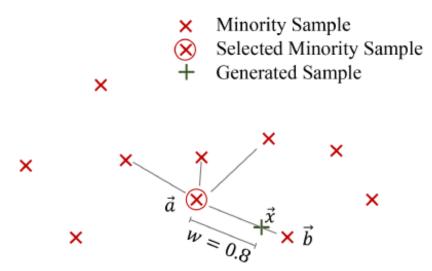


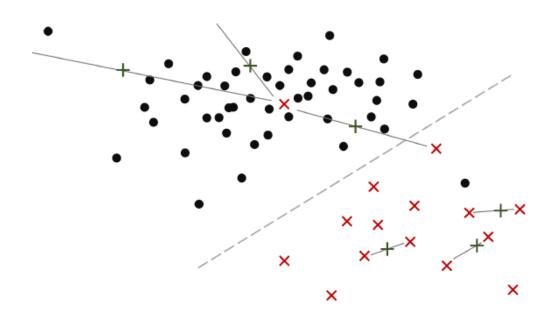
A bit of background: SMOTE



https://arxiv.org/pdf/1711.00837.pdf

SMOTE linearly interpolates a randomly selected minority sample and one of its k = 5 nearest neighbors

SVM, Borderline SMOTE

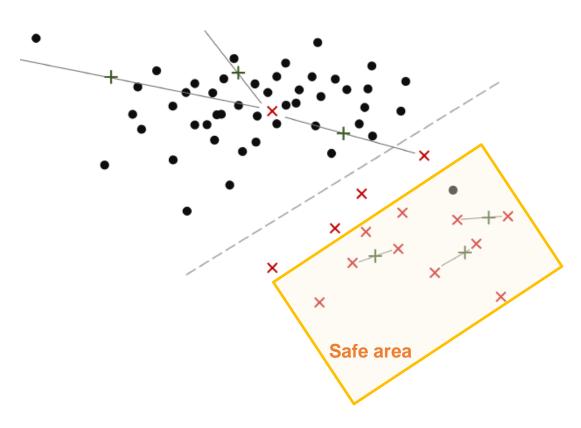


https://arxiv.org/pdf/1711.00837.pdf

- We should not create samples in areas that are safe
- We should create samples at the boundary



SVM, Borderline SMOTE

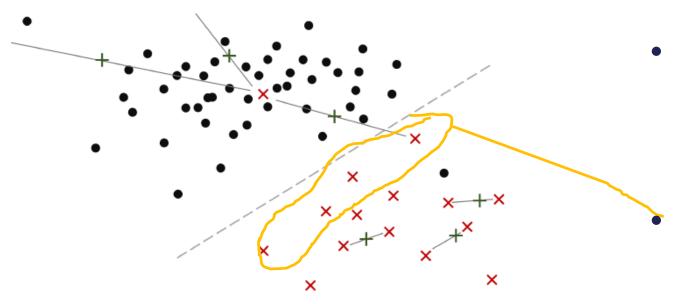


https://arxiv.org/pdf/1711.00837.pdf

- We should not create samples in areas that are safe
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SVM, Borderline SMOTE



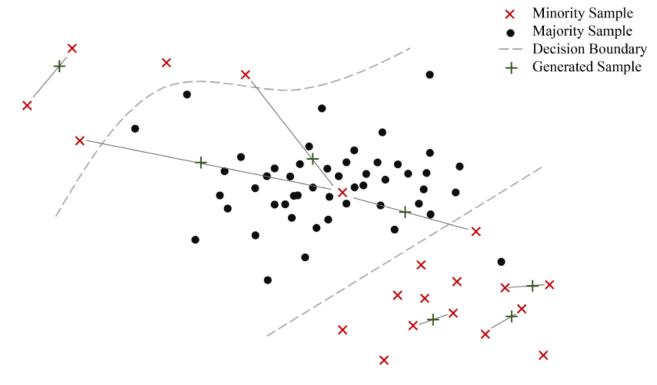
We should not create samples in areas that are safe

We should create samples at the boundary

https://arxiv.org/pdf/1711.00837.pdf



SMOTE, noise and intra-class clusters



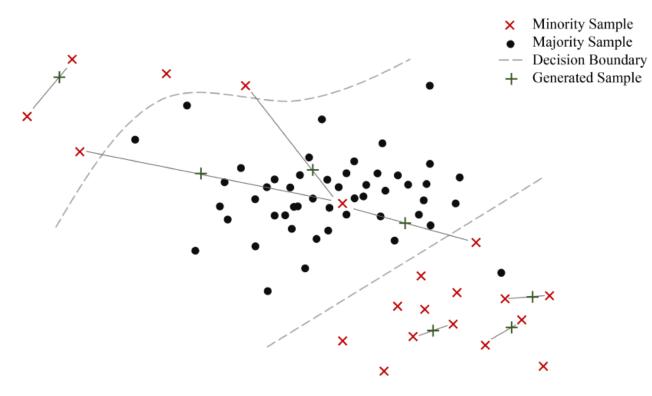
https://arxiv.org/pdf/1711.00837.pdf

 SMOTE might create noisy examples

 SMOTE does not contemplate intra-class clusters



Examples of intra-class clusters



https://arxiv.org/pdf/1711.00837.pdf

Fraudulent credit card transactions:

- Transactions in foreign countries
- High value transactions



K-Means SMOTE – the idea

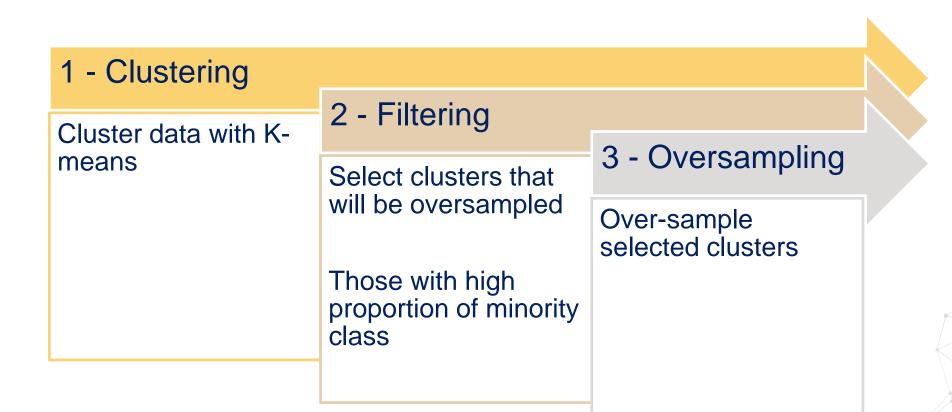
 Boost minority class regions by creating samples within naturally occurring clusters of the minority class.

Contemplate intra-class clusters

Avoid introducing noise



K-Means SMOTE - 3 steps





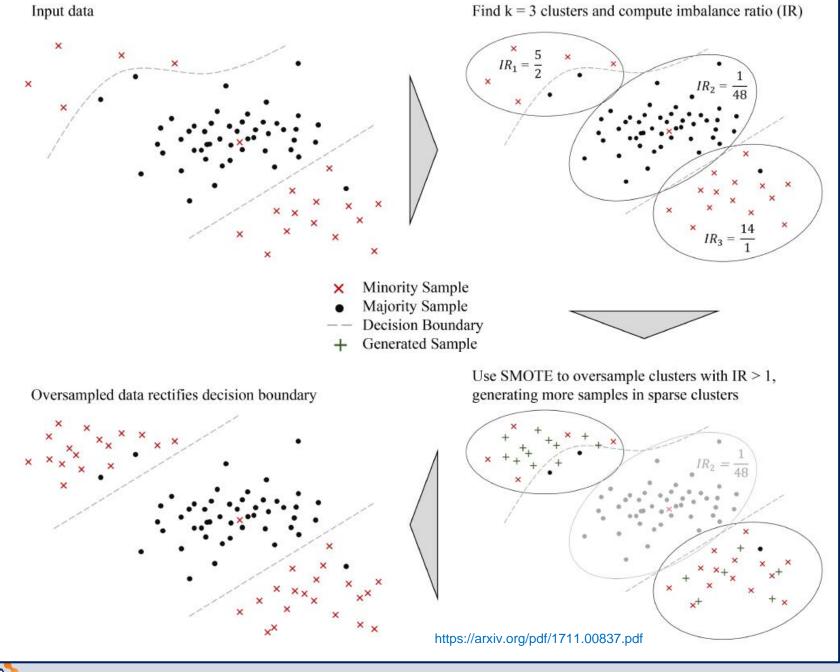
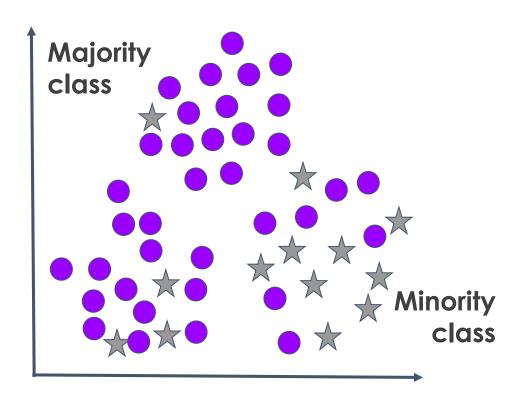


Diagram showing the 3 steps of K-means SMOTE

By default clusters where 50% are minority are selected by default

Clustering step



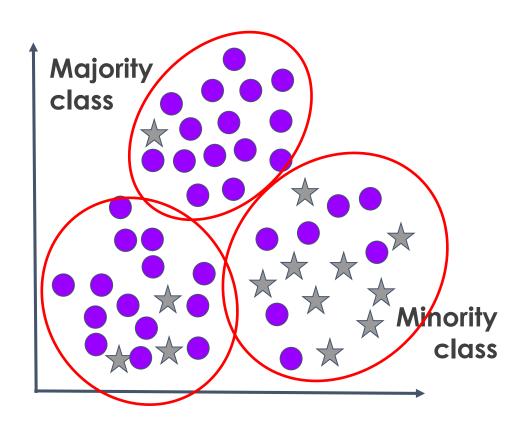
Find clusters → K means algorithm with entire dataset

We need to know K, or treat K as a hyperparameter

K-means may be slow to converge in huge datasets



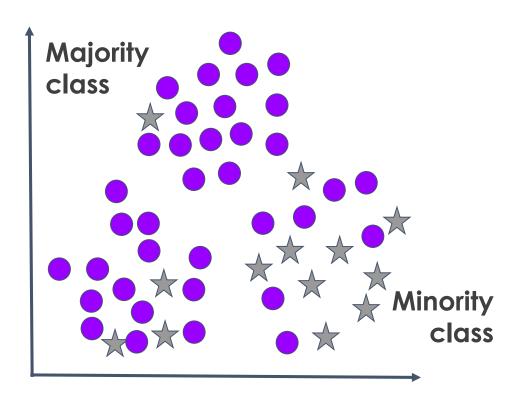
Filtering step



- By default select those clusters where 50% of observations belong to minority
- IR = 1 = # minority / # majority
- We can increase IR, thus we select clusters with higher proportion of minority class
- IR becomes another hyperparameter



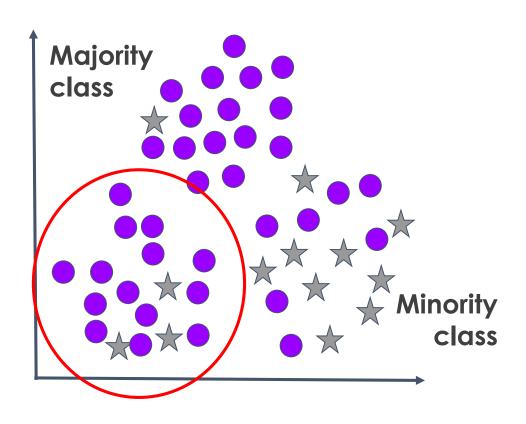
Over-sampling



- Determine how many samples to create in each cluster
- Asign weights to clusters, more weights to clusters with less minority observations.



Over-sampling

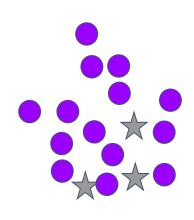


We take this cluster to proceed with the demo

- Determine how many samples to create in each cluster
- Asign weights to clusters, more weights to clusters with less minority observations.



Cluster weight



- 1. Determine Euclidean distance between all samples from minority
- 2. Determine Mean Euclidean distance
- 3. density = $\frac{Xmin}{L2mean^{number\ of\ features}}$
- 3. Sparsity = 1 / density
- 4. Cluster sparsity = Sparsity /sum(Sparsity all clusters)

Cluster weight

- 1. Determine Euclidean distance between all samples from minority
- 2. Determine Mean Euclidean distance



3. density =
$$\frac{Xmin}{L2mean^{number\ of\ features}}$$

- 3. Sparsity = 1 / density
- Cluster sparsity = Sparsity /sum(Sparsity all clusters)



Cluster weight

- 1. Determine Euclidean distance between all samples from minority
- 2. Determine Mean Euclidean distance (L2-mean)



3. density =
$$\frac{\# minority}{L2mean^{number of features}}$$

- 3. Sparsity = 1 / density
- Cluster sparsity = Sparsity /sum(Sparsity all clusters)

K-Means SMOTE

 Calculate the number of synthetic examples that need to be generated for each cluster

$$g_i = cs_i \times G$$

csi = cluster sparsity

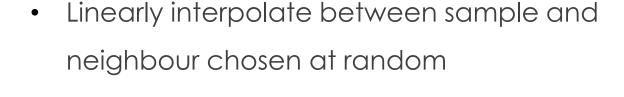
G = total number of samples to generate

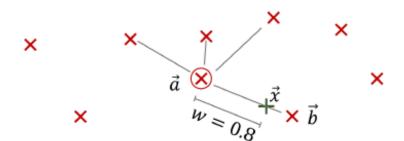
gi = number of samples to generate from cluster i



SMOTE with samples in cluster







 If cluster has few samples, we may not have enough neighbours

https://arxiv.org/pdf/1711.00837.pdf

 The number of neighbours becomes a hyperparameter



K-mean SMOTE – my thoughts

- The rationale is well thought, it makes sense
- The implementation of the algorithm is not super straight forward
- A lot of parameters to adjust
- Potentially some EDA to corroborate those parameters



Imbalanced-learn: KMeansSMOTE

```
sm = KMeansSMOTE(
    sampling strategy='auto', # samples only the minority class
    random state=0, # for reproducibility
    k neighbors=2,
    n jobs=None,
    kmeans estimator=KMeans(n clusters=3, random state=0),
    cluster balance threshold=0.1,
    density exponent='auto'
X res, y res = sm.fit resample(X, y)
```





THANK YOU

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