

## Supplementary Material of ART-ASyn

### 001 Overview

002 This supplementary material complements the main paper  
003 by providing detail explanation of the synthetic anomaly  
004 composition steps of ART-ASyn and quantitative analysis  
005 of several hyperparameters used in the paper.

### 006 §1. Clarification of Synthesis Method

007 In this section, the implementation details of synthesis  
008 method are provided along with the ablation study of the  
009 effect of each synthesis step.

#### 010 §1.1. Implementation Details

011 As introduced in Section 3.2.1 of the main paper, the  
012 synthesis pipeline is built upon three primary transfor-  
013 mations of crystallization, blurring, and intensity scal-  
014 ing. The following subsections provide detailed ex-  
015 planations of each transformation and their composi-  
016 tion. For the full implementation, please refer to  
017 `preprocessing/ART_ASyn.py` in the code.

##### 018 §1.1.1. Crystallization ( $T_{\text{cryst}}$ )

019 In computer graphics, crystallization is a non-photorealistic  
020 rendering (NPR) technique that mimics the appearance of  
021 crystals or stained glass. Instead of preserving all fine de-  
022 tails, the image is decomposed into cells with uniform col-  
023 ors, producing a stylized effect [1]. It follows the imple-  
024 mentation of:

- 025 1. A fixed number of random "seed" points are sampled  
026 across the image grid. These act as nuclei around  
027 which regions will grow.
- 028 2. Each pixel in the image is assigned to the near-  
029 est seed using a spatial nearest-neighbor search uti-  
030 lizing `cKDTree` algorithm from Python module  
031 `scipy.spatial`. This yields a Voronoi diagram  
032 where the plane is divided into non-overlapping cells,  
033 each containing all pixels closest to its seed.
- 034 3. For each Voronoi cell, the algorithm computes the  
035 mean RGB color of all original pixels within the re-  
036 gion. Every pixel in that cell is then replaced by this  
037 mean color.

038 This operation corresponds to the crystallization transfor-  
039 mation mentioned in Section 3.2.1 of the main paper.

##### 040 §1.1.2. Blurring ( $T_{\text{blur}}$ )

041 Gaussian blurring is applied to both the raw  
042 anomaly layer and to crystallized regions utilizing  
043 `ImageFilter.GaussianBlur` from Python module  
044 of `PIL` with radii of 8 and 3, respectively. This simulates  
045 consolidation-like smoothness or the gradual density  
046 transitions observed in real lung pathologies.

##### §1.1.3. Intensity Scaling ( $T_{\text{rib}}$ )

The alpha channel of anomalies is modulated by a rib-  
intensity-weighted mask computed from the original chest  
X-ray. This ensures that synthetic anomalies preserve ra-  
diographic consistency with surrounding rib shadows and  
do not unrealistically occlude high-density bony structures.

##### §1.1.4. Composition of Transformations

The three transformations are applied sequentially and com-  
posed together:

1. A blurred version of brush painted anomaly is gener-  
ated.
2. A crystallized version of brush painted anomaly is  
generated.
3. These layers are alpha-scaled and overlaid to cre-  
ate a composite with both sharp-edged and smoothed  
structures.
4. The anomaly alpha is refined with lung and rib-  
intensity masks, ensuring anatomical plausibility.

This layered application follows the conceptual descrip-  
tion in Fig. 1 of the main paper, an initial painted anomaly  
is progressively refined by blurring, crystallization, and rib-  
aware scaling before being overlaid onto the normal chest  
X-ray to composite the synthetic abnormal image.

### §1.2. Ablation Study

We conducted stepwise experiments to analyze the contri-  
bution of different synthesis transformations. The ablation  
starts with the baseline synthesis ( $A_{\text{base}}$ ), then incorporates  
blurring ( $T_{\text{blur}}$ ) and crystallization ( $T_{\text{cryst}}$ ), and finally inte-  
grates all transformations into the complete pipeline ( $A_{\text{final}}$ ).  
The transformations of blur and crystallization are intro-  
duced jointly, as they are designed to operate in a composite  
manner: blurring smooths structural boundaries while crys-  
tallization introduces irregular granular patterns, together  
forming a realistic density base.

The base of our synthetic abnormality ( $A_{\text{base}}$ ) pro-  
duces texture-based, cocoon-like patterns with irregu-  
lar edges (see Figure S1-1). Quantitatively,  $A_{\text{base}}$   
achieves 0.8315/0.6603 in AP/AUC on CheXpert and  
0.9112/0.8797 on ZhangLab, compared to 0.7394/0.5929  
and 0.7938/0.6842 under the no-synthesis setting, and im-  
proves Dice on QaTa-ZeroShot from 0.0707 to 0.3067.  
Compared to established methods,  $A_{\text{base}}$  already exceeds  
most baselines, although its AUC values of 0.6603 on  
CheXpert and 0.8797 on ZhangLab remain below the Anat-  
Paste baseline of 0.7825 and 0.9103, respectively. This sug-  
gests that the texture painted  $A_{\text{base}}$  alone creates anomaly  
like structures.

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Synthesis Step	CheXpert		ZhangLab		QaTa-ZeroShot
	AP	AUC	AP	AUC	Dice
None	0.7394	0.5929	0.7938	0.6842	0.0707
$A_{\text{base}}$	0.8315	0.6603	0.9112	0.8797	0.3067
$A_{\text{base}} + T_{\text{blur}} + T_{\text{cryst}}$	0.7840	0.5953	0.8536	0.7574	0.0954
$A_{\text{final}}$	<b>0.8639</b>	<b>0.7125</b>	<b>0.9621</b>	<b>0.9350</b>	<b>0.3275</b>

Table S1. Quantitative analysis of the effect of different synthesis steps on anomaly detection performance across CheXpert, ZhangLab, and QaTa-ZeroShot.

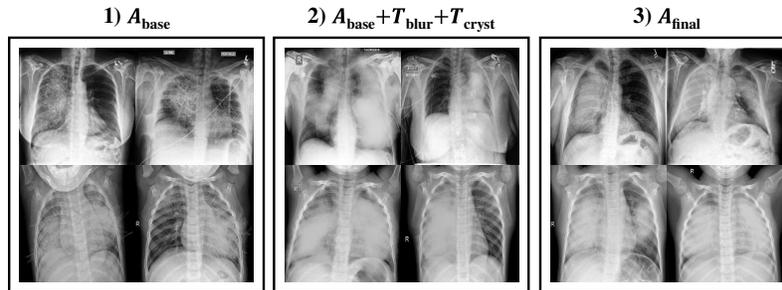


Figure S1. Visual comparison of different synthesis steps.

094 Adding blur and crystallization ( $A_{\text{base}} + T_{\text{blur}} + T_{\text{cryst}}$ )  
095 produces denser structures but tends to obscure anatomical  
096 boundaries (see Figure S1-2). This results in significant  
097 performance degradation compared to  $A_{\text{base}}$ . The  
098 variant lowers AP/AUC to 0.7840/0.5953 on CheXpert  
099 and 0.8536/0.7574 on ZhangLab, with Dice dropping to  
100 0.0954 on QaTa-ZeroShot, which is only slightly better  
101 than the no-synthesis setting. adding an intensity-scaling  
102 step ( $T_{\text{rib}}$ ) to restore anatomical visibility restores our ART-  
103 ASyn method, producing the  $A_{\text{final}}$  synthesis (see Figure S1-  
104 3). Collectively, blur and crystallization provide a neces-  
105 sary density base, and when combined with intensity scaling,  
106 they complement the texture-focused  $A_{\text{base}}$  to achieve  
107 superior anomaly realism as reported in Table 1 of the main  
108 paper and also Table S1.

109 In summary, the ablation demonstrates that texture syn-  
110 thesis alone ( $A_{\text{base}}$ ) already enhances anomaly detection,  
111 but combining blur and crystallization with intensity scaling  
112 in the full  $A_{\text{final}}$  pipeline is essential to balance texture fi-  
113 delity and anatomical consistency, resulting in the strongest  
114 and most consistent performance across datasets.

## 115 §2. Sensitivity of the Binarization Threshold 116 for Anomaly Mask Computation

117 For ease of experimentation in the main paper, we fix the binarization  
118 threshold at  $\tau = 0.2$  when deriving the anomaly mask  $\hat{M}_{\text{anomaly}}$   
119 from the pixel-wise reconstruction error between the synthetic input  
120  $I_{\text{syn}}$  and its reconstruction  $\hat{I}$ . However, as a dataset-dependent  
121 thresholding hyperparameter, it

is important to evaluate the model’s sensitivity to its setting.  
We therefore explore a wide range of  $\tau$  from 0.2 to 0.8 with  
an interval of 0.2 and report their respective anomaly detection  
performance on CheXpert and ZhangLab, and QaTa-ZeroShot as  
reported in Table S2.

On CheXpert, the best scores occur at  $\tau = 0.6$  with AP/AUC  
0.8694/0.7325, while the lowest are at  $\tau = 0.4$  with  
0.8523/0.6981, giving sensitivity ranges of 0.0171 (AP) and  
0.0344 (AUC). On ZhangLab, the highest AP/AUC is at  
 $\tau = 0.2$  with 0.9621/0.9350 and the lowest at  $\tau = 0.6$   
with 0.9348/0.8930, corresponding to ranges of 0.0273 (AP)  
and 0.0420 (AUC). For QaTa-ZeroShot, Dice peaks at  
 $\tau = 0.6$  with 0.3319 and is lowest at  $\tau = 0.4$  with  
0.2963 (range 0.0356). Overall, the metrics are relatively  
insensitive to  $\tau$ , with mild improvements near  $\tau = 0.6$   
for CheXpert and QaTa-ZeroShot, and near  $\tau = 0.2$  for  
ZhangLab.

It is evident from the results that although the model’s scores  
vary with the choice of  $\tau$ , the magnitude of these changes  
is minor, below most of the performance margins by which our  
method exceeds competing baselines. Therefore, the performance  
of the model is relatively insensitive to the exact binarization  
threshold  $\tau$ .

## §3. Quantitative Analysis of the Effect of Loss Components

In Section 3.2.2 of the main paper, the loss is defined as:

$$\mathcal{L}_{\text{total}} = \mathcal{L}_{\text{local}} + \mathcal{L}_{\text{global}} + \mathcal{L}_{\text{feat}} + \mathcal{L}_{\text{dice}}. \quad (1)$$

Threshold ( $\tau$ )	CheXpert		ZhangLab		QaTa-ZeroShot
	AP	AUC	AP	AUC	Dice
0.2	0.8639	0.7125	<b>0.9621</b>	<b>0.9350</b>	0.3275
0.4	0.8523	0.6981	0.9417	0.9019	0.2963
0.6	<b>0.8694</b>	<b>0.7325</b>	0.9348	0.8930	<b>0.3319</b>
0.8	0.8531	0.7270	0.9459	0.9204	0.3133

Table S2. Sensitivity analysis of the effect of different binarization thresholds  $\tau$  on anomaly detection result of CheXpert, ZhangLab and QaTa-ZeroShot.

Loss Configuration	CheXpert		ZhangLab		QaTa-ZeroShot
	AP	AUC	AP	AUC	Dice
$\mathcal{L}_{\text{global}}$	0.8250	0.6671	0.8124	0.7457	0.2771
$\mathcal{L}_{\text{global}} + \mathcal{L}_{\text{local}}$	0.8319	0.6675	0.8788	0.7575	0.3125
$\mathcal{L}_{\text{global}} + \mathcal{L}_{\text{feat}}$	0.8347	0.6742	0.9382	0.9053	0.2527
$\mathcal{L}_{\text{global}} + \mathcal{L}_{\text{dice}}$	0.8362	0.6860	0.9032	0.8297	0.2804
$\mathcal{L}_{\text{global}} + \mathcal{L}_{\text{local}} + \mathcal{L}_{\text{feat}}$	0.8573	0.7043	0.9409	0.9018	0.3227
$\mathcal{L}_{\text{global}} + \mathcal{L}_{\text{local}} + \mathcal{L}_{\text{dice}}$	0.8445	0.6801	0.9058	0.8550	0.3266
$\mathcal{L}_{\text{global}} + \mathcal{L}_{\text{feat}} + \mathcal{L}_{\text{dice}}$	0.8499	0.6762	0.9388	0.9003	0.3240
$\mathcal{L}_{\text{total}}$	<b>0.8639</b>	<b>0.7125</b>	<b>0.9621</b>	<b>0.9350</b>	<b>0.3275</b>

Table S3. Quantitative analysis of the effect of training under different loss component permutations on anomaly detection performance across CheXpert, ZhangLab, and QaTa-ZeroShot.

149 In this section, the effect of each permutation of differ-  
 150 ent loss components on anomaly detection performance  
 151 for datasets CheXpert, ZhangLab, and QaTa-ZeroShot is  
 152 shown in Table S3.

153 With the standard global reconstruction loss  $\mathcal{L}_{\text{global}}$ ,  
 154 the model reaches baseline anomaly detection perfor-  
 155 mance of 0.8250/0.6671 on CheXpert and 0.8124/0.7457 on  
 156 ZhangLab, with a Dice score of 0.2771 on QaTa-ZeroShot.  
 157 This serves as the foundation upon which the effects of ad-  
 158 ditional loss terms can be assessed.

159 Incorporating an extra loss component of the local re-  
 160 construction loss  $\mathcal{L}_{\text{local}}$ , the feature-alignment loss  $\mathcal{L}_{\text{feat}}$ ,  
 161 or the Dice loss  $\mathcal{L}_{\text{dice}}$  consistently improves performance.  
 162 Among these, the feature-alignment loss  $\mathcal{L}_{\text{feat}}$  particularly  
 163 effective on the one-class classification task. It elevates the  
 164 AP/AUC of CheXpert to 0.8347/0.6742 and ZhangLab to  
 165 0.9382/0.9053, representing the largest gains relative to the  
 166 baseline.

#### 167 §4. Efficiency Analysis of PBTSEg

168 In the main paper, we propose a progressive binary thresh-  
 169 olding lung segmentation method (PBTSEg). Table S4 re-  
 170 ports the number of images processed per second on the  
 171 same computer with different number of workers. It demon-  
 172 strates the efficiency limitation of PBTSEg, as the simpler  
 173 lung segmentation method in AnatPaste is approximately

No. of Workers	AnatPaste	PBTSEg
	Image/Second	Image/Second
1	99.47	8.95
2	160.22	17.51
4	286.52	26.83
8	423.99	40.52

Table S4. Efficiency analysis of lung segmentation utilizing Anat-Paste and PBTSEg.

10 times faster. This sacrifice in efficiency is offset by im- 174  
 improved segmentation quality which is critical for ensuring 175  
 that anomalies are synthesized within accurate lung bound- 176  
 aries, which directly impacts the anatomical plausibility and 177  
 effectiveness of the generated samples. Additionally, effi- 178  
 ciency limitation of PBTSEg represents a one-time prepro- 179  
 cessing cost rather than an ongoing bottleneck during train- 180  
 ing as it only need to be computed once for the dataset and 181  
 can then be reused throughout all subsequent experiments. 182

#### References 183

- [1] Narendra Ahuja, Byong An, and Bruce Schachter. Image 184  
 representation using voronoi tessellation. *Computer Vision, 185*  
*Graphics, and Image Processing*, 29(3):286–295, 1985. 1 186