

# Multi-Modal Medical Image Augmentation for Controlled Heterogeneity and Fair Outcomes

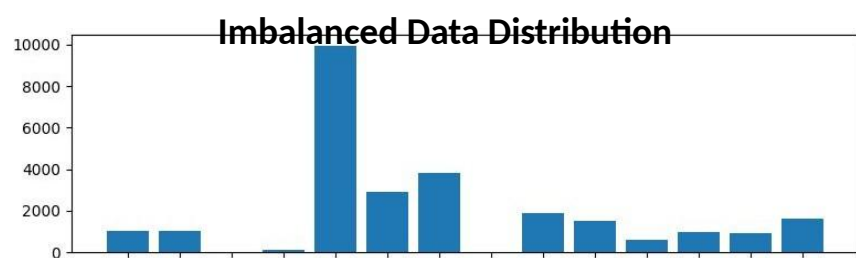
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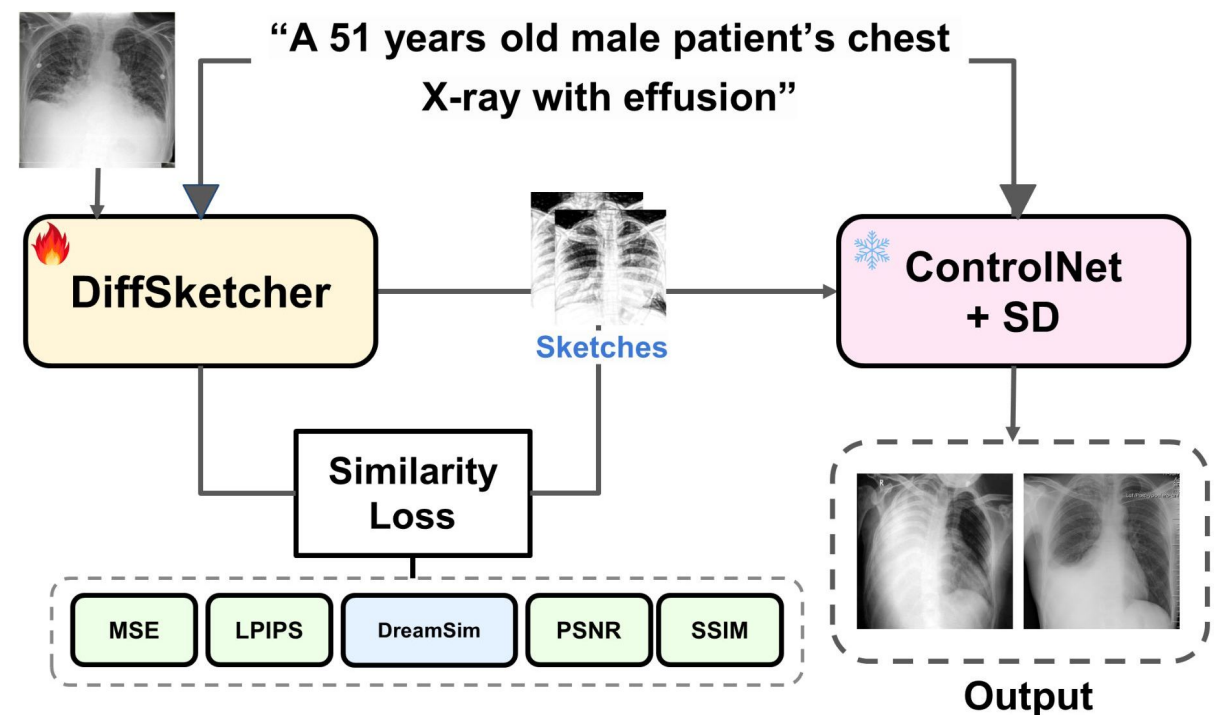
## Motivation

- In medical domain, **data imbalance** among different patient group is critical problem.
- Data augmentation** is a widely used strategy to solve this problem.
- However, generating specific patient groups data with **various conditions** (age, sex, disease) is challenging.



Our study **aims** to augment medical images under specific conditions

## Abstract



## Framework

### (1) Select patient groups for data augmentation

: we propose a **new metric** to measure **necessity of data augmentation** of patient groups

$$M_K = \frac{1}{2} \left( \frac{|D_K|}{|D|} + S_K \right) \quad (1) \quad S_K = \frac{2 * \sum_{1 \leq i \leq j \leq |D_K|} d_{ij}}{|D_K|(|D_K| - 1)} \quad (2)$$

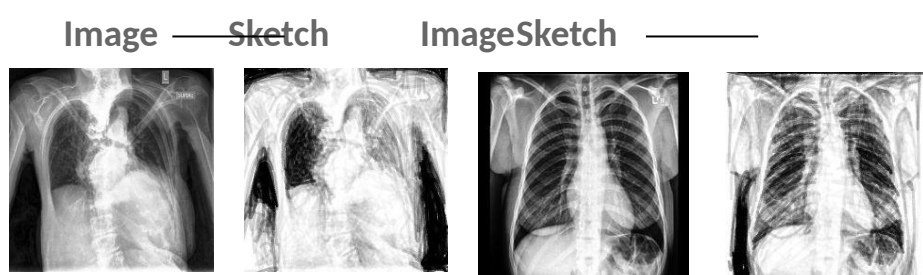
$$d_{ij} = \frac{1}{3} * \left[ \frac{1}{C} \sum_{c=1}^C f(v_c^i, v_c^j) + \frac{|g^i // 10 - g^j // 10|}{g^m // 10} + \delta(s^i, s^j) \right] \quad (3)$$

$$f(v_c^i, v_c^j) = \begin{cases} 1 & \text{if } v_c^i \neq v_c^j \\ 0 & \text{otherwise} \end{cases} \quad (4) \quad \delta(s^i, s^j) = \begin{cases} 1 & \text{if } s^i \neq s^j \\ 0 & \text{otherwise} \end{cases} \quad (5)$$

$M_K$  is the majority score and  $K$  is the index of patient groups.  $|D|$  is the total number of data and  $|D_K|$  is the number of data from  $K$ 'th patient group.  $v$  is the label vector and  $c$  is the class index.  $g$  indicates the age of the patient and  $m$  is the index of the maximum age.  $s$  represents the sex of the patient.

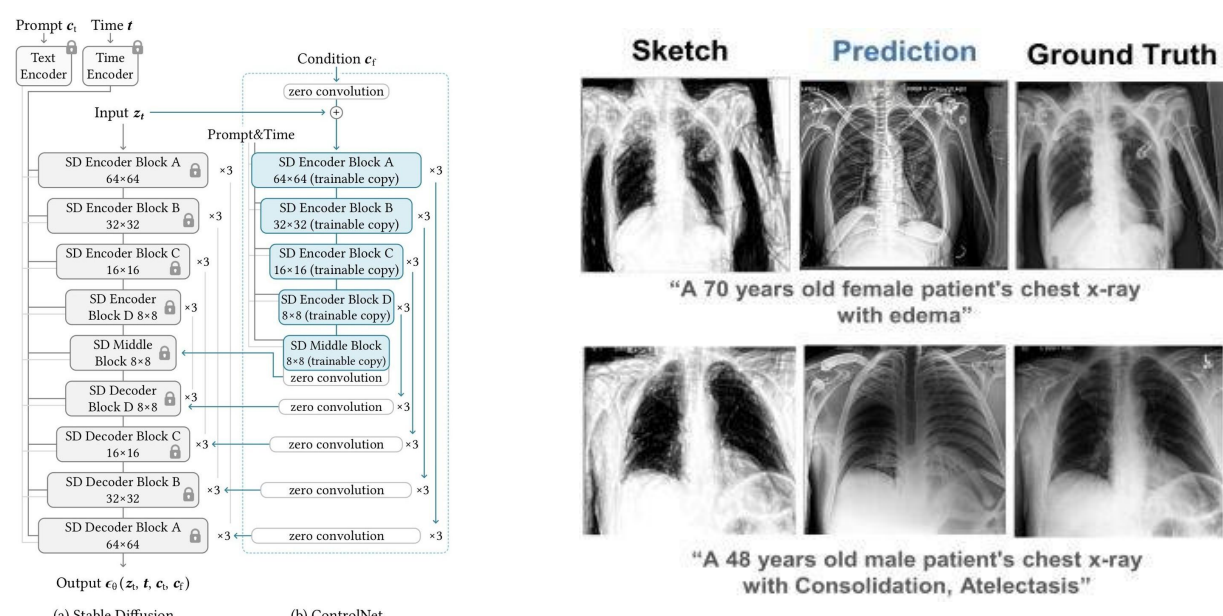
### (2) Collect sketch-image pair using DiffSketcher

: DiffSketcher is vector graphics sketch generator. We pairs to train ControlNet.



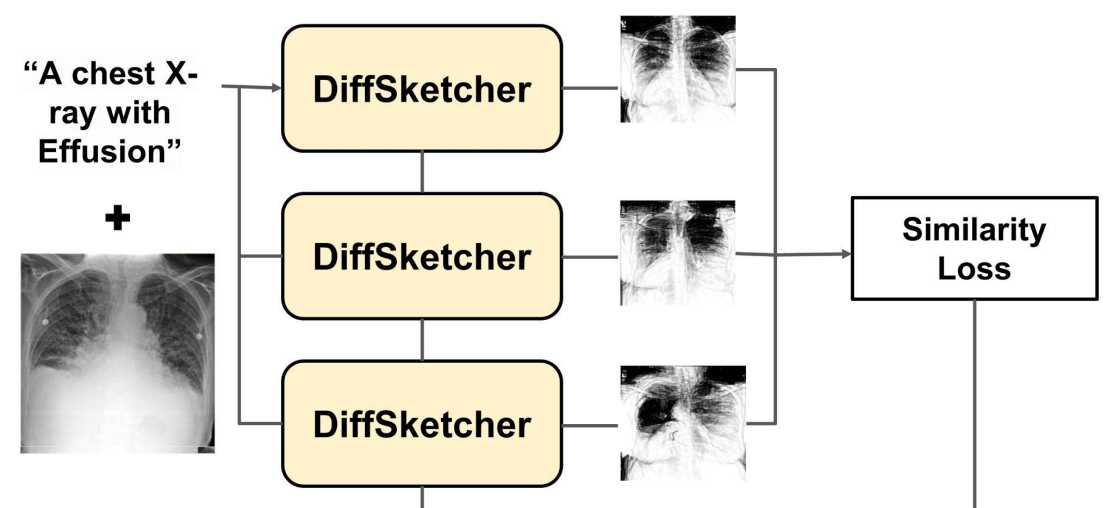
### (3) Fine-tune pre-trained ControlNet

: Below results are obtained from the **training process**.



### (4) Obtain diverse sketch data from DiffSketcher

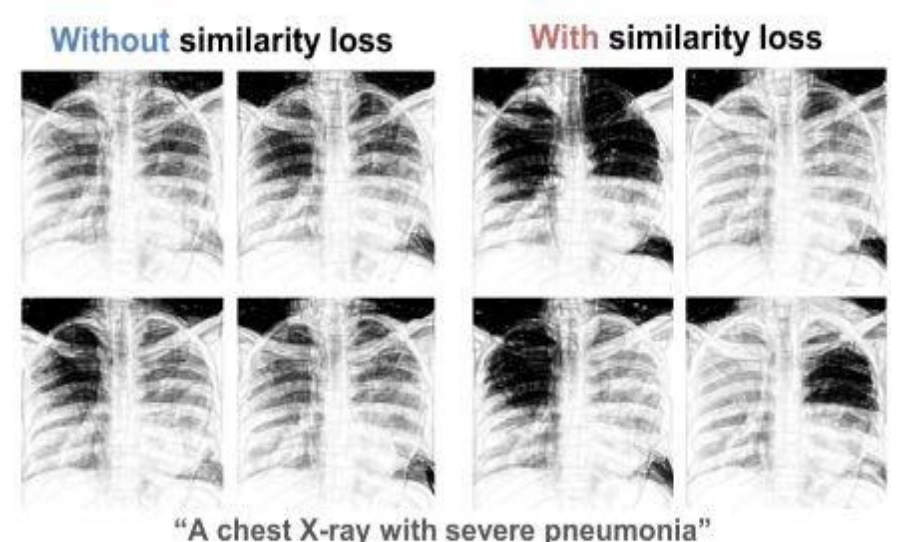
: we obtain multiple various sketched from DiffSketcher with the **similarity loss which we propose**.



**Similarity loss:** consists of both low-level and high-level distance metrics

$$\mathcal{L}_{sim}^i = \alpha * \frac{\sum_{1 \leq j \leq n, i \neq j} (1 - h(x_i, x_j))}{n - 1} + (1 - \alpha) * \frac{\sum_{1 \leq j \leq n, i \neq j} (1 - l(x_i, x_j))}{n - 1} \quad (6)$$

$\alpha$  regulates the contributions of high-level and low-level similarity metrics.  $i$  and  $j$  are the index of the DiffSketcher.  $x$  is the input image.  $h$  represents high-level similarity metric and  $l$  is the low-level similarity metric.  $n$  is the number of sketches we generate at the same time.



### (5) Measure diversity of augmented patient group

: we propose a **new metric** to measure **diversity of dataset using saliency map**.

$$D - score = \frac{2}{N(N - 1)} \sum_{1 \leq i \leq j \leq N} \frac{1}{C} \sum_{c=1}^C \omega(A_c^i, A_c^j) \quad (7)$$

$\omega$  is the Wasserstein distance and  $A$  indicates saliency map.  $c$  represents the index of the convolutional block and  $N$  is the total number of generated samples.