

Self-Supervised Graph Representations of WSIs



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Introduction

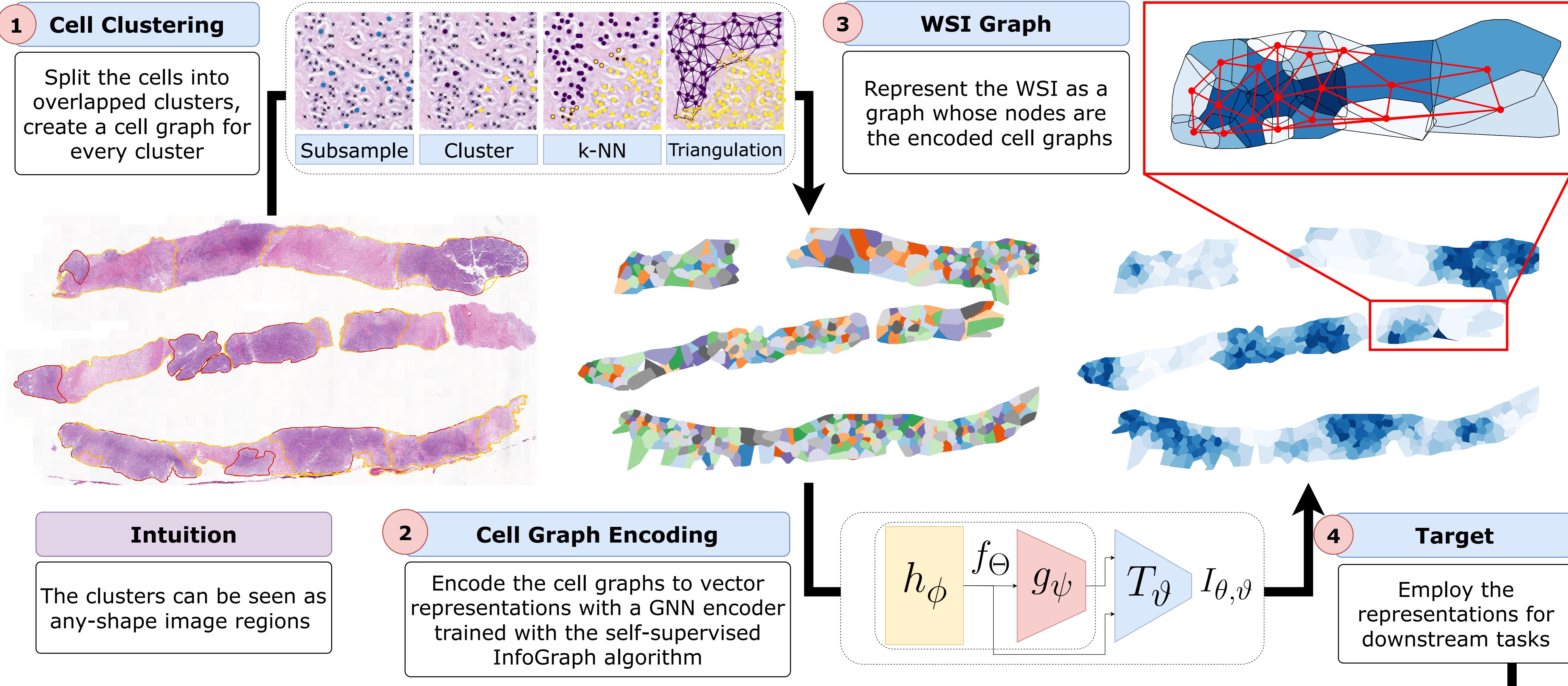
Graph representations of histology images have been proposed in order to leverage the patch positional information with patch graphs (PG) and to model the relationships between biological entities with cell graphs (CG). However, the PG approach relies on a CNN that ignores the entity space whereas the CG can only be applied to small images. We propose a framework based on **self-supervised learning** for the **representation of WSIs as graphs** whose nodes are seen as any-shape regions of the image, extending the PG approach, and their features encode **cell information**, as vector representations of CGs of the regions.

Data

The data is provided by the *Institut Català de la Salut (ICS)*. We have two sets of breast and lung unannotated WSIs, which have been used independently to train two encoders.

Data type	Tissue	Annotations	Quantity	Extracted cell graphs
WSI	Breast	-	13	5948
WSI	Breast	Region	8	2780
WSI	Lung	-	7	4312
Patch	Lung	Cell	20	20

Method

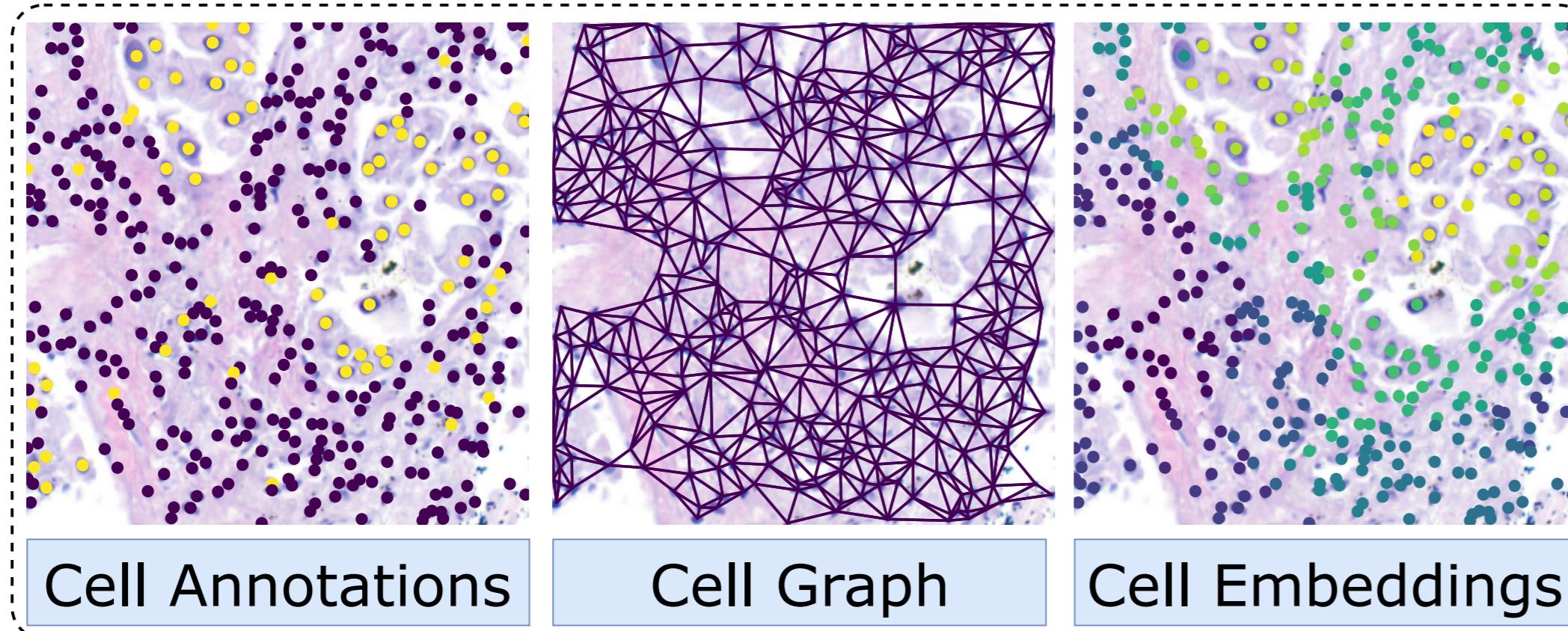


Linear Evaluation Protocol

Linear evaluation protocol at cell and region level

A Cell Level

Encoder trained with unannotated **lung WSIs**. The cell level evaluation is done on patches with labelled cells



B Region Level

Another encoder is trained with unannotated **breast WSIs**. For region level evaluation the annotated WSIs the cell graphs are labelled

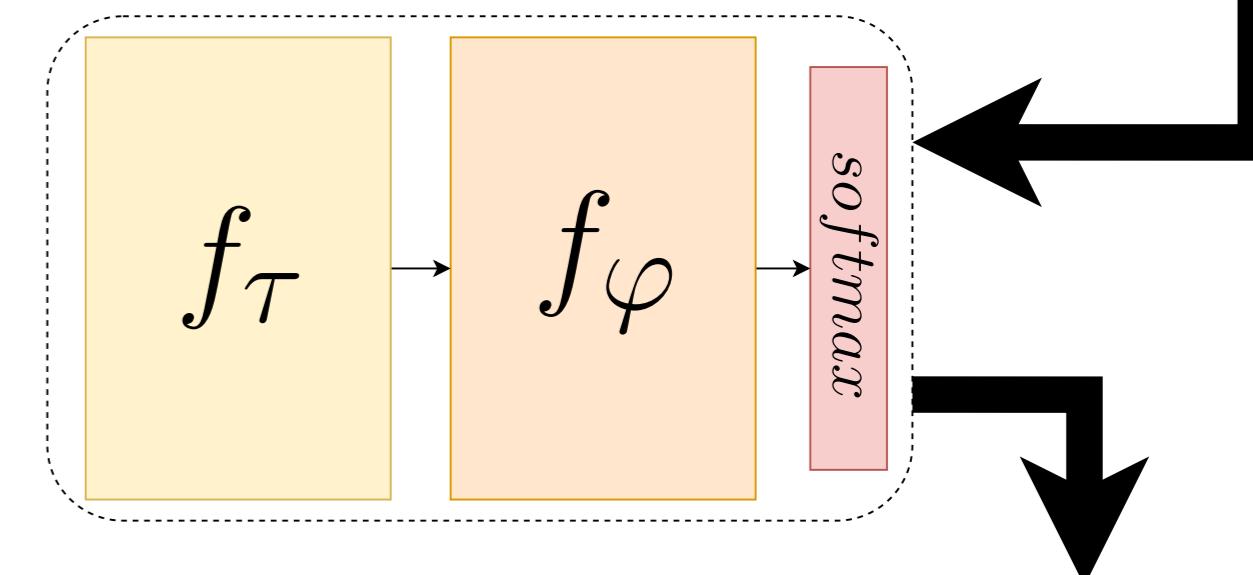
A	Input	Accuracy	Precision	Recall	F1
	Random	50.0 ± 1.0	34.3 ± 1.4	49.5 ± 1.6	40.5 ± 1.4
	x_i	78.6 ± 0.6	76.2 ± 3.1	54.7 ± 1.9	63.7 ± 1.2
	z_i	87.5 ± 0.7	83.3 ± 1.2	79.5 ± 1.5	81.3 ± 0.9

B	Method	Accuracy	Precision	Recall	F1
	Random	50.4 ± 0.6	79.8 ± 0.5	50.5 ± 0.9	61.9 ± 0.7
	$ V_k $	72.0 ± 0.6	72.2 ± 0.6	100.0 ± 0.0	83.7 ± 0.4
	\bar{x}_k	78.6 ± 0.6	79.8 ± 1.3	94.6 ± 2.0	86.4 ± 0.2
	z_k	86.5 ± 0.6	88.7 ± 0.9	92.7 ± 1.2	90.6 ± 0.4

Clustering for ROI detection

5 Graph Clustering

Perform graph clustering with Deep Modularity Networks algorithm for ROI detection



Intuition

Clustering with GNNs includes both graph topology and node embeddings



C Evaluation

Clustering	ARI	NMI	F
Random	0.0 ± 0.0	0.0 ± 0.0	11.6 ± 0.4
k-Means	18.3 ± 6.3	19.5 ± 6.5	30.3 ± 3.6
DMoN	19.8 ± 20.2	21.2 ± 15.7	60.9 ± 10.3

Conclusions

We present a framework for the analysis of WSIs that learns representations for overlapped regions of a slide based on the their cells and creates a graph of regions to represent the WSI. The scarce annotations available for unseen images during training are employed to evaluate the representations at both cell and region level, outperforming the raw input features. We also employ this graph representation for ROI detection addressed as a graph clustering task with GNNs that efficiently includes both graph topology and node embeddings.

References

- [1] Sun, F.-Y., Hoffman, J., Verma, V., & Tang, J. (2019). InfoGraph: Unsupervised and Semi-supervised Graph-Level Representation Learning via Mutual Information Maximization. *International Conference on Learning Representations*.

- [2] Tsitsulin, A., Palowitch, J., Perozzi, B., & Müller, E. (2020). Graph Clustering with Graph Neural Networks.