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Topic 20: Random Forest

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Key points: .

Disclaimer: The note is built on Prof. Jinchi Lv's lectures of the course at USC, DSO 607, High-Dimensional Statistics and Big Data Problems.

20.1 Motivation

Denote by $m(\mathbf{X})$ the measurable nonparametric regression function with p-dimensional random vector \mathbf{X} taking values in $[0,1]^p$. The Random Forest algorithm aims to learn the regression function in a nonparametric way based on the observations $\mathbf{x}_i \in [0,1]^p$, $y_i \in \mathbb{R}$, $i = 1, \dots, n$, from the model

$$y_i = m(\mathbf{x}_i) + \epsilon_i$$

where X, x_i , ε_i , $i = 1, \dots, n$ are independent, and $\{x_i\}$ and $\{\varepsilon_i\}$ are two sequences of identically distributed random variables. x_i is distributed identically as X.

Why Random Forest (RF)? RF has gained significant popularity due to its

- High accuracy: RF consistently rank among the top performer, often surpassing more complex models
- Robustness: RF are less subject to overfitting due to the ensemble nature leveraging multiple decision trees
- Interpretability: RF provide rankings of feature importance

As illustrated in Figure 20.1, in a level-2 tree, each node (cell) defines the point where the current cell split and new cells are produced. The sets of features eligible for splitting cells at level k-1 are denoted as $\Theta_k := \{\Theta_{k,1}, \cdots, \Theta_{k,2^{k-1}}\}$, where $\Theta_{k,s} \subset \{1, \cdots, p\}$.



Figure 20.1: Level-2 Tree Example

Given any T (and the associated splitting criterion) and $\Theta_{1:k}$, the tree estimate denoted as $\hat{m}_{T(\Theta_{1:k})}$ for a test

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point $\mathbf{c} \in [0,1]^p$ is defined as

$$\hat{m}_{T(\Theta_{1:k})}(\mathbf{c}, \mathcal{X}_n) := \sum_{(\mathbf{t}_1, \dots, \mathbf{t}_k) \in T(\Theta_{1:k})} \mathbf{1}_{\mathbf{c} \in \mathbf{t}_k} \left(\frac{\sum_{i \in \{i: \mathbf{x}_i \in \mathbf{t}_k\}} y_i}{\# \{i: \mathbf{x}_i \in \mathbf{t}_k\}} \right)$$

where $X_n := \{\mathbf{x}_i, y_i\}_{i=1}^n$ the fraction is defined as 0 when no sample is in the cell \mathbf{t}_k , and $\mathbf{1}_{\mathbf{c} \in \mathbf{t}_k}$ is an indicator function = 1 if $\mathbf{c} \in \mathbf{t}_k$ and = 0 otherwise.

20.2 Chi et al. (2022): High Dimensional RFs

Following Chi et al. (2022), for a RF model where

- a sequence of distinct $\Theta_{1:k}$ results in a distinct tree
- every set of available features $\Theta_{l,s}$, $l=1,\cdots,k$; $s=1,\cdots,2^{l-1}$

Column subsampling Define a **column subsampling** procedure: $\Theta_{l,s}$, $\forall l,s$ has $[\gamma_0 p]$ distinct integers among $1, \dots, p$, with $[\cdot]$ the ceiling function for some $0 < \gamma_0 \le 1$. γ_0 is the predetermined constant parameter of column subsampling. Introduce the boldface random mappings $\Theta_{1:k}$, which are independent and uniformly distributed over all possible $\Theta_{1:k}$ for all integer k. Then random forests estimate for \mathbf{c} with observations X_n is given by

$$\mathbb{E}\left(\hat{m}_{T(\boldsymbol{\Theta}_{1:k})}\left(\mathbf{c}, \mathcal{X}_{n}\right) \mid \mathcal{X}_{n}\right) = \sum_{\boldsymbol{\Theta}_{1:k}} \mathbb{P}\left(\bigcap_{s=1}^{k} \left\{\boldsymbol{\Theta}_{s} = \boldsymbol{\Theta}_{s}\right\}\right) \hat{m}_{T(\boldsymbol{\Theta}_{1:k})}\left(\mathbf{c}, \mathcal{X}_{n}\right)$$

The expectation is taken over sets of available features.

Observation resampling Let $A = \{a_1, \dots, a_B\}$ be a set of subsamples with each a_i consisting of $\lceil bn \rceil$ observations (indices) drawn without replacement from $\{1, \dots, n\}$ for some positive integer B and $0 < b \le 1$; in addition, each a_i is independent of model training. The default values of B and B are 500 and 0.632 1 . Then the tree estimate using subsample B is define as

$$\hat{m}_{T(\Theta_{1:k}),a}\left(\mathbf{c},\mathcal{X}_{n}\right) \coloneqq \sum_{\left(\mathbf{t}_{1},\cdots,\mathbf{t}_{k}\right)\in T(\Theta_{1:k})} \mathbf{1}_{\mathbf{c}\in\mathbf{t}_{k}} \left(\frac{\sum_{i\in a\cap\left\{i:\mathbf{x}_{i}\in\mathbf{t}_{k}\right\}} y_{i}}{\#\left(a\cap\left\{i:\mathbf{x}_{i}\in\mathbf{t}_{k}\right\}\right)}\right)$$

the random forests estimate given A is then

$$B^{-1} \sum_{a \in A} \mathbb{E} \left[\hat{m}_{T,a} \left(\mathbf{\Theta}_{1:k}, \mathbf{c}, \mathcal{X}_n \right) \mid \mathcal{X}_n \right] := B^{-1} \sum_{a \in A} \mathbb{E} \left[\hat{m}_{T(\mathbf{\Theta}_{1:k}),a} \left(\mathbf{c}, \mathcal{X}_n \right) \mid \mathcal{X}_n \right]$$

CART-split criterion Given a cell t, a subset of observation indices a and a set of available features $\Theta \subset \{1, \dots, p\}$, the CART-split is defined as

$$(\hat{j}, \hat{c}) = \arg \min_{j \in \Theta, c \in \{x_{ij}: \mathbf{x}_i \in \mathbf{t}, i \in a\}} \left[\sum_{i \in a \cap P_L} (\overline{y}_L - y_i)^2 + \sum_{i \in a \cap P_R} (\overline{y}_R - y_i)^2 \right]$$
(20.1)

 $^{{}^{1}\}text{Or}$, b = 1 but observations are drawn with replacement.

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where

$$P_L := \left\{ i : \mathbf{x}_i \in \mathbf{t}, x_{ij} < c \right\}$$

$$\overline{y}_L := \sum_{i \in a \cap P_L} \frac{y_i}{\#(a \cap P_L)}$$

$$P_R := \left\{ i : \mathbf{x}_i \in \mathbf{t}, x_{ij} \ge c \right\}$$

$$\overline{y}_R := \sum_{i \in a \cap P_R} \frac{y_i}{\#(a \cap P_R)}$$

The CART-split criterion conditional on the sample is a deterministic splitting criterion; conditioning on another sample leads to another deterministic splitting criterion. Define \hat{T}_a as the sample tree growing rule that is associated with a splitting criterion following Eq. (20.1), the tree estimates using \hat{T}_a can be similarly defined as

$$\hat{m}_{\hat{T}_a(\Theta_{1:k})}(\mathbf{c}, \mathcal{X}_n) := \sum_{(\mathbf{t}_1, \dots, \mathbf{t}_k) \in \hat{T}_a(\Theta_{1:k})} \mathbf{1}_{\mathbf{c} \in \mathbf{t}_k} \left(\frac{\sum_{i \in \{i: \mathbf{x}_i \in \mathbf{t}_k\}} y_i}{\# \{i: \mathbf{x}_i \in \mathbf{t}_k\}} \right)$$

the definition is the same for $\hat{m}_{\hat{T}_{a,d}}$. Then the random forests estimate for a test point $\mathbf{c} \in [0,1]^p$ is given by

$$B^{-1} \sum_{a \in A} \mathbb{E} \left(\hat{m}_{\hat{T}_a, a} \left(\mathbf{\Theta}_{1:k}, \mathbf{c}, \mathcal{X}_n \right) \mid \mathcal{X}_n \right)$$

where the average and conditional expectation correspond to the sample and column subsamplings respectively, and they are interchangeable.

Bias-variance decomposition For a tree growing rule T and $\Theta_{1:k}$, the population version is defined as

$$m_{T(\Theta_{1:k})}^{*}(\mathbf{c}) := \sum_{(\mathbf{t}_{1}, \dots, \mathbf{t}_{k}) \in T(\Theta_{1:k})} \mathbf{1}_{\mathbf{c} \in \mathbf{t}_{k}} \mathbb{E}\left(m(\mathbf{X}) \mid \mathbf{X} \in \mathbf{t}_{k}\right)$$
(20.2)

for each test point $\mathbf{c} \in [0,1]^p$. And the \mathbb{L}^2 prediction loss for random forests is defined as

$$\mathbb{E}\left[m(\mathbf{X}) - B^{-1} \sum_{a \in A} \mathbb{E}\left(\hat{m}_{\hat{T}_a, a}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}, \mathcal{X}_n\right) \mid \mathbf{X}, \mathcal{X}_n\right)\right]^2$$
(20.3)

if we use the full sample $a = \{1, \dots, n\}$, and denote \hat{T}_a and $\hat{m}_{\hat{T}_a, a}$ as \hat{T} and $\hat{m}_{\hat{T}}$, the sample subsampling and average $B^{-1} \sum_{a \in A} (\cdot)$ in the random forests estimate are no longer needed, then Eq.(20.3) can be simplified as

$$\mathbb{E}\left[m(\mathbf{X}) - \mathbb{E}\left(\hat{m}_{\hat{T}}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}, \mathcal{X}_{n}\right) \mid \mathbf{X}, \mathcal{X}_{n}\right)\right]^{2}$$

By Jensen's inequality and Cauchy-Schwarz inequality,

$$\frac{1}{2}\mathbb{E}\left[m(\mathbf{X}) - \mathbb{E}\left(\hat{m}_{\hat{T}}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}, \mathcal{X}_{n}\right) \mid \mathbf{X}, \mathcal{X}_{n}\right)\right]^{2}$$

$$\leq \mathbb{E}\left[m(\mathbf{X}) - m_{\hat{T}}^{*}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}\right)\right]^{2} + \mathbb{E}\left[m_{\hat{T}}^{*}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}\right) - \hat{m}_{\hat{T}}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}, \mathcal{X}_{n}\right)\right]^{2}$$
approximation error (squared bias) estimation variance

Consistency of RF Models

For a cell t and its two daughter cells t' and t", define

$$\begin{split} (\mathbb{I})_{\mathbf{t},\mathbf{t}'} &:= \mathbb{P}\left(X \in \mathbf{t}' \mid X \in \mathbf{t}\right) \operatorname{Var}\left(m(X) \mid X \in \mathbf{t}'\right) + \mathbb{P}\left(X \in \mathbf{t}'' \mid X \in \mathbf{t}\right) \operatorname{Var}\left(m(X) \mid X \in \mathbf{t}''\right) \\ (\mathbb{II})_{\mathbf{t},\mathbf{t}'} &:= \mathbb{P}\left(X \in \mathbf{t}' \mid X \in \mathbf{t}\right) \left[\mathbb{E}(m(X) \mid X \in \mathbf{t}') - \mathbb{E}(m(X) \mid X \in \mathbf{t})\right]^2 \\ &+ \mathbb{P}\left(X \in \mathbf{t}'' \mid X \in \mathbf{t}\right) \left[\mathbb{E}(m(X) \mid X \in \mathbf{t}'') - \mathbb{E}(m(X) \mid X \in \mathbf{t})\right]^2 \end{split}$$

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and $(\mathbb{I})_{\mathbf{t},\mathbf{t}''}$ and $(\mathbb{II})_{\mathbf{t},\mathbf{t}''}$ are defined similarly.

in this context, we assume the following regularity conditions:

- **Absolutely continuous distribution**: f, the density function of X, is bounded away from 0 and ∞
- Covariates and model errors: assume $p = O(n^{K_0})$ for $K_0 > 0$, and there is a symmetric distribution around 0 for ϵ_1 , s.t. $\mathbb{E}|\epsilon_1|^q < \infty$ for sufficiently large q > 0
- **Bounded regression functions**: $\sup_{\mathbf{c} \in [0,1]^p} |m(\mathbf{c})| \le M_0$, for some $M_0 > 0$
- Sufficient impurity decrease: $\exists \alpha_1 \ge 1 \text{ s.t. } \forall \mathbf{t} = t_1 \times \cdots \times t_p$,

$$\operatorname{Var}\left[m(\mathbf{X})\mid\mathbf{X}\in\mathbf{t}\right]\leq\alpha_{1}\sup_{j\in\left\{1,\cdots,p\right\},c\in t_{j}}\left(\mathbb{II}\right)_{\mathbf{t},\mathbf{t}\left(j,c\right)}$$

where

- $-(\mathbb{I})_{t,t'}$: conditional bias decrease (or conditional impurity decrease)
- Var [m(X) | X ∈ t]: conditional *total* bias, Var [m(X) | X ∈ t] = (\mathbb{I})_{t,t'} + (\mathbb{I})_{t,t'}
- Intuition: having large conditional bias decrease on each cell is a desired property for achieving a good control of the squared bias of random forests estimate

Sufficient impurity decrease (SID) Define the functional class

$$SID(\alpha) := \{m(\mathbf{X}) : m(\mathbf{X}) \text{ satisfies SID with } \alpha_1 \leq \alpha \}$$

the size of $SID(\alpha)$ is **non-decreasing** in $\alpha \ge 1$: if $m(\mathbf{X}) \in SID(\alpha - c)$ for some $\alpha - c \ge 1$ and c > 0, then $m(\mathbf{X}) \in SID(\alpha)^2$.

Under the regularity conditions mentioned above, we have the following theorem

Theorem 20.2.1: Consistency

Let $0 < b \le 1, 0 < \gamma_0 \le 1, \alpha_2 > 1, 0 < \eta < 1/8, 0 < c < 1/4 and <math>\delta > 0, 2\eta < \delta < \frac{1}{4}$. Let $A = \{a_1, \dots, a_B\}$ with $\#a_i = \lceil bn \rceil$ for $i = 1, \dots, B$ and $a \in A$. Then $\exists C > 0$ s.t. for all large n and each $1 \le k \le c \log_2 \lceil bn \rceil$,

$$\mathbb{E}\left[m(\mathbf{X}) - \mathbb{E}\left(\hat{m}_{\hat{T}_{a},a}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}, X_{n}\right) \mid \mathbf{X}, X_{n}\right)\right]^{2}$$

$$\leq C\left[\alpha_{1}\left(\lceil bn \rceil\right)^{-\eta} + \left(1 - \gamma_{0}\left(\alpha_{1}\alpha_{2}\right)^{-1}\right)^{k} + \left(\lceil bn \rceil\right)^{-\delta + c}\right]$$

In addition, when aggregate over row subsamples (over $a \in A$), get

$$\mathbb{E}\left[m(\mathbf{X}) - \frac{1}{B}\mathbb{E}\left(\hat{m}_{\hat{T}_a,a}\left(\mathbf{\Theta}_{1:k}, \mathbf{X}, X_n\right) \mid \mathbf{X}, X_n\right)\right]^2$$

$$\leq C\left[\alpha_1\left(\lceil bn \rceil\right)^{-\eta} + \left(1 - \gamma_0\left(\alpha_1 \alpha_2\right)^{-1}\right)^k + \left(\lceil bn \rceil\right)^{-\delta + c}\right]$$

Here, the feature dimensionality p and tree height k decide the number of all possible cells when growing trees.

²Many popular regression functions belong to this functional class.

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Bias-variance decomposition Under Theorem (20.2.1), both bias and variance depend implicitly on p through n in the upper bounds.

Proposition 20.2.2: Bias-Variance Decomposition

Under Thm (20.2.1), for all large n and $1 \le k \le c \log_2 n$, it holds that

Squared bias :=
$$\underbrace{\mathbb{E}\left[m(\mathbf{X}) - m_{\hat{T}}^*(\mathbf{\Theta}_{1:k}, \mathbf{X})\right]^2}_{\text{approximation error}}$$

$$\leq O\left(n^{-\eta} + \underbrace{\left(1 - \gamma_0(\alpha_1\alpha_2)^{-1}\right)^k}_{\text{Main term of bias}}\right) + \underbrace{O\left(n^{-\delta+c}\right)}_{\text{Uninteresting error}}$$

where $n^{-\eta}$ upper-bounds the error caused by the sample CART-splits. Under theoretical CART-splits, $n^{-\eta}$ vanishes and $\alpha_2 = 1$. and

Estimatino Variance :=
$$\mathbb{E}\left[m_{\hat{T}}^*(\boldsymbol{\Theta}_{1:k}, \mathbf{X}) - \hat{m}_{\hat{T}}(\boldsymbol{\Theta}_{1:k}, \mathbf{X}, \mathbf{X}_n)\right]^2$$

$$\leq O(n^{-\eta}) + \underbrace{O(n^{-\delta+c})}_{\text{Unintersting error}}$$

where the upper bound is conservative since we establish a uniform upper bound for the variances of **individual** trees.

Relevant features Under the regularity conditions, for some cells, only the splits along the relevant feature directions can reduce a sufficient amount of bias. To be precise, introduce a variance of SID with some $S_0 \subset \{1, \dots, p\}$ below

• **Sufficient impurity decrease 2** $\exists \alpha_1 \geq 1 \text{ s.t. for each cell } \mathbf{t} = t_1 \times \cdots \times t_p$

$$\operatorname{Var}(m(\mathbf{X}) \mid \mathbf{X} \in \mathbf{t}) \leq \alpha_1 \sup_{j \in S_0, c \in t_j} (\mathbb{II})_{\mathbf{t}, \mathbf{t}(j, c)}$$

When the regularity conditions on the underlying regression function and SID are assumed, SID2 holds only if S_0 includes all relevant features

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References

Chien-Ming Chi, Patrick Vossler, Yingying Fan, and Jinchi Lv. Asymptotic properties of high-dimensional random forests. *The Annals of Statistics*, 50(6):3415–3438, 2022.