Unsupervised Lung Cancer nodule Representation using Region of Interest

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1 Training objective

Let I be the image of dimension n * n. For now let the scale be fixed to m * m. The task is to train a prediction function g.

Let each block be indexed $x_1, x_2, x_3...x_n$.

$$f_{\theta}(x_i) + e_i \to v_i$$

where,

 $x_i \in m \times m$

 $v_i \in d$

 e_i : positional embedding

 f_{θ} : convnet with params θ

 θ is trainable parameters.

Each block x_i is passed through a function f to get vector representation of each block, where $v_1, v_2, v_3....v_n$ represent vector representations.

Now, we apply a pooling layer p over all these vector representations, except the block to be predicted.

$$p(\{v_1 \dots v_{i-1} v_{i+1} \dots v_n\}) \to V_{\setminus i}$$

This $V_{\setminus i}$ is passed through a regression function g to get predicted vector representation v'_i for the predicted block

$$g_{\phi}\left(v_{\backslash i}\right) \to v_{i}'$$

where,

 g_{ϕ} : a neural network with parameters ϕ

 ϕ is trainable parameter.

Let q be a function that takes in v_i' and converts this vector back into image space.

$$q_{\omega}(v_i') \to I_i'$$

where,

 q_{ω} : a upsampling neural network with parameters ω

is trainable parameter. This generates a block of the predicted Image I. On passing all the block vectors $v_1, v_2, ... v_n$ through d we would get the predicted image I'

Hence when each patch or block v_i tries to predict the original patch x_i Loss in image space can be defined as

$$L_i = \frac{1}{2} \left(I_i' - I_i \right)^2$$

The final objective is defined as

$$J_{\theta,\omega,\phi} = \frac{1}{2} \sum_{i} (q_{\omega}(g_{\phi}(p(\{f_{\theta}(x_1)..f_{\theta}(x_i-1), f_{\theta}(x_i+1)..f_{\theta}(x_n)\}))) - I_i)^2$$

2 Region of Interest

• for each block i in $x_1, x_2...x_n$, r_i is the residuals in vector space, then for higher residuals patch x_i is region of interest.

$$r_i = \frac{1}{2} \left(v_i' - v_i \right)^2$$