

Authors	Pub. Title	info	ID	pdf name
Peter Neidlinger (1,*), Omar S. M. El Nahhas (1, 2, *), Hannah Sophie Campanella, Shengjia Chen, Manbir Singh, Ruchika V	Benchmarking foundation models as feature extractors for weakly-supervised computational pathology	Benchmarking foundation models as feature extractors for weakly-supervised computational pathology	1	neidlinger_benchmarking_foundation_models_as_feature_extractors_for_weakly-supervised_computational_pathology.pdf
Gabriele Campanella, Shengjia Chen, Manbir Singh, Ruchika V	A clinical benchmark of public self-supervised pathology foundation models	A clinical benchmark of public self-supervised pathology foundation models	2	fm_features_mit_benchmark.pdf
Rohan Bareja, Francisco Carrillo-Perez, Yuanning Zheng, Marija Pilić	Evaluating Vision and Pathology Foundation Models for Computational Pathology: A Comprehensive Benchmark Study	Evaluating Vision and Pathology Foundation Models for Computational Pathology: A Comprehensive Benchmark Study	3	path_nonpath_evaluation.pdf
Narmin Ghaffari Laleh a, Hannah Sophie Muti a, Chiara Maria Lavin	Benchmarking weakly-supervised deep learning pipelines for whole slide classification in computational pathology	Benchmarking weakly-supervised deep learning pipelines for whole slide classification in computational pathology	4	cpath_pipeline_evaluation.pdf
Jeaeun Lee, Jeewoo Lim, Keunho Byeon, Jin Tae Kwak	Benchmarking Pathology Foundation Models: Adaptation Strategies and Scenarios	Benchmarking Pathology Foundation Models: Adaptation Strategies and Scenarios	5	peft.pdf
Mingu Kang * Heon Song * Seonwook Park Donggeun Yoo Sérgio	Benchmarking Self-Supervised Learning on Diverse Pathology Datasets	Benchmarking Self-Supervised Learning on Diverse Pathology Datasets	6	ssl.pdf
Shengjia Chen, Gabriele Campanella, Abdulkadir Elmas, Aryeh Stor	Benchmarking Embedding Aggregation Methods in Computational Pathology: A Clinical Data Perspective	Benchmarking Embedding Aggregation Methods in Computational Pathology: A Clinical Data Perspective	7	embeddingaggregation.pdf

name	SSL method	alias(es)	info	paper	used in bench weights	
TANGLE				Transcriptomics-guided Slide Representation Learning	https://huggingface.co/Path-VLM/tangle-768-vlm	
ToPoFM				ToPoFM: Topology-Guided Pathology Foundation		
HistoGPT				Generating dermatopathology reports from gigapath		
CtransPath	contrastive learning		CTransPath (dim:768) integrates a CNN and Swin Transformer,	[1,2,3,5,7]		https://drive.google.com/drive/folders/1UW0tYDQFvTjEgRnXZlHwzCkKdLm3BfV
UNI	DINOv2		UNI (dim:1024)Chen et al. in August 2023	[1,2,3,5,7]		
Phikon	iBOT			Scaling Self-Supervised Learning for Histopathology		[1,2,3,5]
Virchow	DINOv2			Vorontsov et al. in September 2023		[1,2,3]
Virchow2	DINOv2			Zimmermann et al. in August 2024		[1,2,3]
H-optimus-O	DINOv2			July 2024 by Saillard et al.		[1,2,3]
Prov-GigaPath	DINOv2				[1,2,3]	
Kaiko	DINO				[1,3]	
Hibou	DINOv2			Nechaev et al., released in June 2024	[1,3]	
PLIP	CLIP		Path-VLM	Huang et al. in August 2023; A visual–language foundation model for pathology	[1,3] https://huggingface.co/Path-VLM/plip-768-vlm	
BiomedCLIP	CLIP		Path-VLM	Zhang et al. in March 2023	[1,3] https://huggingface.co/Path-VLM/biomedclip-768-vlm	
CONCH	CoCa		Path-VLM; CONCH was trained on proprietary datasets rather than public ones		[1,3]	
DinoSSLPPath	DINOv1			published by Kang et al. in December 2022	[1] https://github.com/kanglab/dinoSSLPPath	
ViT-L14	DINOv2		by Kaiko.ai		[1]	
PRISM			multimodal slide analysis	Shaikovski et al. in May 2024	[1]	
MADELEINE				Jaume et al. in August 2024	[1]	
CHIEF				Wang et al. in September 2024; A pathology foundation model	[1]	
Phikon-v2	DINOv2				[2,3]	
tres50_imagenet		ResNet-50	Truncated ResNet50 (tres50 imagenet, dim:1024), pretrained on ImageNet		[2,4,6,7]	
ViT "Vision Transformers"		SP85M	publicly available on HuggingFace		[2,4]	
Vit-S		SP22M	publicly available on HuggingFace		[2,6]	
Rudolf-V	DINOv2			RudolfV: A Foundation Model by Pathologists for Computational Pathology at Health System Scale	[2]	
Campanella Et al.	DINO			Computational Pathology at Health System Scale	[2]	
Campanella Et al.	MAE			Computational Pathology at Health System Scale	[2]	
Lunit	DINO			Benchmarking Self-Supervised Learning on Dividing Cells	[3,5] https://github.com/lunit-lab/lunit	
HIPT	DINO				[3] https://github.com/HIPT-Histopathology-Image-Pretraining/hipt	
BEPH	BeiT			A foundation model for generalizable cancer diagnosis	[3] https://drive.google.com/drive/folders/1UW0tYDQFvTjEgRnXZlHwzCkKdLm3BfV	
GPFM	DINOv2				[3]	
UNI2	DINOv2				[3]	
QuiltNet-B16	CLIP		Path-VLM		[3] https://huggingface.co/Path-VLM/quiltnet-b16-vlm	
Mizero pubmed	contrastive learning		Path-VLM		[3] https://drive.google.com/drive/folders/1UW0tYDQFvTjEgRnXZlHwzCkKdLm3BfV	
Mizero clinmedbert	contrastive learning		Path-VLM		[3] https://drive.google.com/drive/folders/1UW0tYDQFvTjEgRnXZlHwzCkKdLm3BfV	
TITAN	Ibot + CoCa		Path-VLM		[3] https://huggingface.co/Path-VLM/titan-768-vlm	
Dino-S16	DINO		VM		[3]	
Dino-B16	DINO		VM		[3]	
Dinov2	DINOv2		VM		[3]	
IBOT-B16	iBOT		VM		[3]	
IBOT-L16	iBOT		VM		[3]	
CLIP-B16	CLIP (contrastive learning)		VLM		[3]	
BLIP-B16-14M	BLIP		VLM		[3]	
Align-base	EfficientNet + BERT (Contrastive)		VLM		[3]	
beit3-L16	iBOT		VLM		[3]	
Imagenet efficientnet-b7					[4]	
dinosmall			DINO-ViT small (dinosmall, dim:384), pretrained on 1.6 billion histology images		[7]	
PathDino					https://huggingface.co/Path-VLM/pathdino-768-vlm	