Kernel ADASYN

Kernel Based Adaptive Synthetic Data Generation for Imbalanced Learning

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Content

Over-sampling method

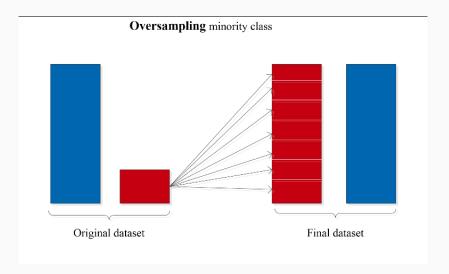
SMOTE

ADASYN

Kernal ADASYN



Over-sampling method



SMOTE

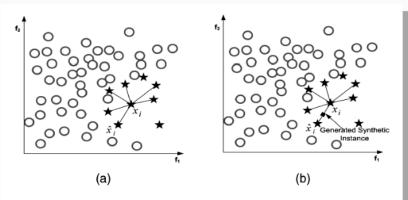


Fig. 3. (a) Example of the K-nearest neighbors for the x_i example under consideration (K = 6). (b) Data creation based on euclidian distance.

ADASYN

Calculate how many minority samples to generate ($N^+ * SR$).

For each minority sample x_i^+ , $i = 1, 2, ..., N^+$, find its K-nearest neighbors, N_i^{maj} of which from the majority.

$$\Gamma_i = \frac{\frac{N_i^{maj}}{K}}{Z}$$
, Z is a standardization factor to make sure $\sum \Gamma_i = 1$

$$g_i = \Gamma_i * N^+ * SR$$



ADASYN

Algorithm ADASYN.

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Input: Training set S = \{(x_i, y_i), i = 1, 2, ..., N, y_i \in \{+, -\}\}, majority N^-, minority N^+, N = N^+ + N^-, SR, K

Output: S' = \{(x_i, y_i), i = 1, 2, ..., N + N^+ * SR SR, y_i \in \{+, -\}\}
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Kernal ADASYN

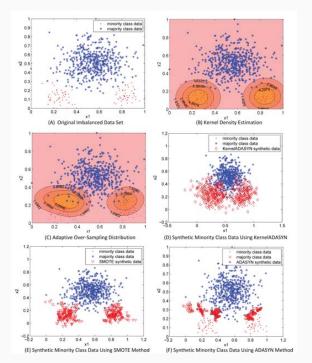
kernal density estimation:

$$\hat{\boldsymbol{\rho}}(\boldsymbol{x}) = \frac{1}{N+h} \sum_{i \in I_{+1}} \hat{r}_i \frac{1}{(\sqrt{2\pi}h)^n} exp(-\frac{1}{2} \frac{|\boldsymbol{x} - \boldsymbol{x}_i|}{h})$$

Not only adaptively shifting the classification decision boundary toward the difficult examples.

But also construct an adaptive over-sampling distribution to generate synthetic minority class data.







Datasets	Methods	OA	Precision	Recall	F-Measure	G-Mean
Pima Indians Diabetes	Decision Tree	0.7049	0.5803	0.5642	0.5696	0.6612
	SMOTE	0.6750	0.5284	0.6627	0.5867	0.6707
	ADASYN	0.6799	0.5374	0.5881	0.5606	0.6535
	KernelADASYN	0.7089	0.5731	0.6560	0.6112	0.6950
ILPD	Decision Tree	0.6515	0.3858	0.3687	0.3741	0.5274
	SMOTE	0.6326	0.3797	0.4518	0.4116	0.5629
	ADASYN	0.6357	0.3841	0.4542	0.4139	0.5642
	KernelADASYN	0.6571	0.4004	0.4598	0.4187	0.5722
Parkinsons	Decision Tree	0.8237	0.6611	0.6417	0.6398	0.7478
	SMOTE	0.8361	0.6627	0.7500	0.6938	0.8012
	ADASYN	0.8216	0.6456	0.6958	0.6611	0.7723
	KernelADASYN	0.8433	0.6672	0.7500	0.7000	0.8059
Vertebral Column	Decision Tree	0.7903	0.6711	0.7040	0.6837	0.7631
	SMOTE	0.7845	0.6491	0.7480	0.6910	0.7720
	ADASYN	0.8013	0.6703	0.7800	0.7195	0.7944
	KernelADASYN	0.7935	0.6559	0.7820	0.7194	0.7984
Breast Cancer	Decision Tree	0.9367	0.9117	0.9058	0.9077	0.9287
	SMOTE	0.9476	0.9171	0.9333	0.9244	0.9439
	ADASYN	0.9401	0.9080	0.9200	0.9133	0.9349
	KernelADASYN	0.9481	0.9115	0.9408	0.9258	0.9463
Breast Tissue	Decision Tree	0.8962	0.7179	0.8600	0.7644	0.8766
	SMOTE	0.9115	0.8530	0.6900	0.7391	0.8074
	ADASYN	0.9288	0.8172	0.8400	0.8196	0.8907
	KernelADASYN	0.8788	0.7313	0.7200	0.7038	0.8070
SPECT	Decision Tree	0.8143	0.5605	0.5037	0.5270	0.6695
	SMOTE	0.7820	0.4759	0.5593	0.5105	0.6830
	ADASYN	0.7902	0.4978	0.5595	0.5227	0.6880
	KernelADASYN	0.8174	0.5513	0.5185	0.5280	0.6755
Winning Time	Decision Tree	0	2	1	0	0
	SMOTE	0	2	2	0	0
	ADASYN	2	0	1	2	2
	KernelADASYN	5	3	4	5	5

Questions

