Questions

1. (4 points) Implementing 2D Convolution Using nn.Linear.

PyTorch's nn.Conv2d uses a weight tensor $W \in \mathbb{R}^{C_{\text{out}} \times C_{\text{in}} \times k \times k}$ and bias $b \in \mathbb{R}^{C_{\text{out}}}$. Given an input $x \in \mathbb{R}^{1 \times C_{\text{in}} \times H_{\text{in}} \times W_{\text{in}}}$, implement a *stride-1*, *no-padding* 2D convolution using only slicing, reshaping, and a single shared nn.Linear layer.

Define:

```
def conv2d_linear(x, W, b):
```

that reproduces the output of nn.Conv2d.

```
Solution:
import torch
import torch.nn as nn
def conv2d_linear(x, W, b):
    # x: (1, C_in, H_in, W_in)
    # W: (C_out, C_in, k, k)
    # b: (C_out,)
   C_{out}, C_{in}, k, _{=} = W.shape
    _, _, H_in, W_in = x.shape
   H_{out} = H_{in} - k + 1
   W_{out} = W_{in} - k + 1
    # ----- RUBRIC (4 marks) -----
    # (1 mark) Correct construction of Linear layer and reshaping:
    #
               linear.weight = W.reshape(C_out, -1)
    #
               linear.bias
                            = b
    # (1 mark) Correct extraction of each (C_in × k × k) patch:
    #
              patch = x[0, :, i:i+k, j:j+k]
    #
              Explanation: convolution uses all channels in patch.
    # (1 mark) Correct flattening into a 1D vector:
    #
              flat = patch.reshape(-1)
    #
              Explanation: Linear expects a 1D input of size C_in*k*k.
    # (1 mark) Correct loop structure and placement of output:
              out[0, :, i, j] = linear(flat)
              Explanation: identical to applying conv filter at (i,j).
    linear = nn.Linear(C_in * k * k, C_out, bias=True)
    linear.weight.data = W.reshape(C_out, -1).clone()
    linear.bias.data = b.clone()
    out = torch.zeros(1, C_out, H_out, W_out)
    for i in range(H_out):
       for j in range(W_out):
            patch = x[0, :, i:i+k, j:j+k]
                                               # (C_in, k, k)
            flat = patch.reshape(-1)
                                               # (C_in*k*k,)
            out[0, :, i, j] = linear(flat)
                                              # shared linear map
    return out
```

2. (4 points) Parameter counting for a modified LeNet-style CNN with two parallel paths.

The input is a grayscale image of shape $(1 \times 32 \times 32)$. The network is defined as follows:

- Conv1: 6 output channels, kernel 5×5 , stride 1, no padding.
- **Pool1:** 2×2 max-pooling with stride 2.
- The resulting feature map is then sent into two parallel branches:
 - Path A:
 - * ConvA1: 10 output channels, kernel 3×3 , stride 1, padding 1.
 - * ConvA2: 12 output channels, kernel 1×1 , stride 1.
 - Path B:
 - * ConvB1: 8 output channels, kernel 1×1 , stride 1.
 - * ConvB2: 16 output channels, kernel 5×5 , stride 1, padding 2.
- The outputs of Path A and Path B are concatenated along the channel dimension (i.e., if Path A produces (C_A, H, W) and Path B produces (C_B, H, W) , the concatenated output is $(C_A + C_B, H, W)$).
- Pool2: 2×2 max-pooling with stride 2 is applied to the concatenated output.
- FC1: Fully connected layer with 50 outputs. Its input is the flattened output of Pool2.
- FC2: Fully connected layer with 10 outputs.

Assume all convolution layers include a bias term.

(a) For each of the following stages, compute the output shape (channels, height, width):

Conv1, Pool1, ConvA1, ConvA2, ConvB1, ConvB2, Concat, Pool2.

- (b) Compute the number of trainable parameters in: Conv1, ConvA1, ConvA2, ConvB1, ConvB2, FC1, FC2.
- (c) Compute the total number of trainable parameters in the entire network.

Solution:

Useful formula:

$$\#\text{params}(\text{conv}) = \text{outC} \cdot (\text{inC} \cdot k_h \cdot k_w) + \text{outC}, \qquad \#\text{params}(\text{FC}) = D_{\text{out}}D_{\text{in}} + D_{\text{out}}.$$

(a) Shapes:

$$(1,32,32) \xrightarrow{\text{Conv1 } (5\times5, \text{ s}=1, \text{ p}=0)} (6,28,28) \xrightarrow{\text{Pool1 } (2\times2, \text{ s}=2)} (6,14,14),$$

$$(6,14,14) \xrightarrow{\text{ConvA1 } (3\times3, \text{ p}=1)} (10,14,14) \xrightarrow{\text{ConvA2 } (1\times1)} (12,14,14),$$

$$(6,14,14) \xrightarrow{\text{ConvB1 } (1\times1)} (8,14,14) \xrightarrow{\text{ConvB2 } (5\times5, \text{ p}=2)} (16,14,14),$$

$$\text{Concat: } (12+16,14,14) = (28,14,14) \xrightarrow{\text{Pool2 } (2\times2, \text{ s}=2)} (28,7,7).$$

(b) Parameter counts:

Conv1:
$$6(1 \cdot 5 \cdot 5) + 6 = 156$$
.
ConvA1: $10(6 \cdot 3 \cdot 3) + 10 = 550$.
ConvA2: $12(10 \cdot 1 \cdot 1) + 12 = 132$.
ConvB1: $8(6 \cdot 1 \cdot 1) + 8 = 56$.
ConvB2: $16(8 \cdot 5 \cdot 5) + 16 = 3216$.
FC1: $D_{in} = 28 \cdot 7 \cdot 7 = 1372$,
 $50 \cdot 1372 + 50 = 68650$.
FC2: $10 \cdot 50 + 10 = 510$.

(c) Total parameters:

$$156 + 550 + 132 + 56 + 3216 + 68650 + 510 = \boxed{73270}.$$

TA Rubric (4 marks):

- 1 Correct shapes for all listed stages (minor off-by-one: -0.5).
- 2 Correct parameter counts for all conv and FC layers (small arithmetic mistakes: -0.5).
- 1 Correct total parameter count with consistent working.

3. (2 points) Feature space dimension for a polynomial kernel.

Consider the polynomial kernel

$$K(\mathbf{x}, \mathbf{z}) = (\mathbf{x}^{\top} \mathbf{z} + 1)^2,$$

where $\mathbf{x}, \mathbf{z} \in \mathbb{R}^2$ with coordinates $\mathbf{x} = (x_1, x_2)$ and $\mathbf{z} = (z_1, z_2)$.

- (a) Expand $K(\mathbf{x}, \mathbf{z})$ explicitly as a polynomial in x_1, x_2, z_1, z_2 .
- (b) Find an explicit feature map $\phi: \mathbb{R}^2 \to \mathbb{R}^m$ such that

$$K(\mathbf{x}, \mathbf{z}) = \langle \phi(\mathbf{x}), \phi(\mathbf{z}) \rangle,$$

and determine m.

Solution:

(a)

$$K(\mathbf{x}, \mathbf{z}) = (\mathbf{x}^{\top} \mathbf{z} + 1)^2 = (x_1 z_1 + x_2 z_2 + 1)^2$$
$$= (x_1 z_1)^2 + (x_2 z_2)^2 + 2x_1 z_1 x_2 z_2 + 2x_1 z_1 + 2x_2 z_2 + 1.$$

(b) We want to write $K(\mathbf{x}, \mathbf{z}) = \langle \phi(\mathbf{x}), \phi(\mathbf{z}) \rangle$.

One valid choice (not unique) is:

$$\phi(\mathbf{x}) = \begin{bmatrix} x_1^2 \\ x_2^2 \\ \sqrt{2} x_1 x_2 \\ \sqrt{2} x_1 \\ \sqrt{2} x_2 \\ 1 \end{bmatrix} \in \mathbb{R}^6.$$

Then

$$\langle \phi(\mathbf{x}), \phi(\mathbf{z}) \rangle = x_1^2 z_1^2 + x_2^2 z_2^2 + 2x_1 x_2 z_1 z_2 + 2x_1 z_1 + 2x_2 z_2 + 1 = K(\mathbf{x}, \mathbf{z}).$$

So the feature space dimension is m = 6.

TA Rubric (2 marks):

- 1 Correct expansion in (a).
- 1 Valid explicit $\phi(\mathbf{x})$ with inner product matching K, and m=6.
- 4. (3 points) Hard-margin SVM: primal, dual, and reason for using the dual.

We are given a linearly separable binary classification dataset $\{(\mathbf{x}_i, y_i)\}_{i=1}^N$ with $\mathbf{x}_i \in \mathbb{R}^d$ and $y_i \in \{-1, +1\}$.

- (a) Write the **primal** optimization problem for the hard-margin SVM.
- (b) Write the corresponding **dual** optimization problem (directly give the final form).
- (c) Give one clear reason why the dual formulation is preferred in practice for nonlinear SVMs.

Solution:

(a) Primal (hard-margin SVM):

$$\min_{\mathbf{w},b} \quad \frac{1}{2} ||\mathbf{w}||^2$$
s.t. $y_i(\mathbf{w}^{\top} \mathbf{x}_i + b) \ge 1, \quad i = 1, \dots, N.$

(b) Dual:

$$\max_{\alpha} \sum_{i=1}^{N} \alpha_i - \frac{1}{2} \sum_{i=1}^{N} \sum_{j=1}^{N} \alpha_i \alpha_j y_i y_j (\mathbf{x}_i^{\top} \mathbf{x}_j)$$
s.t. $\alpha_i \ge 0$, $\sum_{i=1}^{N} \alpha_i y_i = 0$.

(c) The dual depends on the data only through inner products $\mathbf{x}_i^{\top} \mathbf{x}_j$, so we can replace them by a kernel $K(\mathbf{x}_i, \mathbf{x}_j)$ and implicitly work in a high- or infinite-dimensional feature space. This "kernel trick" is not possible in the primal.

TA Rubric (3 marks):

- 1 Correct primal formulation.
- 1 Correct dual formulation with objective and constraints.
- 1 Correct explanation of kernel trick as the reason for preferring the dual.
- 5. (2 points) Backpropagation through a 2×2 max-pooling layer.

Consider a single-channel feature map (activation) before pooling:

$$A = \begin{bmatrix} 1 & -2 & 3 & 0 \\ 4 & 5 & -1 & 2 \\ 0 & 7 & 6 & -3 \\ 8 & -4 & 1 & 9 \end{bmatrix}.$$

We apply a 2×2 max-pooling operation with stride 2 and no padding. This produces a 2×2 output:

$$P = \text{MaxPool}_{2 \times 2, s=2}(A).$$

- (a) Compute the pooled output P explicitly.
- (b) Suppose the gradient of the loss w.r.t. the pooled output is

$$\frac{\partial L}{\partial P} = G = \begin{bmatrix} 1 & 2 \\ 3 & 4 \end{bmatrix}.$$

Compute the gradient of the loss w.r.t. the input activations, $\frac{\partial L}{\partial A}$, as a 4×4 matrix. Briefly explain in one sentence how max-pooling routes gradient back to the input.

Solution:

(a) We have four pooling windows (stride 2):

$$W_{11}: A[1:2, 1:2] = \begin{bmatrix} 1 & -2 \\ 4 & 5 \end{bmatrix} \Rightarrow \max = 5,$$

$$W_{12}: A[1:2, 3:4] = \begin{bmatrix} 3 & 0 \\ -1 & 2 \end{bmatrix} \Rightarrow \max = 3,$$

$$W_{21}: A[3:4, 1:2] = \begin{bmatrix} 0 & 7 \\ 8 & -4 \end{bmatrix} \Rightarrow \max = 8,$$

$$W_{22}: A[3:4, 3:4] = \begin{bmatrix} 6 & -3 \\ 1 & 9 \end{bmatrix} \Rightarrow \max = 9.$$

Thus

$$P = \begin{bmatrix} 5 & 3 \\ 8 & 9 \end{bmatrix}.$$

(b) Max-pool gradient: each G_{ij} is sent back only to the position of the max in window W_{ij} ; all other positions get zero.

Max positions in A:

- W_{11} : value 5 at (2,2) in A.

- W_{12} : value 3 at (1,3) in A.

- W_{21} : value 8 at (4,1) in A.

- W_{22} : value 9 at (4,4) in A.

So

$$\frac{\partial L}{\partial A} = \begin{bmatrix} 0 & 0 & 2 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 3 & 0 & 0 & 4 \end{bmatrix},$$

where each nonzero entry is copied from the corresponding element of G.

Max-pooling routes the gradient only to the input element(s) that achieved the maximum in each pooling window; all other elements in that window receive zero gradient.

TA Rubric (2 marks):

- 1 Correct pooled output $P = \begin{bmatrix} 5 & 3 \\ 8 & 9 \end{bmatrix}$.
- 1 Correct locations of max elements and nonzero entries in $\frac{\partial L}{\partial A}$. Correct value assignments for the gradient (matching G) and brief explanation that only argmax positions receive gradient.
- 6. (1 point) Can cross-validation be used to choose the number of clusters K in K-Means? Briefly justify.

Solution: No: K-Means has no labels, so there is no validation error to compute. Also, its objective (within-cluster sum of squares) always decreases as K increases, making cross-validation meaningless for selecting K.

- 7. (2 points) Q-learning update.
 - (a) Write the **Bellman optimality Q-learning update equation** and briefly explain what each term represents (reward, discount, bootstrap target, learning rate, and current Q-value).
 - (b) You are given the Q-table:

$$Q = \begin{array}{c|cc} & L & R \\ \hline A & 1.0 & 0.5 \\ B & 0.2 & 0.4 \end{array}$$

The agent is in state A, takes action R, receives reward r=2, and transitions to state B. Use learning rate $\alpha=0.5$ and discount $\gamma=0.9$.

Compute the updated value of Q(A, R).

Solution:

(a)

$$Q(s,a) \leftarrow Q(s,a) + \alpha \Big(r + \gamma \max_{a'} Q(s',a') - Q(s,a) \Big).$$

- r: immediate reward
- γ : discount factor
- $\max_{a'} Q(s', a')$: bootstrap estimate of optimal future value
- Q(s, a): current estimate
- α : learning rate controlling how much new information overrides old

(b)

$$\max_{a'} Q(B,a') = 0.4,$$

$$Q(A,R) \leftarrow 0.5 + 0.5 \left(2 + 0.9 \cdot 0.4 - 0.5\right) = 0.5 + 0.93 = 1.43.$$

TA Rubric (2 marks):

- 1 Correct Bellman update equation and each term identified clearly.
- 1 Correct numerical update: Q(A, R) = 1.43.