Transformers, Vision Transformers and SAMJ

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Content

Transformers and Vision Transformers

Segment Anything Model (SAM) and SAM-like models

SAMJ

Hands on activities

Transformers and Vision Transformers



The Transformer

Introduced in 2017 by Vaswani et al, from Google



New architecture "just" for language translation

Currently is the cornerstone of the Artificial Intelligence revolution





ChatGPT



Music generation



Protein folding



Figure 1: The Transformer - model architecture.

3 key contributions

Sef-attention

Multi-head attention

Positional encoding

Attention Is All You Need

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Embeddings locate similar ideas together



Attention blocks

self-attention + multihead attention



change the "meaning" of words given the context











Jay Alammar Visualizing machine learning one concept at a time. @JayAlammar on Twitter, YouTube Channel

Blog About

The Illustrated Transformer

Discussions: Hacker News (65 points, 4 comments), Reddit r/MachineLearning (29 points, 3 comments) Translations: Arabic, Chinese (Simplified) 1, Chinese (Simplified) 2, French 1, French 2, Italian, Japanese, Korean, Persian, Russian, Spanish 1, Spanish 2, Vietnamese Watch: MIT's Deep Learning State of the Art lecture referencing this post

Featured in courses at Stanford, Harvard, MIT, Princeton, CMU and others

3blue1brown videos on Transformers

The Illustrated Transformer

Generative Pre-trained Transformer (GPT)



Trained for next token prediction

Emerging capabilities

Works for translation, question answering... (!!!!)



Generative Pre-trained Transformer



Generative Pre-trained Transformer

Scaling the model works

Prompt : Python code to find the smallest factor of a number

GPT-1:

the lack of a body in the room before me. after several moments of silence, he spoke again. " you are my daughter. the two of us are one. and in time you will

GPT-2

let $p = \&[5 - 3] \Rightarrow (1, 2, 3)$ The example above uses the "P" module to print the smallest factor of the number:

GPT-3

import math def lfact (n): factors = [1] factors . append (n) while n % factor <> 0: factor = factor * 2 - 1 while int (factor) > n % factor : factors . append (factor) return factors

What the code does:

Imports math module for math functions. [..]

Vision Transformer (ViT)

Using transformers for vision

AN IMAGE IS WORTH 16x16 WORDS: TRANSFORMERS FOR IMAGE RECOGNITION AT SCALE

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> *equal technical contribution, [†]equal advising Google Research, Brain Team {adosovitskiy, neilhoulsby}@google.com

Required huge amounts of data and params to outperform CNNs



Vision Transformer (ViT)





Transformers vs CNNs

BiT -Big Transfer (CNN) ImageNet - 1.2m images ImageNet21k - 14m images ViT-H > ViT-L > ViT-B



Transformers in Vision - Useful resources

Overview of ViTs with one of the authors

ViT explanation with code

ViTs for small datasets

ViTs for small datasets (the whole channel is quite good)

Foundational models for Vision: <u>SAM</u> and <u>Dino</u>





Transformers are data hungry

Transformers need a LOT of data

The bigger they are and the more data they see the more they learn about it

Transformers can learn relationships between anything

aminoacids -> protein folding



images





Everything to everything models -Multimodal transformers













Natural images		Object detection	
Trend	Dataset	Best Model	
	COCO test-dev	Co-DETR	ViT
	COCO minival	Co-DETR	ViT
	сосо-о	EVA	ViT
:	PASCAL VOC 2007	Cascade Eff-B7 NAS-FPN (Copy Paste pre-training, single-scale)	CNN



Colonoscopy			Medical Image segmentation		
images	Trend	Dataset	1	Best Model	
		Kvasir-SEC	G	DUCK-Net	CNN
		CVC-Clinic	cDB	DUCK-Net	CNN
	·	CVC-Color	nDB	DUCK-Net	CNN
		ETIS-LARI	BPOLYPDB	DUCK-Net	CNN



		Medical Image segmentation		
CT scans	* . * . * .	Synapse multi-organ CT	Swin UNETR	CNN
MRI cardiac images	an an an an an	Automatic Cardiac Diagnosis Challenge (ACDC)	FCT	CNN
Tissue images		MoNuSeg	Hi-gMISnet	CNN
Nuclei images		2018 Data Science Bowl	DuAT	ViT
Gland segmentation in Colon Histology images	10 AL AL AL AL	GlaS	Hi-gMISnet	CNN

Tasks with **millions of images available** are dominated by **transformers**

Specific tasks with **more difficult data acquisition** are still dominated by **CNNs**



Transformers in Microscopy - Cell segmentation

Transformers still **underperform** CNN methods for cell segmentation

Cellpose (CNN) method is still the king

Analysis | Published: 26 March 2024

The multimodality cell segmentation challenge: toward universal solutions

Jun Ma, Ronald Xie, Shamini Ayyadhury, Cheng Ge, Anubha Gupta, Ritu Gupta, Song Gu, Yao Zhang, Gihun Lee, Joonkee Kim, Wei Lou, Haofeng Li, Eric Upschulte, Timo Dickscheid, José Guilherme de Almeida, Yixin Wang, Lin Han, Xin Yang, Marco Labagnara, Vojislav Gligorovski, Maxime Scheder, Sahand Jamal Rahi, Carly Kempster, Alice Pollitt, ... Bo Wang 🗠 + show authors

Nature Methods (2024) Cite this article

13k Accesses | 65 Altmetric | Metrics

Debunked by

Cellpose with transformer backbone underperforms CNN backbone

Transformers do not outperform Cellpose

Carsen Stringer[†], Marius Pachitariu[†]

HHMI Janelia Research Campus, Ashburn, VA, USA

[†] correspondence to (stringerc, pachitarium) @ janelia.hhmi.org

CNN

ViT

Transformers in Microscopy - Cell segmentation

Article | Published: 14 December 2020

Cellpose: a generalist algorithm for cellular segmentation

Carsen Stringer, Tim Wang, Michalis Michaelos & Marius Pachitariu 🖾

Nature Methods 18, 100-106 (2021) | Cite this article

82k Accesses | 990 Citations | 176 Altmetric | Metrics

Transformers do not outperform Cellpose

Carsen Stringer[†], Marius Pachitariu[†] HHMI Janelia Research Campus, Ashburn, VA, USA [†] correspondence to (stringerc, pachitarium) @ janelia.hhmi.org

Cellpose authors claim that ViTs success may not translate to biological images It may be impossible to collect millions of diverse biological images for training



Transformers in Microscopy - Cell segmentation

SAM (Segment Anything model) performs well on cell data

Training data of natural images, cell images were a small percentage

There might be hope for ViTs in cell images



https://segment-anything.com/demo#

29

The story of Uncle SAM





Segment-Anything Model (SAM)

Foundation model from MetaAl



Transform: encoding / decoding



PROMPT

The ChatGPT of the Computer Vision



Model SA-1B

BIG DATA

- Natural photographies
- Huge model (~1GB)
- 11M diverse, high-res. images
- 1.1B segmentation masks
- Open, privacy







Alexander Kirillov et al. IEEE/CVF, 2023, 2700 citations



dation

Models

SAM for Science?



Variants of SAM Models

- MicroSAM
- MedSAM
- CellSAM
- EfficientSAM
- MobileSAM
- ...







MicroSAM C. Pape CellSAM

ARCGIS

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Acceleration of annotations Megakaryocotes on human biopsis SAM Large model SAM extension of WuPath SAM on server R. Sarkis, CHUV, L-F. Celma, EPF April 2024

Web interface Python package QuPath Napari Fiji

SAMJ Annotation with SAM on FIJI (CPU)

SAMJ

- FIJI Plugin and ICY plugin
- Model Efficient SAM (run on CPU)
- Automatic installation of the Python environment
- Smart strategy for tiling





SAMJ Team: Carlos, Caterina, Arrate, Vladimir Ulman, Adrian Ines, Jonathan Heras, Curtis Rueden, Jean-Christophe, Daniel



Segment Anything Model (SAM) and SAM-like models



Segment Anything Model

https://segment-anything.com

by Meta Al



Segment Anything Model



Promptable Segmentation (bounding box and points)



Real-time interaction (~50 miliseconds)



1 Billion masks, 11 Million images



Manual to automatic annotation process



Real-time web browser interaction



Real-world scenarios



Diverse and high-resolution images



Vision Transformer-based Architecture (ViT)



Zero-Shot Capabilities

Ethical and fairness focus



Image		Prompt points, box, masks
Ļ		↓
		Prompt encoder
		Ļ
	\rightarrow Embedding \rightarrow	Mask decoder
Image encoder		Ļ
		Final Mask
GPU @.15s		Web-browser (CPU) @.055s





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Segment Anything Data Engine

1

Model-assisted manual annotation stage (2)

Semi-automatic stage with a mix of automatically predicted masks and model-assisted annotation

3

Fully automatic stage, model generates masks without annotator input



SAM's Zero-Shot transfer capabilities on image types





SAM for Science

- MicroSAM
- CellSAM
- MedSAM

• • • •



MicroSAM C. Pape



CellSAM



ARCGIS



EfficientSAM

https://yformer.github.io/efficient-sam/

by Y. Xiong et al.



EfficientSAM

Develop SAMI, a masked image pretrained framework to reconstruct features from SAM ViT-H image encoder

SAMI-pretrained backbone generalize to many tasks including classification Deliver EfficientSAM, a light-weight SAM model for practical deployment



EfficientSAM









EfficientSAM: points



Input (left), SAM (middle), EfficientSAM (right)



EfficientSAM: ROIs



Input (left), SAM (middle), EfficientSAM (right)



EfficientSAM: Everything



Input (left), SAM (middle), EfficientSAM (right)



SAMJ

https://github.com/segment-anything-models-java/SAMJ-IJ











SAMJ Functionalities

SAM-like models implementation

Semi-automatic annotation

Adaptable to different use cases

No need of GPU

Different models based on SAM (e.g. EfficientSAM) available for your annotations Annotation of objects through **different prompts** (bounding box, points, etc) in seconds Capable of performing annotations over **different images**, cell types, morphologies, etc

Using different SAM-like models with a CPU as using lighter versions



SAMJ usage example



Original Image*





Generated Mask

SAMJ Annotations

*Original Image obtained from the Cell Tracking Challenge



SAMJ usage example



Original Image*





Generated Mask

SAMJ Annotations

*Original Image obtained from doi.org/10.1016/j.compbiomed.2021.104673



Hands on activities



Counting nuclei!



- 1. Download the image from the GitHub repository.
- 2. Open Fiji and SAMJ plugin.
- 3. Open the image and encode it with the SAMJ plugin.
- 4. Start annotating nuclei for 20 seconds.

Who can annotate more?

Annotation of myelin sheaths on Fiji!



- 1. Segment by using few preprocessing set and then threshold
- 2. Annotate by hand (mouse).
- 3. Annotate using the magic wand of Fiji (select the tolerance) and then interpolate the selection (menu Edit>Selection>Interpolate)
- 4. Annotate using SAMJ
- 5. Comment these 4 methods in term of speed of annotation, accuracy of segmentation, and required resources.



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