

# Note: Building trees and splitting algorithms

Note: Oct 28, 2019

#### **▼** Reference:

- 1. <u>Paper</u> Liu, F. T., Ting, K. M., & Zhou, Z. H. (2008, December). Isolation forest. In *2008 Eighth IEEE International Conference on Data Mining* (pp. 413-422). IEEE.
- 2. <u>Paper Hariri, S., Kind, M. C., & Brunner, R. J. (2018)</u>. Extended Isolation Forest. \_arXiv preprint arXiv\_:1811.02141.
- 3. Paper: MORF, USPORF

#### **▼** Goals

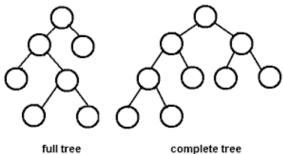
- Apply Algo2-3 from EIF paper to USPORF
- ▼ Paper1: Isolation Forest

#### **▼** 1. Definition: Isolation Tree (iTree)

- T is divided into 2 daughters nodes T\_I,T\_r
- stop when reach (i) height limit, (ii) |X| = 1 in external nodes (iii) data in X have same value

Fig 1: iTree is proper binary tree= each internal node has 2 daughters

- external (ending) nodes= n = |X|
- dimension = N
- internal (starting and latent) nodes = *n-1*
- tot nodes = 2n-1
- Thus Tree grow linearly



# **▼ 2. Definition: Pathlength** h(x)

- h(x) = number of edge from root node to terminal node ~ avg. depth unsuccessful search in binary tree?
- $max h(x) \sim n, E[h(x)] = log n$
- ▼ Paper2: Extended Isolation Forest

# ▼ Algo 1:

# **Algorithm 1** : $iForest(X, t, \psi)$

**Inputs**: X - input data, t - number of trees,  $\psi$  - subsampling size

**Output**: a set of *t iTrees* 

1: **Initialize** Forest

2: set height limit  $l = ceiling(\log_2 \psi)$ 

3: **for** i = 1 to t **do** 

 $X' \leftarrow sample(X, \psi)$ 

 $Forest \leftarrow Forest \cup iTree(X', 0, l)$ 

6: end for

7: **return** Forest

- I = average tree height. We are only interested length <I</li>
- \psi ~ 256 is enough, \t ~ 100 is enough
- Forest = {iTree}, i = 1,..,t-1 → see Algo 2
- X' (subset of X with sample size \psi)

complxity  $\sim O(t\psi \log \psi)$ 

# ▼ Algo 2: Splitting algorithm

#### Algorithm 2 iTree(X, e, l)

**Input:** X - input data, e - current tree height, l - height limit

#### Output: an iTree

- 1: if  $e \ge l$  or  $|X| \le 1$  then
- 2: **return**  $exNode\{Size \leftarrow |X|\}$
- 3: **else**
- 4: randomly select a normal vector  $n \in \mathbb{R}^{|X|}$  by drawing each coordinate of  $\vec{n}$  from a uniform distribution.
- 5: randomly select an intercept point  $p \in \mathbb{R}^{|X|}$  in the range of X
- 6: set coordinates of n to zero according to extension level
- 7:  $X_l \leftarrow filter(X, (X p) \cdot n \le 0)$
- 8:  $X_r \leftarrow filter(X, (X-p) \cdot n > 0)$
- 9: **return** inNode{  $Left \leftarrow iTree(X_l, e+1, l)$ ,

$$Right \leftarrow iTree(X_r, e+1, l),$$

 $Normal \leftarrow n$ ,

 $Intercept \leftarrow p$ 

10: **end if** 

- 6: extension level ={0,1} → {Standard IF, Extended IF}
- **▼** Splitting Algorithm
  - 1. normal vector (gradient), choose from uniform distribution in R^N

$$\hat{n} \in \mathbb{R}^N$$
, N-dimension

2. random intercept for the cut choose from uniform distribution in  $\mathbf{X}$ 

$$ec{p} \in \mathcal{X}, \mathcal{X} ext{ current node member} \}$$

1. and 2. create the splitting criteria

$$(ec{x}-ec{p})\cdot\hat{n}\leq 0,\;\;ec{x}\in\mathcal{X}$$

- ≤ go left daughter node
- > go right daughter node
- ▼ Return

- While {if e < I or |X| > 1}: inNode{ Left-daughter iTree, Right-daughter iTree, Normal, Intercept }
  - e← e+1, depth level in the decision tree
- {if  $e \ge 1$  or  $|X| \le 1$ }: exNode{Size  $\leftarrow |X|$ }, end

#### ▼ Algo 3:

```
Algorithm 3 PathLength(x,T,e)

Input: x - an instance, T - an iTree, e - current path length; to be initialized to zero when first called Output: path length of x

1: if T is an external node then

2: return e+c(T.size)\{c(.) is defined in Equation (2)}

3: end if

4: n \leftarrow T.Normal

5: p \leftarrow T.Intercept

6: if (x-p) \cdot n \leq 0 then

7: return PathLength(x,T.left,e+1)

8: else if (x-p) \cdot n > 0 then

9: return PathLength(x,T.rigth,e+1)

10: end if
```

#### ▼ Parameters

- x =one data in the tree, xin X'
- e = path length, set to 0 at first
- *T = iTree(X',e,l)* from **Algo 2** 
  - T = { exNode{ pop |X| in the node}, InNode{Normal, Intercept, left, right } }

#### ▼ PathLength algorithm

eqn (2): H(i), harmonic number = In(i) +0.5772156649 ?

$$c(n) = 2H(n-1) - (2(n-1)/n)$$

- While {T = inNode}: PathLength(x, T.left or T.right, e ← e+1)
- {if T=exNode}: h(x)=e+c(T.size) end

### Anomoly score s(n)

$$s(x,n)=2^{-E[h(x)]/c(n)},\; E[h(x)]pprox \log(n)$$

## ▼ Paper3 MORF

- ▼ Goal:
  - In structured data lying on manifold (easily searchable i.e. spreadsheets, image, text, NN), the indices is as important as magnitude
  - · build distribution in manifold
- **▼** Background:
  - 1. Classification
    - data  $D_n \sim F_XY$  distribution
  - 2. Random Forest
    - D\_n ← {S^L\_theta\*, S^R\_theta\*}
    - theta\* =(e\*\_j=feature basis, tau\*= split): tend to do halves split
    - $I(S) = Gini impurity = 1-p_i^2$
  - 3. SPORF
- ▼ Algo 1: Manifold Decision Tree

$$\frac{\text{function}}{\text{function}} T = \text{GROWTREE}(X, y, fA, \Theta)$$

▼ Algo 2: best node split

$$\frac{\text{function}}{\text{function}} (j*,\tau*) = \text{FINDBESTSPLIT}(X,y)$$

best split = minimizing Gini impurity / entropy

- ▼ Paper4 USPORF (URerF)
  - ▼ Goal:
    - estimate Geodesic distance btw 2 points
  - ▼ Parameters:

$$x=\{x_1,..,x_N\}\in\mathbb{R}^p$$

- each tree,  $t = \{1,..,T\}$  has subset of sample X', |X'| = m < N
- use subset of feature d<p,
- transform X'∈ R^P into X~ ∈R^d

$$ilde{X}_{d imes 1} = A^T X_{p imes 1}', \ \ A_{d imes p}$$

- ▼ Spliting Criteria
  - 2-GMM Splitting with Fast-BIC
    - = split with global maximum log likelihood, s = cluster number
- ▼ Algo 1: build an unsupervised random decision tree
  - function t = BUILDTREE(X' subset,d allowed dimension,Θ split criteria)
  - if satisfy Θ: split X' into {X'I,X'r} #note: optimal split ~ middle
  - if not satisfy Θ: LeafNode(X)