



# Gaussian Soft Decision Trees for Interpretable Feature-Based Classification

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# Outline

- **Introduction**
- Previous Works
- Proposed Method
- Experiments
- Conclusion

# Black Box

- Deep neural network is a **black box**
  - Its decision process is not interpretable
  - Difficult to trust decisions even with high accuracy

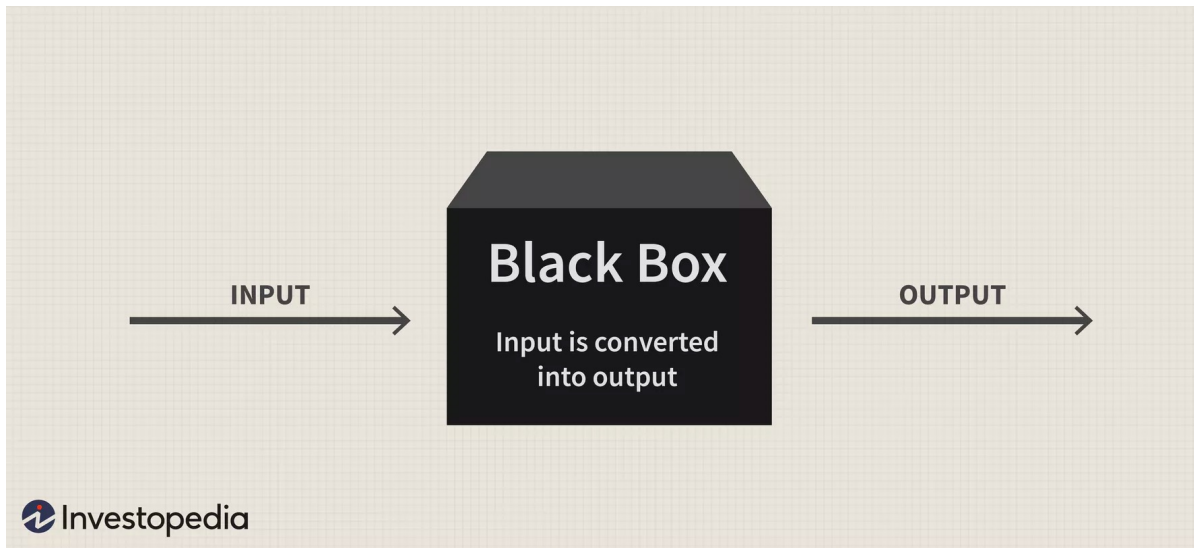




Image from <https://www.investopedia.com/terms/b/blackbox.asp>



# Interpretable ML (1)

- Research to interpret a model's decisions
  - Important especially in bio or medical domains
- **Global interpretability**
  - A model's decision process is itself interpretable
    - Linear models or decision trees
- **Local interpretability**
  - To explain decisions made by black box models
    - Recent works for deep neural networks

# Interpretable ML (2)

- Research to interpret a model's decisions
  - Important especially in bio or medical domains
- **Global interpretability** 
  - A model's decision process is itself interpretable
  - Global interpretability makes **reliable decisions**
- **Feature-based classification** 
  - Simple models can be better than neural networks
  - Generalizability is more important the capability

# Tree Models

- Tree models provide global interpretability
  - Each decision is represented as a path in the tree, which has its own meaning

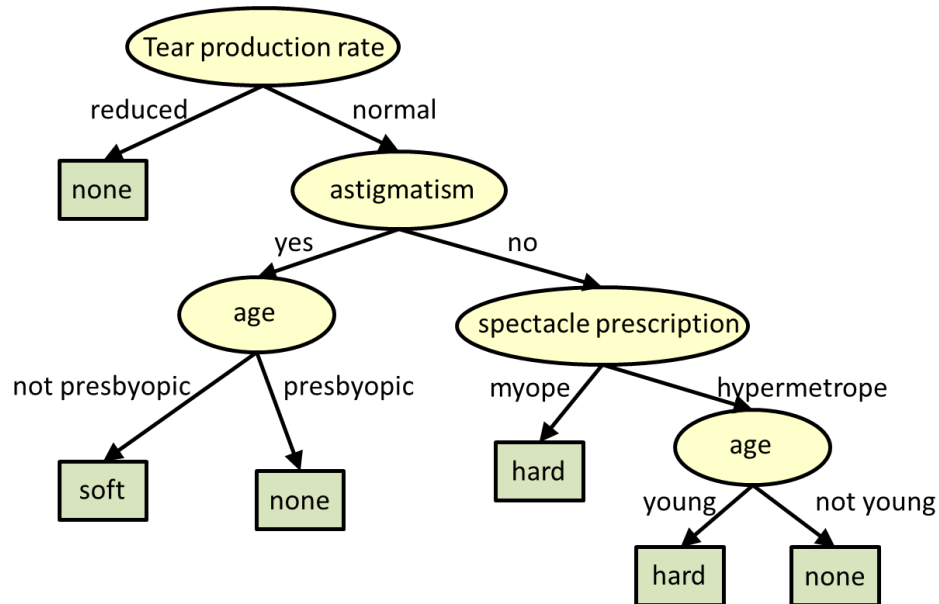


Image from <https://www.cs.cmu.edu/~bhiksha/courses/10-601/decisiontrees/>



# Limitations of Tree Models

- **Linear decisions**

- Restrict the overall representation power
- Make it difficult to learn complex decision rules

- **Large tree depth**

- Limits the interpretability of models
- Tree depth means the complexity of interpretation
- *Is a tree still interpretable with large depth  $d > 10$ ?*



# Problem Definition

- **Given** a feature-based dataset  $\mathcal{D} = \{(\mathbf{x}_i, y_i)\}_i$ 
  - No structural information exists in  $\mathbf{x}$
  - Each element in  $\mathbf{x}$  is itself meaningful
- **Train** an interpretable tree classifier  $f$
- **Maximizing** its accuracy and interpretability
  - Addressing the limitations of previous models





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# Decision Trees

- One of the most popular tree models
  - Has been used for decades
  - Learns an explicit decision rule at each branch
    - For instance, to pass  $x$  to the left child if  $x_3 > 3$
- **Strength**
  - Its decision process is clear and interpretable
- **Weakness**
  - It easily overfits, making limited performance



# Soft Decision Trees (1)

- Improve the representation power of DTs
  - Perform a soft decision with all features
  - Learn a soft target distribution at each leaf
- **Strength**
  - Larger capability to learn complex decision rules
- **Weakness**
  - Less interpretability due to the soft decisions



# Soft Decision Trees (2)

- SDTs are characterized by **soft decisions**
- The probability  $f_i$  at node  $i$  to pass  $\mathbf{x}$  to the right child is

$$f_i = \sigma(\mathbf{w}_i^\top \mathbf{x} + b_i)$$

- $\mathbf{w}_i$  and  $b_i$  are learnable parameters at node  $i$
- $\sigma$  is the sigmoid function for the split
- The probability to the left is  $1 - f_i$  thanks to  $\sigma(\cdot)$



# Soft Decision Trees (3)

- The interpretability is worse than that of DTs
  - Because all features are used for every decision
  - Each decision path involves  $O(dm)$  parameters
    - $d$  is the depth, and  $m$  is the number of features
- EDiT (ICDM 2019) focused on decreasing  $m$ 
  - It learns a sparse weight vector at each branch
  - However, the large depth  $d$  remains the same



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# Overview (1)

- **Gaussian Soft Decision Trees (GSDT)**
  - Tree model having a multi-branched structure
  - Decisions are modeled as Gaussian mixtures
  - Address the limitations of previous tree models
- **Main ideas**
  - Gaussian decisions
  - Low-rank perturbation
  - Path regularization
  - Post-optimization



# Overview (2)

- GSDT first computes the arrival probability  $\mathbf{r}(\mathbf{x})$
- Then, the prediction is done by a single leaf:

$$\hat{y}(\mathbf{x}) = \mathbf{p}_i \quad \text{where } i = \operatorname{argmax}_k r_k(\mathbf{x})$$

- $\mathbf{p}_i$  is the class distribution learned by leaf  $i$
- The training is done by a gradient-based way
  - All parameters are updated at the same time
  - We minimize the hinge loss for classification



# Gaussian Decisions (1)

- We make all decisions as Gaussian mixtures
- This enables us to preserve the interpretability even with the nonlinearity of decisions

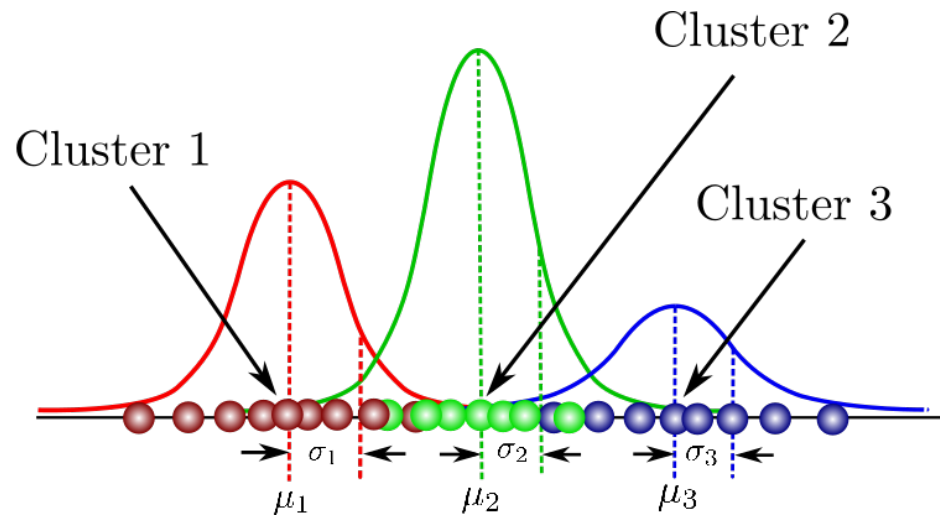
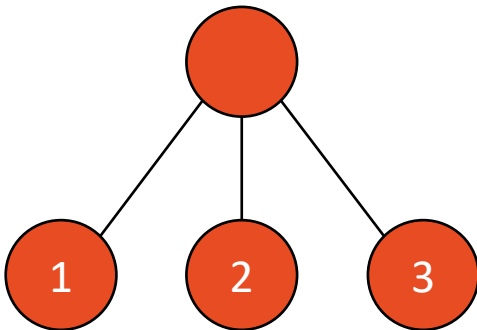


Image from <https://towardsdatascience.com/gaussian-mixture-models-explained-6986aaf5a95>

# Gaussian Decisions (2)

- The probability  $f_{ij}(\mathbf{x})$  of  $\mathbf{x}$  from node  $i$  to  $j$  is

$$f_{ij}(\mathbf{x}) = \frac{\exp(\mathcal{L}(\theta_j | \mathbf{x}))}{\sum_k \exp(\mathcal{L}(\theta_k | \mathbf{x}))},$$

- $\mathcal{L}$  is the log likelihood of  $\mathbf{x}$ , which is defined as

$$\mathcal{L}(\theta_j | \mathbf{x}) = -\frac{1}{2} \left( (\mathbf{x} - \boldsymbol{\mu}_j)^\top \boldsymbol{\Sigma}_j^{-1} (\mathbf{x} - \boldsymbol{\mu}_j) + \log \det(\boldsymbol{\Sigma}_j) + d \log(2\pi) \right).$$

- $\boldsymbol{\mu}_j$  and  $\boldsymbol{\Sigma}_j$  are learned through backpropagation



# Gaussian Decisions (3)

- Gaussian decisions make several advantages
  - **Nonlinearity**
    - Each branch can learn a complex decision function
  - **Interpretability of decisions**
    - $f_{ij}(\mathbf{x})$  is itself interpretable as a probability
  - **Interpretability of parameters**
    - $\mu_i$  summarizes the examples arriving at node  $i$
    - $\Sigma_i$  gives insights about the given features
      - E.g., which feature is more important than others?



# Gaussian Decisions (4)

- What if we apply multiple branches directly to soft decision trees?
  - It makes multiple children at each branch as

$$\mathbf{p}(\mathbf{x}) = \text{softmax}(\mathbf{W}\mathbf{x} + \mathbf{b})$$

- However,  $\mathbf{p}$  becomes no longer interpretable
  - $w_{ij} \neq$  the correlation between  $x_j$  and  $p_i$



# Low-Rank Perturbation (1)

- It is burdensome to learn a full matrix  $\Sigma_i$ 
  - Because of the  $\log \det(\Sigma_i)$  and  $\Sigma_i^{-1}$  operations
- Diagonal covariance is a simple choice
  - But it ignores the correlations between features
- We propose **low-rank perturbation**
  - Strengthen the diagonal  $\Sigma_i$  with correlations
  - Involve only  $O(m)$  additional parameters

# Low-Rank Perturbation (2)

- Our covariance matrix at each node  $i$  is

$$\Sigma_i = \text{diag}(\log(1 + \exp(\boldsymbol{\sigma}_i))) + \mathbf{U}\mathbf{U}^\top$$

- $\boldsymbol{\sigma}_i \in \mathbb{R}^m$  is a learnable vector
- $\mathbf{U} \in \mathbb{R}^{m \times k}$  is a learnable matrix
- $k$  is the target rank
  - We set  $k$  to 1 or 2 in experiments



# Path Regularization

- How to encourage GSDT to utilize all leaves?
  - GSDT is prone to use only a few leaf nodes
- We add the **path regularizer** to the objective

$$l_{\text{lr}}(\mathcal{B}) = \sum_{j \in \mathcal{N}_d} r_j(\mathcal{B}) \log r_j(\mathcal{B}) \quad \text{where} \quad \mathbf{r}(\mathcal{B}) = \frac{1}{|\mathcal{B}|} \sum_{\mathbf{x} \in \mathcal{B}} \mathbf{r}(\mathbf{x}),$$

- $l_{\text{lr}}(\mathcal{B})$  calculates the negative entropy of  $\mathbf{r}(\mathcal{B})$
- $\mathbf{r}(\mathcal{B})$  is the mean arrival probability for batch  $\mathcal{B}$

# Post-Optimization (1)

- Each leaf  $i$  corresponds to a set of examples
  - That arrive at the leaf node  $i$  at the inference
  - $\mathcal{X}_i = \{x \in \mathcal{D} \mid \operatorname{argmax}_k r_k(\mathbf{x}) = i\}$
- **Post-optimization**
  - Our technique to maximize the correspondence
  - We make the dist.  $\mathcal{N}_i$  represent the examples  $\mathcal{X}_i$
  - The interpretability of leaves further improves



# Post-Optimization (2)

- The algorithm is given as follows:
  - $\mu_j$  and  $\Sigma_j$  are updated to represent the set  $\mathcal{X}_j$

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**Algorithm 1:** Post-optimization of the leaf Gaussians of GSDT.

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**Input:** A trained GSDT  $M$ , a set  $\mathcal{D}$  of training features, a learning rate  $\alpha$  for the covariances, and the number  $n$  of iterations

- 1: **for** leaf node  $j$  in  $M$  **do**
  - 2:    $\mathcal{X}_j \leftarrow \{\mathbf{x} \in \mathcal{D} \mid \arg \max_k r_k(\mathbf{x}) = j\}$
  - 3:    $\boldsymbol{\mu}_j \leftarrow \sum_{\mathbf{x} \in \mathcal{X}_j} \mathbf{x}$
  - 4:   **for**  $i \in [1, n]$  **do**
  - 5:      $l \leftarrow \text{sum}((\boldsymbol{\Sigma}_j - \text{cov}(\mathcal{X}_j))^2)$
  - 6:      $\boldsymbol{\Sigma}_j \leftarrow \boldsymbol{\Sigma}_j - \alpha \cdot \partial l / \partial \boldsymbol{\Sigma}_j$
  - 7:   **end for**
  - 8: **end for**
  - 9: Fine-tune the whole parameters of  $M$  for a fixed number of epochs
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# Experimental Setup

- **Datasets**

- We use six public feature-based datasets
- Taken from UCI Repository or Kaggle
- All of them are bio and medical domains
  - Interpretability is a crucial factor

- **Baselines**

- Interpretable models: LR, SVM, DT, SDT, EDiT
- Black box models: RF, MLP

# Classification Accuracy

- GSDT shows the best accuracy in five datasets
  - GSDT outperforms even strong black box models

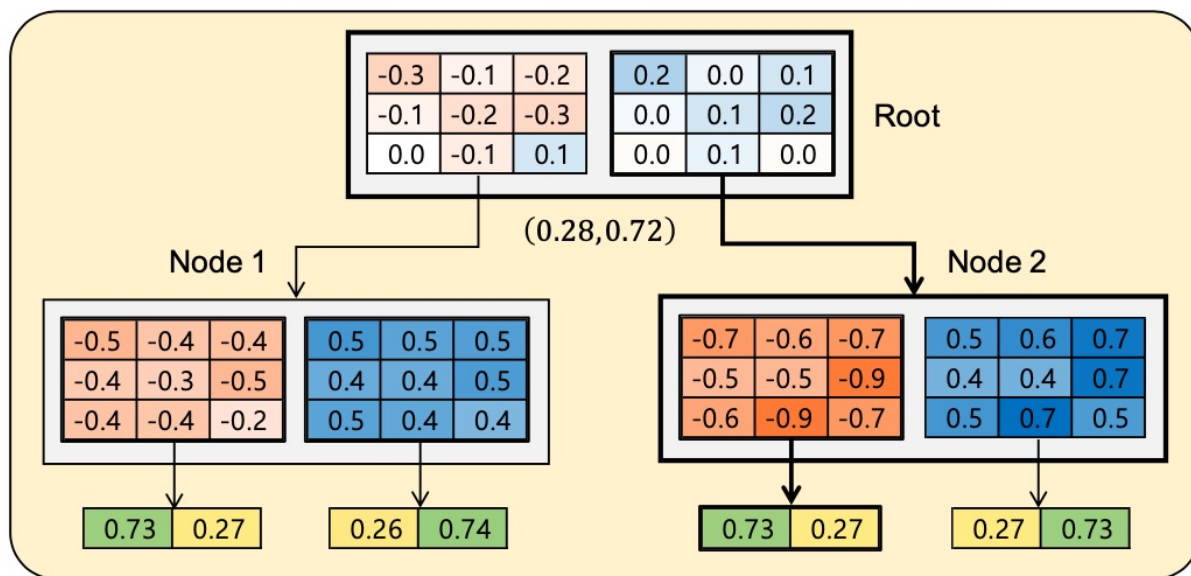
Model	Brain	Breast	Breast-wis	Diabetes	Heart	Hepatitis
LR	63.4 ± 0.0	65.5 ± 0.0	97.1 ± 0.0	<u>76.0 ± 0.0</u>	<b>86.9 ± 0.0</b>	77.4 ± 0.0
SVM-lin	61.0 ± 0.0	62.1 ± 0.0	97.1 ± 0.0	<b>76.6 ± 0.0</b>	83.6 ± 0.0	77.4 ± 0.0
SVM-rbf	58.5 ± 0.0	70.7 ± 0.0	97.1 ± 0.0	<u>76.0 ± 0.0</u>	<b>86.9 ± 0.0</b>	77.4 ± 0.0
DT	70.5 ± 0.7	68.8 ± 1.6	96.0 ± 0.9	69.7 ± 1.6	67.2 ± 1.6	70.0 ± 6.9
SDT	66.8 ± 5.0	73.3 ± 5.2	97.9 ± 0.0	<u>76.0 ± 0.7</u>	80.7 ± 2.7	67.3 ± 4.7
EDiT	58.5 ± 0.0	75.0 ± 2.6	97.1 ± 0.2	74.6 ± 1.5	85.2 ± 2.3	<u>77.8 ± 3.8</u>
MLP	<u>73.4 ± 1.7</u>	73.3 ± 2.3	<u>98.6 ± 0.2</u>	75.0 ± 0.8	80.5 ± 1.5	64.2 ± 3.0
RF	68.0 ± 2.3	<u>76.6 ± 0.8</u>	98.1 ± 0.3	73.4 ± 0.7	84.8 ± 0.8	70.3 ± 2.4
<b>GSDT</b>	<b>73.5 ± 1.5</b>	<b>77.2 ± 1.7</b>	<b>98.8 ± 0.6</b>	<u>76.0 ± 0.9</u>	<b>86.9 ± 1.2</b>	<b>78.2 ± 3.1</b>

# Structure Visualization

- GSDT provides a clear decision process
  - Each mean vector is a representative of the path

Feature  $x =$ 

0.54	-0.72	-0.09	-0.31	-0.56	-0.73	-0.99	-0.62	-0.35
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## Breast-cancer-wisconsin

### Features

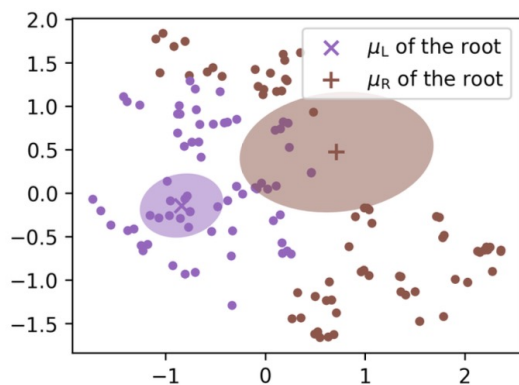
- $x_1$ : Clump thickness
- $x_2$ : Uniformity of cell size
- $x_3$ : Uniformity of cell shape
- $x_4$ : Marginal adhesion
- $x_5$ : Single epithelial cell size
- $x_6$ : Bare nuclei
- $x_7$ : Bland chromatin
- $x_8$ : Normal nucleoli
- $x_9$ : Mitoses

### Diagnosis (as labels)

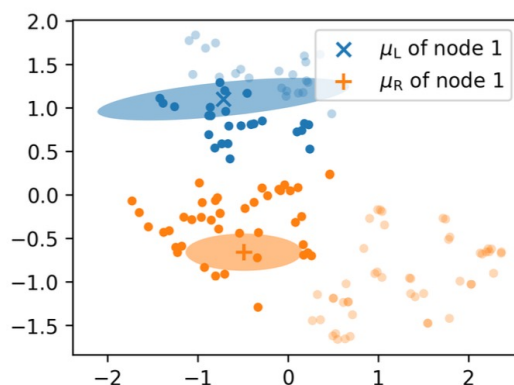
- $y_1$ : Benign
- $y_2$ : Malignant

# Learned Distributions

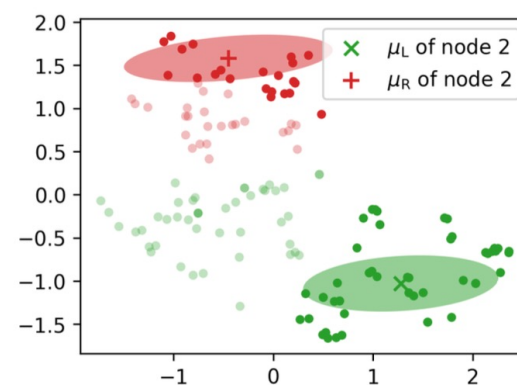
- GSDT learns meaningful node distributions
  - The root node splits examples horizontally
  - The internal nodes split examples vertically
    - Nodes 1 and 2 take different sets of examples



(b) Decision by the root.



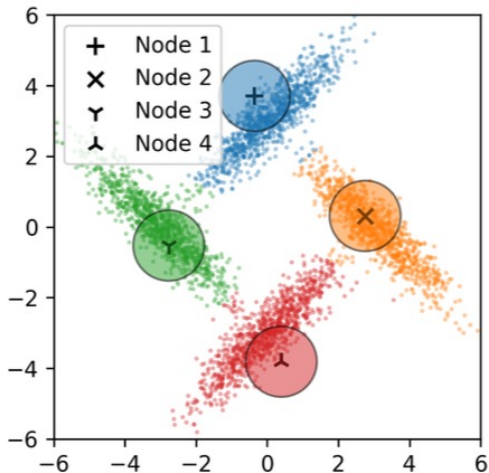
(c) Decision by node 1.



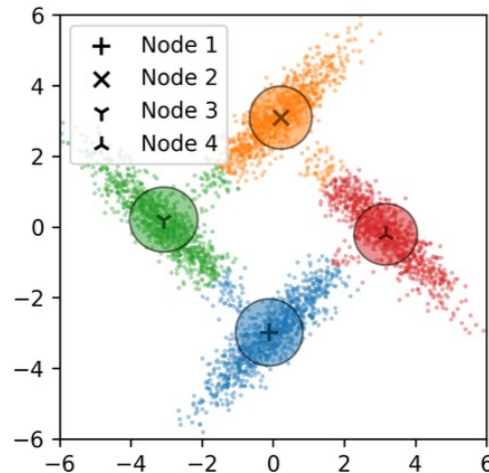
(d) Decision by node 2.

# Covariance Matrix

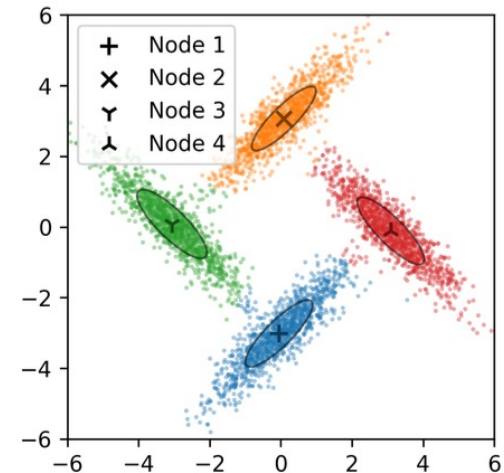
- Our low-rank perturbation makes the best fit
  - The identity and diagonal covariances are simple but fail to model the given distributions



(a) Identity.



(b) Only diagonal.



(c) Low-Rank Perturbed.



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# Conclusion

- **Gaussian Soft Decision Trees (GSDT)**
  - Our novel tree model for interpretable learning
  - Multi-branched structure with nonlinear decisions
- Main ideas
  - Gaussian decisions with low-rank perturbation
  - Path regularization
  - Post-optimization
- Experiments
  - GSDT outperforms baselines with interpretability



# Thank you!

Code and datasets:

<https://github.com/leesael/GSDT>