LMP 1210 - Basic principles of machine learning in biomedical research

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Announcements

- Next week, we will start project presentations.
- 6 teams on March 28
 - Your project does not need to be 100% done, report what remains to be done for the upcoming week
- 7 teams on April 4
- Each team will have 15 minutes total 12 minutes for presentation
- Remember:
 - Most of your grade is based on the presentation + project report.
 - Reports will be graded by me and the TAs, presentations by me
- Bring:
 - A *practiced* presentation
 - Laptop
 - HDMI Adaptor

Grading rubric

- Scope of the project
- How much of the initial project proposal you have completed
- How well you present the core idea:
 - What problem do you care about
 - Why do you care about solving it?
 - How did you solve it?
 - When you implemented your method, what happened?
- What have you learned through doing the project?

Overview

- Limitations of supervised learning
- Self supervised learning
- [Time permitting: Transformers]
- Imaging in computational histopathology

Imaging in medicine

- <u>History of Medical Imaging, Bradley et. al, 2008</u>
- Nuclear medicine: Using radiation to see inside the human body
 - X-ray discovered in 1895 (won the Nobel in 1901)
 - CT, PET discovered thereafter
- Magnetic resonance imaging: Mapping resonance in the body to images
- Ultrasound imaging: Mapping high-frequency sound waves to images
- Histopathological imaging: Images of stained tissue samples



 Given a dataset, the model parameters are learned via maximum likelihood estimation

$$\mathcal{L}(y, x) = \log p(y|x; \theta)$$

N

Score function (high is good, low is bad)

 $\theta = rg\max_{\theta} \sum_{i=1}^{\mathcal{L}} \mathcal{L}(y_i, x_i)$ Solve this optimization problem to learn of the negative of the log-likelihood function

Deep neural networks typically learned using tools that leverage automatic differentiation



Computer vision

Computer vision has had a front row seat to the advances in deep learning
 ImageNet Challenge

Deng, Jia, et al. "Imagenet: A large-scale hierarchical image database." *2009 IEEE conference on computer vision and pattern recognition*. leee, 2009.

IM A GENET

- 1,000 object classes (categories).
- Images:
 - \circ 1.2 M train
 - **100k test.**



Error rates on Imagenet over time



Neural networks in a slide

- Simplest neural network is a multi-layer perceptron
- Neural networks are known to be universal function approximators



Convolutional neural networks

Capture the fact that we may want representations that are spatially invariant



Deep residual neural networks



Researchers found that deep networks had a hard time learning the identity function.

They added a skip-connections between layers:

$$h_k = \phi(conv(h_{k-1})) + \sum_{j < k-1} h_{k-j}$$

Deep Residual Learning for Image Recognition, He et. al, 2015

Limitations of supervised learning

- Deep neural networks have proven very successful in learning useful representations of image data from large datasets
- Models like AlexNet, ResNet trained on imagenet capture features useful for multiple different tasks
- For a new task:
 - Need fine-grained labels associated with each example
 - Standard approach: Use a pre-trained imagenet model and fine-tune on new dataset
- Self-supervised learning:
 - What if we do not need labels to learn good representations?







$$\theta = \arg \max_{\theta} \sum_{i=1}^{3} \mathcal{L}(x;\theta) + \mathcal{L}(y_1|x_1;\theta_2) + \mathcal{L}(y_3|x_3;\theta_2)$$

 Have a combination of labelled and un-labelled data in your dataset

Unsupervised and semi-supervised learning of high-dimensional images is hard

- Even if there is a small space of concepts unsupervised models of image data are challenging to build
- Need a good model of each pixel in the image.
- Recently there has been a lot of work in leveraging generative adversarial networks for this problem
- Idea: Can we build representations without labels and without modeling each pixel as a random variable?

Self-supervised learning

- Recent (last 4-5 years) development in machine learning
- **Principle**: Leverage domain knowledge about what kinds of information the representation should contain when building it
- Learn about self-supervised learning by examples

Notation

$$\mathcal{T}: x \to \tilde{x}$$

 ${\mathcal O}$

- Transformation of an image [random crop, rotation, jittering, color normalization]
 - Preserves the identity of the image

- sim(k, k')
- Similarity function
 - Measure of similarity of two vectors
 - Mean squared error, cosine similarity

SSL 1 - Learning with contrastive examples

- <u>A Simple Framework for Contrastive Learning of Visual</u> <u>Representations, Chen et. al, ICML 2020</u>
- Builds upon earlier work: <u>Unsupervised Feature Extraction by</u> <u>Time-Contrastive Learning and Nonlinear ICA, Hyvarinen et. al</u>

SIMCLR: Self-supervised learning with contrastive examples



Randomly sample a mini-batch of datapoints.

Minimize loss below

Goal: Learn representations that recognize that the class of transformations in T preserve identity.

Note: No labels used.

$$i_{i,j} = -\log P \frac{\exp(\sin(z_i, z_j))}{\sum_{k=1}^{2N} \mathbb{I}_{[k \in i]} \exp(\sin(z_i, z_k))} , \quad (1)$$

How good are the representations?

	Food	CIFAR10	CIFAR100	Birdsnap	SUN397	Cars	Aircraft	VOC2007	DTD	Pets	Caltech-101	Flowers
Linear evaluation	on: 76.9	95.3	80.2	48.4	65.9	60.0	61.2	84.2	78.9	89.2	93.9	95.0
Supervised	75.2	95.7	81.2	56.4	64.9	68.8	63.8	83.8	78.7	92.3	94.1	94.2
SimCLR (ours)	89.4	98.6	89.0 88.7	78.2	68.1	92.1	87.0 88.0	86.6	77.8 78 8	92.1	94.1 94.2	97.6
Random init	88.3	96.0	81.9	77.0	53.7	91.4	84.8	69.4	64.1	82.7	72.5	92.5

A Simple Framework for Contrastive Learning of Visual Representations

Table 8. Comparison of transfer learning performance of our self-supervised approach with supervised baselines across 12 natural image classification datasets, for ResNet-50 (4×) models pretrained on ImageNet. Results not significantly worse than the best (p > 0.05, permutation test) are shown in bold. See Appendix B.8 for experimental details and results with standard ResNet-50.

SSL - Learning without contrastive examples

- In the above examples, the quality of representations will depend on the choice of negative examples used.
- Can we learn without negative examples?
- <u>DINO: Emerging Properties in Self-Supervised Vision</u> <u>Transformers, Caron et. al, 2021</u>
 - Key idea: Instead of comparing the representations with respect to random negative examples, compare the representation to a different crop of itself

DINO

Decision making with images

- Ultrasound:
 - Echocardiograms
 - Visualize beating of the heart to assess normal function
 - Abdominal ultrasounds
 - Assess healthy function of abdominal organs
- X-rays:
 - Breast cancer screening
 - Guiding surgery to remove blood clots, insert catheters
 - Friday: Hear from Ruizhi Liao on combining text and chest x-ray data

Technical issues in machine learning for medical imaging

- The general setup is almost always as follows:
 - Collect a large set of images [X]
 - Use notes/clinical variables/expert annotation to come up with labels [Y]
 - Use a deep learning model predict Y from X
- Fairness:
 - <u>Reading Race: AI Recognises Patient's Racial Identity In Medical</u> Images, Banerjee et. al, 2021
- Selection bias:
 - Causality matters in medical imaging, Castro et. al, 2019

Case study 1: Deep learning for echocardiograms

- Sound waves to image the heart
- Why:
 - Check for problems with your valves or chambers
 - Check if heart problems are causing shortness of breath
 - Assess congenital heart defects



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A taxonomy of echocardiograms

- Most common: Transthoracic echocardiogram
- Transesophageal echocardiogram
 - Transducer guided down patient's throat
 - Records sound waves bouncing off the heart pumping and interprets them as images
- Doppler echocardiogram
 - Used to assess bloodflow
- Stress echocardiogram
 - Ultrasound after excercise

Case study 1: Predicting cardiac amyloidosis

- Artificial intelligence-enabled fully automated detection of cardiac amyloidosis using electrocardiograms and echocardiograms, Goto et. Al, Nature Communications, 2021
- Cardiac amyloidosis
 - deposition of protein in the heart muscle, can result in heart failure
 - believed to be rare but likely underdiagnosed
 - manifests in both ECGs and echo-cardiography but features are not highly specific and difficult to spot
 - Gold standard: biopsy (costly and risky to patient)

Where machine learning can help

- How can we design a method that:
 - Fits into the clinical workflow for cardiac patients
 - If used, improve underdiagnosis of disease?
- Key-idea: Two-stage approach
 - Step 1: Build ML models from ECG data (readily available at most care providers)
 - Finding: Models have decent accuracy but not enough for conclusive diagnosis
 - Step 2: Build ML models from echocardiogram data
 - Finding: Models outperform human experts
 - Use step 1 to decide which patients should undergo an echocardiogram and apply model from step 2

A multi-center study

Table 1 Study-level demographic information (ECG cohort).												
	BWH		MGH		UCSF							
	Case	Control	Case	Control	Case	Control						
Number of studies	2249	8684	405	437	372	731						
Age, years ± SD	69.9 ± 10.4	62.3 ± 13.2	72.9 ± 9.0	73.8 ± 8.8	67.7 ± 12.9	67.5 ± 11.7						
Age Groups												
≤30, n (%)	2 (0.1)	97 (1.1)	1 (0.2)	1 (0.2)	2 (0.5)	0 (0.0)						
30-50, n (%)	78 (3.5)	1,370 (15.8)	7 (1.7)	6 (1.4)	36 (9.7)	69 (9,4))						
50-70, n (%)	901 (40.1)	4548 (52.4)	143 (35.3)	135 (30.9)	136 (36.6)	278 (38.0)						
70-90, n (%)	1242 (55.2)	2606 (30.0)	254 (62.7)	295 (67.5)	198 (53.2)	384 (52.5)						
>90, n (%)	26 (1.2)	63 (0.7)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)						
HR, bpm ± SD	76.4 ± 16.7	75.9 ± 18.5	78.6 ± 16.6	75.1 ± 19.8	79.6 ± 18.7	72.2 ± 16.3						
Sinus rhythm, n (%)	1,736 (77.2)	8,072 (93.0)	283 (69.9)	371 (84.9)	365 (98.1)	729 (99.7)						

Step 1: ECG model

 Results Ok but not considered good enough for evaluating interventions for a rare diagnosis since it will result in a large number of false positives





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Step 2: Echocardiogram model

 Performance significantly better when using a richer (but more expensive) data modality



Score: 0.17

Score: 0.04

Score: 0.99

Score: 1.00

Recall: Metrics

- Positive Predictive Value (PPV): TP/(TP+FP)
 - A high PPV will indicate that a positive result is likely correct
- Sensitivity: TP/(TP+FN)
 - A highly sensitive test will have few-false negatives

Analyzing the combined approach

- ECG model:
 - MGH: PPV 3.9% with Sensitivity 71%
 - BWH: PPV 3.4% with Sensitivity 52.4%
- Echo model:
 - MGH: PPV 32.7% with Sensitivity 66.9%
 - BWH: PPV: 33.4% with Sensitivity 67%
- Combined:
 - MGH: PPV: 76.6% with Sensitivity 47.5%
 - BWH: PPV: 73.9% with Sensitivity 34.8%

Transformers

- Built on the attention mechanism
- Many modern tools with deep learning are based off the transformer architecture (ChatGPT, Claude, Vision Transformers, OpenAI Whisper)
- General purpose neural network that works well on images, speech, text....
- Resources:
 - Roger and Jimmy Ba's lecture notes
 - <u>https://nlp.seas.harvard.edu/2018/04/03/attention.html</u>
 - <u>https://lilianweng.github.io/posts/2020-04-07-the-transformer-family/</u>

Case study 2: Deep learning for histopathological image data

 <u>Scaling Vision Transformers to Gigapixel</u> <u>Images via Hierarchical Self-Supervised</u> <u>Learning</u>, Chen et. al, CVPR 2022

Histopathological images in the clinical workflow

- Histopathology: Microscopic examination of tissue to study diseases and their different presentations,
- Pipeline:
 - Surgery, biopsy or autopsy for excision of tissue
 - Placed in a fixative to stabilize tissue
 - Investigated under a microscope
- Histopathological images are routinely used for clinical diagnoses of cancer
 - Key question: How can we use machine learning to build representations of histopathological image data?

Slide-Level Supervised Learning (Weak Supervision)



Lipkova et al. 2021, In Review

Weakly-Supervised Learning: Finding Needles in Haystacks via Attention



Lu et al. 2021, Nature BME

Current Paradigm is limited by: Clinical Domain Knowledge

- Requires clinical domain knowledge to:
 - label image regions in WSIs with known morphological phenotypes (patch-level tasks)
 - 2. Make prognostic decisions from subjective interpretation of the entire WSI (slide-level tasks)
- How can we identify new phenotypic biomarkers?
- What are we missing in current decision-making that can guide prognosis?



Current Paradigm is limited by: Clinical Domain Knowledge



Current pipelines for creating representations of whole slide images make use of ResNet50 architectures pretrained on imagenet.

Self-Supervised Learning: Pixel-Level Annotations are Not Needed!



Lipkova et al. 2021, In Review, Ciga et. al

We build upon recent work [Resource and data efficient self supervised learning, Ciga et. al, 2021] who show that self-supervision yields general purpose representations of histopathological images

DINO-based Knowledge Distillation for Patch-based Representations

Loss Function: p1 log (p2)



Chen et al. 2021, In Preparation

DINO

- We wanted to study the use of non-contrastive self-supervised learning for creating representations
- Input:
 - Two crops with color contrasts from the same image
- Goal of self-supervised learning:
 - Teach the network that these two crops are from the same image
 - Output of student network is trained to match the distribution of teacher network via minimizing cross-entropy loss
 - Avoid network collapse by having two networks
 - Train the student via gradient descent
 - Teacher is not trained, weights are updated via exponential moving average from students
- Does not require negative samples
 - Data inductive biases in natural images may not hold in H&E pathology slides

DINO: Emerging properties in self-supervised vision transformers, Caron et. Al,

DINO-based Knowledge Distillation for Patch-based Representations

Loss Function: p1 log (p2)



DINO

- Output of student network is trained to match the distribution of teacher network via:
 - minimizing cross-entropy loss
 - EMA to update teacher network
- Does not require negative samples
 - Data inductive biases in natural images may not hold in H&E pathology slides
- Vision Transformer (ViT) used as encoder
 - 256 x 256 H&E tissue patches are further patched as 16 x 16 patch embeddings

Chen et al. 2021, In Preparation

Study Design



Adenocarcinoma (40%)

This is the most prevalent form of lung cancer and usually arises in the cells lining the alveoli. It is a common form of lung cancer in people who have never smoked, but is also seen in smokers.

• Experiments:

- Organ-specific vs. pan-cancer training
 - TCGA Lung (1033 WSIs) vs Entire TCGA (~8788 WSIs)
- Comparisons with SOTA methods
 - SimCLR, SimSiam

Squamous cell carcinoma (30%)

These tumours appear in the flat cells that line the inside of the airways, usually near the bronchi. This form of the disease is usually caused by smoking and is more common in men than women. The tumours tend to grow slowly. Slide-Level Tasks:

- LUAD vs. LUSC Subtyping
- LUAD + LUSC Survival Analysis
- TP53 + KRAS Mutation Prediction

Results [1]: TCGA Lung Subtyping (LUAD vs. LUSC)



Method							
ImageNet Transfer	ResNet-50	ImageNet	100	0.945 ± 0.018	0.943 ± 0.019	0.917 ± 0.024	0.888 ± 0.031
SimCLR SimSiam DINO	ResNet-50 ResNet-50 ViT	Lung Only Lung Only Lung Only	100 100 100	$\begin{array}{c} 0.950 \pm 0.026 \\ 0.952 \pm 0.017 \\ 0.948 \pm 0.021 \end{array}$	$\begin{array}{c} 0.947 \pm 0.017 \\ 0.944 \pm 0.018 \\ 0.942 \pm 0.019 \end{array}$	$\begin{array}{c} 0.934 \pm 0.025 \\ 0.935 \pm 0.026 \\ 0.937 \pm 0.021 \end{array}$	$\begin{array}{c} 0.897 \pm 0.028 \\ 0.897 \pm 0.029 \\ 0.928 \pm 0.024 \end{array}$
SimCLR SimSiam DINO	ResNet-50 ResNet-50 ViT	Pan-Cancer Pan-Cancer Pan-Cancer	100 100 100	$\begin{array}{c} 0.951 \pm 0.016 \\ 0.493 \pm 0.085 \\ \textbf{0.957} \pm \textbf{0.019} \end{array}$	$\begin{array}{c} 0.948 \pm 0.017 \\ 0.534 \pm 0.072 \\ 0.949 \pm 0.019 \end{array}$	$\begin{array}{c} 0.930 \pm 0.023 \\ 0.508 \pm 0.085 \\ \textbf{0.941} \pm \textbf{0.022} \end{array}$	$\begin{array}{c} 0.898 \pm 0.026 \\ 0.603 \pm 0.040 \\ \textbf{0.931} \pm \textbf{0.024} \end{array}$

• Self-supervised feature extractors from DINO are more data-efficient than pretrained ResNet-50 on ImageNet for subtyping

Results [2]: TCGA Lung Subtyping (LUAD vs. LUSC) + Mutation Prediction (TP53 + KRAS)

Lung Adenocarcinoma (LUAD)

Lung Squamous Cell Carcinoma (LUSC)

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Method	Model Architecture	Training Source	Epochs	100%	75%	50%	25%	TP53	KRAS
ImageNet Transfer	ResNet-50	ImageNet	100	0.945 ± 0.018	0.943 ± 0.019	0.917 ± 0.024	0.888 ± 0.031	$\textbf{0.756} \pm \textbf{0.053}$	0.761 ± 0.073
SimCLR SimSiam DINO	ResNet-50 ResNet-50 ViT	Lung Only Lung Only Lung Only	100 100 100	$\begin{array}{c} 0.950 \pm 0.026 \\ 0.952 \pm 0.017 \\ 0.948 \pm 0.021 \end{array}$	$\begin{array}{c} 0.947 \pm 0.017 \\ 0.944 \pm 0.018 \\ 0.942 \pm 0.019 \end{array}$	$\begin{array}{c} 0.934 \pm 0.025 \\ 0.935 \pm 0.026 \\ 0.937 \pm 0.021 \end{array}$	$\begin{array}{c} 0.897 \pm 0.028 \\ 0.897 \pm 0.029 \\ 0.928 \pm 0.024 \end{array}$	$\begin{array}{c} 0.694 \pm 0.073 \\ 0.698 \pm 0.084 \\ 0.751 \pm 0.041 \end{array}$	$\begin{array}{c} 0.737 \pm 0.044 \\ 0.681 \pm 0.117 \\ \textbf{0.771} \pm \textbf{0.059} \end{array}$
SimCLR SimSiam DINO	ResNet-50 ResNet-50 ViT	Pan-Cancer Pan-Cancer Pan-Cancer	100 100 100	$\begin{array}{c} 0.951 \pm 0.016 \\ 0.493 \pm 0.085 \\ \textbf{0.957} \pm \textbf{0.019} \end{array}$	$\begin{array}{c} 0.948 \pm 0.017 \\ 0.534 \pm 0.072 \\ 0.949 \pm 0.019 \end{array}$	$\begin{array}{c} 0.930 \pm 0.023 \\ 0.508 \pm 0.085 \\ \textbf{0.941} \pm \textbf{0.022} \end{array}$	$\begin{array}{c} 0.898 \pm 0.026 \\ 0.603 \pm 0.040 \\ \textbf{0.931} \pm \textbf{0.024} \end{array}$	$\begin{array}{c} 0.687 \pm 0.100 \\ 0.516 \pm 0.073 \\ 0.746 \pm 0.051 \end{array}$	$\begin{array}{c} 0.711 \pm 0.127 \\ 0.612 \pm 0.051 \\ 0.740 \pm 0.052 \end{array}$

• Self-supervised feature extractors from DINO are more data-efficient than pretrained ResNet-50 on ImageNet for subtyping

• No difference found in gene mutation prediction

Results [3]: DINO Attentions to Cellular Identities



Questions?