DSV: An Alignment Validation Loss for Self-supervised Outlier Model Selection



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Overview

Motivation: How can we search hyperparameters for self-supervised anomaly detection (SSAD)?

Why is it difficult? No labels are given at training

- \rightarrow No validation data can be used for HP search
- \rightarrow Unsupervised model selection problem

Proposed method: DSV

Validation loss for unsupervised model selection Focus on HPs in a data augmentation function E.g., the patch size in CutOut

Proposed Method

Discordance: The distance between ℓ and $Z_{\text{test}}^{(a)}$ **Separability:** The dist. between Z_{trn} and $Z_{\text{test}}^{(a)}$ on ℓ

Surrogate losses:

$$\mathcal{L}_{\text{dis}} = \frac{d(\mathcal{Z}_{\text{trn}} \cup \mathcal{Z}_{\text{aug}}, \mathcal{Z}_{\text{test}})}{d(\mathcal{Z}_{\text{trn}}, \mathcal{Z}_{\text{aug}})} \qquad \mathcal{L}_{\text{sep}} = \frac{\text{std}(\text{proj}(\mathcal{Z}_{\text{test}}))}{d(\mathcal{Z}_{\text{trn}}, \mathcal{Z}_{\text{aug}})}$$

Simplified final loss: $\mathcal{L}_{DSV} = \mathcal{L}_{dis} - \mathcal{L}_{sep}$ / \mathcal{L}_{dis}



Problem Definition

Problem: Unsupervised model selection for SSAD

Given

- 1. Data augmentation function f_{aug}
- 2. Set Φ of different detector models that are trained with f_{aug} with different HPs
- 3. Training data \mathcal{D}_{trn} (w/o labels)
- 4. Test data \mathcal{D}_{test} (as a transductive setting)

Find

Detector $\phi^* \in \Phi$ with best accuracy on $\mathcal{D}_{\text{test}}$

Main Ideas

Proposition: Functional alignment

Let f_{gen} be the anomaly-generating function Detector ϕ will be accurate if it is trained with f_{aug} that is aligned w/ f_{gen}

Q: How can we define the alignment? Idea 1: Compute the embedding distance btw. data augmented by $f_{aug}(\mathcal{D}_{trn})$ and true anomalies in \mathcal{D}_{test}

Experiments

Task: Industrial image anomaly detection
Detector: ResNet18-based classifier model
Trained to minimize the cross entropy between normal and augmented data
Augmentation functions: CutOut, CutPaste, etc.

Q: We don't know which are the true anomalies... Idea 2: Use \mathcal{D}_{test} as a whole, not only anomalies, comparing $\mathcal{D}_{trn} \cup \mathcal{D}_{aug}$ and \mathcal{D}_{test}

Steps for Approximation

Step 1: Decompose alignment into two terms: **Discordance** h_d and **Separability** h_s

Step 2: Design approximate **surrogate losses** $h_d \rightarrow \mathcal{L}_{dis}$ and $h_s \rightarrow \mathcal{L}_{sep}$

Step 3: Propose the DSV loss w/ \mathcal{L}_{dis} and \mathcal{L}_{sep}

Target HPs: Patch size of augmentation

| $f_{ m aug}$ | Avg. | Rand. | Base | MMD | STD | MC | SEL | HITS | DSV |
|--------------------------|-------|-------|-------|-------|-------|-------|-------|-------|--------------|
| CutOut | 0.739 | 0.776 | 0.741 | 0.735 | 0.739 | 0.749 | 0.727 | 0.757 | 0.813 |
| CutAvg | 0.739 | 0.817 | 0.721 | 0.692 | 0.745 | 0.751 | 0.744 | 0.742 | <u>0.806</u> |
| $\operatorname{CutDiff}$ | 0.743 | 0.711 | 0.739 | 0.730 | 0.744 | 0.747 | 0.741 | 0.777 | 0.811 |
| CutPaste | 0.788 | 0.841 | 0.694 | 0.756 | 0.818 | 0.862 | 0.830 | 0.850 | 0.884 |

