



26TH INTERNATIONAL CONFERENCE ON MEDICAL IMAGE COMPUTING
AND COMPUTER ASSISTED INTERVENTION
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Additional Positive Enables Better Representation Learning for Medical Images

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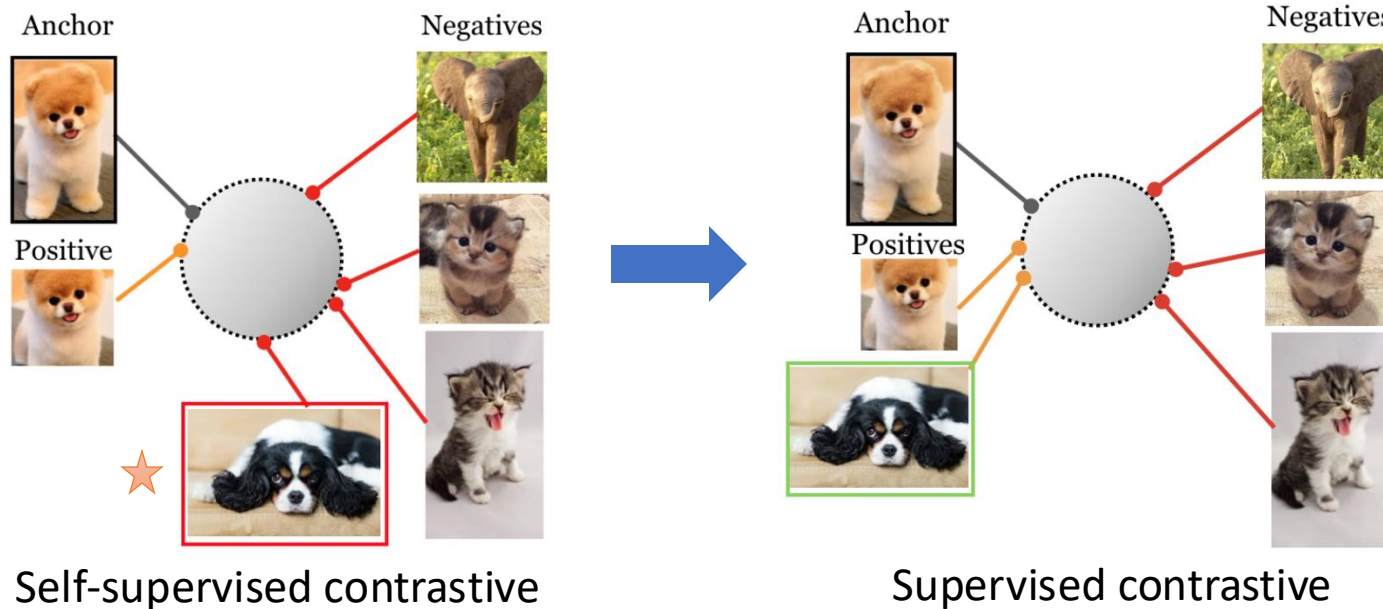
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Pittsburgh



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Guangdong Cardiovascular Institute

Limitations in SOTA self-supervised learning framework

- ❑ Positive pairs are generated using data augmentations
- ❑ Instance with the same label from other images are not utilized



Can we identify such positives without label?

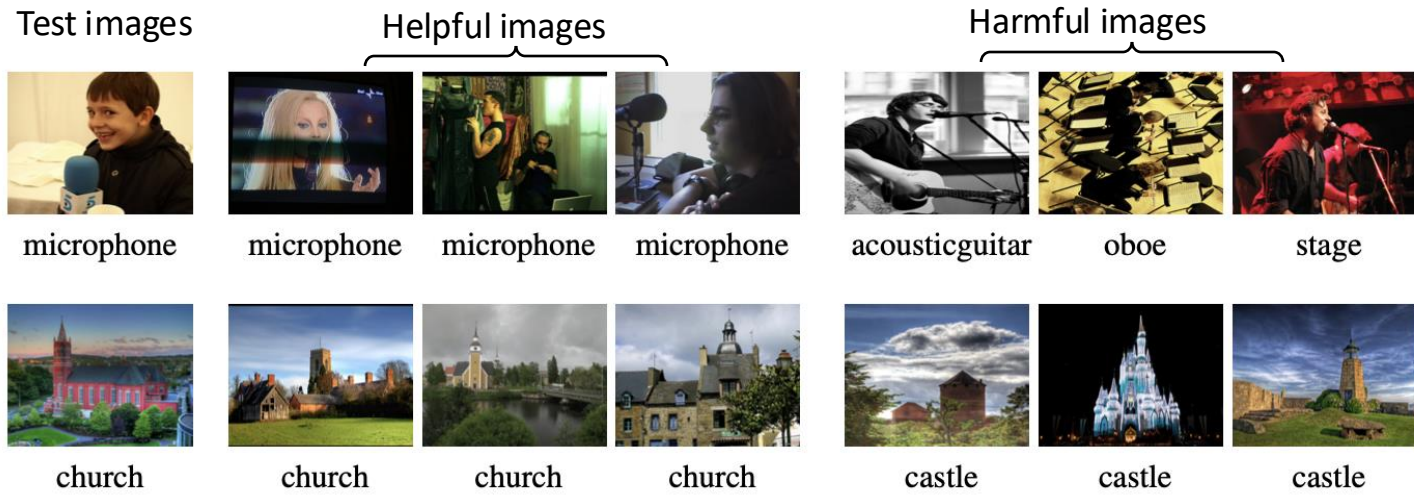
Influence Function: TracIn

- ❑ Measure how the loss of a test sample changes with the training process of a training sample
- ❑ A **positive** TracIn value means **helpful** example that reduces loss, a **negative** TracIn value means **harmful** example that increases loss

$$\text{TracIn}(x_i, x_k) = \sum_{t: x_t = x_i}^T \eta_t \nabla \ell(w_t, x_k) \cdot \nabla \ell(w_t, x_i).$$

total number of iterations → T
learning rate at iteration t → η_t
gradient of sample x_k at iteration t → $\nabla \ell(w_t, x_k)$

Model interpretation

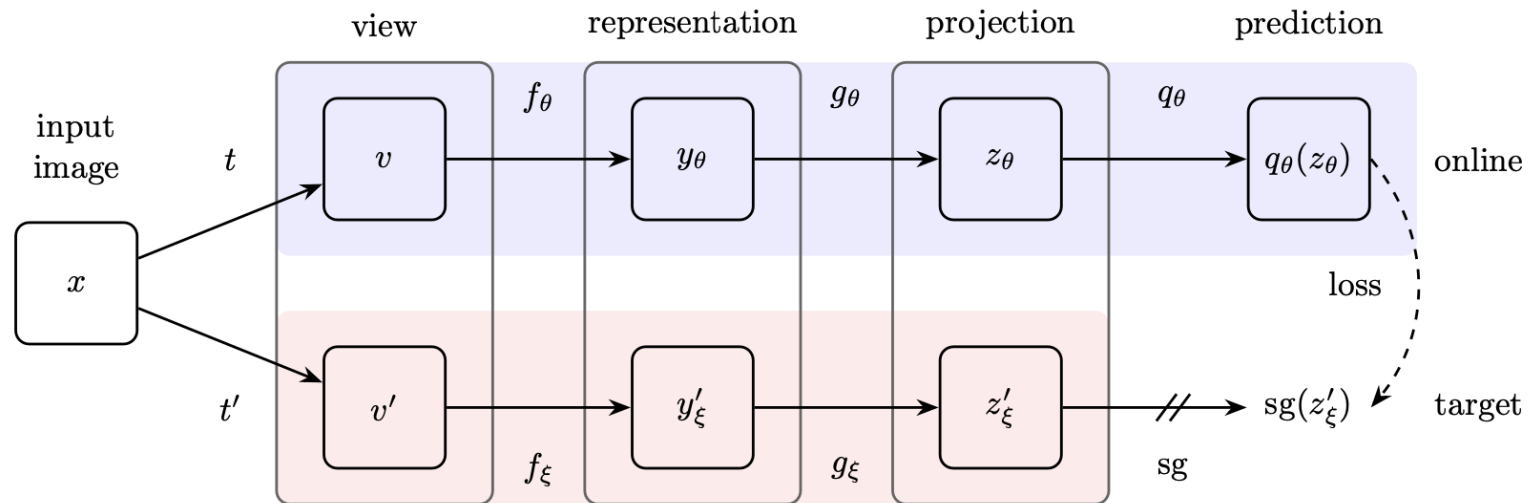


Mislabel identification (self influence)



BYOL: Bootstrap your own latent

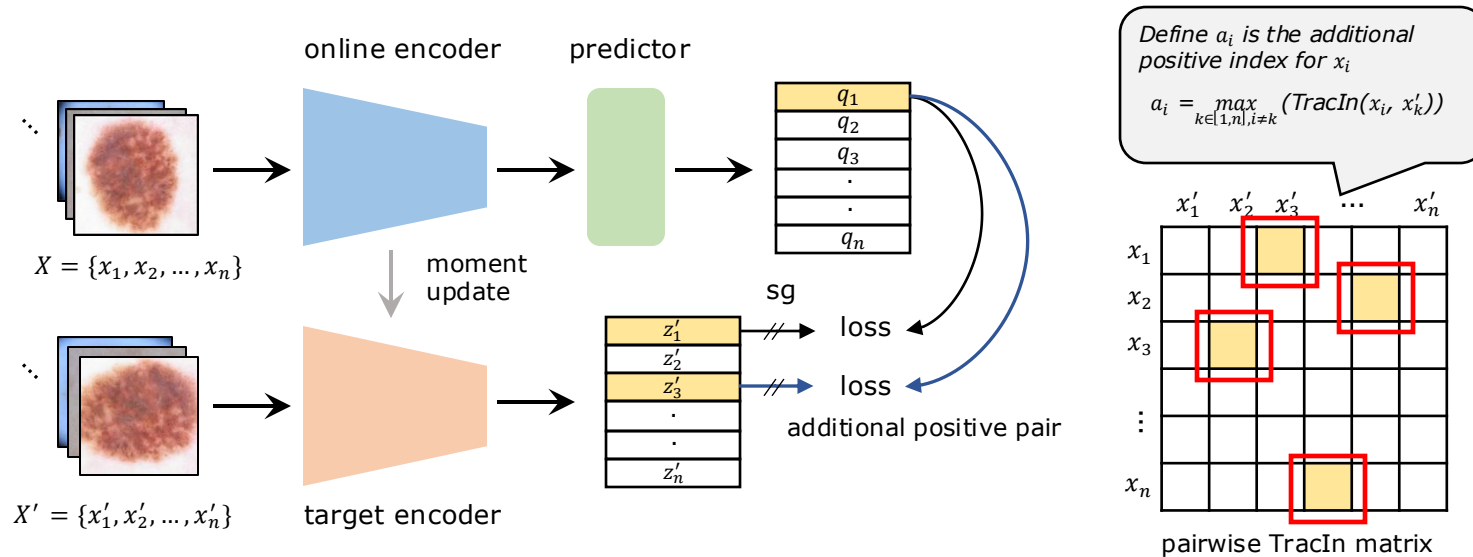
- ❑ Only positive pair is required (no negatives)
- ❑ Easy to compute per-sample gradient



$$\mathcal{L}_{\theta, \xi} \triangleq \|\bar{q}_\theta(z_\theta) - \bar{z}'_\xi\|_2^2 = 2 - 2 \cdot \frac{\langle q_\theta(z_\theta), z'_\xi \rangle}{\|q_\theta(z_\theta)\|_2 \cdot \|z'_\xi\|_2}$$

BYOL-TracIn

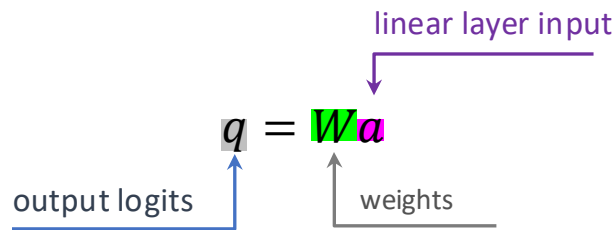
- ❑ Use BYOL loss function without label
- ❑ Selecting another sample with the largest TracIn value
- ❑ Use the gradient of the last linear layer to estimate (save resources)



Efficient per-sample gradient

- ❑ TracIn requires the gradient of each sample.
- ❑ PyTorch or Tensorflow do not support batch-wise per-sample gradient

For linear layer:



loss function

$$\nabla_W f(q) = \nabla_q f(q) \cdot a^T$$

target network output (constant)

$$f(q) = 2 - 2 \cdot \frac{\langle q, z \rangle}{(\|q\|_2 \cdot \|z\|_2)}$$

gradient

$$\begin{aligned} \text{TracIn}(x_i, x_k) &\approx \eta_t \nabla_W f(q_i) \cdot \nabla_W f(q_k) \\ &= \eta_t (\nabla_q f(q_i) a_i^T) \cdot (\nabla_q f(q_k) a_k^T) \\ &= \eta_t (\nabla_q f(q_i) \cdot \nabla_q f(q_k)) \cdot (a_i \cdot a_k) \end{aligned}$$



$$\nabla_q f(q) = 2 \cdot \left(\frac{\langle q, z \rangle \cdot q}{(\|q\|_2^3 \cdot \|z\|_2)} - \frac{z}{\|q\|_2 \cdot \|z\|_2} \right)$$

We only need to save the inputs and outputs of the linear layer in one forward operation

Semi-supervised results

- ❑ Dataset: ISIC 2019 (25,331 images), ChestX-ray (108,948 images)
- ❑ Our method: **BYOL-TracIn**, upper bound: **BYOL-Sup**
- ❑ BYOL-TracIn-pretrained: using a pretrained model for computing TracIn

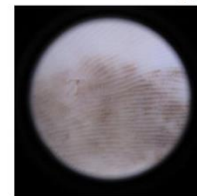
Method	ISIC 2019			ChestX-ray		
	10%	50% BMA ↑	100%	10%	50% AUC ↑	100%
Random	0.327(.004)	0.558(.005)	0.650(.004)	0.694(.005)	0.736(.001)	0.749(.001)
BYOL [12]	0.399(.001)	0.580(.006)	0.692(.005)	0.699(.004)	0.738(.003)	0.750(.001)
FNC [17]	0.401(.004)	0.584(.004)	0.694(.005)	0.706(.001)	0.739(.001)	0.752(.002)
FT [30]	0.405(.005)	0.588(.008)	0.695(.005)	0.708(.001)	0.743(.001)	0.751(.002)
FS	0.403(.006)	0.591(.003)	0.694(.004)	0.705(.003)	0.738(.001)	0.752(.002)
FS-pretrained	0.406(.002)	0.596(.004)	0.697(.005)	0.709(.001)	0.744(.002)	0.752(.002)
BYOL-TracIn	0.403(.003)	0.594(.004)	0.694(.004)	0.705(.001)	0.742(.003)	0.753(.002)
✓ BYOL-TracIn-pretrained	0.408(.007)	0.602(.003)	0.700(.006)	0.712(.001)	0.746(.002)	0.754(.002)
BYOL-Sup	0.438(.006)	0.608(.007)	0.705(.005)	0.714(.001)	0.748(.001)	0.756(.003)

Transfer learning results

- ❑ Dataset: ISIC 2016 (900 images), Shenzhen (662 images)
- ❑ FS: Feature Similarity

Method	ISIC 2016 Precision \uparrow	Shenzhen AUC \uparrow
Random	0.400(.005)	0.835(.010)
BYOL [12]	0.541(.008)	0.858(.003)
FNC [17]	0.542(.007)	0.862(.006)
FT [30]	0.559(.011)	0.876(.005)
FS	0.551(.003)	0.877(.004)
FS-pretrained	0.556(.004)	0.877(.006)
BYOL-TracIn	0.555(.012)	0.880(.007)
BYOL-TracIn -pretrained	0.565(.010)	0.883(.001)
BYOL-Sup	0.592(.008)	0.893(.006)

Anchor image



Label:NV

Top-3 most similar images in a mini-batch ←



Label:NV

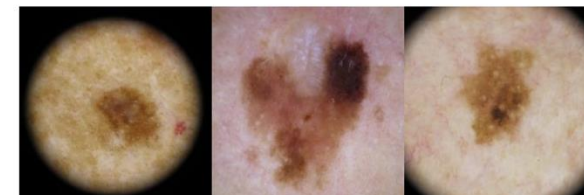
Label:NV

Label:NV

TracIn:0.023

TracIn:0.018

TracIn:0.016



Label:MEL

Label:MEL

Label:NV

FS:0.907

FS:0.894

FS:0.892



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Conclusions

1. BYOL-TrackIn provides a way to select additional positive pair for self-supervised pre-training, which could increase the diversity of features in positive pairs.
2. BYOL-TrackIn shows significant improvements in both semi-supervised and transfer learning settings