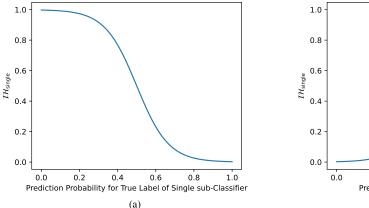
# Supplementary Materials for Paper: Dual Granular Balanced Deep Forest for Drug Combination Prediction

August 15, 2024



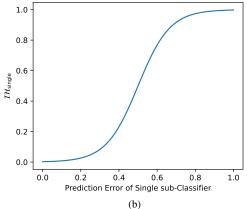


Figure S1. (a) Visualization of the IH function varying with the instance's prediction probability for the correct label. (b) Visualization of the IH function varying with the instance's prediction error.

$$Sensitivity = \frac{TP}{TP + FN}$$
 (S1)

$$Specificity = \frac{TN}{TN + FP}$$
 (S2)

$$\text{f1-macro} = \frac{1}{2} \sum_{i=1}^{2} 2 \times \frac{\text{Precision}_{i} \times \text{Recall}_{i}}{\text{Precision}_{i} + \text{Recall}_{i}} \tag{S3}$$

$$Gmean = \sqrt{Sensitivity \times Specificity}$$
 (S4)

$$AUROC = \int_0^1 TPR \, d(FPR) \tag{S5}$$

$$AUPR = \int_{0}^{1} Precision d(Recall)$$
 (S6)

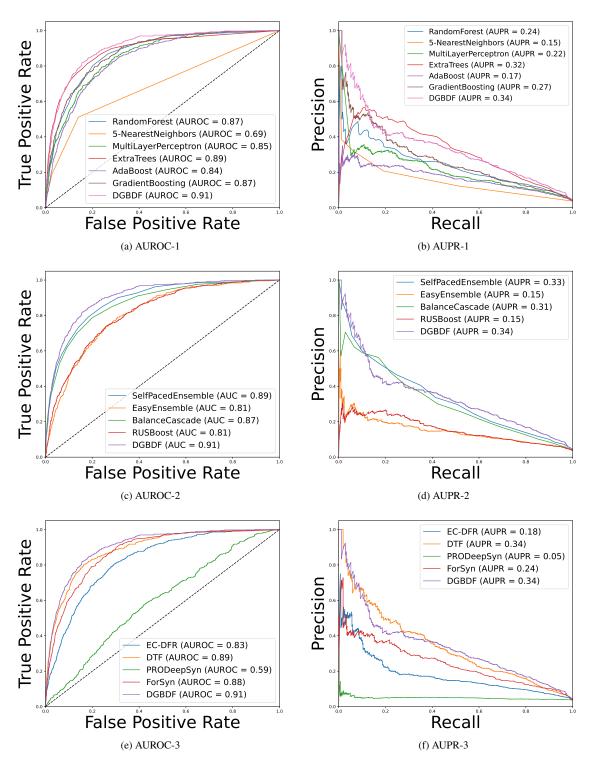


Figure S2. Comparison of AUROC and AUPR between DGBDF and various Canonical, Imbalanced Learning, and Existing Drug Combination Prediction Algorithms. DGBDF consistently outperforms other models in AUROC and AUPR.

TABLE SI Summary of Compared Algorithms

Algorithm	Description	
Self-Paced Ensemble (SPE)	Under-samples by coordinating data hardness through self-paced learning during the process of base classifier training, suitable for highly imbalanced and extremely skewed datasets.	
Easy Ensemble (EE)	Under-samples the majority class $(\mathcal{N})$ in the dataset during the training process of each base classifier and integrates the results.	
Balance Cascade (BC)	Guides the resampling process each time based on the existing trained base classifiers, based on EE.	
RUSBoost (RUSB)	Combines random under-sampling and boosting techniques, randomly under-sampling the majority class instances during each iteration.	
EC-DFR	An enhanced cascade deep forest model for drug combination prediction.	
DTF	Extracts latent representations of drugs and cell lines from drug combinations, synergizing data through tensor factorization, then predicts the synergistic state of drug combinations through a neural network.	
ProDeepSyn	Integrates PPI networks with omics data, constructing low-dimensional embeddings through GCN to predict anticancer synergistic drug combinations (SDCs).	
ForSyn	An improved deep forest method that includes two embedded units: the RF-CSU unit for handling data imbalance and the ETF-DR unit for handling high-dimensional features, to predict synergistic drug combinations in different cells.	

TABLE SII
Summary of Drug Combinations and Evidence

No	Drug 1	Drug 2	Evidence (PMID)	Description
1	PD173074	Erlotinib	37173994	Dual inhibition of FGFRs and EGFR by PD173074 and EGFR inhibitor erlotinib was synergistic in CCA
2	Entinostat	Paclitaxel	27177222	Entinostat significantly enhanced paclitaxel-mediated anti-proliferative/anti-survival effects on NSCLC cells in vitro and in vivo
3	Entinostat	Irinotecan	34258881	Combinations of irinotecan plus entinostat synergistically kill p53-positive CRC cells
4	Entinostat	Sorafenib	25371323	A combination of entinostat and sorafenib resulted in an additive to supra-additive growth inhibition of human bile duct adenocarcinoma cell lines
5	Bortezomib	Vinorelbine	25833390	Bortezomib increased the efficacy of ifosfamide and vinorelbine in pediatric Hodgkin lymphoma

#### **Algorithm S1** DGBDF

```
Input:
  1: \mathcal{D}: training set
Output:
  2: F: DGBDF
Function:
  3: \mathcal{F} \leftarrow \emptyset, \mathcal{IH} \leftarrow \emptyset, l \leftarrow 1
  4: while l \le d do // d is the max depth of the cascade
             if 1 = 1 then
                   \mathcal{L}^l, \mathcal{A}^l, \mathcal{IH}^l \leftarrow \text{Algorithm2}(\mathcal{D}, \mathcal{D}, 4)
  6:
  7:
                   \mathcal{L}^l, \mathcal{A}^l, \mathcal{IH}^l \leftarrow \text{Algorithm2}(\mathcal{D}', \mathcal{D}'_s, 4)
  8:
  9:
             end if
             Put \mathcal{L}^l into \mathcal{F}, \mathcal{IH}^l into \mathcal{IH}
10:
             Calculate performance M^l for \mathcal{F} by Eq. (13) with equal layer weights
11:
             if l \neq 1 and M^l - M^{l-1} < \epsilon then
12:
                   \mathcal{F} \leftarrow \mathcal{F} - \mathcal{L}^l
13:
                   break
14:
             end if
15:
             \mathcal{D}' \leftarrow concatenate(X, \mathcal{A}^l), Y
16:
             \mathcal{D}_s' \leftarrow \text{Algorithm3}(\mathcal{D}', \mathcal{IH}, nl, 5)
17:
             l \leftarrow l+1
19: end while
20: Weight per layer by Eq. (11)-(12)
21: return final cascade model F(x) = \frac{1}{l} \sum_{j=1}^{l} w^{j} * \mathcal{L}^{j}(x)
```

# **Algorithm S2** the generation of DGBDF's *l*-th layer

```
Input:
```

19: **return**  $\mathcal{L}^l$ ,  $\mathcal{A}^l$ ,  $\mathcal{IH}^l$ 

```
1: \mathcal{D}: training set
  2: \mathcal{D}_s: sampled training set
  3: n: number of ensemble classifiers per layer
Output:
  4: the DF's l-th layer \mathcal{L}^l
  5: the augmented features A^l
  6: the \mathcal{IH}^l of per ensemble classifier
Function:
  7: \mathcal{L}^l \leftarrow \emptyset, \mathcal{A}^l \leftarrow \emptyset, \mathcal{IH}^l \leftarrow \emptyset
  8: for i = 1 to n do
               Split \mathcal{D} by the 5-fold stratified cross validation to get \mathcal{T}_j, \mathcal{V}_j where 1 \leq j \leq 5
                \begin{aligned} & \mathcal{T}_{j}' \leftarrow \mathcal{T}_{j} \cap \mathcal{D}_{s} \\ & \mathcal{L}_{i} \leftarrow \emptyset, \mathcal{A}_{i} \leftarrow \emptyset, \mathcal{IH}_{i} \leftarrow \emptyset \end{aligned} 
 10:
 11:
               for j = 1 to 5 do
 12:
                      Train \mathcal{E}_{i,j} by Algorithm4(\mathcal{T}'_i, \mathcal{T}_j, 20) and get its prediction \hat{y}_{i,j} for \mathcal{V}_j
 13:
                      Calculate \mathcal{IH}_{i,j} for instances in \mathcal{V}_j by Eq. (3)
 14:
                      Put \mathcal{E}_{i,j} into \mathcal{L}_i, \hat{y}_{i,j} into \mathcal{A}_i, \mathcal{IH}_{i,j} into \mathcal{IH}_i
 15:
 16:
               Put \mathcal{L}_i into \mathcal{L}^l, \mathcal{A}_i into \mathcal{A}^l, \mathcal{IH}_i into \mathcal{IH}^l
 17:
 18: end for
```

# Algorithm S3 Balancing Module

# **Input:**

- 1:  $\mathcal{D}$ : training set
- 2:  $\mathcal{IH}$ : IH values set for all instances
- 3: K: number of all classifiers
- 4: b: number of bins

# **Output:**

5:  $\mathcal{D}_s$ : sampled training set

#### **Function:**

- 6:  $\mathcal{P} \leftarrow \text{positive in } \mathcal{D}, \mathcal{N} \leftarrow \text{negative in } \mathcal{D}, \mathcal{N}_s \leftarrow \emptyset, \mathcal{D}_s \leftarrow \emptyset$
- 7: Calculate  $\overline{\mathcal{IH}}$  of per instance in  $\mathcal{N}$  by Eq. (4)
- 8: Calculate per bin's capacity c by Eq. (8)
- 9: Calculate the percentiles  $p_0, p_1, \ldots, p_b$  of  $\overline{\mathcal{IH}}$  w.r.t. c
- 10: Cut majority set  $\mathcal{N}$  into b bins by Eq. (9) w.r.t.  $p_0, p_1, \ldots, p_b$
- 11: **for** i = 1 to b **do**
- 12: Calculate uncertainty U of instances in Bin-i by Eq. (7)
- 13: Calculate sampling probability pf per instance in Bin-i by Eq. (10)
- 14: Sample  $|\mathcal{P}|/b$  instances  $B_i'$  from Bin-i w.r.t. sampling probability
- 15:  $\mathcal{N}_s \leftarrow \mathcal{N}_s \cup B_i'$
- 16: Put  $B_i'$  into  $\mathcal{N}_s$
- 17: **end for**
- 18:  $\mathcal{D}_s \leftarrow \mathcal{N}_s \cup \mathcal{P}$
- 19: return  $\mathcal{D}_s$

# Algorithm S4 Balanced Ensemble Classifier

# **Input:**

- 1:  $\mathcal{T}$ : training set
- 2:  $\mathcal{T}_s$ : initial sampled training set
- 3:  $n\_estimators$ : number of decision trees

#### **Output:**

4:  $\mathcal{E}$ : balanced ensemble classifier

#### **Function:**

- 5:  $\mathcal{E} \leftarrow \emptyset$ ,  $\mathcal{IH} \leftarrow \emptyset$
- 6: **for** i=1 to  $n\_estimators$  **do**
- 7: Train decision tree  $t_i$  on  $\mathcal{T}_s$
- 8: Calculate  $\mathcal{IH}_i$  for instances in  $\mathcal{T}$  by Eq. (3)
- 9: Put  $\mathcal{IH}_i$  into  $\mathcal{IH}$
- 10:  $\mathcal{T}_s \leftarrow Algorithm3(\mathcal{T}, \mathcal{IH}, i, 5)$
- 11: Put  $t_i$  into  $\mathcal{E}$
- 12: **end for**
- 13: return  $\mathcal{E}$