moves until a leaf node is achieved

**Backup:** backprogress the value to each node in trajectory.

[1] *Mastering the game of Go with deep neural networks and tree search*, David Silver, Aja Huang, 2016

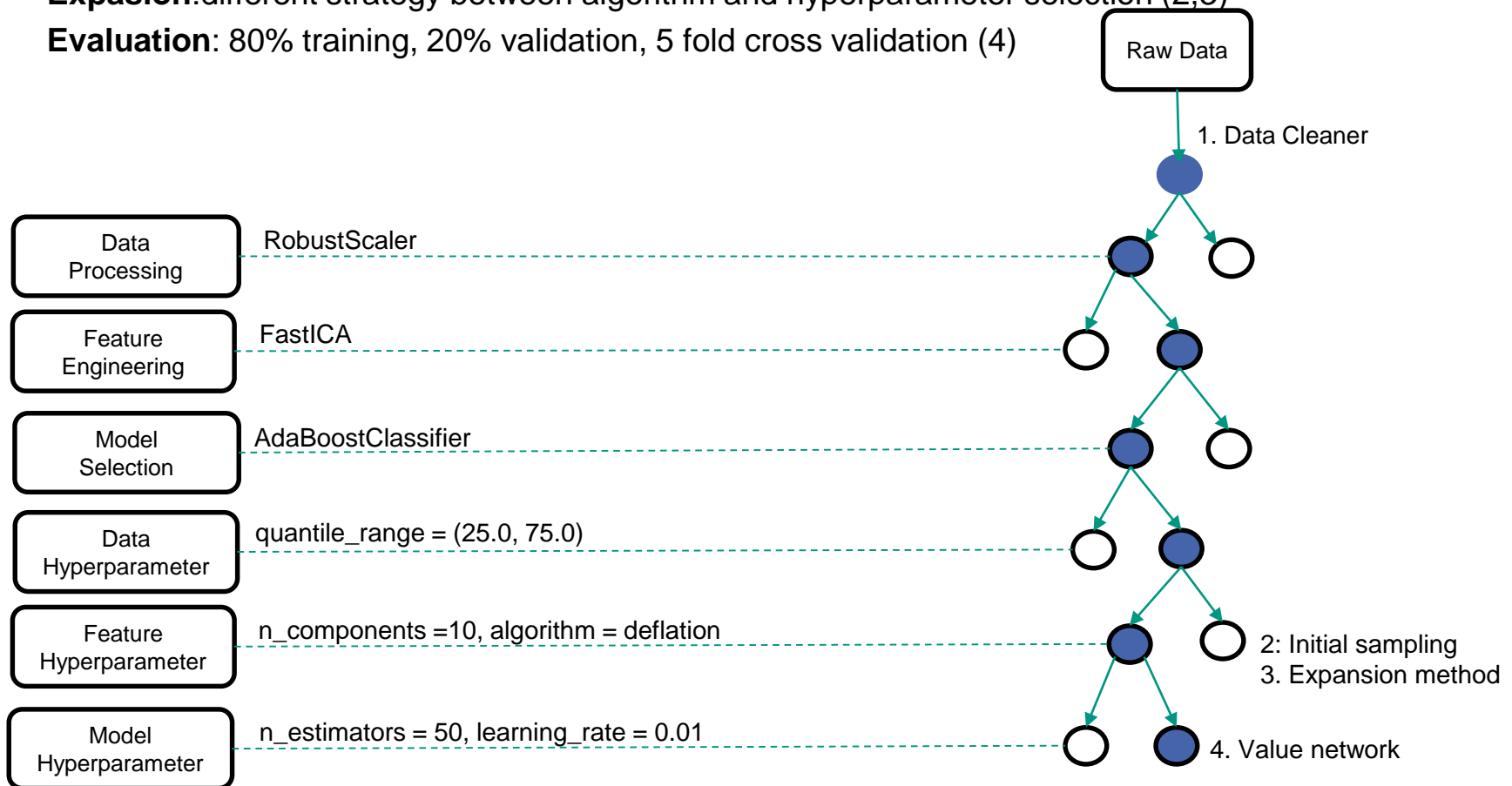
# 3.1. MCTML

## ■ Structure

### Data Cleaning (1)

**Expansion:** different strategy between algorithm and hyperparameter selection (2,3)

**Evaluation:** 80% training, 20% validation, 5 fold cross validation (4)



## 3.2. Detail of MCTML(1)

### ■ 1. Data cleaner

- missing value imputation: NaN, ?, -
- conver non-numerical feature to numerical feature

### ■ 2. Initial sampling

hyperparameter name	value range	sampled values
n_estimators	(50,500)	[50,100,200,300,400,500]
learning_rate	(0.01,2.0)	[0.01,0.05,0.1,0.5,1.0,1.5,2.0]

*Table 2 : Example of hyper-parameter samples using initial state sampling method*

## 3.3. Detail of MCTML(2)

### 3. Expansion algorithm

New parameter:  $K = \lceil C_{pw} N^\alpha \rceil$

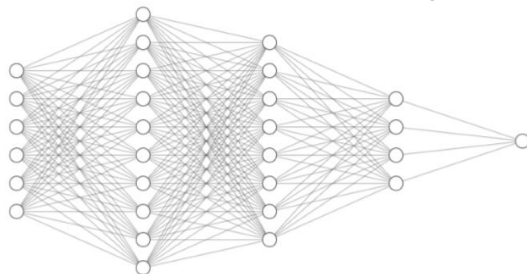
When to expand node? All node visited at least 10, and K increase by one

Which node to expand? Selection between neighbors and random nodes.

### 4. Value network

Parameter Name	Value
Neuron Number of input layer	6
Neuron Number of hidden layer 1	10
Neuron Number of hidden layer 2	8
Neuron number of hidden layer 3	4
number of Neuron output layer	1
Optimizer	Adam
Number of epochs	100
Performance function	MSE
Training Goal	0.01

**Table 3: Value network parameter setting**



**Figure 3.2: Value network structure**

#### Algorithm 2 Expansion algorithm

**Input:** Node  $N$

**Output:** new action  $A_{sug}$

```

1: function PROGRESSIVE WIDENING( $N$ )
2:   for  $t = 0, 1, 2, 3 \dots$  do
3:     Let  $k = \lceil C_{pw} t^\alpha \rceil$ ,  $A_{best} = \max_n Q_n$ ,  $Q_{mean} = \frac{\sum_{n=1}^N Q_n}{N}$ 
4:     if  $N_{Node} > 10$  and  $k$  increase by 1 then
5:       Find 2 neighbors  $A_{n1}, A_{n2}$ , 2 random samples  $A_{s1}, A_{s2}$ .
6:       choose the best evaluated sample  $A_{sug}$  and its evaluation  $Q_{A_{sug}}$ .
7:       if  $Q_{A_{sug}} > Q_{mean}$  then
8:         return  $A_{sug}$ 
9:       end if
10:    else
11:      return  $A_{best}$ 
12:    end if
13:  end for
14: end function

```

**Figure 3.1: Expansion Algorithm**



## 4.1. Evaluation Environment

■ Smart Data Innovation Lab Platform: 1 core with 4GB RAM.

■ Search space: 5/5/14 (53)

Data processor: 5

Feature Engineering : 5

Model: 14

Hyperparameter: 53

■ Dataset: OpenML CC18 suite

Name	Sample	Features	Class	Missing value
texture	5500	40	11	-
segment	2310	19	7	-
wilt	4839	6	2	-
car	1728	6	4	X
PhishingWebsites	11055	30	2	-

*Table 3: Example of datasets properties from CC18 suite.*

Name	Step	# $\lambda$
quantile	Data	2
norm	Data	1
minmax	Data	1
robust	Data	1
stand	Data	-
-----		
Fastica	Feature	3
PCA	Feature	2
Polynomial	Feature	3
Agglomeration	Feature	1
KernelPCA	Feature	1
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AdaBoost	Model	3
Bernoulli	Model	2
decision tree	Model	3
extra tree	Model	2
Gaussian naive Bayes	Model	1
Gradient boosting	Model	4
KNN	Model	3
LDA	Model	3
Linear SVC	Model	2
SVC	Model	3
Multinomial naive Bayes	Model	2
Passive aggressive	Model	4
QDA	Model	2
random forest	Model	4

*Table 4 : Default search space, # means number of hyperparameters*



## 4.2. Evaluation Results

### ■ Default setting

	Baseline	AutoSklearn	MCTML
Candidate models	SGD	Search Space	Search Space
Default parameter	learning rate = 0.0001 penalty = l2 loss function = hinge	ensemble size=1 meta learning = 0	c = 2 c <sub>pw</sub> = 4 α = 0.3
Variable parameter	—	total time = 30min	total time = 30min

Table 5 : Default setting of methods Baseline, AutoSklearn and MCTML

- Performance:  $F1 = 2 * \frac{Precision * Recall}{Precision + Recall}$
- Stability: 5 runs variance
- Evaluation summary

	Baseline	AutoSklearn	MCTML
Performance	0	14	16
Stability	—	12	22

Table 6: Statistical summary of performance and stability

Dataset	MCTML	AutoSklearn
car	<b>0.986</b>	0.938
churn	<b>0.987</b>	0.335
climate	<b>0.975</b>	0.842
cnae-9	<b>0.792</b>	0.569
cylinder-bands	0.632	<b>0.831</b>
balance-scale	0.944	0.962
fourier	<b>0.921</b>	0.802
karhunen	<b>0.950</b>	0.658
satimage	<b>0.899</b>	0.766
morphological	<b>0.840</b>	0.752
eucalyptus	<b>0.905</b>	0.662
zernike	<b>0.818</b>	0.735
optdigits	<b>0.980</b>	0.948
pendigits	<b>0.982</b>	0.750
diabetes	<b>0.952</b>	0.463
spambase	<b>0.934</b>	0.851
splice	<b>0.095</b>	0.629
tic-tac-toe	<b>0.972</b>	0.488
letter	<b>0.889</b>	0.748
dna	<b>0.588</b>	0.331
first-order	<b>0.518</b>	0.490
jm1	0.466	<b>0.518</b>
kc1	0.326	<b>0.413</b>
kc2	0.419	0.435
plxel	<b>0.892</b>	0.811
MiceProtein	<b>0.991</b>	0.877
phpJNxH0q	<b>0.954</b>	0.897
ozone-level-8hr	<b>0.818</b>	0.666
pc1	0.286	0.306
pc3	0.195	<b>0.425</b>
PhishingWebsites	<b>0.925</b>	0.905
qsar-biodeg	<b>0.868</b>	0.631
segment	<b>0.957</b>	0.900
semeion	0.812	<b>0.836</b>
wall-robot	<b>0.923</b>	0.839
wdbc	0.963	0.941
wilt	<b>0.975</b>	0.857

Table 7: F1 score on all dataset of methods MCTML and Autosklearn

## 5. Future Work

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- Regression tasks --- change task type
- Initial state sampling --- meta learning
- Value network --- pipeline conversion and structure
- More evaluation methods & Bigger search space

# Thanks

