

VARIATIONAL AUTOENCODERS FOR LUNG CANCER DIAGNOSIS

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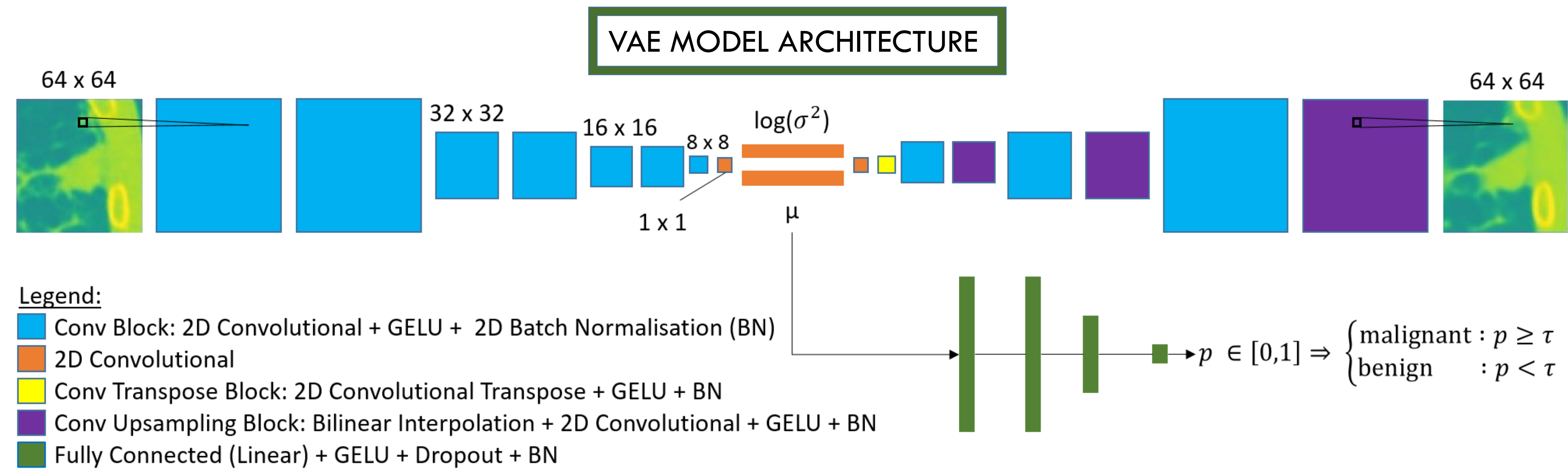
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MOTIVATION

Recent advances in generative methods have brought them to the forefront of AI research. VAEs offer many avenues of potential research:

- **EXPLAINABLE FEATURES AND CLASSIFICATION**
- **SYNTHETIC DATA**
- **LATENT TRAVERSALS**

This work aims to demonstrate these methods on a dataset of lung cancer lesions.



MATERIALS AND METHODS

DATASET:

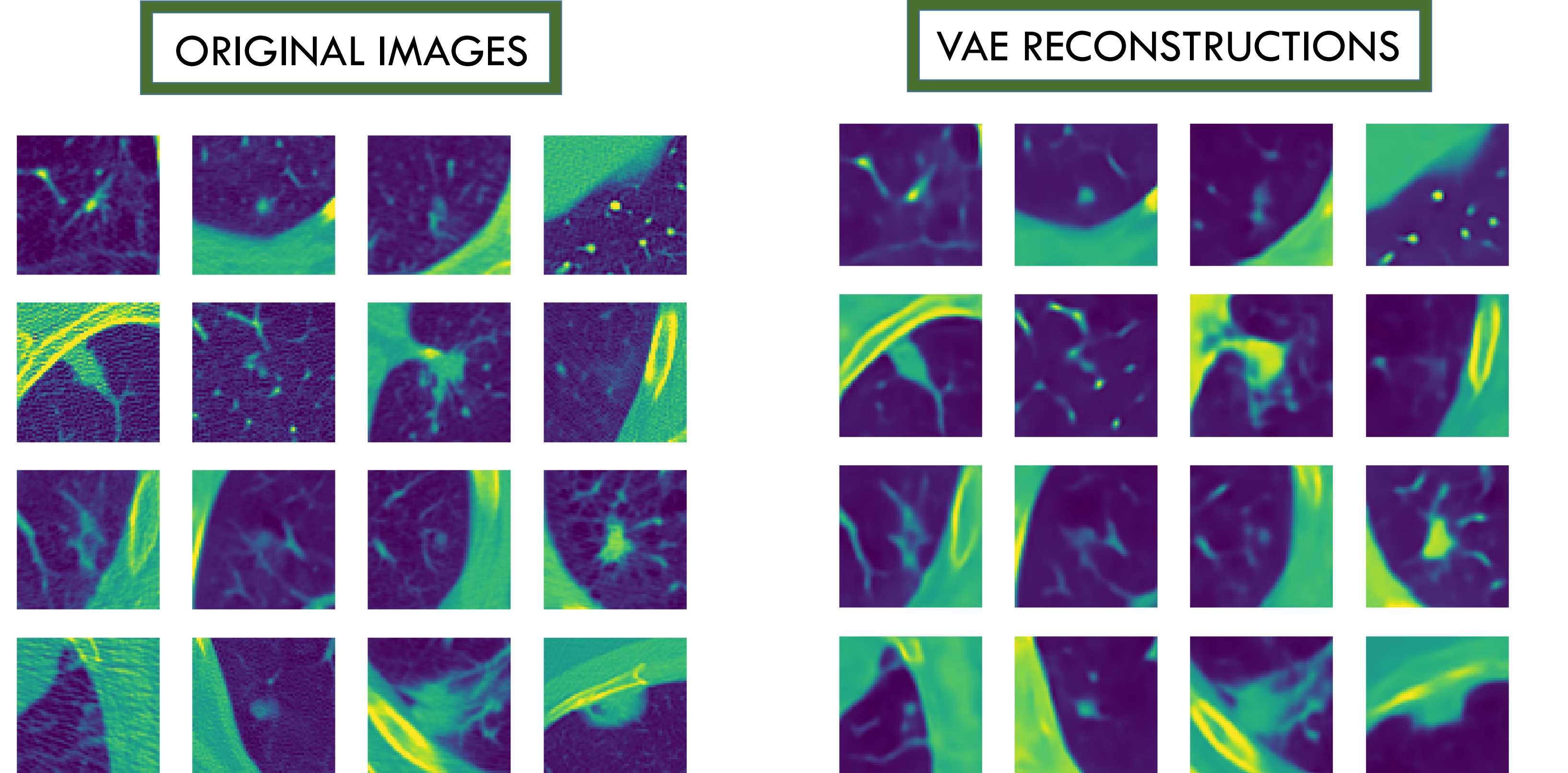
Results are produced from the LIDC-IDRI dataset which holds 875 annotated CT scans [2]. Malignancy labels and segmentation masks were agreed by 4 radiologists.

METHODS:

- Gaussian VAE and Dirichlet VAE (encourages disentanglement).
- Extract latent vectors and use as feature representations in Multi-Layered Perceptron (MLP) diagnostic model.
- K-Means and CLASSIX clustering.

WHY VAEs?

- **unsupervised:** do not require labelled data.
- **generative:** can produce (realistic) synthetic data



LATENT EXPLORATION

CLUSTERING:

Shows lesions are separated by shape, patient and class in latent space:

- **58%** of patients had over 50% of slices in a single cluster.
- **70%** of clusters had over 75% of one class (malignant/non-mal).
- See samples from highest proportion malignant/benign clusters.

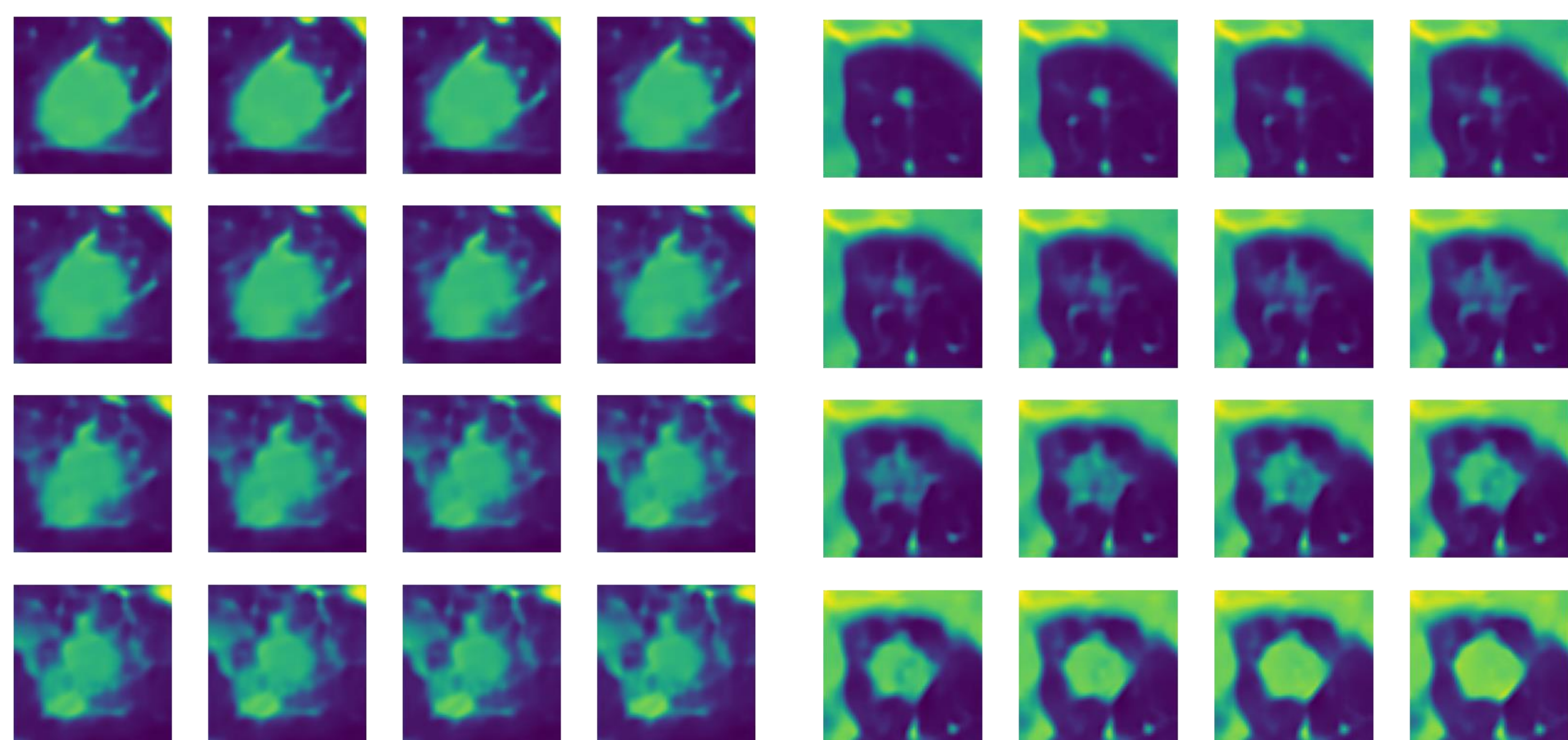
LATENT TRAVERSALS:

- Initial work shows its possible to find **generalisable and clinically meaningful feature directions**.
- Uses average direction vector between centre of two groups of similar images with/without a given feature. Multiples of the vector are applied to a new image to generate a smooth transition.

LATENT TRAVERSAL / FEATURE DIRECTIONS

IRREGULAR BORDER

TUMOUR GROWTH



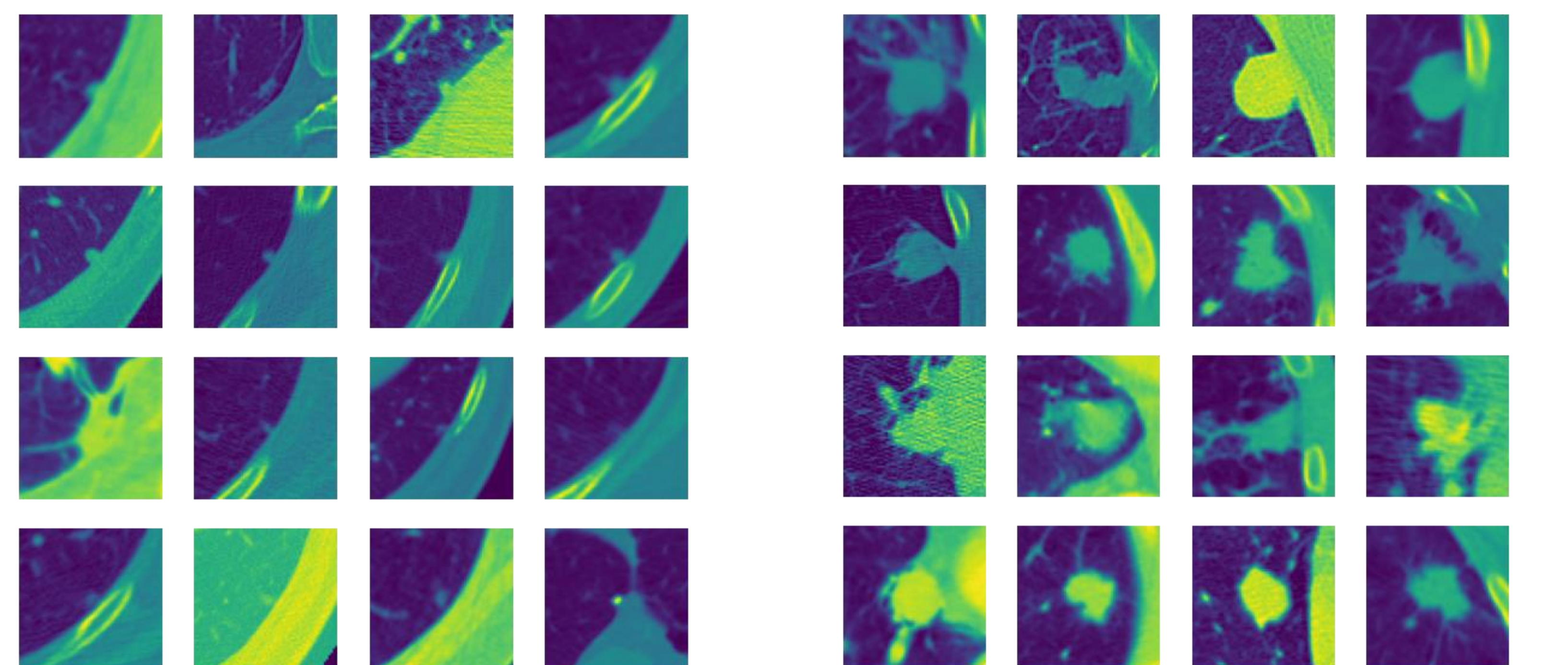
LUNG CANCER DIAGNOSIS

- Results presented show that VAE + MLP combination achieves state-of-the-art performance for lung cancer diagnosis: **0.98 AUC** and **93% Accuracy**.
- Best model uses fine-tuning of encoder with MLP loss to produce classification-optimised latent vectors.
- Direct comparison with Silva et al. [4] who used VAE latent vectors for same task.
- VAE feature vectors are resistant to noise/small changes unlike traditional CNN-derived features.

	VAE _{EM}	VAE	Systematic Review [3]	Silva et al. [4]	Radiologist [1]
AUC	0.98	0.89	0.7 - 0.97	0.94	0.85
Accuracy	0.93	0.82	0.88 - 0.99	0.90	

BENIGN CLUSTER

MALIGNANT CLUSTER



NEXT STEPS

- Use clustering to generate **pseudo-labels** for weakly supervised classification.
- See how latent traversals affect classifications for **diagnostic feature discovery**.
- Create **synthetic data** to help reduce overfitting.
- **Segmenting bone and fat** to remove this impact from the latent space.
- Find direction corresponding to time – **Temporal VAE** for time to event data e.g. pre/post treatment scans. Evolve lesions in time to look for **response to treatment**.

REFERENCES:

- [1] Al Mohammad, S. L. Hillis, W. Reed, M. Alakhras, and P. C. Brennan. Radiologist performance in the detection of lung cancer using CT. *Clinical Radiology*, 74(1):67–75, 2019.
- [2] Armato, S. G., McLennan, G., Bidaut, L., McNitt-Gray, M. F., Meyer, C. R. et al. [2011]. The Lung Image Database Consortium (LIDC) and Image Database Resource Initiative (IDRI): A completed reference database of lung nodules on CT scans *Medical Physics* 38(2) 915–931.
- [3] Jassim, M. M. and Jaber, M. M. Systematic review for lung cancer detection and lung nodule classification: Taxonomy, challenges, and recommendation future works. *Journal of Intelligent Systems*, 31(1):944–964, 2022.
- [4] Silva, F., Pereira, T., Frade, J., Mendes, J., Freitas, C. et al. [2020]. Pre-training autoencoder for lung nodule malignancy assessment using CT images. *Applied Sciences* 10(21).