

Exercise Session 1

Learning at Scale: Supervised, Self-Supervised, and Beyond

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Additional Reading Materials. We recommend following three papers:

- [1] Oord, Aaron van den, Yazhe Li, and Oriol Vinyals. “**Representation Learning with Contrastive Predictive Coding.**” arXiv preprint arXiv:1807.03748 (2018).
- [2] Chen, Ting, et al. “**A Simple Framework for Contrastive Learning of Visual Representations.**” International Conference on Machine Learning (ICML). PMLR, 2020.
- [3] Radford, Alec, et al. “**Learning Transferable Visual Models from Natural Language Supervision.**” International Conference on Machine Learning. PMLR, 2021.

Task 1. InfoNCE in Contrastive Learning.

In contrastive learning, the goal is to learn useful representations without labels. For each condition c , the model is trained to identify the single positive sample among K distractors. For example, in instance discrimination for images, c is one augmented view of an image, x^+ is another independent augmentation of the same image, and the negatives $\{x_i^-\}$ are views of other images (e.g., from the same minibatch or from a memory queue). In vision language tasks, c is an image, x denotes captions, x^+ is the matched caption, and the negatives are captions of other images. A common objective for this is the InfoNCE loss. We will investigate it from a probabilistic perspective in this task.

Suppose that, given a condition c , we form a candidate set $X = \{x_0, \dots, x_K\}$ where $x_0 = x^+$ is the positive sample and x_1, \dots, x_K are K negative samples drawn i.i.d. from $q_{\text{noise}}(x)$. The InfoNCE objective is defined as

$$\mathcal{L}_{\text{InfoNCE}} = - \sum_{(X,c) \in D} \left[\log \frac{\frac{p_{\text{data}}(x^+ | c)}{q_{\text{noise}}(x^+)}}{\sum_{j=0}^K \frac{p_{\text{data}}(x_j | c)}{q_{\text{noise}}(x_j)}} \right],$$

where the term inside the logarithm represents the probability of correctly identifying x^+ as the positive sample among all candidates.

In practice, we typically set $q_{\text{noise}}(x) = p_{\text{data}}(x)$, so the numerator simplifies to $\frac{p_{\text{data}}(x^+ | c)}{p_{\text{data}}(x^+)}$, whose logarithm equals the pointwise mutual information. Then, we can encode x and c with

neural networks to obtain embeddings, define a score to approximate this density ratio. Consequently, the InfoNCE loss encourages the model to learn effective representations that captures mutual information between x and c , by learning a higher estimated ratio (or score) for true positive pairs than for negative samples.

- (a) **Connection to cross-entropy loss.** Choose logits whose exponentials match the relative weights of candidates in the InfoNCE numerator/denominator; then build connection between InfoNCE and cross-entropy loss.
- (b) **Relation to NCE.** The local NCE loss is defined as

$$\mathcal{L}_{\text{NCE}} = \sum_{(X,c) \in D} \left[-\log \frac{p_{\text{data}}(x^+ | c)}{p_{\text{data}}(x^+ | c) + K q_{\text{noise}}(x^+)} - \sum_{j=1}^K \log \frac{K q_{\text{noise}}(x_j^-)}{p_{\text{data}}(x_j^- | c) + K q_{\text{noise}}(x_j^-)} \right].$$

Show that this objective can be seen as optimizing the same logits as the InfoNCE but with a binary cross-entropy loss.

- (c) **Effect of K .** In the case $q_{\text{noise}}(x) = p_{\text{data}}(x)$, analyze the effect of the number of negative samples K on the InfoNCE loss.

Hint: In this case, the logarithm of the numerator equals the pointwise mutual information. Optimizing the InfoNCE loss corresponds to maximizing a lower bound on the mutual information.

Solution

- (a) **Connection to cross-entropy loss.** Define logits $s_j = \log p_{\text{data}}(x_j | c) - \log q_{\text{noise}}(x_j)$. Then

$$\mathcal{L}_{\text{InfoNCE}} = - \sum_{(X,c) \in D} \log \frac{\exp(s^+)}{\sum_{j=0}^K \exp(s_j)} = - \sum_{(X,c) \in D} \sum_{j=0}^K y_j \log \text{softmax}(s)_j,$$

where y is the one-hot label for the true sample x^+ (i.e., for s^+). Thus this is exactly the multi-class softmax cross-entropy between y and the model distribution $P(\cdot | X, c)$.

In practice for contrastive learning, one example is $s_j = \langle f(x_j), f(c) \rangle / \tau$ (with temperature τ , network f), giving

$$- \sum_{(X,c) \in D} \log \frac{\exp(\langle f(x^+), f(c) \rangle / \tau)}{\sum_{j=0}^K \exp(\langle f(x_j), f(c) \rangle / \tau)}.$$

- (b) **Relation to NCE.** In (local) NCE, each pair (x, c) is classified as *data* vs. *noise*. With K noises per positive and using the logits s_j defined in (a), we have:

$$\frac{p_{\text{data}}(x^+ | c)}{p_{\text{data}}(x^+ | c) + K q_{\text{noise}}(x^+)} = \sigma(s^+ - \log K), \quad (1)$$

$$\frac{K q_{\text{noise}}(x^-)}{p_{\text{data}}(x^- | c) + K q_{\text{noise}}(x^-)} = \sigma(-s^- + \log K) = 1 - \sigma(s^- - \log K), \quad (2)$$

with σ as the sigmoid function. Then, we have

$$\mathcal{L}_{\text{NCE}} = -\log \sigma(s^+ - \log K) - \sum_{j=1}^K \log (1 - \sigma(s_j^- - \log K)) \quad (3)$$

$$= -(y \log \sigma(z) + (1 - y) \log (1 - \sigma(z))), \quad (4)$$

with $z = s - \log K$, $y \in \{0, 1\}$ denote the binary label ($y = 1$ for positive, $y = 0$ for negative). Thus, NCE and InfoNCE optimize the same logits s_j ; they differ only in how the likelihood is factored (sum of binary log-losses vs. one multi-class log-loss).

(c) **Effect of K .** For the case where $q_{\text{noise}}(x) = p_{\text{data}}(x)$, the mutual information is

$$I(x^+; c) = \sum_{x^+, c} p(x^+, c) \log \frac{p(x^+ | c)}{p(x^+)}.$$

If the labels are fully reliable, the InfoNCE estimator yields the lower bound (see Oord et al., 2018, Appendix A for proof)

$$I(x^+; c) \geq \log(K+1) - \mathcal{L}_{\text{InfoNCE}}.$$

By plugging the loss in the right-hand side, we can see that as K increases, the lower bound becomes tighter, which can often lead to better performance for models trained with the InfoNCE loss. However, in practice a very large K may introduce too much noise in the negatives and increases compute/memory cost, so K should be balanced with its temperature factor and batch design.

Task 2. Masked Language Modeling as Pseudo-Likelihood and -Perplexity.

Consider a sequence $x = (x_1, \dots, x_T)$ from a data distribution \mathcal{D} . For masked language modeling (MLM), let us draw a random mask set $M \subseteq \{1, \dots, T\}$ by sampling each position independently with probability $q \in (0, 1)$. Let $x_{\setminus t}$ denote x with the token in position t hidden (or replaced) and other tokens visible. Let x_t be an actual token in the position t . The MLM training objective is

$$\mathcal{L}_{\text{MLM}}(\theta) = -\mathbb{E}_{x \sim \mathcal{D}} \mathbb{E}_M \left[\sum_{t \in M} \log p_{\theta}(x_t | x_{\setminus t}) \right]. \quad (5)$$

(a) **Connection to pseudo-likelihood.** Define the (negative) pseudo log-likelihood (NPLL):

$$\text{NPLL}(\theta) = -\mathbb{E}_{x \sim \mathcal{D}} \left[\sum_{t=1}^T \log p_{\theta}(x_t | x_{\setminus t}) \right]. \quad (6)$$

Show that under independent Bernoulli masking at rate q , i.e., $M \stackrel{iid}{\sim} \text{Bern}(q)$:

$$\mathbb{E}_M \left[\sum_{t \in M} \log p_{\theta}(x_t | x_{\setminus t}) \right] = q \sum_{t=1}^T \log p_{\theta}(x_t | x_{\setminus t}), \quad (7)$$

and hence $\mathcal{L}_{\text{MLM}}(\theta) = q \cdot \text{NPLL}(\theta)$.

Hint: Use indicators $\mathbf{1}_{\{t \in M\}}$ and linearity of expectation.

- (b) **Pseudo-perplexity and an unbiased estimator.** Define the pseudo-perplexity (PPPL) for a sequence x by

$$\text{PPPL}(x) = \exp\left(\frac{1}{T} \sum_{t=1}^T -\log p_{\theta}(x_t | x_{\setminus t})\right). \quad (8)$$

- (i) Show that $\log \text{PPPL}(x)$ equals the average token-wise NPLL.
 (ii) Propose a practical *unbiased* single-pass estimator of $S(x) = \sum_{t=1}^T -\log p_{\theta}(x_t | x_{\setminus t})$ by sampling one index $U \sim \text{Unif}\{1, \dots, T\}$ and evaluating only $-\log p_{\theta}(x_U | x_{\setminus U})$. Then, prove that it is indeed unbiased.
 (c) **Relation to autoregressive maximum likelihood.** An autoregressive (AR) model maximizes

$$\log p_{\theta}(x) = \sum_{t=1}^T \log p_{\theta}(x_t | x_{<t}). \quad (9)$$

- (i) Explain why AR likelihood can be evaluated exactly, whereas MLM/pseudo-likelihood generally cannot yield a normalized joint $p_{\theta}(x)$.
 (ii) **Bonus:** State a modeling assumption(s) under which minimizing $\text{NPLL}(\theta)$ is statistically consistent. That is, if θ_n denotes the estimator from n samples, then $\text{NPLL}(\theta_n) \rightarrow \text{NPLL}(\theta^*)$ and $\theta_n \rightarrow \theta^*$ as $n \rightarrow \infty$. Sketch the proof of consistency.
Hint: Consider applying results from M-estimation theory.
 (iii) Give one advantage and one limitation of MLM vs. AR for downstream tasks.

Bonus (BERT 80/10/10). In BERT, of the selected tokens, 80% are replaced by [MASK], 10% by a random token, and 10% are left unchanged. Argue how this reduces train-test mismatch and prevents over-reliance on [MASK]; predict qualitative effects of using 100% or 0% [MASK].

Solution

- (a) **Connection to pseudo-likelihood.** Let $I_t := \mathbf{1}_{\{t \in M\}}$ with $I_t \stackrel{iid}{\sim} \text{Bern}(q)$, independent of x . Then

$$\begin{aligned} \mathbb{E}_M \left[\sum_{t \in M} \log p_{\theta}(x_t | x_{\setminus t}) \right] &= \mathbb{E}_M \left[\sum_{t=1}^T I_t \log p_{\theta}(x_t | x_{\setminus t}) \right] = \\ &= \sum_{t=1}^T \mathbb{E}[I_t] \log p_{\theta}(x_t | x_{\setminus t}) = q \sum_{t=1}^T \log p_{\theta}(x_t | x_{\setminus t}). \end{aligned}$$

Taking expectation over $x \sim \mathcal{D}$ yields

$$\begin{aligned} \mathcal{L}_{\text{MLM}}(\theta) &= -\mathbb{E}_{x \sim \mathcal{D}} \mathbb{E}_M \left[\sum_{t \in M} \log p_{\theta}(x_t | x_{\setminus t}) \right] \\ &= -q \mathbb{E}_{x \sim \mathcal{D}} \left[\sum_{t=1}^T \log p_{\theta}(x_t | x_{\setminus t}) \right] = q \cdot \text{NPLL}(\theta). \end{aligned}$$

- (b) **Pseudo-perplexity and an unbiased estimator.**

(i) By definition,

$$\log \text{PPPL}(x) = \log \exp \left(\frac{1}{T} \sum_{t=1}^T -\log p_{\theta}(x_t | x_{\setminus t}) \right) = \frac{1}{T} \sum_{t=1}^T -\log p_{\theta}(x_t | x_{\setminus t}),$$

which is the average token-wise NPLL.

(ii) A single-pass unbiased estimator of $S(x) = \sum_{t=1}^T -\log p_{\theta}(x_t | x_{\setminus t})$ is

$$\widehat{S}(x) = T \cdot (-\log p_{\theta}(x_U | x_{\setminus U})), \quad \text{where } U \sim \text{Unif}\{1, \dots, T\}.$$

Intuition: We estimate the sum of T terms by sampling one uniformly at random and scaling by T .

Unbiased-ness follows by linearity of expectation:

$$\begin{aligned} \mathbb{E}_U [T \cdot (-\log p_{\theta}(x_U | x_{\setminus U}))] &= \sum_{t=1}^T \Pr(U = t) T \cdot (-\log p_{\theta}(x_t | x_{\setminus t})) = \\ &= \sum_{t=1}^T \frac{1}{T} T \cdot (-\log p_{\theta}(x_t | x_{\setminus t})) = S(x). \end{aligned}$$

(c) **Relation to autoregressive maximum likelihood.**

(i) For an autoregressive (AR) model,

$$\log p_{\theta}(x) = \sum_{t=1}^T \log p_{\theta}(x_t | x_{<t}) \Rightarrow p_{\theta}(x) = \prod_{t=1}^T p_{\theta}(x_t | x_{<t}),$$

where each factor is directly given by the model, so we can compute likelihood exactly.

For the MLM/pseudo-likelihood, if the conditionals came from a joint normalized $p_{\theta}(x)$, they would satisfy:

$$p_{\theta}(x_t | x_{\setminus t}) = \frac{p_{\theta}(x)}{\sum_{x'_t} p_{\theta}(x_1, \dots, x_{t-1}, x'_t, x_{t+1}, \dots, x_T)} \quad (*)$$

(i.e., marginalizing out position t from the joint). But there is no guarantee that such a joint exists. We model the conditional distributions directly without constraints to ensure compatibility.

Here's a counterexample showing incompatibility. Consider $T = 2$ with tokens u, v . Suppose the MLM model defines:

$$p_{\theta}(x_1 = u | x_2 = v) = 1 \quad (10)$$

$$p_{\theta}(x_2 = v | x_1 = u) = 0 \quad (11)$$

If these came from a joint $p_{\theta}(x_1, x_2)$, we would need:

$$p_{\theta}(u, v) = p_{\theta}(x_1 = u | x_2 = v) \cdot p_{\theta}(x_2 = v) = 1 \cdot p_{\theta}(v)$$

and also:

$$p_{\theta}(u, v) = p_{\theta}(x_2 = v | x_1 = u) \cdot p_{\theta}(x_1 = u) = 0 \cdot p_{\theta}(u) = 0$$

This gives $p_{\theta}(v) = 0$, but then the first conditional wouldn't be well-defined (division by zero in the marginal). Therefore, no joint distribution exists that yields these conditionals.

- (ii) For statistical consistency of minimizing $\text{NPLL}(\theta)$, we need different assumptions than for joint distributions.

Key point: We do NOT assume the conditionals come from a joint distribution (which would contradict part (i)).

Sufficient assumptions for consistency:

- The data is generated i.i.d. from some true distribution with conditionals $p^*(x_t | x_{\setminus t})$
- The model class is correctly specified: $\exists \theta^* \in \Theta$ such that $p_{\theta^*}(x_t | x_{\setminus t}) = p^*(x_t | x_{\setminus t})$ for all t
- Identifiability: if $p_{\theta}(x_t | x_{\setminus t}) = p_{\theta'}(x_t | x_{\setminus t})$ for all t and almost all x , then $\theta = \theta'$
- Standard regularity conditions (e.g., compact parameter space, continuous likelihood)

Sketch of consistency proof: Define the empirical estimator from n samples $x^{(1)}, \dots, x^{(n)}$:

$$\theta_n = \arg \max_{\theta \in \Theta} \frac{1}{n} \sum_{i=1}^n \sum_{t=1}^T \log p_{\theta}(x_t^{(i)} | x_{\setminus t}^{(i)})$$

The population objective is:

$$\mathcal{Q}(\theta) = \mathbb{E}_{x \sim p^*} \left[\sum_{t=1}^T \log p_{\theta}(x_t | x_{\setminus t}) \right]$$

For each position t , by the KL divergence inequality:

$$\mathbb{E}_{x \sim p^*} [\log p_{\theta}(x_t | x_{\setminus t})] \leq \mathbb{E}_{x \sim p^*} [\log p^*(x_t | x_{\setminus t})]$$

with equality if and only if $p_{\theta}(x_t | x_{\setminus t}) = p^*(x_t | x_{\setminus t})$ almost surely.

Therefore $\mathcal{Q}(\theta) \leq \mathcal{Q}(\theta^*)$ with equality only at $\theta = \theta^*$ (by identifiability).

By the law of large numbers, the empirical objective converges to the population objective uniformly over Θ . Since θ^* is the unique maximizer of $\mathcal{Q}(\theta)$, standard M-estimation theory gives us $\theta_n \rightarrow \theta^*$ in probability as $n \rightarrow \infty$.

Remark: The assumption of correct specification can be relaxed. Even if no θ^* exactly matches the true conditionals, consistency still holds for the θ^* that minimizes the KL divergence to the true conditionals, making pseudo-likelihood robust to model misspecification.

Note: Knowledge of M-estimation theory is not relevant for the exam.

- (iii) *One advantage of MLM:* It supports bidirectional context, making it natural for tasks that require information from both directions (e.g., fill-in-the-blank or fill-in-the-middle). Training with bidirectional context often yields strong representations for fill-in-the-blank and encoder-style downstream tasks. One limitation: to enable generation, an MLM typically needs an encoder-decoder architecture, whereas an autoregressive (AR) model requires only a decoder. Moreover, under a strict MLM objective, only the masked tokens contribute to the loss in each forward pass, so token usage is less efficient than in AR models.

Bonus (BERT 80/10/10). Among the tokens chosen for prediction, replacing 80% with [MASK], 10% with a random token, and leaving 10% unchanged reduces train–test mismatch and discourages over-reliance on the [MASK] because:

- The 10% unchanged positions force the model to sometimes predict when the true token is visible, mitigating the fact that [MASK] never appears at test time.
- The 10% random replacements inject noise so the model cannot treat [MASK] as the sole prediction signal.

Qualitatively, using 100% [MASK] increases train–test mismatch and risks overfitting to the special token; using 0% [MASK] makes many targets trivially copyable (identity shortcut), weakening the learning signal, harming generalization, and reducing model understanding of the language.

Task 3. Exploring Contrastive Learning with SimCLR.

See Task A in the Jupyter notebook.

Task 4. Exploring the Scaling Behaviour of LMs with a Series of Pythia Models.

See Task B in the Jupyter notebook.
