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SOoD: Self-Supervised Out-of-Distribution Detection Under Domain Shift for Multi-Class Colorectal Cancer Tissue Types

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Abstract

The goal of out-of-distribution (OoD) detection is to identify unseen categories of inputs different from those seen during training, which is an important requirement for the safe deployment of deep neural networks in computational pathology. Additionally, to make OoD detection applicable in clinical applications, one may encounter image data distribution shifts. This paper argues that practical OoD detection should handle both semantic shift and data distribution shift simultaneously. We propose a new selfsupervised OoD detector for colorectal cancer tissue types based on a clustering scheme. Our work's central tenet benefits from multi-view consistency learning with a supplementary view based on style augmentation to mitigate domain shift. The learned representation is then adapted to minimize images' predictive entropy to segregate indistribution examples from OoDs on the target data domain. We evaluated our method on two public colorectal tissue types datasets. Our method achieved state-of-the-art OoD detection performance over various self-supervised baselines. The code, data, and models are available at https: //github.com/BehzadBozorgtabar/SOoD.

1. Introduction

Colorectal Cancer (CRC) is considered one of the most occurring cancers worldwide, and early-stage CRC diagnosis can significantly improve the chances for therapy of patients [6]. In CRC, the Tumor MicroEnvironment (TME) analysis plays an essential role in cancer grading, and prognostication [23]. Thus, developing automatic tissue phenotyping in Whole Slide Images (WSIs) is of great importance. In recent years, deep learning models have been widely developed for multi-class tissue type classification [24, 38, 2]. While these deep models implicitly assume that the datasets are independent and identically distributed (i.i.d), in practice, collected datasets are typically far from

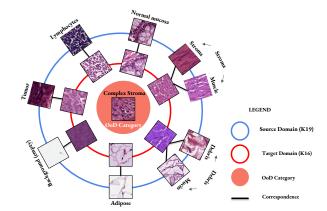


Figure 1: Motivation of the proposed OoD detection under domain shift. SOoD seeks to handle both data distribution shift and semantic shift. Histological images of different tissue types present high appearance variability between the source domain (K19) and the target domain (K16), and the target domain contains an additional unknown class (complex stroma).

the i.i.d assumption. Histological images present high appearance variability in a real-world scenario due to acquiring data in various conditions, including different scanners or staining procedures. To mitigate this issue, domain adaptation techniques [14, 13, 1, 36] disclose the inference-time data (target domain) to model for adapting the representation from the training data (source domain). Nevertheless, most domain adaptation methods [14, 13] assume a closed-set scenario, where the source and target domains share the same distribution of classes (label set).

In clinical routine, a model is often exposed to new data with unknown categories, e.g., tissues from specific cancer subtypes. Thus, making a model robust to the presence of out-of-distribution (OoD) samples and sidestep potentially inaccurate predictions is crucial for the model's safe deployment. Although the task of OoD detection has seen considerable progress [17, 34, 40], developing practical OoD

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detection for computational pathology has been a particularly challenging problem for two reasons. *First*, deep neural networks (DNNs) often make overconfident predictions to unknown inputs [31]. *Second*, due to the domain discrepancy mentioned earlier, OoD detectors may mistakenly detect a test sample from known categories but have a different style/domain as an OoD. Fig. 1 shows the correspondence of different tissue types across two CRC datasets, Kather-19 (K19) [24] and Kather-16 (K16) [25], as the source domain and target domain. Histological images present high appearance variability between two data domains, and the target domain contains an additional unknown class (complex stroma).

To address the limitations of current OoD detection methods, we propose **SOoD**, short for **S**elf-supervised **O**ut**o**f-**D**istribution detection under domain shift for multi-class colorectal cancer tissue types, a new self-supervised OoD detector to mitigate both semantic shift and data distribution shift. We illustrate the pre-training stage of SOoD in Fig. 2 and present a pseudo-code implementation in Algorithm 1. In summary, we highlight the contributions below:

- Our method (SOoD) is the first work to consider the problem of multi-class OoD detection under domain shift for clinical applications to the best of our knowledge;
- The proposed self-supervised OoD method builds upon multi-view consistency paradigm with complementary style augmentation to mitigate domain shift, as opposed to current OoD detections, which focus on a single image domain;
- We propose a new self-training scheme for OoD detection via minimizing images' predictive entropy of unlabeled images to segregate in-distribution examples from OoDs on the target data domain. Our method does not require OoD samples during training and is capable of working with unlabeled source datasets alleviating costly annotations;
- Experimental results show consistent improvement of proposed OoD detection performance over state-of-the-art (SOTA) self-supervised methods [8, 7, 42, 3] on two hematoxylin & eosin (H&E) stained CRC tissue types datasets [24, 25].

2. Related Work

The problem of OoD detection has seen considerable progress in computer vision and medical image analysis, as OoD detection is crucial for the safe deployment of deep learning systems. The related work in this area is sizable. Thus, we mainly focus on the recent deep learning-based methods in supervised [15] and unsupervised [44] settings. Most of the current OoD detection solutions presume access to the OoD datasets during training [22, 47] or validation steps [27, 26, 34] that are not well suited for the general use of OoD detection in real-world applications. Some interesting methods [16, 27, 26] benefit from adversarial samples via perturbation of the training samples to improve the robustness of their network, which results in higher training time and suboptimal solutions.

On the other hand, recent methods [17, 34, 40] rely on generative or reconstruction-based training schemes [39, 49], deep one-class classifiers [35, 29], and, more recently, self-supervised approaches [15, 5, 30, 3]. Overall, the underlying rationale behind those methods is modeling the representation of in-distribution data either using a oneclass [35] or multi-class setup [4], and then a detection function is usually defined to detect OoDs. However, most previous OoD detection methods assume that the training and test data would follow a similar distribution (style/domain). This assumption can negatively impact OoD detection as OoD detectors may erroneously detect a test sample from known classes but have a different style as an OoD class. A possible solution would be to use additional data from the new target domain and formulate this problem as openset domain adaptation [37, 32], where the source domain contains in-distribution labeled data and the target domain contains novel classes in addition to the classes present in the source domain. Nevertheless, it would require labeled source data and costly annotations by domain experts. Recent studies have shown that contrastive training [8, 19, 9]significantly improves OoD detection [45, 42]. These methods attempt to learn representation based on attracting similar views of a sample and repelling disagreeing views from each other. However, current contrastive training methods are incentivized to learn features from a single image domain.

3. Method

We start by motivating our approach before explaining the methodological details. The main goal of OoD detection under domain shift is to learn domain-invariant representation between one specific domain (i.e., source domain) and a testing domain (i.e., target domain) so that an OoD model can robustly leverage such invariances to a new unlabeled target domain. The domain invariance is often ignored or not formulated in previous OoD detection methods. As a result, *OoD detectors may mistakenly detect a new test example from known classes but have a different style as an OoD class.*

This problem setting is also different from typical unsupervised domain adaptation (UDA) approaches as a new target domain contains an additional unknown class. Besides, unlike UDA methods, we formulate our proposed OoD to deal with the unlabeled source data, which is highly demanded in practical applications. In this work, we revisit current state-of-the-art contrastive learning-based selfsupervised methods [10, 7, 8, 18] for the OoD task using only positive pair samples. In particular, we extend the two-view consistency learning paradigm based on selfaugmentation to a multi-view version with style augmentation as the new complementary view.

The objective of the pre-training stage is to simultaneously learn domain-invariant features and consistent cluster assignments between multiple views of the same tissue image in an entirely unsupervised setting (Sect. 3.1). The learned representation is then adapted using a *self-training* scheme on the unlabeled target domain images. Typically, one can use the most probable cluster predicted by the network as pseudo labels. Since the pseudo labels are often noisy, we propose to segregate known and unknown samples using the entropy of the clustering output and opt for only highly confident (lower entropy) target images for pseudo labeling. More specifically, we perform entropy minimization on selected unlabeled target images w.r.t source prototypes from (Sect. 3.1) to segregate known categories from OoDs (Sect. 3.2). Finally, we define the OoD function to detect OoDs.

3.1. Multi-View Consistency

We revisit the recent self-supervised clustering scheme [7], which clusters the data while imposing an agreement between cluster assignments obtained from different augmentations of the same image. Specifically, we address the limitation of the current self-augmentation consistency learning paradigm in the presence of data domain shift. To do so, we extend typical two-view consistency learning to a multi-view version with style augmentation of the target domain as the new complementary view. Each source domain image x_s is transformed into two in-domain augmented views, including weakly augmented view x_{sw} and heavily augmented view x_{sh} , and the encoder output of the weakly augmented view provides a pseudo label for the predictions on heavily augmented view. Furthermore, we add an additional view based on style augmentation x_{style} to make the model robust against domain shift, especially in the absence of labeled data. As for the style augmentations, weakly augmented images are mapped from the source domain to the target domain via a pre-trained CycleGAN model [48]. Such a new view makes the model invariant to the image style by further covering the target data distributions and adding the regularization effect through multi-view consistency learning.

More precisely, we apply a non-linear mapping f_{θ} to the multi-view augmented images to match their representations to K dimensional features. The non-linear encoder f_{θ} includes the convolutional neural network (CNN) backbone followed by a 2-layer MLP network. Given an image

Algorithm 1: SOoD PyTorch pseudocode w/o multi-crop (pre-training stage).

_ I	nulli-crop (pre-training stage).
	Input: S : unlabeled or partially labeled source samples, $trslt$:
	pre-trained style transformer on $\mathcal S$ and unlabeled target
	samples \mathcal{T}, f_{θ} : encoder network
	Output: updated f_{θ}
	Parameter : tp: temperature, λ_1, λ_2 : weights for the loss
	terms, sinkhorn: Sinkhorn-Knopp function
1	for x in loader do // load a minibatch with n
	samples from ${\cal S}$
2	<pre>x_sw = weak_augment(x)// augmented views</pre>
3	$x_{sh} = heavy_{augment}(x)$
4	$x_style = trslt(x_sw)$
5	scores_sw, scores_sh, scores_style = $f_{\theta}(x_sw)$, $f_{\theta}(x_sh)$,
	$f_{ heta}(x_style)//$ output n-by-K
6	pseudo_sw, pseudo_sh, pseudo_style = sinkhorn(scores_sw),
	<pre>sinkhorn(scores_sh), sinkhorn(scores_style)// apply</pre>
	sinkhorn to generate pseudo label
7	$\ell_{heavy} = H(pseudo_sw, scores_sh)/2 + H(pseudo_sh,$
	scores_sw)/2
8	ℓ_{style} = H(pseudo_sw, scores_style)/2 + H(pseudo_style,
	scores_sw)/2
9	$\ell_{mv} = \lambda_1 \ell_{heavy} + \lambda_2 \ell_{style}$
10	$\ell_{mv}.backward()$ // back-propagate
11	$update(f_{ heta}) // encoder update$
12	def H(pseudo, score):
13	<pre>pseudo = pseudo.detach()// stop gradient</pre>
14	pred = softmax(score / tp, dim=1)
15	return -(pseudo * log(pred)).sum(dim=1).mean()

x from one of the three different augmentations of the input source image, we compute its cluster assignments (codes) by matching its feature representations to a set of K trainable prototypes $\{c_1, \dots, c_K\}$. These soft assignments are in the form of probability distributions over K dimensions. Then, the probability P is obtained by normalizing the output of the encoder f_{θ} with a softmax function:

$$P(x)^{(i)} = \frac{\exp\left(f_{\theta}(x)^{(i)} / \tau\right)}{\sum_{k=1}^{K} \exp\left(f_{\theta}(x)^{(k)} / \tau\right)}$$
(1)

where τ is a temperature parameter [46]. As in [7], features before last linear layer of f_{θ} and prototypes are ℓ_2 normalized. We optimize the multi-view consistency loss ℓ_{mv} w.r.t. the parameters of the encoder θ . Thus, the encoder output of the weakly augmented view provides a pseudo label for predictions of two other augmented views from the source image based on heavy and style augmentation through the cross-entropy losses ℓ_{heavy} and ℓ_{style} ,

$$\min_{\theta} \ell_{mv} = \lambda_1 \ell_{heavy} + \lambda_2 \ell_{style} \tag{2}$$

where λ 's denote the weights for the heavily augmented view loss ℓ_{heavy} and style augmented view loss ℓ_{style} . The pseudo label is obtained by applying the iterative *Sinkhorn*-

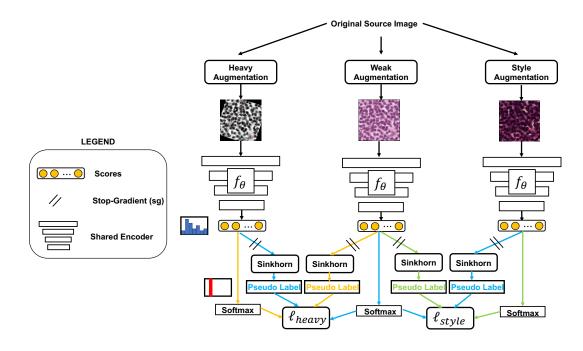


Figure 2: **The overview of the pre-training stage of SOoD**. Augmented views for input source images are generated, and the multi-view consistency loss is optimized. The encoder output of the weakly augmented view provides a pseudo label for predictions of two other augmented views from the source image based on heavy and style augmentation.

Knopp algorithm [12] on the output of the encoder f_{θ} to select all prototypes the same amount of time.

Details of the loss terms. We first describe the loss term ℓ_{heavy} for the heavily augmented view, and a similar formula holds for the loss term of the style augmented view ℓ_{style} . For the heavily augmented view, we use RandAugment [11] that mainly deals with color intensity and geometrical transformations. We first compute for an unlabeled weakly augmented view its pseudo label $\hat{P}(x_{sw}) \in$ $\{1, \dots, K\}$ w.r.t the K prototypes. This is achieved by first applying a stop-gradient (sg) operator on the encoder output and then using iterative Sinkhorn-Knopp algorithm [12], $P(x_{sw}) = \operatorname{sinkhorn}(\operatorname{sg}(P(x_{sw}))))$. Then, we optimize the encoder to match the heavily augmented view prediction $P(x_{sh})$ to the pseudo label $\hat{P}(x_{sw})$ using the cross-entropy loss. In practice, we additionally benefit from a multi-crop data augmentation strategy [7] such that from a given image, we generate a set V of different positive views. This set contains two anchor views and several local image crops of smaller resolution. The predictions of all crops are attracted to the anchor views to further improve the quality of the learned embeddings. We minimize the loss ℓ_{heavy} with stochastic gradient descent:

$$\ell_{heavy} = \min_{\substack{\theta \\ x_{sw} \in \left\{x_{1}^{a}, x_{2}^{a}\right\}}} \sum_{x'_{sh} \in V, x'_{sh} \neq x_{sw}} H\left(\hat{P}\left(x_{sw}\right), P\left(x'_{sh}\right)\right)$$
(3)

where $H(a, b) = -a \log b$, and x_1^a and x_2^a denote the anchor views. Similar formula holds for ℓ_{style} to align the style augmented view prediction $P(x_{style})$ to the pseudo label $\hat{P}(x_{sw})$ using the cross-entropy loss:

$$\ell_{style} = \min_{\substack{\theta \\ x_{sw} \in \left\{x_1^a, x_2^a\right\}}} \sum_{x'_{style} \in V, x'_{style} \neq x_{sw}} H\left(\hat{P}\left(x_{sw}\right), P\left(x'_{style}\right)\right)$$
(4)

The style augmented view loss ℓ_{style} complements heavily augmented view loss ℓ_{heavy} by making the encoder robust to style variation present in the target domain. Following [18], we use a symmetrized loss for both loss terms (Eq. 3 & Eq. 4) as symmetrization helps boost accuracy (see Algorithm 1).

3.2. Self-Training via Entropy Minimization

We incorporate an additional self-training criterion on the target domain into our model to further facilitate OoD detection. For this purpose, the pre-trained encoder f_{θ} from the previous step is applied on unlabeled target images to generate the pseudo-label for target samples, which are then used to fine-tune the encoder. Since we are not using label information on the target domain, we use the entropy of the cluster assignment to draw a boundary between indistribution and OoDs such that we expect that the entropy of OoDs is larger than entropy for the in-distribution samples. To determine the optimal threshold for the entropy,

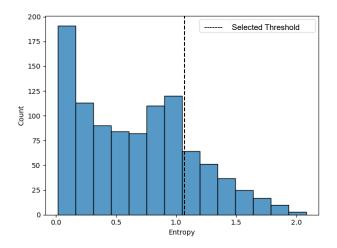


Figure 3: The entropy histogram of cluster assignments for the style augmented images. We set the threshold ρ to 1.07 such that the entropy of $\simeq 80\%$ of training examples will be lower than ρ . This threshold is used to select highly confident target examples for pseudo labeling.

we first compute the entropy of the style augmented images used for training and select a threshold ρ such that the majority of the style augmented images ($\simeq 80\%$)¹ have an entropy lower than ρ . Then we apply this threshold to the unlabeled images from the target domain to select highly confident samples for pseudo labeling. We perform an analysis of ρ in Fig. 3. For the self-training, we perform predictive entropy minimization on pseudo-labeled target data to make them tighter clustered around the source prototypes $\{c_1, \dots, c_K\}$. This increases the confidence of cluster predictions and identifies OoDs if they have different characteristics compared to in-distribution samples. The prototypes are kept fixed during self-training, and only the parameters of f_{θ} are updated. We minimize the entropy loss for the self-training step ℓ_{st} as follows:

$$\ell_{st} = \mathbb{E}_{x_t \sim \mathcal{T}} \left[\sum_{k=1}^{K} -P\left(x_t\right)^{(k)} \log P\left(x_t\right)^{(k)} \right]$$
(5)

where $P(x_t)^{(k)}$ is the probability obtained by the encoder shows unlabeled target sample x_t matches with cluster prototype c_k .

Inference: At inference time, a test image x_{test} is passed through the trained encoder f_{θ} to obtain its feature representation $v_{test} = f_{\theta}(x_{test})$. v_{test} is then compared with the top M similar features $\{v_m\}$ of the target domain's training samples based on the cosine similarity. An OoD

detection score $\mathcal{S}(\cdot, \cdot)$ is computed as follows:

$$\mathcal{S}\left(v_{test}; \{v_m\}\right) := -\frac{1}{M} \sum_{m=1}^{M} sim(v_m, v_{test}) \qquad (6)$$

where $sim(\mathbf{a}, \mathbf{b}) = \frac{\mathbf{a}^T \mathbf{b}}{\|\mathbf{a}\| \|\mathbf{b}\|}$ and $\mathcal{S}(\cdot, \cdot)$ is normalized using the maximum and minimum scores of the set such that $\mathcal{S}(\cdot, \cdot) \in [0, 1]$. Intuitively, the scores of OoD samples should be larger than the scores from in-distribution ones.

4. Experiments

Datasets and Evaluation Metrics. We evaluate SOoD on two H&E stained publicly available CRC datasets, Kather-19 (K19) [24] and Kather-16 (K16) [25], as the source domain and target domain. K16 dataset contains 5,000 images patches of 150×150 pixels each $(74\mu m \times 74\mu m)$ from H&E WSIs, while K19 dataset contains 100,000 H&E stained patches at (0.5 μ m/pixel). There is a data distribution shift across two image domains together with a semantic shift of tissue phenotypes. Incorporating expert pathologists' feedback [1], we group debris/mucus and stroma/muscle as debris and stroma, respectively, to correspond between the two datasets. As a result, we end up with seven tissue categories shared between two domains, including (tumor, stroma, lymphocytes, debris, normal mucosa, adipose, and background or empty class). The target domain contains an additional tissue category of complex stroma that is not present in the source domain, and we consider this tissue type as OoD class. In total, we end up with 11,495 training images (7,995 from the source domain and 3,500 from the target domain) without using OoDs. For the validation set, we use 997 images from the source domain (pre-training and self-training), 621 images from the target domain for t-SNE visualization purpose. For the test set, we use 879 images from the target domain, including 438 OoD images. The rest of the test images are equally distributed between seven in-distribution classes. We use OoD detection metrics: area under the ROC curve (AUC) and area under the precision-recall curve (AUPRC) and present mean \pm std on the test set for all experiments over three runs. Our experiments follow the setting for multi-crop using two anchor views at resolution 144 \times 144 pixels and multiple small crops (local views) of resolution 96 \times 96 pixels.

Implementation Details. Our implementations are based on PyTorch 1.9 [33]. We adopt the ResNet18 [20] as the backbone network for SOoD. All networks are trained using SGD optimizer (*momentum* = 0.9), with a weight decay of 1e - 6 and a learning rate of 0.06. A cosine scheduler is used during the training. A hyper-parameter search was conducted to find the optimal batch size (64), τ (0.1), prototypes (16), λ_1 (1) and λ_2 (1). Also, we found

¹This ratio is determined based on the distribution of unknown samples.

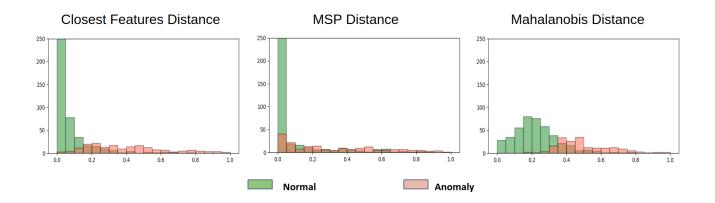


Figure 4: **The histograms of OoD detection scores** for in-distribution (normal) and OoDs (anomalies) on the target set (K16). We compare our OoD detection score (left) to other anomaly scores (middle-right). Our OoD detection score clearly discriminates in-distribution and OoD test images.

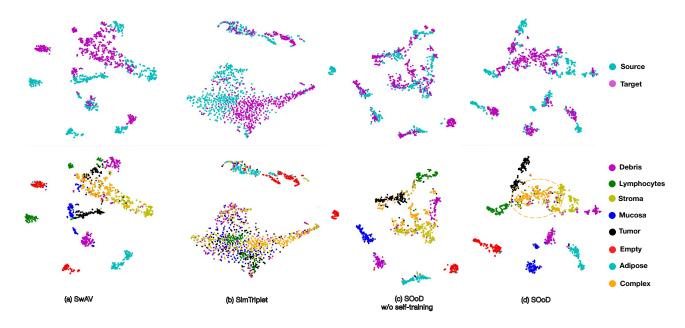


Figure 5: **The t-SNE [43] visualization** of the feature representations extracted by the encoder trained on the source (K19) and target sets (K16) for domain alignment (top row) and the different classes' representations (bottom row). We compare our method (c-d) to other SOTA self-supervised methods (a-b). Further fine-tuned with self-training, our model learns domain-invariant representation and can separate OoDs from in-distribution test samples (highlighted with dashed **orange** contour).

the optimal number of nearest neighbors for k-NN and $M \in \{5, 10, 20, 50\}$ in Eq. 6 and set it to 10 for all baselines. We conduct ablation studies for the chosen λ 's and the number of prototypes. We apply random resize crops and horizontal flips augmentation for the weakly augmented view. Besides, we use RandAugment [11] for obtaining the heavily augmented view. Our model has been pre-trained for 300 epochs using ℓ_{mv} and then fine-tuned by minimizing ℓ_{st} for additional 20 epochs with a lower learning rate of

0.001. For a fair comparison with [7], we set the Sinkhorn regularization parameter ϵ to 0.05 and use three iterations for all runs. To ensure consistent and comparable comparisons, we use the same experimental setup for all baselines.

Comparison with SOTA Methods. Since no prior work has been done for our specific setup, we provide our baselines for OoD detection under domain shift based on stateof-the-art self-supervised learning methods. We first validate the SOoD framework used in this study with state-

* p < 0.001; a bilateral Welch t-test with respect to the top result.

[†] We compute the OoD detection score before self-training, using only the pre-trained model.

(A)		Source Only			Target Only		Both Domains		
Method		AUROC		JPRC	AUROC	AUPRC	AUROC	AUPR	С
SimCLR [9]		72.79 \pm	1.64 66	$.84 \pm 1.93$	87.05 ± 1.24	81.15 ± 2.10	88.75 ± 1.1	4 83.51	± 1.97
SwAV [7] [†]		70.70 \pm	1.76 60	$.80 \pm 1.80$	84.42 ± 1.32	76.73 ± 2.13	76.84 ± 1.6	6 69.68 :	± 2.17
SwAV [7]	‡	$78.50~\pm$	1.55 72	$.57 \pm 2.09$	87.34 ± 1.25	80.81 ± 2.12	85.73 ± 1.2	8 81.96 :	± 1.88
CDMSAD	• [41]	72.85 \pm	1.65 66	$.85 \pm 1.96$	84.03 ± 1.34	78.64 ± 2.03	69.34 ± 1.8	6 58.47 :	± 1.70
(B)	CSI	[42]	GOAD [3]	SimTriplet [28]	SwAV [7]	DINO [8]	Our model w/o sel	lf-training [†]	SOoD [‡]
AUROC	89.64	± 1.95	70.56 ± 2.42	69.53 ± 1.32	67.58 ± 2.77	52.56 ± 0.43	88.38 ± 1	.30 92	$\textbf{2.77} \pm \textbf{0.48}$
AUPRC	86.32	± 2.21	63.11 ± 1.84	60.01 ± 1.42	60.44 ± 2.81	46.52 ± 0.29	80.43 ± 2	.84 90	0.90 ± 1.00
C)	Sup.	Tr100%	Sup. Src10	0% SwAV-100	% DINO-100%	SOoD-100%	SOoD-20%	SOoD-10%	SOoD-1%
near (ACC)	78.3	1 ± 5.98	65.13 ± 3.12	57 48.83 ± 1 .	83 42.03 \pm 8.51	$\textbf{73.24} \pm \textbf{0.39}$	$\textbf{73.39} \pm \textbf{0.69}$	73.24 ± 0.82	62.59 ± 1.42
near (F1)	78.2	7 ± 5.86	$62.89 \pm 1.$	15 45.67 ± 2 .	31 37.17 \pm 8.49	$\textbf{69.88} \pm \textbf{0.92}$	$\textbf{70.29} \pm \textbf{0.72}$	70.22 ± 0.92	60.29 ± 0.94
NN (ACC)	77.4	8 ± 1.61	41.50 ± 3.5	84 41.72	32.20	83.45	-	-	-
NN (F1)	78.0	1 ± 1.48	37.28 ± 3.9	97 37.11	28.46	83.25	-	-	-

[‡] We compute the OoD detection score after self-training.

Table 1: Evaluation of the proposed method and baselines. Section-wise best scores are in Bold.

of-the-art OoD detection methods, including contrastive training-based methods and self-supervised approaches. We also evaluate the representation quality by following common practice in self-supervised learning with a linear classifier on top of frozen features from the pre-training stage. The baselines include:

- state-of-the-art contrastive training based OoD detection methods, including CSI [42], open-set OoD detection (GOAD) [3], and CDMSAD [41],
- self-supervised learning-based methods [9, 7] pretrained on *source domain*, *target domain*, and *both domains* [28, 8, 7], respectively,
- supervised linear classification on frozen features from different self-supervised methods [8, 7].

Results and Discussion. In Table 1 A, we first report the results of state-of-the-art self-supervised methods [9, 7, 41] using the same backbone architecture as in SOoD, a ResNet18 on K16 and K19 datasets. Similar to SOoD, these methods are concerned with the scenarios where the source data are not labeled [9, 7] except for [41], where the source data is partially labeled. We use the same OoD detection score based on the closest feature distance for all the baselines for a fair comparison. The contrastive training-based methods are strong baselines [9, 7, 41] and can generalize reasonably well on the target data with the model pre-trained on the source data only, target data only, or both domains. The models that take advantage of the target distribution gain superior performance improvement than the same models trained on the source data. In addition, combining both data domains for training can further improve performance.

In Table 1 B, regarding the models trained on both domains, the GOAD method [3] often incorrectly detects a known class of target domain as an OoD due to the domain shift. CSI [42] benefits from contrastive learning to contrast each image with distributionally-shifted augmentations of itself. The methods in [7, 8] are based on a clustering scheme, and we use the same self-training approach as in SOoD for a fair comparison. SimTriplet [28] is the only method that incorporates multiple instances, but it is not formulated to address domain shift. For the input views of [28], we use the same three augmented views as in SOoD. Unlike these baselines, our method learns generalizable semantic properties in the feature space via clustering. Our designed domain-invariant formulation gains huge improvements under domain shift, and our results outperform other state-of-the-art self-supervised methods. Additionally, self-training used in our method further enhances the OoD detection performance. In Table 1 C, we also evaluate the quality of frozen features from the pre-training stage (pre-trained with ℓ_{mv}) via training a linear classifier on the frozen features. The objective is to show the effectiveness of SOoD to classify in-distribution target images correctly. Furthermore, we use a nearest neighbors classifier (k-NN) without any finetuning to vote for the label of in-distribution test images from the target domain. In Table 1, we report average F1 and accuracy (ACC) scores for seven in-distribution classes (all) using both schemes. We use the fully-supervised trained model (ResNet18) by utilizing all labeled style augmented images as the upper bound (Sup. Tr.-100%)². Self-supervised features from pre-

²One can consider a trained, supervised model on fully labeled real target images as the upper bound, but this setting is not realistic as we do not use label information from the target domain.

	K Sen	sitivity		Loss Weights	Sensitivity		Final Model (SOoD)
Metric	K=8	K=24	$\ell_1 = 3 \ell_2 = 1$	$\ell_1 = 1 \ell_2 = 3$	w/o ℓ_{heavy}	w/o ℓ_{style}	K=16, $\ell_1 = 1 \ell_2 = 1$
AUROC (A) AUROC (B)	$ \begin{array}{ } 82.52 \pm 1.69 \\ 88.85 \pm 0.57 \end{array} $	$\begin{array}{c} 86.70 \pm 4.33 \\ 89.24 \pm 0.40 \end{array}$	$\begin{array}{c} 85.18 \pm 2.12 \\ 90.77 \pm 0.47 \end{array}$	$\begin{array}{c} 82.76 \pm 0.86 \\ 90.99 \pm 0.28 \end{array}$	$\begin{array}{c} 84.63 \pm 0.43 \\ 91.60 \pm 0.59 \end{array}$	$\begin{array}{c c} 83.95 \pm 1.85 \\ 85.87 \pm 3.25 \end{array}$	$\begin{array}{c} 88.38 \pm 1.30 \\ 92.77 \pm 0.48 \end{array}$
AUPRC (A) AUPRC (B)	$ \begin{vmatrix} 71.44 \pm 2.03 \\ 85.30 \pm 0.22 \end{vmatrix} $	$\begin{array}{c} 79.60 \pm 7.56 \\ 84.71 \pm 0.43 \end{array}$	$\begin{array}{c} 76.41 \pm 3.22 \\ 87.95 \pm 0.33 \end{array}$	$\begin{array}{c} 71.80 \pm 0.25 \\ 88.64 \pm 0.39 \end{array}$	$\begin{array}{c} 75.76 \pm 1.07 \\ 90.62 \pm 0.54 \end{array}$	$\begin{array}{c c} 75.63 \pm 2.40 \\ 84.56 \pm 2.26 \end{array}$	$\begin{array}{c} 80.43 \pm 2.84 \\ 90.90 \pm 1.00 \end{array}$

Table 2: Ablation studies for the different number of prototypes K and loss weight values. We evaluate the models for both before self-training (A) and after the self-training stage (B).

Metric	Color Jittering	SOoD
AUROC (A) AUROC (B)	$\begin{array}{c} 82.44 \pm 1.20 \\ 88.58 \pm 0.73 \end{array}$	
AUPRC (A) AUPRC (B)	$\begin{array}{c} 75.23 \pm 1.12 \\ 86.54 \pm 0.57 \end{array}$	$\begin{vmatrix} 80.43 \pm 2.84 \\ 90.90 \pm 1.00 \end{vmatrix}$

Table 3: **Ablation studies** for different augmentation techniques. We evaluate the models for both before self-training (A) and after the self-training stage (B).

Checkpoint	AUROC	AUPRC		
	Mahalanobis Distance [26]			
Before Self-Training After Self-Training	$\begin{array}{ } 79.08 \pm 0.98 \\ 92.36 \pm 0.44 \end{array}$	69.13 ± 1.95 90.22 ± 0.74		
6	MSP Distance [21]			
Before Self-Training After Self-Training		$\begin{array}{c} 64.56 \pm 4.13 \\ 79.05 \pm 0.85 \end{array}$		
	Closest Features Distance			
Before Self-Training After Self-Training	$\begin{array}{ } 88.38 \pm 1.30 \\ 92.77 \pm 0.48 \end{array}$	$\begin{array}{c} 80.43 \pm 2.84 \\ 90.90 \pm 1.00 \end{array}$		

Table 4: The evaluation of the proposed method with different **OoD detection techniques** before and after selftraining.

trained SOoD perform particularly well with either learning a linear classifier or *k*-NN and surpass SOTA selfsupervised methods and supervised baseline. For example, SOoD trained with 10% labeled source data outperforms the fully supervised model trained from scratch on the source domain (Sup. Src.-100%), reducing the gap with full-label training (Sup. Tr.-100%). Finally, compared to [7, 28], the t-SNE [43] visualization of extracted features from the encoder f_{θ} shows a better alignment of the source and target domains, representations of the known classes, and a better separation of OoDs (see Fig. 5).

Ablations. We provide ablation studies to analyze the key factors that lead to the success of SOoD. These ablations concerning various aspects of SOoD's design, including loss terms, loss weight values, number of proto-types, augmentation techniques, and OoD detection scores (see Table 2). We sweep over a different number of pro-

totypes (16-24) and find that our method is not very sensitive to the number of prototypes, but using fewer prototypes $(< 2 \times \text{classes})$ leads to performance degradation. As argued, the model trained with additional style augmented view achieved a significant performance boost compared to baselines without this complementary view (w/o ℓ_{stule}). The sensitivity test for the loss weight values also shows each loss term for augmented views is equally important, improving the regularization effect of multi-view learning. Table 3 compares the OoD detection performance of our style augmentation with other augmentation (color jittering) and shows that the performance is significantly improved as we learn style-invariant representation. For all ablation experiments in Table 2 and Table 3, self-training the model on the target domain yields a better separation of OoDs from known classes and higher accuracy than the model trained only with optimizing ℓ_{mv} . This is also indicated by the t-SNE [43] visualization in Fig. 5. Finally, we compare our proposed OoD detection score with popular techniques (see Table 4), including Maximum over softmax probabilities (MSP) [21] and Mahalanobis distance [26]. The comparison demonstrates that OoD detection in these baselines might be failing due to the semantic ambiguity of some tissue categories, while ours achieves superior performance.

5. Conclusion

Our method is the first self-supervised OoD detection for CRC tissue types under domain shift in a zero-labeled data regime, yielding a more realistic and practical setting and alleviating costly annotations. It is also critical to safely deploy DNNs in computational pathology to generalize to a new clinical site with new categories not presented in a source dataset. We show that our designed multi-view consistency learning together with a self-training scheme gains substantial performance improvements in both OoD detection and classification of in-distribution samples compared to SOTA self-supervised methods. SOoD can be easily adjusted to be applied to different organs and histology tasks. In future work, we plan to design new formulations to improve the accuracy of pseudo labels and the generalizability of our method to unseen datasets.

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