

DEEP LEARNING ASSIGNMENT 1

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▼ TASK 1

- Purpose: set up NumPy and a small show() helper to inspect shapes/values during development.
- Why it matters: In neural nets, most bugs come from shape mismatch. Printing the shapes early avoids silent errors!!

Ref:

- [Flattening Arrays](#)
- [Datatype Conversion](#)

```
import numpy as np
np.random.seed(0)

def show(name, arr, max_items=5):
    arr = np.array(arr)
    print(f"{name}: shape={arr.shape}, first={arr.reshape(-1)[:max_items]}")

def to_col(x):
    """
    To convert feature vector x into col vector of shape (n,1), it accepts x
    """
    x = np.array(x, dtype=float)
    if x.ndim == 1:
        return x.reshape(-1, 1)
    if x.ndim == 2:
        if x.shape[1] == 1:      # (n,1)
            return x
        if x.shape[0] == 1:      # (1,n)
            return x.reshape(-1, 1)
    raise ValueError(f"to_col: ### EXPECTED (n,), (n,1), or (1,n). Got {x.sh

def to_1d_labels(y):
    """
    To convert labels into shape (m,) that has values 0/1, it accepts y as --
    """
    y = np.array(y)
    if y.ndim == 0:
        return y.reshape(1).astype(int)
    if y.ndim == 1:
        return y.astype(int)
```

```

if y.ndim == 2:
    return y.reshape(-1).astype(int)
raise ValueError(f"to_1d_labels: unexpected shape {y.shape}")

```

Activation Function

The sigmoid activation:

$$\text{sig}(z) = 1 / (1 + e^{-z})$$

Why it matters?

-> LR outputs a prob for binary classification, -> Sigmoid maps the linear score z to a number between 0 and 1, -> Quick tests like: $\text{sig}(0)=0.5$ help verify correctness.

Reference: [Usage of sigmoid](#)

```

# Squeezeesss any num into 0 to 1

# If z is large positive ---> sigmoid is near 1
# If z is large negative ---> sigmoid is near 0
# If z = 0 ---> sigmoid is 0

def sigmoid(z):
    return 1 / (1 + np.exp(-z))

print("sigmoid(0):", sigmoid(0))
print("sigmoid(5):", sigmoid(5))
print("sigmoid(-5):", sigmoid(-5))

```

```

sigmoid(0): 0.5
sigmoid(5): 0.9933071490757153
sigmoid(-5): 0.0066928509242848554

```

Forward pass

A) Single sample forward

- Input: one sample x with n features
- Computation: $z = w^T \cdot x + b$ and $y' = \text{sig}(z)$
- Output: scalar z and scalar y'

B) Multi-sample forward

- Input: dataset x with m samples and n features
- Computation: $z = w^T \cdot x + b$ and $y' = \text{sig}(z)$
- Output: vectors $z \rightarrow R^m$ and $y' \rightarrow R^m$

1. -> Both are exactly the same model, the transpose just makes the dot product dims right.

2. -> SGD trains on 1 random sample @ a time, so we need the single sample forward,
3. -> For evaluation and later tasks, it's convenient to compute predictions for all samples at once, so my ffn also support multi sample forward.

Ref:

- [Blog on Logistic Regression from Scratch](#)

```
def forward(x, w, b, debug=False):
    """
    I've written a little flexible frwd pass;
    => if x is 1 sample: (n,) or (n,1) or (1,n) ---> returns scalar y_hat a
    => if x is multiple samples: (m,n) -> returns (m,1) y_hat and (m,1) z
    """
    w = to_col(w) # juz to ensure (n,1)

    A = np.array(x, dtype=float)

    # Case 1: single sample
    if A.ndim == 1 or (A.ndim == 2 and (A.shape[0] == 1 or A.shape[1] == 1)):
        x = to_col(A) # (n,1)
        z_mul = np.dot(w.T, x) + b # (1,1)
        z = float(z_mul)
        y_prob = float(sigmoid(z))

        if debug:
            print("---- forward (single) ----")
            show("x", x)
            show("w", w)
            print("b:", b)
            print("z:", z)
            print("y_prob:", y_prob)
            print("-----")

    return z, y_prob

    # Case 2: multiple samples
    if A.ndim == 2:
        X = A # (m,n)
        if X.shape[1] != w.shape[0]:
            raise ValueError(f"OOPSS!!! forward: X has {X.shape[1]} features")

        z = np.dot(X, w) + b # (m,1)
        y_prob = sigmoid(z) # (m,1)

        if debug:
            print("---- forward (many) ----")
            show("X", X)
            show("w", w)
            print("b:", b)
            show("z", z)
            show("y_prob", y_prob)
```

```

        print("-----")
        return z, y_prob

    raise ValueError(f"forward: unexpected input shape {A.shape}")

```

▼ Loss Function

A) Single sample loss

- For one sample: $J = -[y \log(y') + (1 - y) \log(1 - y')]$

B) Multi-sample average loss

- For m samples: $J = -\frac{1}{m} \sum_{i=1}^m [y^i \log(y'^i) + (1 - y^i) \log(1 - y'^i)]$

Tiny epsilon (clipping):

- In floating-point arithmetic, y' can become numerically 0 / 1 for extreme values of z causing $\log(0)$ which is undefined.
- Clipping ensures $y' \rightarrow [\epsilon, 1-\epsilon]$ so the loss remains finite.
- risk getting inf/nan if weights get large or learning rate is high.
- This is just a numerical safeguard, it doesn't change the core algorithm.

Sanity checks:

- If prediction matches label well \rightarrow loss small
- If prediction is confidently wrong \rightarrow loss large

Ref:

- [Log loss for Binary Classification](#)
- [Clipping in log-loss](#)

```

def bin_ce_loss(y, y_prob, eps=1e-12, debug=False):
    """
    binary cross entropy loss:
    => if y_prob is scalar -> returns scalar loss
    => if y_prob is (m,1) -> returns avg loss over m
    """
    y = to_1d_labels(y)

    if np.isscalar(y_prob):
        yh = float(min(max(y_prob, eps), 1 - eps))
        yy = int(y[0])
        J = -(yy * np.log(yh) + (1 - yy) * np.log(1 - yh))
        if debug:
            print("loss fxn(single):", float(J))
        return float(J)

```

```

y_prob = np.array(y_prob, dtype=float).reshape(-1) # (m,)
y_prob = np.clip(y_prob, eps, 1 - eps)

if y_prob.shape[0] != y.shape[0]:
    raise ValueError(f"bin_ce_loss: y has {y.shape[0]} items but y_prob

m = y.shape[0]
J = -(1.0 / m) * np.sum(y * np.log(y_prob) + (1 - y) * np.log(1 - y_prob

if debug:
    print("loss (many):", float(J))

return float(J)

```

▼ Gradients for one sample (partial derivatives)

- Goal: Compute gradients for one training example (SGD).
- Key idea: For logistic regression + log loss + sigmoid, the error term simplifies nicely: For a single training example, it gives the partial derivatives:
 - $\text{der}(w) = (y' - y) \cdot x$, $\text{der}(b) = (y' - y)$
- Sanity checks:
 - If $y' > y$ for a negative example, gradient pushes weights down.
 - Gradients match weight dimensions.

Why?:

- SGD updates parameters using one sample's gradient at a time.
- learning signal: if the prediction is too high/low compared to y , the error term $(y' - y)$ tells us how to adjust w and b .

```

def gradients_one(x, y, y_prob, debug=False):
    x = to_col(x)          # (n,1)
    y = to_1d_labels(y)   # (1,)
    y = int(y[0])
    y_prob = float(y_prob)

    err = (y_prob - y)    # scalar
    dw = err * x          # (n,1)
    db = err                # scalar

    if debug:
        print("---- gradients_one ----")
        print("error (y_prob - y):", err)
        show("dw", dw)
        print("db:", float(db))
        print("-----")

```

```
return dw, db
```

▼ Training loop (SGD + stopping)

What the algorithm does?

- Initialize w and b (e.g., zeros)
- Repeat:
 - Pick a random index i
 - Take training pair ($x(i), y(i)$)
 - Forward pass: compute $y'(i)$
- Compute loss $J(i)$
- Compute gradients $\text{der}(w)$, $\text{der}(b)$
- Update:
 - $w \rightarrow w - \alpha * \text{der}(w)$, $b \rightarrow b - \alpha * \text{der}(b)$
- Accumulate running cost over N iterations:
 - $J_{\text{running}} := \text{Summation}(k=1 \text{ to } N) J^k$
- Every N steps, compare to previous running cost and stop when:
 - $|J_{\text{running}} - J_{\text{running-prev}}| < \text{threshold}$
- Stopping logic in my code:
 - Tracking “running loss” every N steps, Stop if improvement becomes tiny (threshold) Or stop at max_iterations
- Sanity checks:
 - Loss trend generally decreases over time
 - Accuracy should improve on simple data

Ref:

- [GD vs SGD](#)
- [GD from Scratch using numpy](#)
- [SGD](#)
- [SGD from Scratch](#)

```
def fit_logreg_sgd(X, y, alpha=0.1, max_iterations=20000, threshold=1e-4, N=
    X = np.array(X, dtype=float)
    if X.ndim != 2:
        raise ValueError("fit_logreg_sgd: X must be shape (m,n)")

    y = to_1d_labels(y) # (m,)
```

```
m, n = X.shape
if y.shape[0] != m:
    raise ValueError(f"fit_logreg_sgd: X has {m} rows but y has {y.shape[0]} rows")

if N is None:
    N = m

w = np.zeros((n, 1))
b = 0.0

stopping = False
J_running = 0.0
J_running_prev = 0.0
iteration = 0

rng = np.random.default_rng(0)
history_batches = []

while not stopping:
    i = rng.integers(0, m)
    x_i = X[i] # (n,)
    y_i = y[i] # scalar 0/1

    _, y_hat = forward(x_i, w, b) # single-sample forward
    J_current = bin_ce_loss(y_i, y_hat) # single-sample loss
    dw, db = gradients_one(x_i, y_i, y_hat)

    w = w - alpha * dw
    b = b - alpha * db

    iteration += 1
    J_running += J_current

    if iteration > max_iterations:
        stopping = True
        print("Stopped: reached max_iterations.")

    if (iteration % N) == 0:
        change = abs(J_running - J_running_prev)
        history_batches.append((iteration, J_running, change))

        if debug_every_N:
            print(f"After {iteration} iterations: J_running={J_running:.4f}, change={change:.4f}")

        if change < threshold:
            stopping = True
            print("Stopped: convergence reached (change < threshold).")

    J_running_prev = J_running
    J_running = 0.0

return w, b, history_batches
```

```
# helper fxn to produce probs from frwd pass for many samples.

def predict_prob(X, w, b):
    # uses many sample forward: z = Xw + b, y_hat = sigmoid(z)
    _, y_hat = forward(X, w, b)
    return y_hat # (m,1)

def predict_class(X, w, b, threshold=0.5):
    y_hat = predict_prob(X, w, b).reshape(-1)
    return (y_hat >= threshold).astype(int)

def accuracy(y_true, y_pred):
    y_true = to_1d_labels(y_true)
    y_pred = to_1d_labels(y_pred)
    return (y_true == y_pred).mean()
```

- Goal: Create a small synthetic dataset to confirm training works end-to-end.

What to verify:

- Shapes print correctly
- Training reduces loss and improves accuracy

```
m = 200
X = np.random.randn(m, 2)
true_w = np.array([[2.0], [-3.0]])
true_b = 0.5

scores = np.dot(X, true_w) + true_b
y = (scores.reshape(-1) > 0).astype(int)

# Train
w, b, hist = fit_logreg_sgd(
    X, y,
    alpha=0.2,
    max_iterations=20000,
    threshold=1e-2,
    N=m,
    debug_every_N=True
)

# this fxn evaluates (many samples)
y_pred = predict_class(X, w, b)
acc = accuracy(y, y_pred)
print("Training accuracy (sanity check):", acc)

# also testing 1 sample frwd
x0 = X[0] # shape (n,)
z0, yhat0 = forward(x0, w, b, debug=True)
pred_class = 1 if yhat0 >= 0.5 else 0
print("True:", y[0], "| Pred class:", pred_class, "| Pred prob:", yhat0)
```

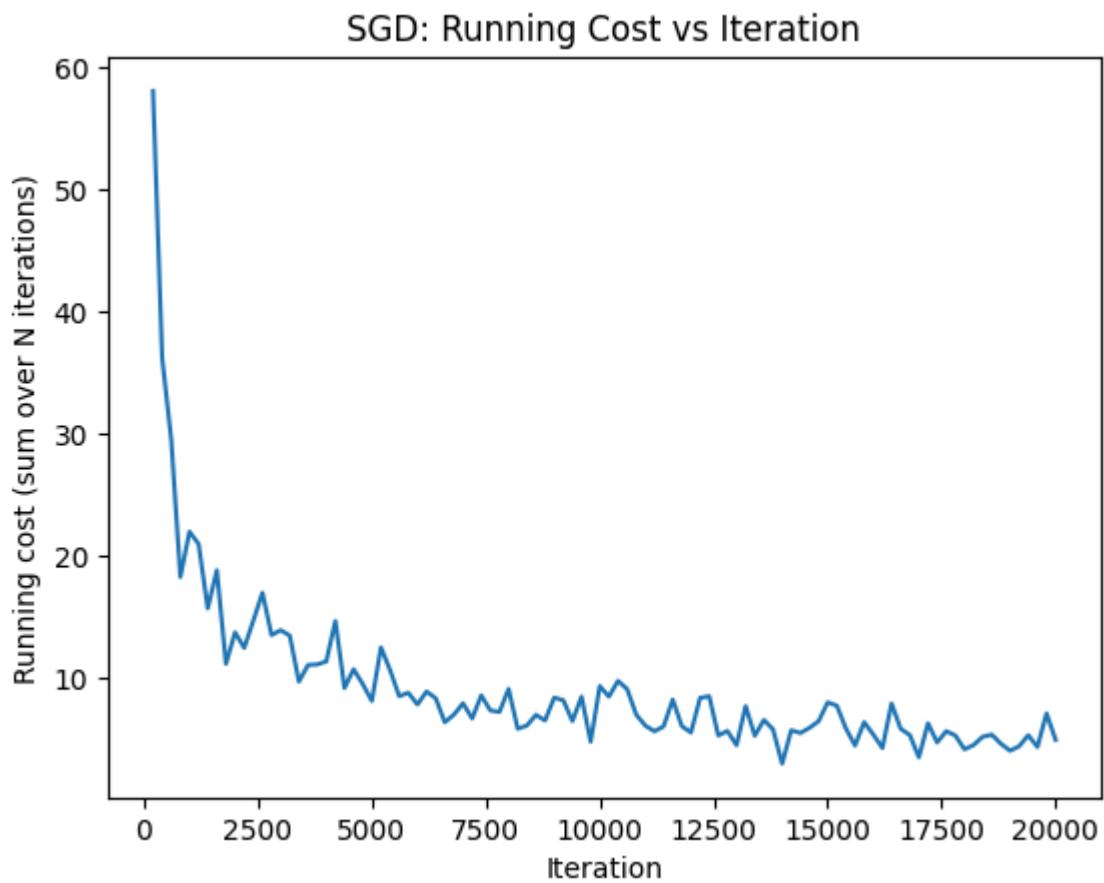
```
After 10800 iterations: J_running=6.9909, change=2.096626
After 11000 iterations: J_running=6.0907, change=0.900239
After 11200 iterations: J_running=5.6304, change=0.460332
After 11400 iterations: J_running=6.0319, change=0.401560
After 11600 iterations: J_running=8.2285, change=2.196546
After 11800 iterations: J_running=6.0540, change=2.174500
After 12000 iterations: J_running=5.5257, change=0.528311
After 12200 iterations: J_running=8.3640, change=2.838293
After 12400 iterations: J_running=8.5057, change=0.141684
After 12600 iterations: J_running=5.2870, change=3.218603
After 12800 iterations: J_running=5.6575, change=0.370497
After 13000 iterations: J_running=4.4921, change=1.165423
After 13200 iterations: J_running=7.6959, change=3.203772
After 13400 iterations: J_running=5.2691, change=2.426762
After 13600 iterations: J_running=6.5581, change=1.288993
After 13800 iterations: J_running=5.7966, change=0.761564
After 14000 iterations: J_running=2.9770, change=2.819550
After 14200 iterations: J_running=5.7166, change=2.739557
After 14400 iterations: J_running=5.4866, change=0.229971
After 14600 iterations: J_running=5.9142, change=0.427651
After 14800 iterations: J_running=6.4567, change=0.542443
After 15000 iterations: J_running=7.9962, change=1.539473
After 15200 iterations: J_running=7.7300, change=0.266134
After 15400 iterations: J_running=5.8535, change=1.876502
After 15600 iterations: J_running=4.4662, change=1.387284
After 15800 iterations: J_running=6.3792, change=1.912973
After 16000 iterations: J_running=5.3599, change=1.019359
After 16200 iterations: J_running=4.2616, change=1.098265
After 16400 iterations: J_running=7.9090, change=3.647446
After 16600 iterations: J_running=5.8367, change=2.072349
After 16800 iterations: J_running=5.3265, change=0.510155
After 17000 iterations: J_running=3.5040, change=1.822559
After 17200 iterations: J_running=6.2868, change=2.782781
After 17400 iterations: J_running=4.7162, change=1.570521
After 17600 iterations: J_running=5.6373, change=0.921034
After 17800 iterations: J_running=5.2895, change=0.347740
After 18000 iterations: J_running=4.1516, change=1.137956
After 18200 iterations: J_running=4.5038, change=0.352200
After 18400 iterations: J_running=5.1852, change=0.681425
After 18600 iterations: J_running=5.3525, change=0.167265
After 18800 iterations: J_running=4.6092, change=0.743226
After 19000 iterations: J_running=4.0375, change=0.571725
After 19200 iterations: J_running=4.4052, change=0.367665
After 19400 iterations: J_running=5.3119, change=0.906742
After 19600 iterations: J_running=4.3482, change=0.963674
After 19800 iterations: J_running=7.0927, change=2.744412
After 20000 iterations: J_running=4.9223, change=2.170358
Stopped: reached max_iterations.
Training accuracy (sanity check): 0.99
---- forward (single) ----
x: shape=(2, 1), first=[1.76405235 0.40015721]
w: shape=(2, 1), first=[ 11.16502405 -17.85126649]
b: 2.25920701378886
z: 14.811580914645154
y_prob: 0.9999996306722162
-----
True: 1 | Pred class: 1 | Pred prob: 0.9999996306722162
```

```
"""
Visual verification says:
- If loss decreases and then flattens -> convergence
- If accuracy increases -> model is learning
"""
```

```
import matplotlib.pyplot as plt

iters = [h[0] for h in hist]
Jrun = [h[1] for h in hist]

plt.figure()
plt.plot(iters, Jrun)
plt.xlabel("Iteration")
plt.ylabel("Running cost (sum over N iterations)")
plt.title("SGD: Running Cost vs Iteration")
plt.show()
```

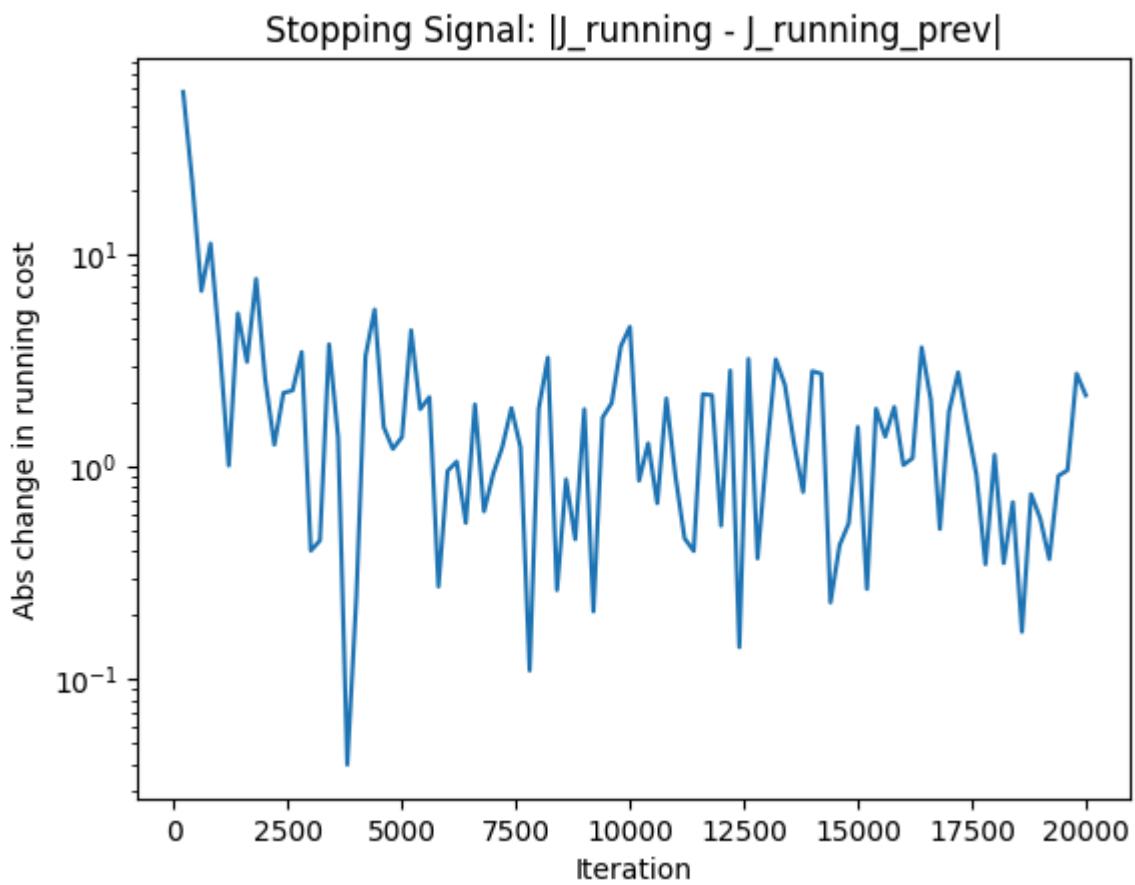


- At the start, the cost drops really fast. That's a good sign, it means the model is learning the main pattern in the data quickly.
- After a while, the line starts going down more slowly and kind of levels off. That usually means the model is almost done learning and further updates are only making small improvements.
- You can also see the line is a bit bumpy instead of perfectly smooth. That's normal for SGD, because I update the weights using one random training example at a time, so the cost can jump a little up and down.

- Overall, since the graph keeps trending downward, training is working properly and the model is improving over time.

```
change = [h[2] for h in hist]

plt.figure()
plt.plot(iters, change)
plt.xlabel("Iteration")
plt.ylabel("Abs change in running cost")
plt.title("Stopping Signal: |J_running - J_running_prev|")
plt.yscale("log") # optional, makes it easier to see shrinking values
plt.show()
```



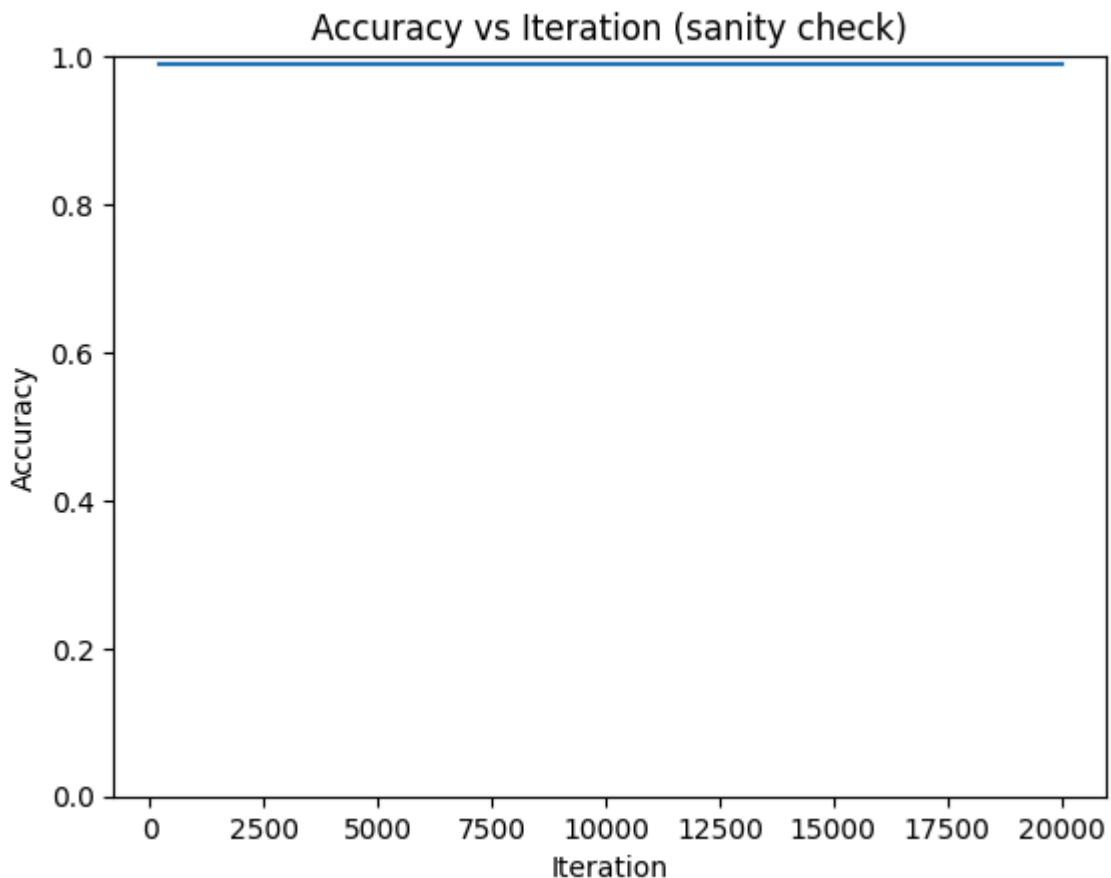
This graph is showing how much the running cost changes from one checkpoint to the next:

- At the beginning, the change is very large which means the model is still learning a lot and the cost is improving quickly.
- After some iterations, the values drop and stay much smaller overall means the model is no longer making big improvements -> it's starting to settle.
- The line still jumps up and down instead of smoothly shrinking to zero. That's normal because I'm using SGD, so each update is based on a random training sample & things can be noisy.

- Even though it's noisy, the important point is: compared to the start, the changes are generally smaller, which suggests the model is getting closer to convergence. If we set a small threshold, training would stop once these changes consistently fall below it.

```
accs = []
for _iter, _, _ in hist:
    y_pred = predict_class(X, w, b) # for sanity test you used same X
    accs.append(accuracy(y, y_pred))

plt.figure()
plt.plot(iters, accs)
plt.xlabel("Iteration")
plt.ylabel("Accuracy")
plt.title("Accuracy vs Iteration (sanity check)")
plt.ylim(0, 1)
plt.show()
```



▼ TASK 2

- Goal: Load blobs and circles datasets and visually inspect class separation.
- Inference:
 - blobs looks (almost) linearly separable -> logistic regression should do well

- circles is non-linear -> logistic regression will struggle

```
# Package imports
import matplotlib
import matplotlib.pyplot as plt
import sklearn
import sklearn.datasets
import pandas as pd
import numpy as np

%matplotlib inline
```

▼ DATASET 1

```
# Use pandas to read the CSV file as a dataframe
df1 = pd.read_csv("blobs600.csv")

# The y values are those labelled 'Class': extract their values
y1 = df1['Class'].values

# The x values are all other columns
del df1['Class'] # drop the 'Class' column from the dataframe
X1 = df1.values # convert the remaining columns to a numpy array

# Check its dimensions

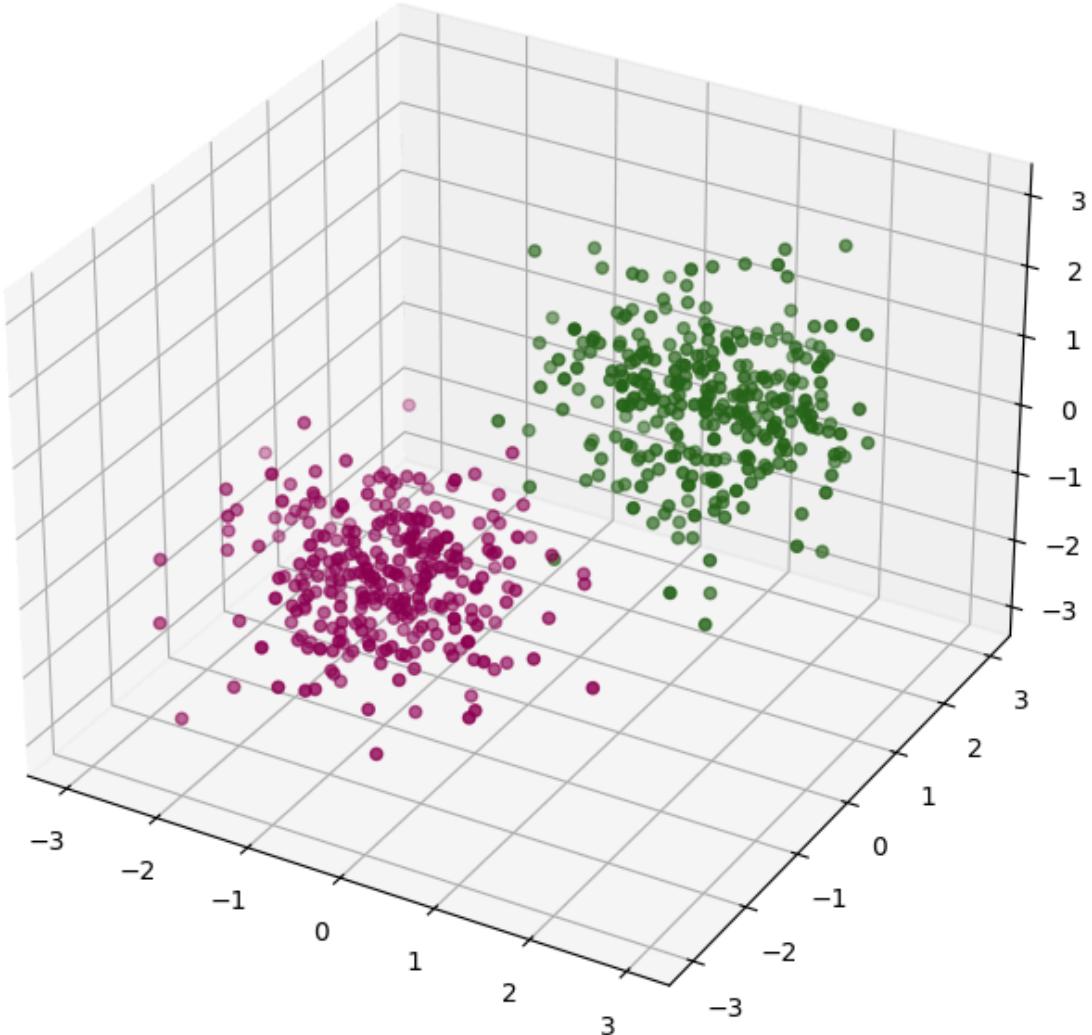
print(f"The dimensions of the dataset are: {np.shape(X1)}")
```

The dimensions of the dataset are: (600, 3)

```
# Plot the dataset in 3D, with colours according to the class label

fig = plt.figure(figsize=(8, 8)) # set the size to 8x8
ax = fig.add_subplot(111, projection="3d")
ax.scatter(X1[:,0], X1[:,1], X1[:,2], c=y1, cmap="PiYG") # changed the colou

plt.show()
plt.close(fig)
```



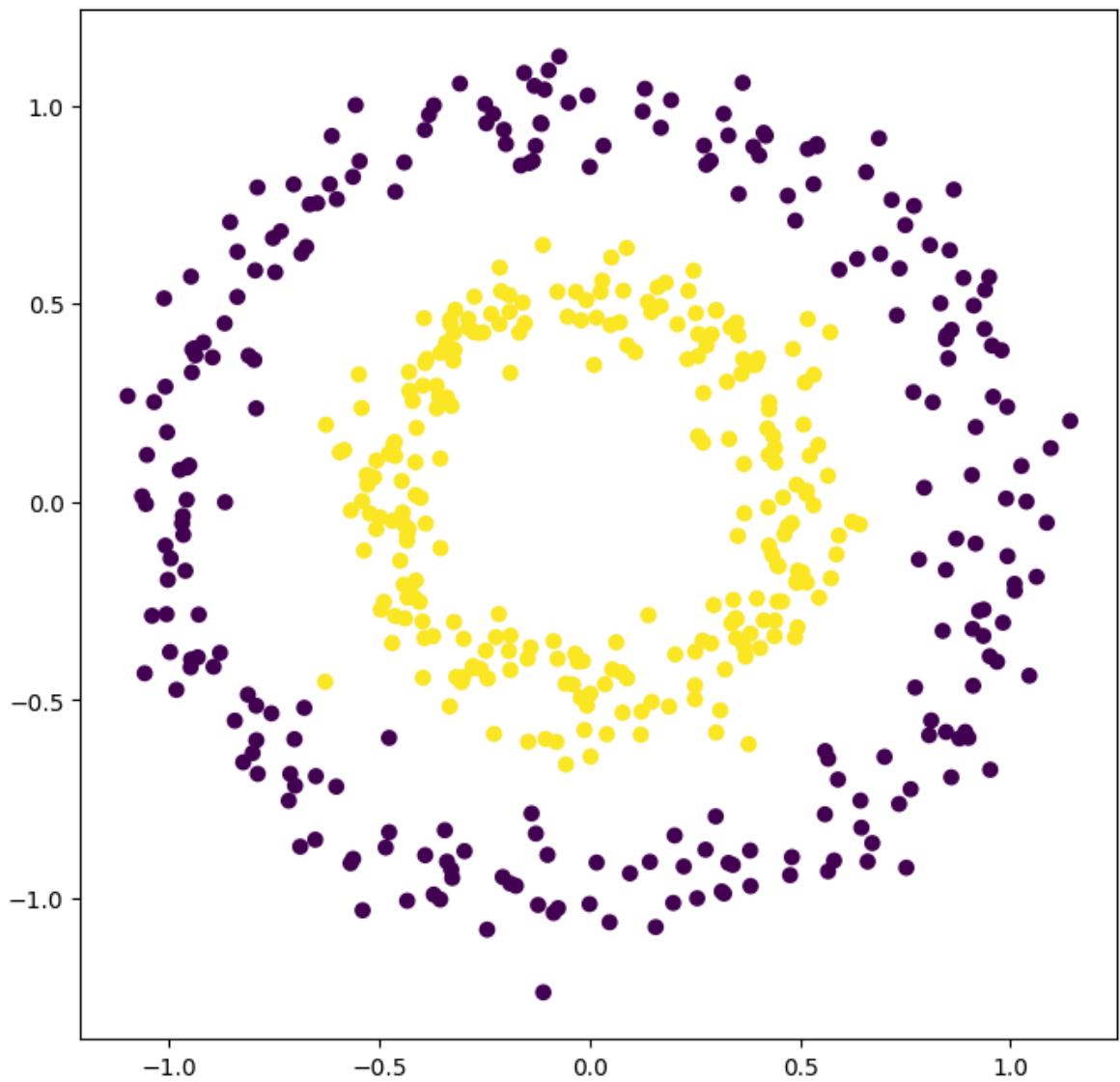
▼ DATASET 2

```
# Use pandas to read the CSV file as a dataframe  
df2 = pd.read_csv("circles500.csv")  
  
# The y values are those labelled 'Class': extract their values  
y2 = df2['Class'].values  
  
# The x values are all other columns  
del df2['Class'] # drop the 'Class' column from the dataframe  
X2 = df2.values # convert the remaining columns to a numpy array
```

```
print(f"The dimensions of Dataset 2 are: {np.shape(X2)}")
```

```
The dimensions of Dataset 2 are: (500, 2)
```

```
# plot X[0] vs X[1] and colour points according to the class, y  
  
fig, ax = plt.subplots(figsize=(8, 8))  
  
ax.scatter(X2[:,0], X2[:,1], c=y2)  
  
plt.show()  
plt.close(fig)
```



```
# Some examples of working with the data, to look at rows/columns  
print ("len(X2):", len(X2))          # outer array: one per sample  
print ("len(X2[0]):", len(X2[0]))    # each inner array is the attributes  
print ("len(X2[:,0]):", len(X2[:,0])) # select column 0 from array  
  
# np.shape returns all dimensions of the array  
(nsamples, nattributes) = np.shape(X2)  
print ("X2: nsamples =", nsamples, ", nattributes =", nattributes)
```

```
# Now example the y vector (1D array)
print ("len(y2)", len(y2))
print ("np.shape(y2):", np.shape(y2))

# You can transpose the y data using 'reshape'
yt = np.reshape(y2, (len(y2),1))
print ("np.shape(yt):", np.shape(yt))
(nsamples, nattribs) = np.shape(yt)
print ("y2 transpose: nsamples =", nsamples, ", nattribs =", nattribs)
```

```
len(X2): 500
len(X2[0]): 2
len(X2[:,0]): 500
X2: nsamples = 500 , nattribs = 2
len(y2) 500
np.shape(y2): (500,)
np.shape(yt): (500, 1)
y2 transpose: nsamples = 500 , nattribs = 1
```

```
print("blobs shape:", df1.shape)
print("circles shape:", df2.shape)

print("blobs columns:", list(df1.columns))
print("circles columns:", list(df2.columns))
```

```
blobs shape: (600, 3)
circles shape: (500, 2)
blobs columns: ['X1', 'X2', 'X3']
circles columns: ['X0', 'X1']
```

▼ split 70/15/15

- Goal: Create train/validation/test splits:
 - Train (70%): fit parameters
 - Validation (15%): choose hyperparameters (learning rate)
 - Test (15%): final evaluation
- Why this matters: prevents cheating by tuning on test set.

Ref : [Splitting using Manual method](#)

```
def split_70_15_15(X, y, seed=0):
    X = np.array(X, dtype=float)
    y = np.array(y).reshape(-1).astype(int)

    m = X.shape[0]
    rng = np.random.default_rng(seed)
    idx = np.arange(m)
    rng.shuffle(idx)

    X = X[idx]
    y = y[idx]
```

```

n_train = int(0.70 * m)
n_val   = int(0.15 * m)

X_train, y_train = X[:n_train], y[:n_train]
X_val, y_val     = X[n_train:n_train+n_val], y[n_train:n_train+n_val]
X_test, y_test   = X[n_train+n_val:], y[n_train+n_val:]

return X_train, y_train, X_val, y_val, X_test, y_test

```

Standardization

Goal:

- standardize features using train mean/std
- train logistic regressor on train set
- evaluate on val/test
- tune learning rate α using validation accuracy

Inference:

- Too small alpha \rightarrow slow learning
- Too big alpha \rightarrow unstable learning / noisy convergence

Ref [Implementing Standardization in Python](#)

```

def standardize_fit(X_train):
    mu = X_train.mean(axis=0, keepdims=True)
    sigma = X_train.std(axis=0, keepdims=True) + 1e-12
    return mu, sigma

def standardize_apply(X, mu, sigma):
    return (X - mu) / sigma

```

Goal:

- Train LR on both datasets
- Plot classification results
- Print prediction distributions
- Compute confusion matrix metrics (TP/FP/FN/TN)

Ref: [Train and Evaluate SGD](#)

```

def train_and_eval_lr(X_train, y_train, X_val, y_val, X_test, y_test,
                      alpha, max_iterations=20000, threshold=1e-4, seed=0, v
                      # Train (SGD)
                      w, b, hist = fit_logreg_sgd(
                          X_train, y_train,

```

```

        alpha=alpha,
        max_iterations=max_iterations,
        threshold=threshold,
        N=len(X_train),
        debug_every_N=verbose
    )

    # Evaluate
    y_tr_pred = predict_class(X_train, w, b)
    y_va_pred = predict_class(X_val, w, b)
    y_te_pred = predict_class(X_test, w, b)

    tr_acc = accuracy(y_train, y_tr_pred)
    va_acc = accuracy(y_val, y_va_pred)
    te_acc = accuracy(y_test, y_te_pred)

    return w, b, hist, tr_acc, va_acc, te_acc

```

- This function simply tries a list of learning rates (alphas) one by one. For each alpha, it trains the logistic regression model for many iterations, then checks how well it performs on the train, validation, and test sets.
- After testing all options, it picks the alpha that gives the highest validation accuracy, because validation performance is the best indicator of how well the model will generalize. In short: I experimented with multiple learning rates and long training runs, and selected the one that performed best on the validation set.

```

def tune_learning_rate(X_train, y_train, X_val, y_val, X_test, y_test, alphas):
    results = []

    for a in alphas:
        w, b, hist, tr_acc, va_acc, te_acc = train_and_eval_lr(
            X_train, y_train, X_val, y_val, X_test, y_test,
            alpha=a, max_iterations=300000, threshold=1e-4, verbose=False
        )
        results.append((a, tr_acc, va_acc, te_acc, w, b, hist))
        print(f"alpha={a:<8} train_acc={tr_acc:.3f} val_acc={va_acc:.3f} test_acc={te_acc:.3f}")

    # best alpha by validation accuracy
    best = max(results, key=lambda t: t[2])
    best_alpha, best_tr, best_va, best_te, best_w, best_b, best_hist = best

    print("\nBest alpha chosen by validation accuracy:", best_alpha)
    print(f"Best: train_acc={best_tr:.3f}, val_acc={best_va:.3f}, test_acc={best_te:.3f}")

    return best_alpha, best_w, best_b, best_hist, results

```

▼ BLOBS600

```
X1_train, y1_train, X1_val, y1_val, X1_test, y1_test = split_70_15_15(X1, y1

print("\nSplit sizes:")
print("Train:", X1_train.shape, y1_train.shape)
print("Val: ", X1_val.shape, y1_val.shape)
print("Test: ", X1_test.shape, y1_test.shape)

mu1, sigma1 = standardize_fit(X1_train)
X_train_s1 = standardize_apply(X1_train, mu1, sigma1)
X_val_s1   = standardize_apply(X1_val, mu1, sigma1)
X_test_s1  = standardize_apply(X1_test, mu1, sigma1)

alphas = [0.001, 0.01, 0.05, 0.1, 0.2]
best_alpha1, w1, b1, hist1, all_results1 = tune_learning_rate(X_train_s1, y1
```

```
Split sizes:
Train: (420, 3) (420,)
Val: (90, 3) (90,)
Test: (90, 3) (90,)
C:\Users\PS3\AppData\Local\Temp\ipykernel_14448\3127467275.py:15: Deprecation
    z = float(z_mul)
Stopped: reached max_iterations.
alpha=0.001      train_acc=0.995 val_acc=0.989 test_acc=1.000
Stopped: reached max_iterations.
alpha=0.01       train_acc=0.995 val_acc=0.989 test_acc=1.000
Stopped: reached max_iterations.
alpha=0.05       train_acc=0.995 val_acc=0.989 test_acc=1.000
Stopped: reached max_iterations.
alpha=0.1        train_acc=0.995 val_acc=0.989 test_acc=1.000
Stopped: reached max_iterations.
alpha=0.2        train_acc=0.998 val_acc=0.989 test_acc=1.000

Best alpha chosen by validation accuracy: 0.001
Best: train_acc=0.995, val_acc=0.989, test_acc=1.000
```

```
y_pred1 = predict_class(X_test_s1, w1, b1).reshape(-1)
correct = (y_pred1 == y1_test)

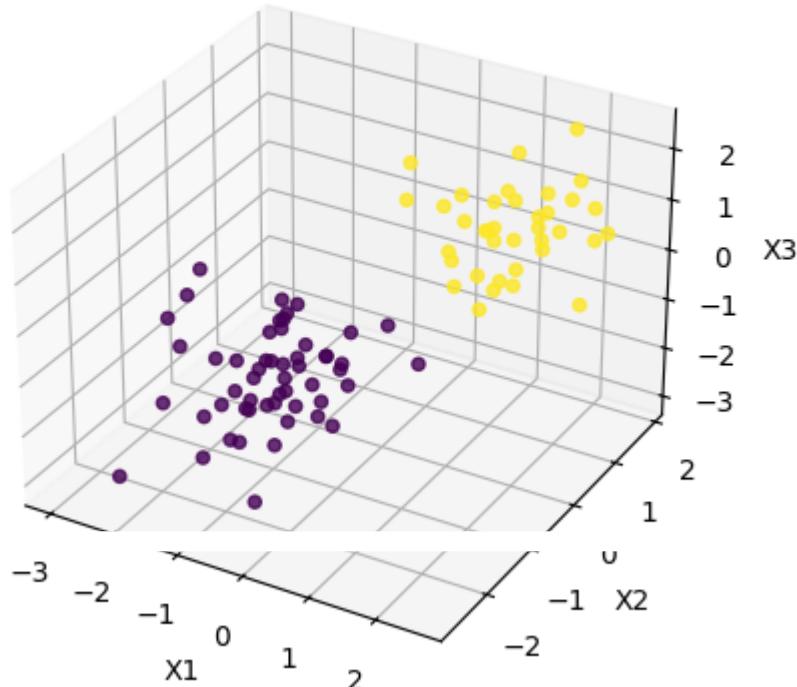
fig = plt.figure()
ax = fig.add_subplot(111, projection="3d")

ax.scatter(X1_test[:, 0], X1_test[:, 1], X1_test[:, 2], c=y_pred1, marker="o")
ax.scatter(X1_test[~correct, 0], X1_test[~correct, 1], X1_test[~correct, 2],
           facecolors="none", edgecolors="k", s=60)

ax.set_xlabel("X1")
ax.set_ylabel("X2")
ax.set_zlabel("X3")
ax.set_title(f"blobs600: Logistic Regression predictions (alpha={best_alpha1})
plt.show()
```

```
print("blobs600 test accuracy:", accuracy(y1_test, y_pred1))
```

blobs600: Logistic Regression predictions (alpha=0.001)



`blobs600 test accuracy: 1.0`

INFERENCE:

- The dataset was split into Train (420 samples), Validation (90), and Test (90) with 3 input features each. I tried multiple learning rates alpha from 0.001 to 0.2, and trained each model for a large number of iterations so every option had a fair chance to learn.
- All learning rates achieved almost the same validation accuracy (0.989) and perfect test accuracy (1.000). This suggests the dataset is very easy to separate, so logistic regression can learn it well regardless of the learning rate.
- Since the validation scores were tied, the code chooses the first best option by default, so it picked alpha = 0.001.

Conclusion: Logistic regression performs extremely well on this dataset, and learning rate choice is not very sensitive here because the classes are strongly separable.

▼ CIRCLES500

```
X2_train, y2_train, X2_val, y2_val, X2_test, y2_test = split_70_15_15(X2, y2)

print("\nSplit sizes:")
print("Train:", X2_train.shape, y2_train.shape)
print("Val:  ", X2_val.shape, y2_val.shape)
```

```
print("Test: ", X2_test.shape, y2_test.shape)

mu, sigma = standardize_fit(X2_train)
X2_train_s = standardize_apply(X2_train, mu, sigma)
X2_val_s   = standardize_apply(X2_val, mu, sigma)
X2_test_s  = standardize_apply(X2_test, mu, sigma)

alphas = [0.001, 0.01, 0.05, 0.1, 0.2]
# alphas = [0.05, 0.1, 0.15, 0.2, 0.25, 0.3]
best_alpha2, w2, b2, hist2, all_results2 = tune_learning_rate(X2_train_s, y2
```

```
Split sizes:
Train: (350, 2) (350,)
Val:   (75, 2) (75,)
Test:  (75, 2) (75,)

C:\Users\PS3\AppData\Local\Temp\ipykernel_14448\3127467275.py:15: Deprecation
    z = float(z_mul)
Stopped: reached max_iterations.
alpha=0.001    train_acc=0.423 val_acc=0.267 test_acc=0.427
Stopped: reached max_iterations.
alpha=0.01     train_acc=0.514 val_acc=0.360 test_acc=0.573
Stopped: reached max_iterations.
alpha=0.05     train_acc=0.369 val_acc=0.373 test_acc=0.387
Stopped: reached max_iterations.
alpha=0.1      train_acc=0.443 val_acc=0.520 test_acc=0.507
Stopped: reached max_iterations.
alpha=0.2      train_acc=0.460 val_acc=0.520 test_acc=0.520

Best alpha chosen by validation accuracy: 0.1
Best: train_acc=0.443, val_acc=0.520, test_acc=0.507
```

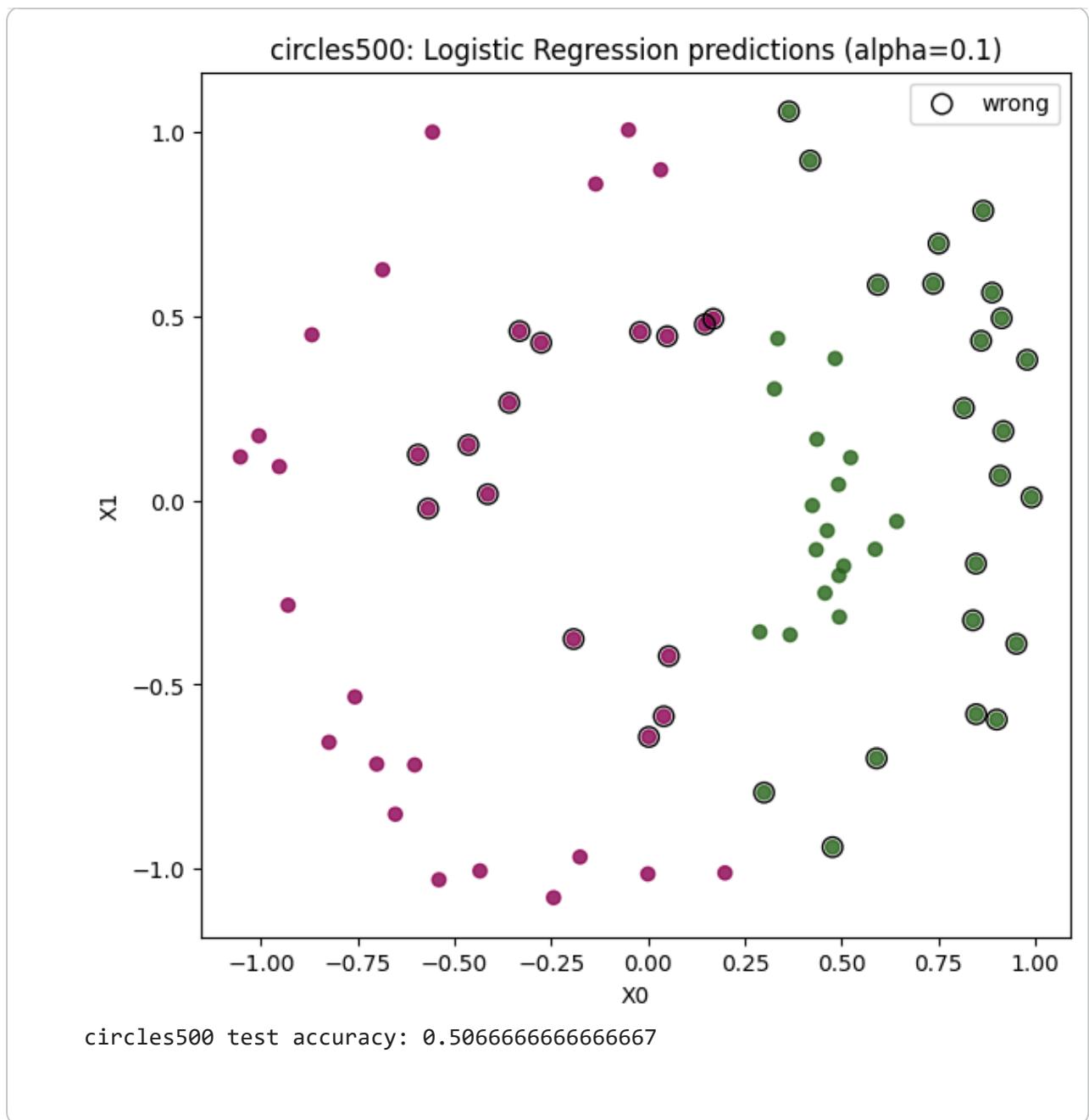
```
import matplotlib.pyplot as plt

y_pred2 = predict_class(X2_test_s, w2, b2).reshape(-1)
correct2 = (y_pred2 == y2_test)

plt.figure(figsize=(7,7))
plt.scatter(X2_test[:, 0], X2_test[:, 1], c=y_pred2, cmap="PiYG", alpha=0.8)
plt.scatter(X2_test[~correct2, 0], X2_test[~correct2, 1],
            facecolors="none", edgecolors="k", s=80, label="wrong")

plt.xlabel("X0")
plt.ylabel("X1")
plt.title(f"circles500: Logistic Regression predictions (alpha={best_alpha2}")
plt.legend()
plt.show()

print("circles500 test accuracy:", accuracy(y2_test, y_pred2))
```



- The dataset was split into Train (350), Validation (75), and Test (75) with 2 input features. I tried different learning rates alpha from 0.001 to 0.2, and trained each model for many iterations to see which one generalises best.
- The best validation accuracy I could get was only around 0.52, and test accuracy is about 0.51. This is close to random guessing (0.50), which tells us logistic regression is struggling on this dataset.
- The main reason is that circles500 is not linearly separable (points form inner vs outer circles), but logistic regression can only learn a straight-line decision boundary in the original feature space.
- Between the learning rates tested, alpha = 0.1 gave the highest validation score (tied with 0.2), so the model chose 0.1.

Conclusion: Logistic regression is not suitable for the circles dataset because the boundary is non-linear. This motivates using a neural network with a hidden layer (Task 3), which can learn non-linear decision boundaries.

```
print("Predicted classes:", np.unique(y_pred2, return_counts=True))
print("True classes:", np.unique(y2_test, return_counts=True))
```

```
Predicted classes: (array([0, 1]), array([36, 39]))
True classes: (array([0, 1]), array([43, 32]))
```

```
import matplotlib.pyplot as plt

iters = [h[0] for h in hist]
Jrun = [h[1] for h in hist]

plt.figure()
plt.plot(iters, Jrun)
plt.xlabel("Iteration")
plt.ylabel("Running cost (sum over N iterations)")
plt.title("Training progress: Running cost vs iteration")
plt.show()
```



At the beginning, the cost drops really fast, this means the model quickly learns the basic pattern in the data. After a while, the curve goes down more slowly and starts to level off, which usually means the model is close to its best solution and further updates only make small improvements. The curve is a bit bumpy because am usingg

SGD (updating using one random sample at a time), so small ups and downs are normal. Overall, the downward trend shows the training process is working properly.

```

tp = np.sum((y2_test == 1) & (y_pred2 == 1))
tn = np.sum((y2_test == 0) & (y_pred2 == 0))
fp = np.sum((y2_test == 0) & (y_pred2 == 1))
fn = np.sum((y2_test == 1) & (y_pred2 == 0))

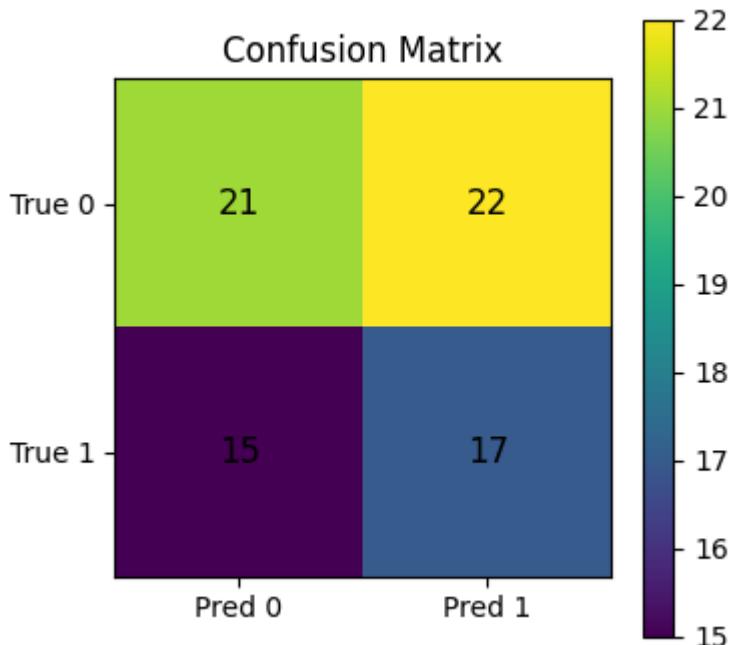
cm = np.array([[tn, fp],
               [fn, tp]])

plt.figure(figsize=(4,4))
plt.imshow(cm)
plt.xticks([0,1], ["Pred 0", "Pred 1"])
plt.yticks([0,1], ["True 0", "True 1"])

for i in range(2):
    for j in range(2):
        plt.text(j, i, cm[i, j], ha="center", va="center", fontsize=12)

plt.title("Confusion Matrix")
plt.colorbar()
plt.show()

```



- This confusion matrix shows exactly how many test samples the model got right and wrong for each class. The diagonal numbers (TN and TP) are the correct predictions, and the other off diagonal numbers (FP and FN) are the mistakes.
- In this (circles) test set, the model is making a lot of mistakes in both directions, which means it's struggling to separate the two classes.
 - True 0 predicted as 0 (TN) = 21 → 21 class-0 points were correctly classified.

- True 0 predicted as 1 (FP) = 22 → 22 class-0 points were wrongly labelled as class-1 (this is high).
- True 1 predicted as 0 (FN) = 15 → 15 class-1 points were wrongly labelled as class-0.
- True 1 predicted as 1 (TP) = 17 → 17 class-1 points were correctly classified.

So out of 75 test samples, correct predictions are $21 + 17 = 38$, giving accuracy:

$$\text{Accuracy} = 38/75 \rightarrow \sim 0.507$$

That's almost the same as guessing, which matches what we saw earlier for the circles dataset.

Conclusion: logistic regression isn't able to learn a good boundary here because the circles data needs a non-linear decision boundary, but logistic regression can only draw a straight line in the feature space. I hope this is exactly why task shifts to a neural network with a hidden layer in Task 3.

▼ TASK3

```
import numpy as np
import matplotlib.pyplot as plt

np.random.seed(0)

def to_col(x):
    x = np.array(x, dtype=float)
    if x.ndim == 1:
        return x.reshape(-1, 1)
    if x.ndim == 2:
        if x.shape[1] == 1:
            return x
        if x.shape[0] == 1:
            return x.reshape(-1, 1)
    raise ValueError(f"to_col: expected (n,), (n,1), or (1,n). Got {x.shape}")

def to_1d_labels(y):
    y = np.array(y)
    if y.ndim == 0:
        return y.reshape(1).astype(int)
    return y.reshape(-1).astype(int)

def accuracy(y_true, y_pred):
    y_true = to_1d_labels(y_true)
    y_pred = to_1d_labels(y_pred)
    return (y_true == y_pred).mean()

def sigmoid(z):
    return 1 / (1 + np.exp(-z))
```

```
def sigmoid_grad_from_a(a):
    # if a = sigmoid(z), then sigmoid'(z) = a(1-a)
    return a * (1 - a)
```

▼ shallow NN implementation

- Goal: Extend LR to 1 hidden layer:
 - Hidden layer: $z1 = w1 \cdot x + b1$, $a1 = \sigma(z1)$
 - Output: $z2 = w2 \cdot a1 + b2$, $y' = \sigma(z2)$
- Backprop idea (single sample SGD):
 - Output error: $\delta_2 = y' - y$
 - Hidden error: $\delta_1 = (w2(\text{transpose}) \cdot \delta_2) * \sigma'(z1)$
- Why this solves circles: hidden layer gives non-linear representation.

Ref :

- [1 HL Shallow NN Arch](#)
- [Backpropagation in NN](#)

```
def forward_nn(x_or_X, W1, b1, W2, b2, return_cache=False, debug=False):
    """
    1-hidden-layer NN forward pass (supports single sample / many samples)

    Parameters:
        W1: (h, n), b1: (h, 1)
        W2: (1, h), b2: (1, 1)

    Input:
        - single sample: (n,), (n,1), (1,n)
        - many samples: (m,n)

    Output:
        - single sample: z2 scalar, y_prob scalar
        - many samples: z2 (m,1), y_prob (m,1)
    """
    A = np.array(x_or_X, dtype=float)

    # ----- MANY SAMPLES (m,n) -----
    if A.ndim == 2 and A.shape[0] > 1:
        X = A # (m,n)
        m, n = X.shape

        if W1.shape[1] != n:
            raise ValueError(f"forward_nn: X has {n} features but W1 expects")

        # z1 = W1 X^T + b1 -> (h,m)
        z1 = np.dot(W1, X.T) + b1
```

```

a1 = sigmoid(z1) # (h,m)

# z2 = W2 a1 + b2 -> (1,m)
z2 = np.dot(W2, a1) + b2
y_prob = sigmoid(z2) # (1,m)

# return as (m,1)
z2_out = z2.T
y_out = y_prob.T

if debug:
    print("forward_nn MANY:")
    print("X:", X.shape, "z1:", z1.shape, "a1:", a1.shape, "z2:", z2)

if return_cache:
    return z2_out, y_out, (z1, a1, z2, y_prob)
return z2_out, y_out

# ----- SINGLE SAMPLE -----
x = to_col(A) # (n,1)
n = x.shape[0]

if W1.shape[1] != n:
    raise ValueError(f"forward_nn: x has {n} features but W1 expects {W1.shape[1]}")

z1 = np.dot(W1, x) + b1 # (h,1)
a1 = sigmoid(z1) # (h,1)
z2 = np.dot(W2, a1) + b2 # (1,1)
y_prob = sigmoid(z2) # (1,1)

z2_out = z2.item()
y_out = y_prob.item()

if debug:
    print("forward_nn SINGLE:")
    print("x:", x.shape, "z1:", z1.shape, "a1:", a1.shape, "z2:", z2_out)

if return_cache:
    return z2_out, y_out, (z1, a1, z2, y_prob, x)
return z2_out, y_out

```

```

def bce_loss(y, y_prob, eps=1e-12):
    """
    log loss:
    - y can be scalar or (m,)
    - y_prob can be scalar or (m,1) or (m,)
    returns scalar loss (avg if many)
    """
    y = to_1d_labels(y)

    # scalar prob
    if np.isscalar(y_prob):
        p = float(np.clip(y_prob, eps, 1 - eps))
        yy = int(y[0])

```

```

        return float(-(yy*np.log(p) + (1-yy)*np.log(1-p)))

# vector prob
p = np.array(y_prob, dtype=float).reshape(-1)
p = np.clip(p, eps, 1 - eps)
if p.shape[0] != y.shape[0]:
    raise ValueError("bce_loss: y and y_prob size mismatch.")
m = y.shape[0]
return float(-(1/m) * np.sum(y*np.log(p) + (1-y)*np.log(1-p)))

```

```

def backprop_one(cache, y, W2):
    """
    Backprop for ONE example using cached values from forward_nn(..., return
    cache for single sample contains: (z1, a1, z2, y_prob, x)
    """
    z1, a1, z2, y_prob, x = cache
    y = int(to_1d_labels(y)[0])

    # delta2 = y' - y
    delta2 = (y_prob - y) # (1,1)

    # dW2 = delta2 * a1(transpose)
    dW2 = np.dot(delta2, a1.T) # (1,h)
    db2 = delta2 # (1,1)

    # delta1 = (W2^T delta2) * sigmoid'(z1)
    delta1 = np.dot(W2.T, delta2) * sigmoid_grad_from_a(a1) # (h,1)

    # dW1 = delta1 * x(transpose)
    dW1 = np.dot(delta1, x.T) # (h,n)
    db1 = delta1 # (h,1)

    return dW1, db1, dW2, db2

```

```

def init_params(n_inputs, n_hidden, seed=0):
    rng = np.random.default_rng(seed)

    # small random weights
    W1 = rng.normal(0, 0.1, size=(n_hidden, n_inputs))
    b1 = np.zeros((n_hidden, 1))

    W2 = rng.normal(0, 0.1, size=(1, n_hidden))
    b2 = np.zeros((1, 1))

    return W1, b1, W2, b2

```

```

def fit_shallow_nn_sgd(X, y, n_hidden=8, alpha=0.1, max_iterations=300000,
                       threshold=1e-4, N=None, seed=0, debug_every_N=True):
    """
    training shallow NN (1 hidden layer, 1 output node) using single-example
    """

```

```
X = np.array(X, dtype=float)
y = to_1d_labels(y)
m, n = X.shape

if N is None:
    N = m

W1, b1, W2, b2 = init_params(n, n_hidden, seed=seed)
rng = np.random.default_rng(seed)

J_running = 0.0
J_prev = 0.0
history = []
iteration = 0

while True:
    # here it picks 1 random sample
    i = rng.integers(0, m)
    x_i = X[i]      # (n,)
    y_i = y[i]      # 0/1

    # forward (single sample) + cache
    _, y_prob, cache = forward_nn(x_i, W1, b1, W2, b2, return_cache=True)

    # loss (single)
    J_current = bce_loss(y_i, y_prob)

    # backprop grad
    dW1, db1g, dW2, db2g = backprop_one(cache, y_i, W2)

    # updation of w and b
    W1 = W1 - alpha * dW1
    b1 = b1 - alpha * db1g
    W2 = W2 - alpha * dW2
    b2 = b2 - alpha * db2g

    iteration += 1
    J_running += J_current

    # stop if max iterations
    if iteration >= max_iterations:
        if debug_every_N:
            print("Stopped: reached max_iterations.")
        break

    # checks for convergence every N steps
    if iteration % N == 0:
        change = abs(J_running - J_prev)
        history.append((iteration, J_running, change))

        if debug_every_N:
            print(f"After {iteration} iters: J_running={J_running:.4f},")

        if change < threshold:
            if debug_every_N:
```

```

        print("Stopped: convergence reached.")
        break

    J_prev = J_running
    J_running = 0.0

return W1, b1, W2, b2, history

```

```

def predict_proba_nn(X, W1, b1, W2, b2):
    _, y_prob = forward_nn(X, W1, b1, W2, b2) # many-sample path
    return y_prob.reshape(-1)

def predict_class_nn(X, W1, b1, W2, b2, threshold=0.5):
    probs = predict_proba_nn(X, W1, b1, W2, b2)
    return (probs >= threshold).astype(int)

```

```

def evaluate_hidden_sizes(X_train, y_train, X_val, y_val, X_test, y_test,
                         hidden_list, alpha=0.1, max_iterations=300000,
                         threshold=1e-4, seed=0):
    results = []
    for h in hidden_list:
        W1, b1, W2, b2, hist = fit_shallow_nn_sgd(
            X_train, y_train,
            n_hidden=h,
            alpha=alpha,
            max_iterations=max_iterations,
            threshold=threshold,
            N=len(X_train),
            seed=seed,
            debug_every_N=False
        )

        ytr = predict_class_nn(X_train, W1, b1, W2, b2)
        yva = predict_class_nn(X_val, W1, b1, W2, b2)
        yte = predict_class_nn(X_test, W1, b1, W2, b2)

        tr_acc = accuracy(y_train, ytr)
        va_acc = accuracy(y_val, yva)
        te_acc = accuracy(y_test, yte)

        results.append((h, tr_acc, va_acc, te_acc, W1, b1, W2, b2, hist))
        print(f"h={h:<3} train={tr_acc:.3f} val={va_acc:.3f} test={te_acc:.3f}")

    best = max(results, key=lambda r: r[2]) # choose best by val accuracy
    print("\nBest hidden size by validation:", best[0], "| val_acc:", best[2])
    return results, best

```

▼ Test different hidden sizes

- Goal: Experiment with $h \rightarrow \{2, 4, 8, 16\}$ & compare train/val/test.

- Inference:

- circles improves dramatically with more hidden units
- blobs already easy; hidden size doesn't matter much

```
hidden_list = [2, 4, 8, 16]

print("== circles500 (2D) shallow NN ==")
circles_results, circles_best = evaluate_hidden_sizes(
    X2_train_s, y2_train,
    X2_val_s, y2_val,
    X2_test_s, y2_test,
    hidden_list,
    alpha=0.1,
    max_iterations=50000,
    threshold=1e-4,
    seed=0
)

print("\n== blobs600 (3D) shallow NN ==")
blobs_results, blobs_best = evaluate_hidden_sizes(
    X_train_s1, y1_train,
    X_val_s1, y1_val,
    X_test_s1, y1_test,
    hidden_list,
    alpha=0.1,
    max_iterations=50000,
    threshold=1e-4,
    seed=0
)
```

```
== circles500 (2D) shallow NN ==
h=2  train=0.757 val=0.667 test=0.667
h=4  train=0.994 val=0.973 test=1.000
h=8  train=1.000 val=1.000 test=1.000
h=16 train=1.000 val=0.987 test=1.000
```

Best hidden size by validation: 8 | val_acc: 1.0

```
== blobs600 (3D) shallow NN ==
h=2  train=0.998 val=0.989 test=1.000
h=4  train=0.998 val=0.989 test=1.000
h=8  train=0.998 val=0.989 test=1.000
h=16 train=0.998 val=0.989 test=1.000
```

Best hidden size by validation: 2 | val_acc: 0.9888888888888889

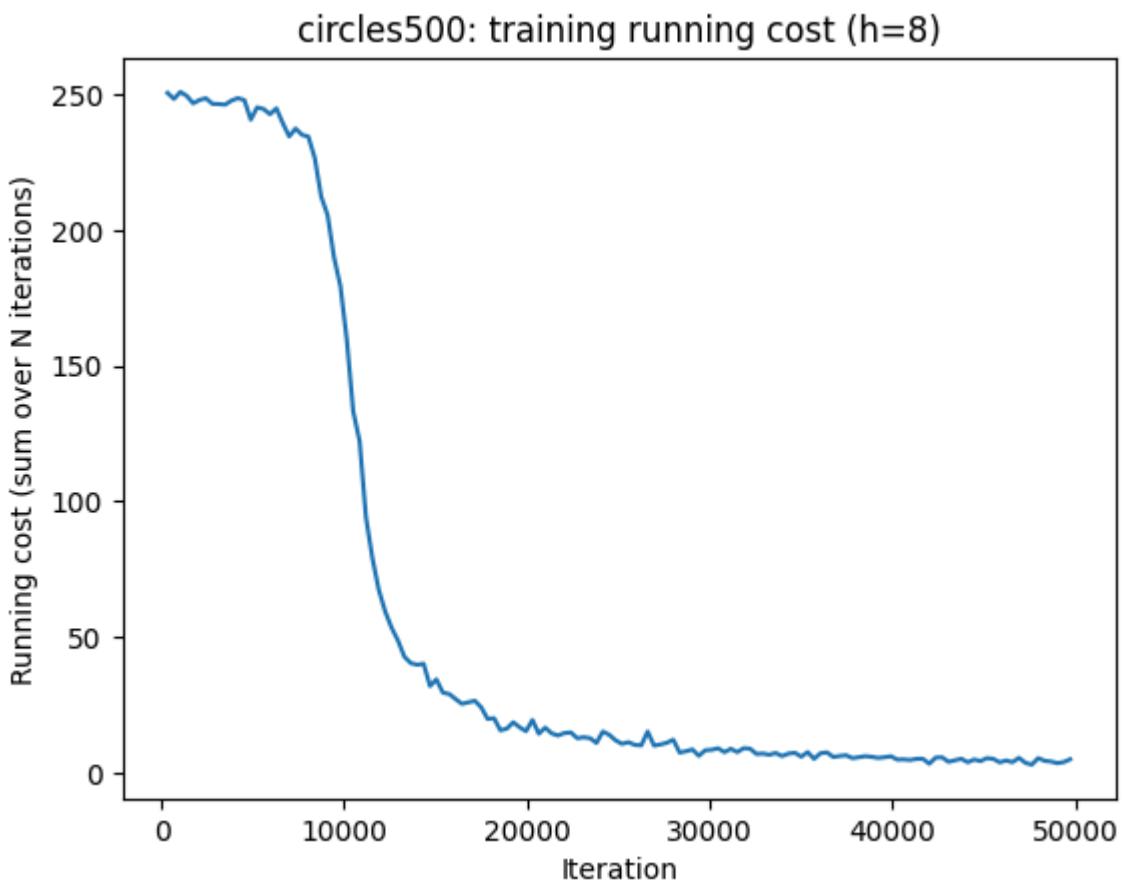
- Adding one hidden layer completely changes what the model can learn. On circles500, logistic regression struggled because it can only draw a straight line boundary, but the shallow NN learns a non-linear boundary, so accuracy jumps to ~100% when hidden nodes are enough (best at h=8).

- On blobs600, the data is already easy and almost linearly separable, so even a small hidden layer ($h=2$) achieves near-perfect performance and increasing hidden nodes doesn't really help.

```
best_h, tr_acc, va_acc, te_acc, W1, b1, W2, b2, hist = circles_best

iters = [t[0] for t in hist]
Jrun = [t[1] for t in hist]

plt.figure()
plt.plot(iters, Jrun)
plt.xlabel("Iteration")
plt.ylabel("Running cost (sum over N iterations)")
plt.title(f"circles500: training running cost (h={best_h})")
plt.show()
```



The cost stays high at the start, then suddenly drops sharply, which shows the network finally finds a good non-linear separation after enough updates. After that, it slowly flattens out near zero, meaning the model has basically learned the pattern and is only making tiny improvements.

```
import matplotlib.pyplot as plt

def plot_loss_history(hist, title_prefix=""):
    iters = [h[0] for h in hist]
```

```
Jrun    = [h[1] for h in hist]
change = [h[2] for h in hist]

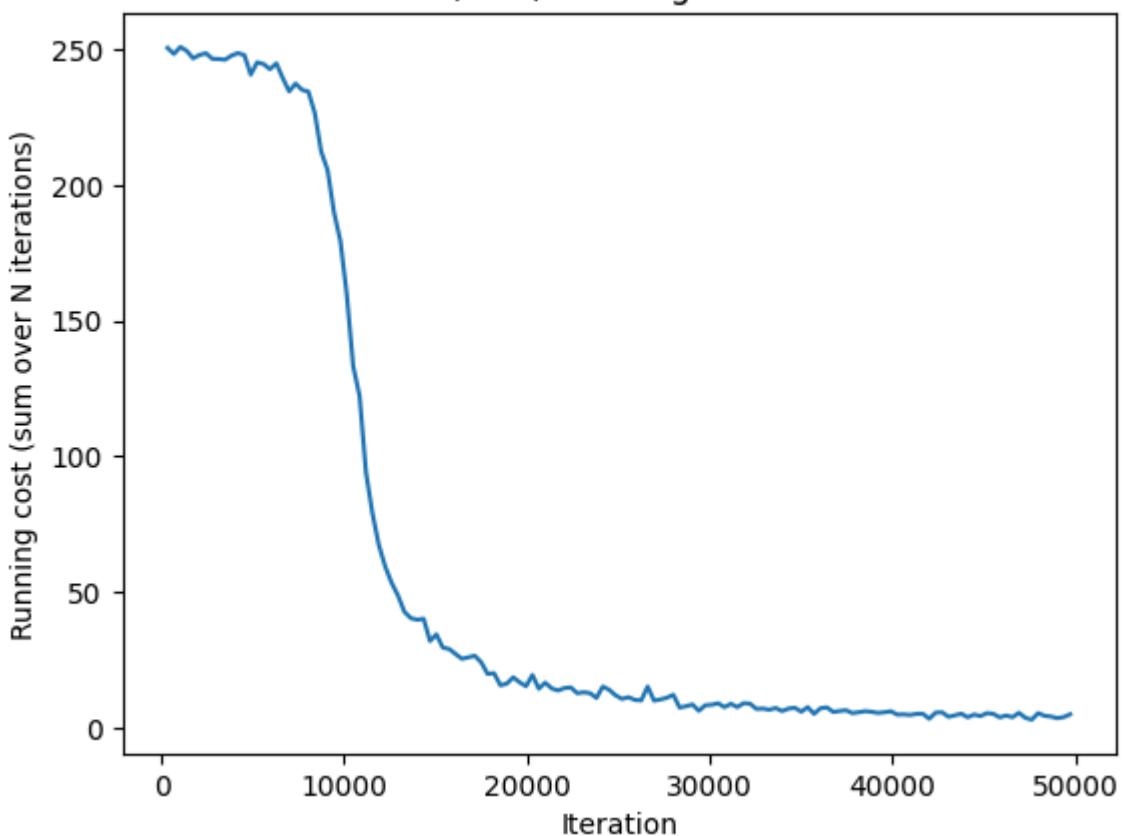
# Running cost plot
plt.figure()
plt.plot(iters, Jrun)
plt.xlabel("Iteration")
plt.ylabel("Running cost (sum over N iterations)")
plt.title(f"{title_prefix} Running cost vs iteration")
plt.show()

# Stopping signal plot
plt.figure()
plt.plot(iters, change)
plt.xlabel("Iteration")
plt.ylabel("|\u033_running - \u033_prev|")
plt.title(f"{title_prefix} Stopping signal vs iteration")
plt.yscale("log")
plt.show()

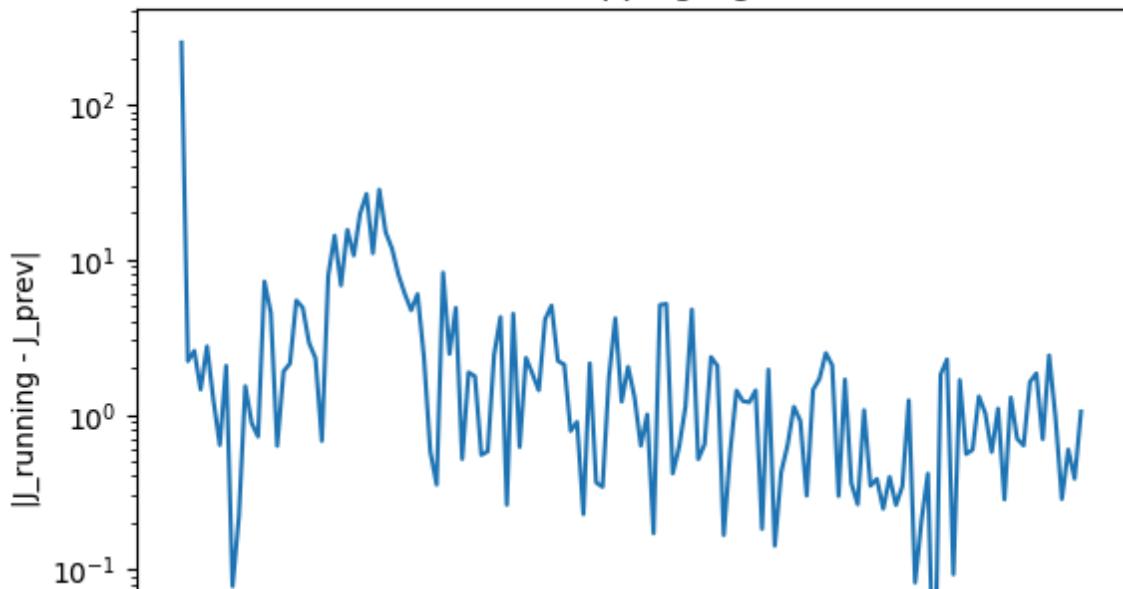
# --- circles ---
best_hc, tr_acc_c, va_acc_c, te_acc_c, W1c, b1c, W2c, b2c, hist_c = circles_
plot_loss_history(hist_c, title_prefix=f"circles500 (h={best_hc})")

# --- blobs ---
best_tb, tr_acc_b, va_acc_b, te_acc_b, W1b, b1b, W2b, b2b, hist_b = blobs_be
plot_loss_history(hist_b, title_prefix=f"blobs600 (h={best_tb})")
```


circles500 (h=8) Running cost vs iteration



circles500 (h=8) Stopping signal vs iteration



1. circles500: running cost vs iteration ($h=8$) (similar cost curve)

This shows the same story, the model improves a lot after a certain point and then settles. The small bumps are normal because the use of SGD, so each update is based on a random sample and the cost can wiggle a bit.

2. circles500: stopping signal vs iteration ($h=8$) (log-scale change plot)

At the beginning the change is huge (model learning fast), but later the changes become much smaller overall. The curve is noisy because SGD is random, but the general trend tells us training is stabilising and getting close to convergence.

3. blobs600: running cost vs iteration (h=2) (cost curve)

The cost drops extremely fast and becomes tiny early on, showing blobs are easy to separate. After that it stays low and flat, which means the model converged quickly.

4. blobs600: stopping signal vs iteration (h=2) (log-scale change plot)

The change starts very large then quickly becomes small, which means training reaches a **stable point** early. Any remaining spikes are just SGD noise and don't affect the fact that the model has already learned the dataset well.

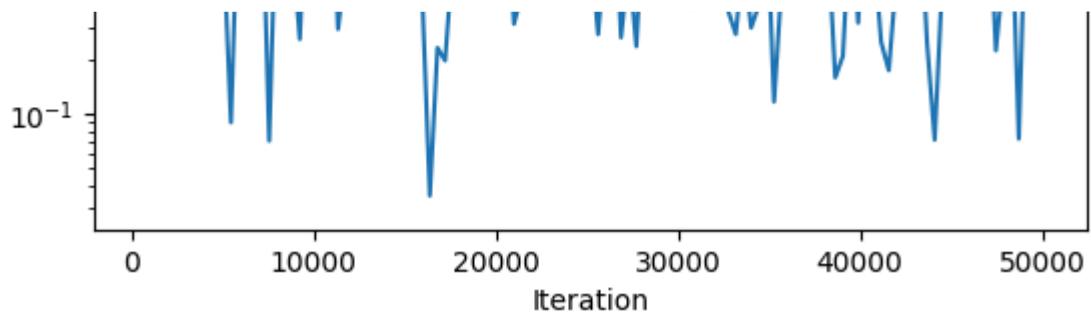
```
import numpy as np
import matplotlib.pyplot as plt

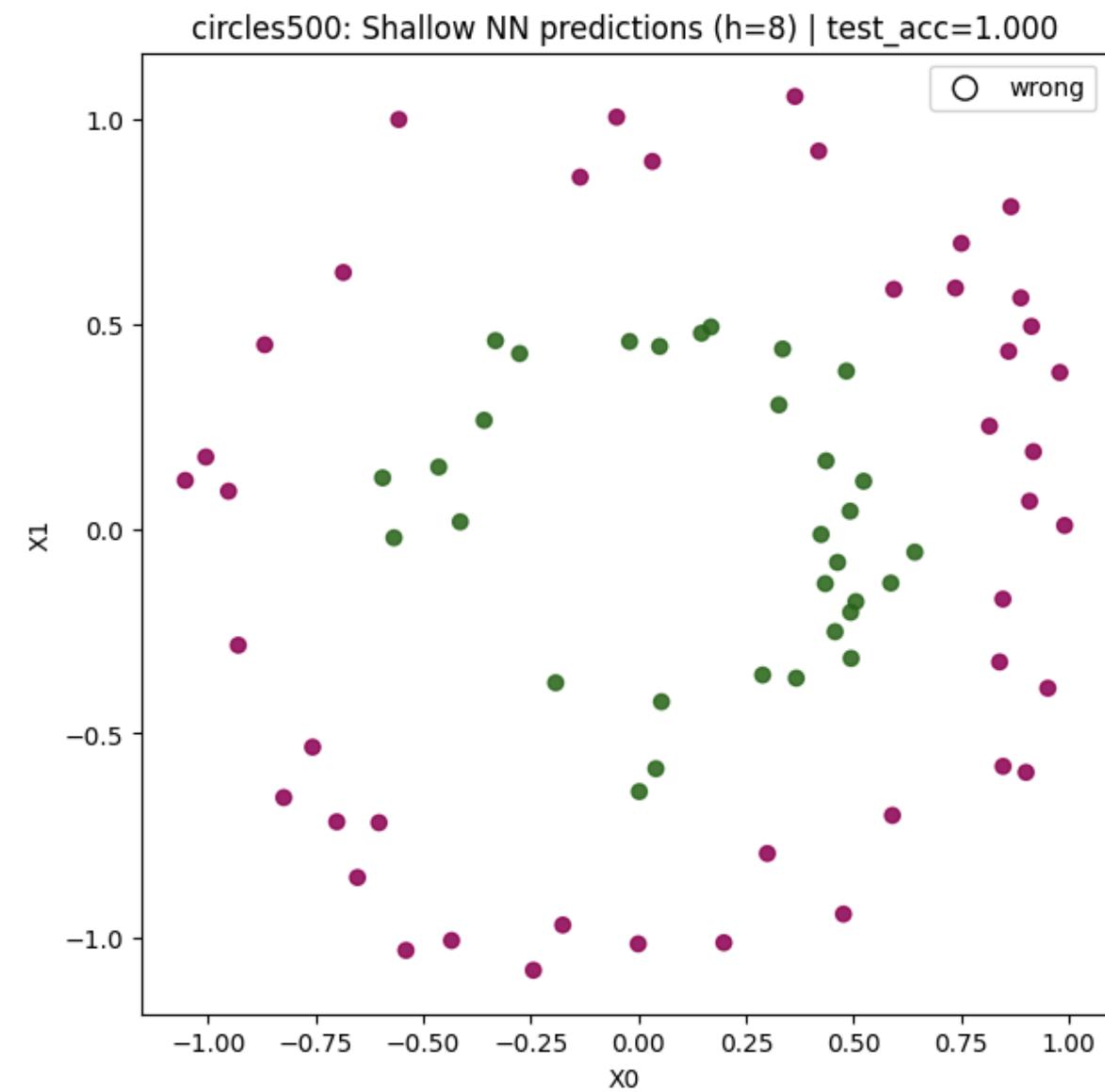
y_pred_c = predict_class_nn(X2_test_s, W1c, b1c, W2c, b2c).reshape(-1)
correct_c = (y_pred_c == y2_test)

plt.figure(figsize=(7,7))
plt.scatter(X2_test[:, 0], X2_test[:, 1], c=y_pred_c, cmap="PiYG", alpha=0.8
plt.scatter(X2_test[~correct_c, 0], X2_test[~correct_c, 1],
            facecolors="none", edgecolors="k", s=90, label="wrong")

plt.xlabel("X0")
plt.ylabel("X1")
plt.title(f"circles500: Shallow NN predictions (h={best_hc}) | test_acc={te_
plt.legend()
plt.show()

print("circles500 predicted class counts:", np.unique(y_pred_c, return_count
print("circles500 test accuracy:", accuracy(y2_test, y_pred_c))
```





circles500 scatter: Shallow NN predictions (h=8), test_acc=1.0

The predicted colors form a clear inner vs outer circle split, which is exactly what we expect for circles data. The wrong markers are basically absent, matching the 100% test accuracy.

```
y_pred_b = predict_class_nn(X_test_s1, W1b, b1b, W2b, b2b).reshape(-1)
correct_b = (y_pred_b == y1_test)

fig = plt.figure(figsize=(8,8))
ax = fig.add_subplot(111, projection="3d")

ax.scatter(X1_test[:, 0], X1_test[:, 1], X1_test[:, 2], c=y_pred_b, cmap="Pi
ax.scatter(X1_test[~correct_b, 0], X1_test[~correct_b, 1], X1_test[~correct_
    facecolors="none", edgecolors="k", s=70, label="wrong")

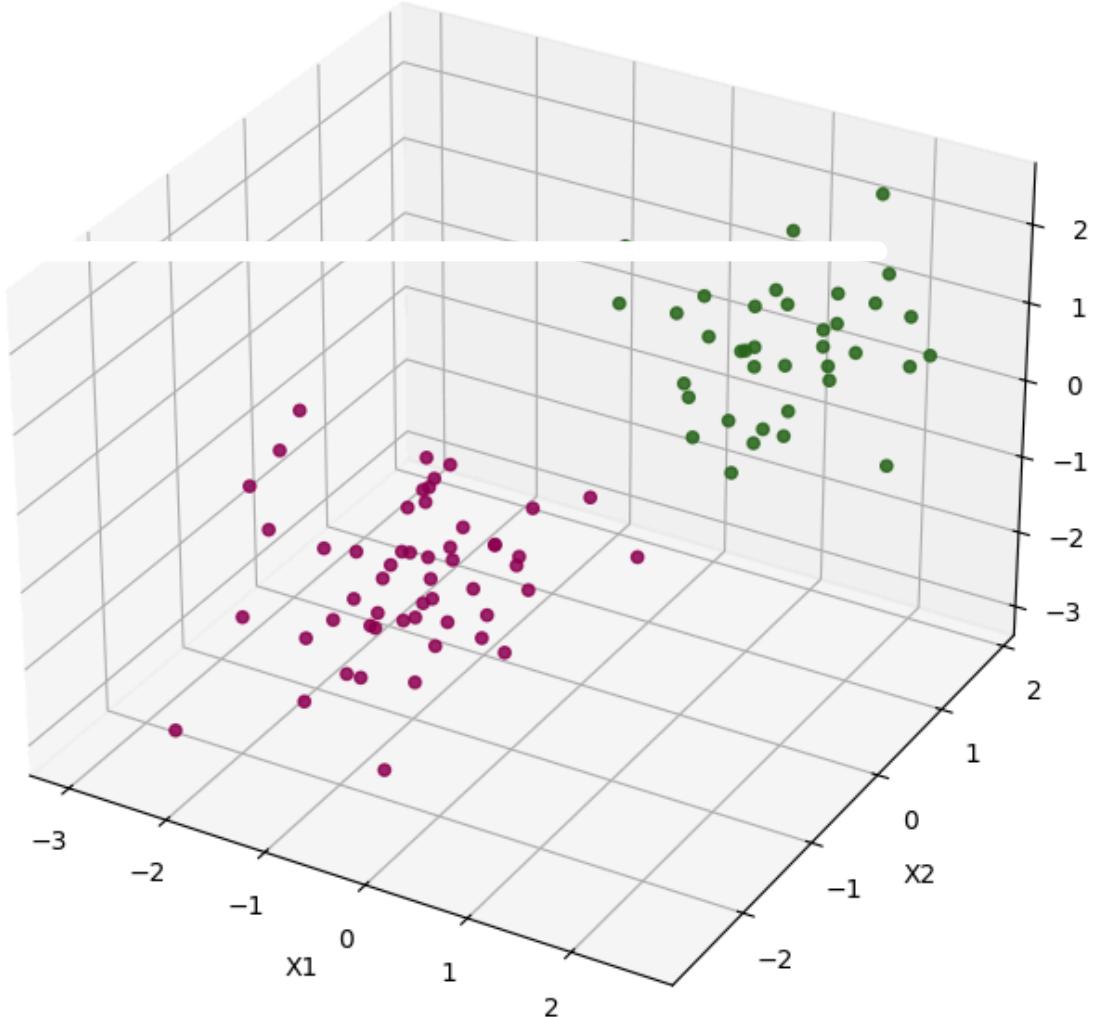
ax.set_xlabel("X1")
```

```
ax.set_ylabel("X2")
ax.set_zlabel("X3")
ax.set_title(f"blobs600: Shallow NN predictions (h={best_hb}) | test_acc={te
ax.legend()
plt.show()

print("blobs600 test accuracy:", accuracy(y1_test, y_pred_b))
```

blobs600: Shallow NN predictions (h=2) | test_acc=1.000

wrong



blobs600 test accuracy: 1.0

blobs600 3D scatter: Shallow NN predictions (h=2), test_acc=1.0

The two clusters are cleanly separated in 3D space, so the model can classify them almost perfectly even with a small hidden layer. This explains why increasing hidden nodes doesn't improve accuracy, it's already solved.

REF: [Confusion matrix from scratch](#)

```

def confusion_counts(y_true, y_pred):
    y_true = np.array(y_true).reshape(-1).astype(int)
    y_pred = np.array(y_pred).reshape(-1).astype(int)

    TN = np.sum((y_true == 0) & (y_pred == 0))
    FP = np.sum((y_true == 0) & (y_pred == 1))
    FN = np.sum((y_true == 1) & (y_pred == 0))
    TP = np.sum((y_true == 1) & (y_pred == 1))
    return TN, FP, FN, TP

def plot_confusion_matrix(TN, FP, FN, TP, title="Confusion Matrix", class0="0",
                         cm = np.array([[TN, FP],[FN, TP]]))

    plt.figure(figsize=(4,4))
    plt.imshow(cm)
    plt.xticks([0,1], [f"Pred {class0}", f"Pred {class1}"])
    plt.yticks([0,1], [f"True {class0}", f"True {class1}"])

    for i in range(2):
        for j in range(2):
            plt.text(j, i, cm[i,j], ha="center", va="center", fontsize=12)

    plt.title(title)
    plt.colorbar()
    plt.show()

```

```

TN, FP, FN, TP = confusion_counts(y2_test, y_pred_c)
print("circles500 confusion matrix:")
print("TN:", TN, "FP:", FP)
print("FN:", FN, "TP:", TP)

plot_confusion_matrix(TN, FP, FN, TP, title="circles500: Shallow NN Confusion
                      class0='0', class1='1'")

TN, FP, FN, TP = confusion_counts(y1_test, y_pred_b)
print("blobs600 confusion matrix:")
print("TN:", TN, "FP:", FP)
print("FN:", FN, "TP:", TP)

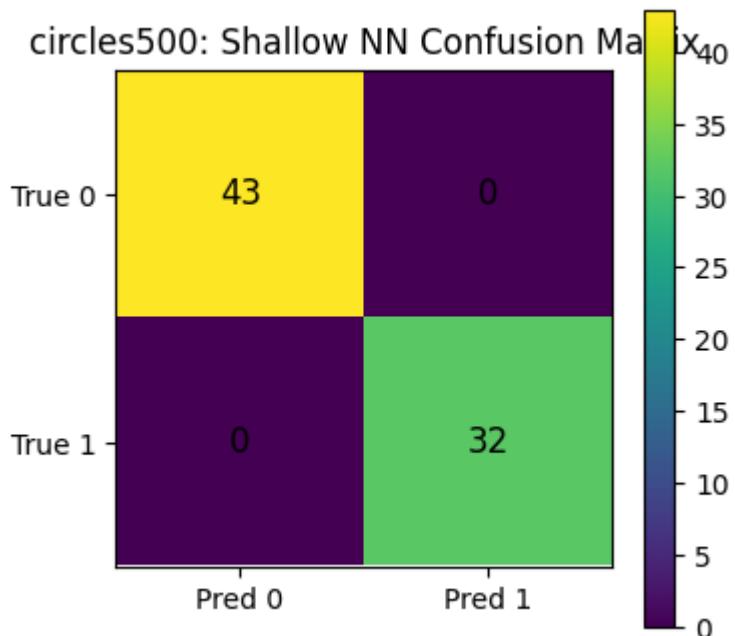
plot_confusion_matrix(TN, FP, FN, TP, title="blobs600: Shallow NN Confusion
                      class0='0', class1='1'")

```

circles500 confusion matrix:

TN: 43 FP: 0

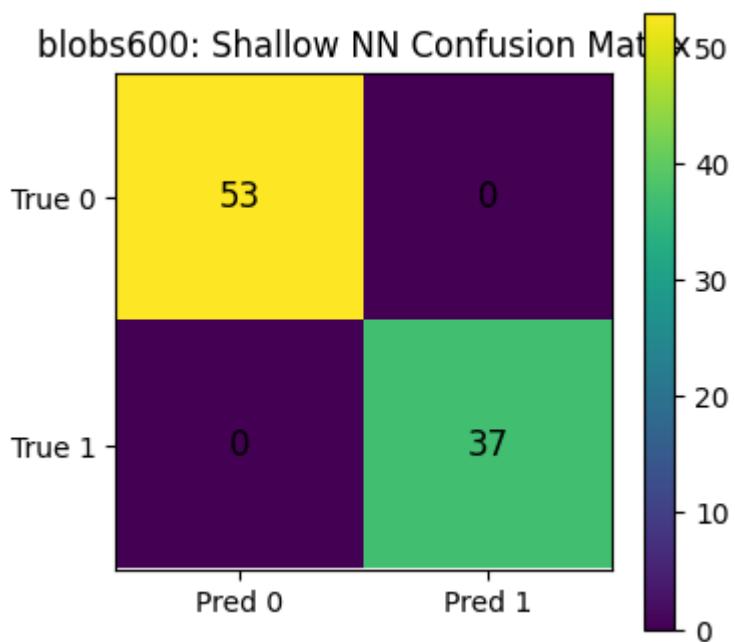
FN: 0 TP: 32



blobs600 confusion matrix:

TN: 53 FP: 0

FN: 0 TP: 37



CONCLUSION:

- circles500 hidden-size:
 - With h=2, the network doesn't have enough capacity, so accuracy is only moderate. Once we reach h=4 or h=8, the model has enough hidden units to learn a strong non-linear boundary, and performance becomes nearly perfect. h=8 is chosen as best because it gives the highest validation accuracy without unnecessary extra complexity.

- blobs600 hidden-size:
 - Even h=2 already achieves near perfect performance because blobs are simple and close to linearly separable. Adding more hidden nodes doesn't change results because the dataset doesn't need extra complexity.

▼ TASK 4

- Goal: Convert 26-class EMNIST letters into a binary dataset:
 - select only F and L
 - relabel to {0,1}: 0 = F ; 1 = L
- Why flatten:
 - 28×28 image → 784 input features for NN.
- Sanity checks:
 - counts near 3300 per class
 - sample plots look like F and L

```
import numpy as np
import matplotlib.pyplot as plt

data = np.load("emnist_letters_85800.npz")

x_data = data["x"]
y_data = data["y"]
```

```
# Do some data checks ...

# Check how many different classes we have: should be 26
n_classes = len(np.unique(y_data))
print(f"The number of unique classes is {n_classes} (should be 26)."

# check that images are scaled: min should be 0, max should be 1
img = x_data[0]
print(f"\nFor a single image, the min value is {img.min()} and the max is {i

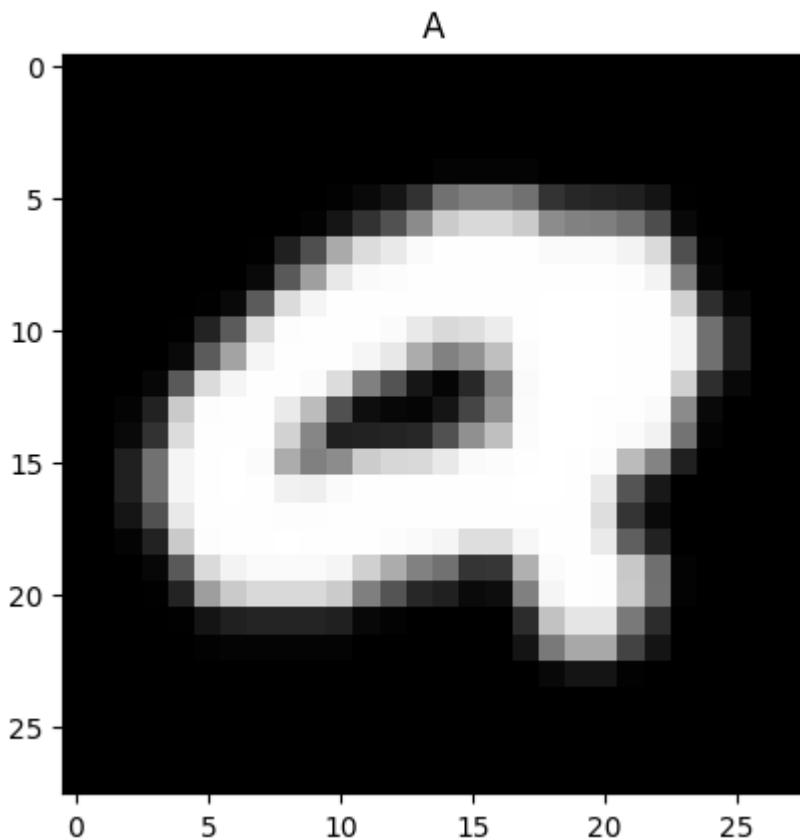
# Check the shape of the two classes
print(f"\nShape of x_data is {x_data.shape}, shape of y_data is {y_data.shap
```

The number of unique classes is 26 (should be 26).

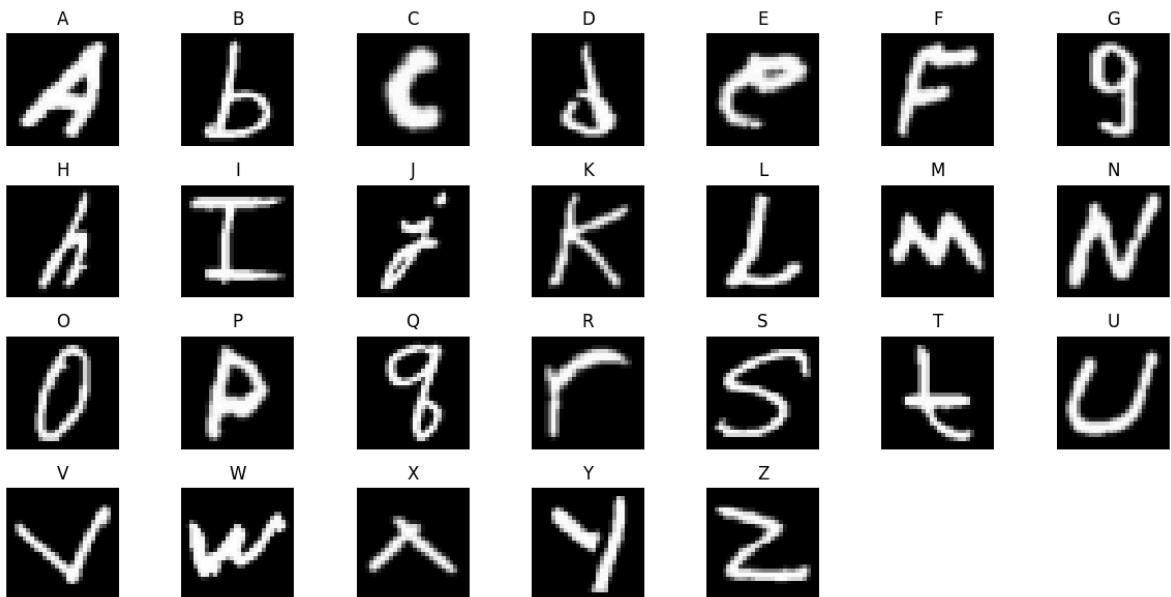
For a single image, the min value is 0.0 and the max is 1.0 (should be 0.0 and 1.0).

Shape of x_data is (85800, 28, 28, 1), shape of y_data is (85800,) (should have shape (85800,))

```
# Plot the first image  
  
plt.imshow(x_data[0].squeeze(), cmap="gray")  
plt.title(chr(int(y_data[0]) + 64))  
plt.show()
```



```
# Plot one item per class  
n = 1000 # plot the n-th item, starting from 0  
  
# Get unique labels  
classes = np.unique(y_data)  
  
plt.figure(figsize=(12, 6))  
  
for i, cls in enumerate(classes):  
    # Find first index of this class  
    idx = np.where(y_data == cls)[0][n]  
  
    plt.subplot(4, 7, i + 1)  
    plt.imshow(x_data[idx].squeeze(), cmap="gray")  
    plt.title(chr(int(cls) + 64)) # 1->A, 2->B, ...  
    plt.axis("off")  
  
plt.tight_layout()  
plt.show()
```



```
# Extract just two classes from the dataset

# PUT YOUR OWN CLASS NUMBERS HERE: remember that A=1, z=26.
c1 = 6    # F
c2 = 12   # L

mask = (y_data == c1) | (y_data == c2)

x_binary = x_data[mask]
y_binary = y_data[mask]

# Now change labels to 0 and 1
y_binary = (y_binary == c2).astype(int)

print("Binary X shape:", x_binary.shape)
print("Binary y shape:", y_binary.shape)
print("Counts (0=F, 1=L):", np.unique(y_binary, return_counts=True))

Binary X shape: (6600, 28, 28, 1)
Binary y shape: (6600,)
Counts (0=F, 1=L): (array([0, 1]), array([3300, 3300]))
```

```
# To help you examine your data, here is a graph to plot 48 images in a grid
```

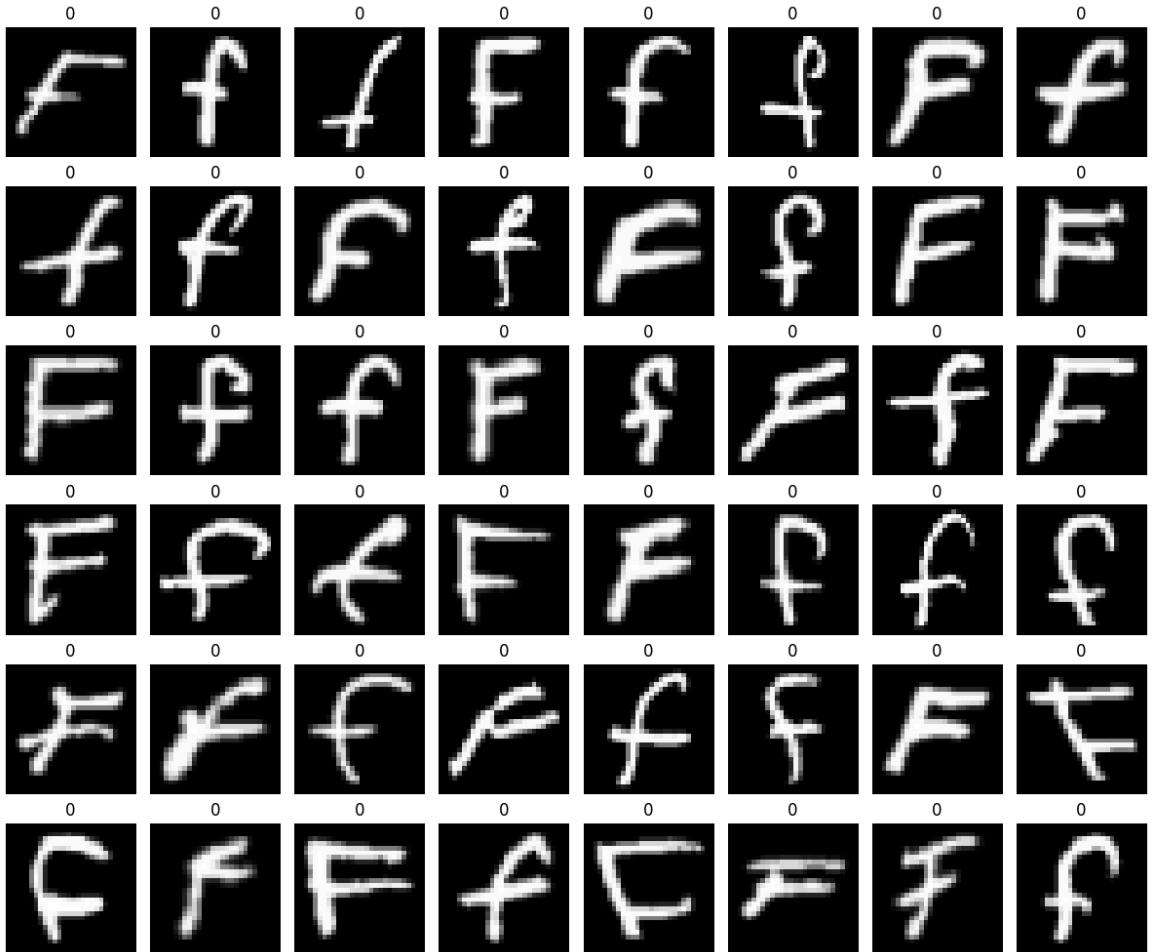
```
def plot_grid(x, y, n):
    plt.figure(figsize=(12, 10))

    for i in range(48):
        idx = n + i

        plt.subplot(6, 8, i + 1)
        plt.imshow(x[idx].squeeze(), cmap="gray")
        plt.title(int(y[idx]))
        plt.axis("off")

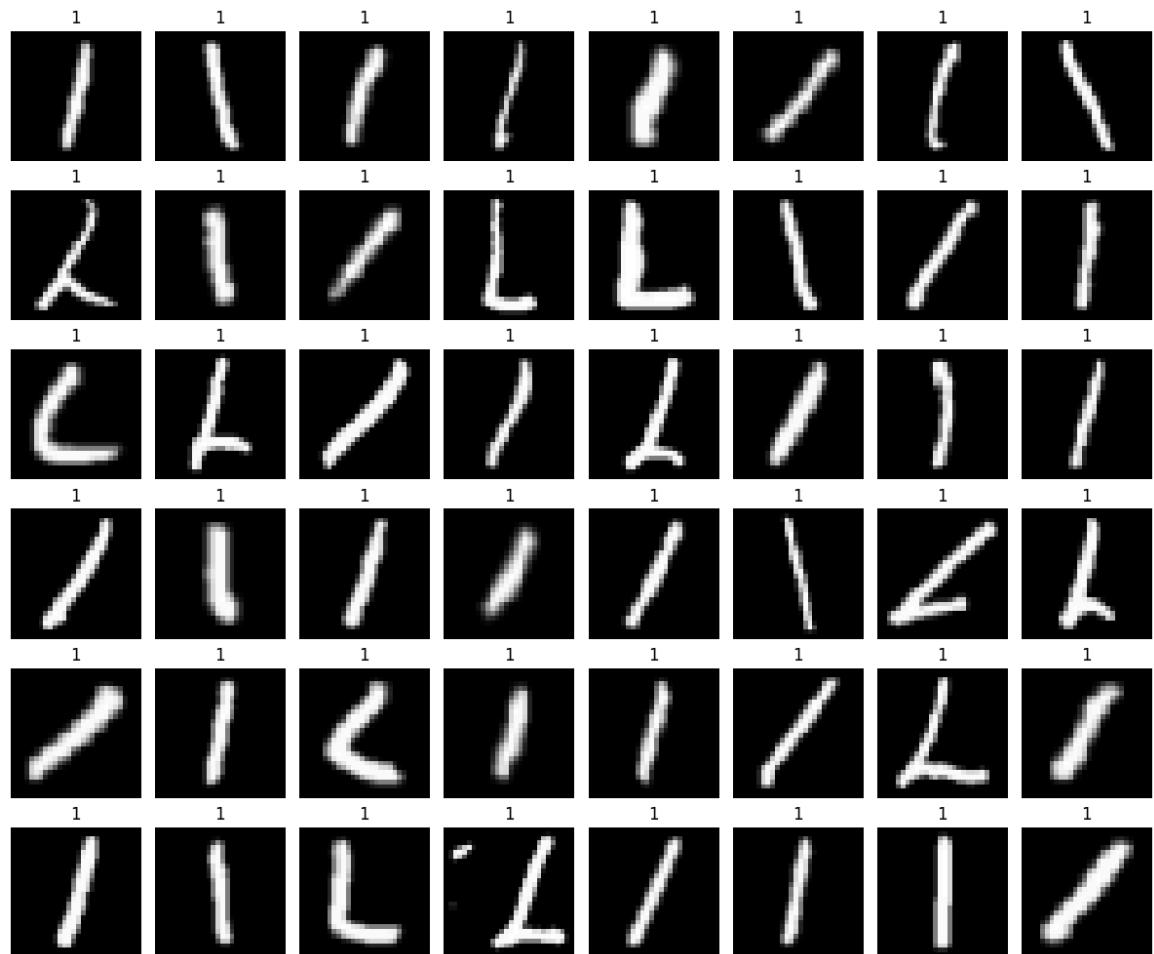
    plt.tight_layout()
    plt.show()
```

```
plot_grid(x_binary, y_binary, n=0)
```



```
# Now check images after 3300 - will all have label 1
```

```
plot_grid(x_binary, y_binary, n=3300)
```



```
X = x_binary.reshape(x_binary.shape[0], -1)    # (m, 784)
y = y_binary.reshape(-1)                      # (m,)

print("X shape:", X.shape)
print("y shape:", y.shape)

X shape: (6600, 784)
y shape: (6600,)
```

▼ Split + standardize + train

- Goal: Same pipeline as before:
 - 70/15/15 split
 - standardize using train only
 - train with bigger hidden nodes (32/64/128)
- Why bigger hidden layer: more inputs (784) → needs more capacity than 2D datasets.

Ref: [Optimal size of hidden layer](#)

```
def split_70_15_15(X, y, seed=0):
    X = np.array(X, dtype=float)
    y = np.array(y).reshape(-1).astype(int)

    m = X.shape[0]
    rng = np.random.default_rng(seed)
    idx = np.arange(m)
    rng.shuffle(idx)

    X = X[idx]
    y = y[idx]

    n_train = int(0.70 * m)
    n_val   = int(0.15 * m)

    X_train, y_train = X[:n_train], y[:n_train]
    X_val, y_val     = X[n_train:n_train+n_val], y[n_train:n_train+n_val]
    X_test, y_test   = X[n_train+n_val:], y[n_train+n_val:]

    return X_train, y_train, X_val, y_val, X_test, y_test

X_train, y_train, X_val, y_val, X_test, y_test = split_70_15_15(X, y, seed=0)

print("Train:", X_train.shape, y_train.shape)
print("Val:  ", X_val.shape, y_val.shape)
print("Test: ", X_test.shape, y_test.shape)

Train: (4620, 784) (4620,)
Val:  (990, 784) (990,)
Test: (990, 784) (990,)
```

```

def standardize_fit(X_train):
    mu = X_train.mean(axis=0, keepdims=True)
    sigma = X_train.std(axis=0, keepdims=True) + 1e-12
    return mu, sigma

def standardize_apply(X, mu, sigma):
    return (X - mu) / sigma

mu, sigma = standardize_fit(X_train)
X_train_s = standardize_apply(X_train, mu, sigma)
X_val_s   = standardize_apply(X_val, mu, sigma)
X_test_s  = standardize_apply(X_test, mu, sigma)

hidden_list = [32, 64, 128]
alpha_list = [0.05, 0.1, 0.2]

best = None # (val_acc, h, alpha, W1,b1,W2,b2,hist)

for h in hidden_list:
    for a in alpha_list:
        W1, b1, W2, b2, hist = fit_shallow_nn_sgd(
            X_train_s, y_train,
            n_hidden=h,
            alpha=a,
            max_iterations=100000,
            threshold=1e-4,
            N=len(X_train_s),
            seed=0,
            debug_every_N=False
        )

        y_val_pred = predict_class_nn(X_val_s, W1, b1, W2, b2)
        val_acc = accuracy(y_val, y_val_pred)

        print(f"h={h:<4} alpha={a:<4} val_acc={val_acc:.3f}")

        if (best is None) or (val_acc > best[0]):
            best = (val_acc, h, a, W1, b1, W2, b2, hist)

best_val_acc, best_h, best_alpha, W1b, b1b, W2b, b2b, hist_best = best
print("\nBest model by validation:")
print("hidden:", best_h, "alpha:", best_alpha, "val_acc:", best_val_acc)

```

```

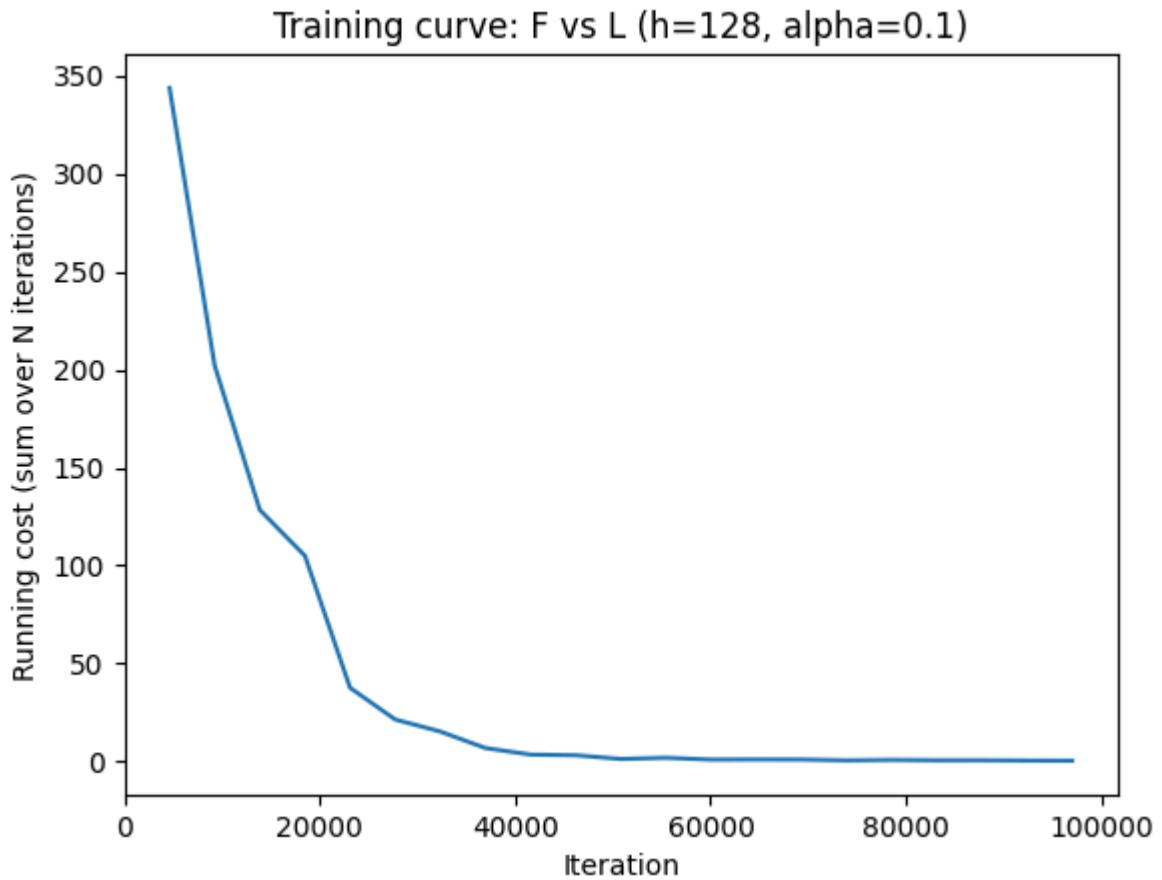
C:\Users\PS3\AppData\Local\Temp\ipykernel_14448\3500915864.py:2: RuntimeWarning:
      return 1 / (1 + np.exp(-z))
h=32  alpha=0.05 val_acc=0.979
h=32  alpha=0.1  val_acc=0.981
h=32  alpha=0.2  val_acc=0.987
h=64  alpha=0.05 val_acc=0.981
h=64  alpha=0.1  val_acc=0.984
h=64  alpha=0.2  val_acc=0.977
h=128 alpha=0.05 val_acc=0.982
h=128 alpha=0.1  val_acc=0.988
h=128 alpha=0.2  val_acc=0.981

```

```
Best model by validation:  
hidden: 128 alpha: 0.1 val_acc: 0.9878787878787879
```

Tried different hidden sizes and learning rates shows performance changes slightly, but the best validation result came from h=128, alpha=0.1. That combination gives enough capacity to learn the 784 pixel input without making training unstable.

```
iters = [t[0] for t in hist_best]  
Jrun = [t[1] for t in hist_best]  
  
plt.figure()  
plt.plot(iters, Jrun)  
plt.xlabel("Iteration")  
plt.ylabel("Running cost (sum over N iterations)")  
plt.title(f"Training curve: F vs L (h={best_h}, alpha={best_alpha})")  
plt.show()
```



The cost falls very quickly and then becomes almost flat near zero. That means the model learned the main pattern early and then just fine-tuned small details as training continued basically it has converged.

```
y_train_pred = predict_class_nn(X_train_s, W1b, b1b, W2b, b2b)  
y_val_pred = predict_class_nn(X_val_s, W1b, b1b, W2b, b2b)
```

```
y_test_pred = predict_class_nn(X_test_s, W1b, b1b, W2b, b2b)

print("Train acc:", accuracy(y_train, y_train_pred))
print("Val acc: ", accuracy(y_val, y_val_pred))
print("Test acc: ", accuracy(y_test, y_test_pred))

TN = np.sum((y_test == 0) & (y_test_pred == 0))
FP = np.sum((y_test == 0) & (y_test_pred == 1))
FN = np.sum((y_test == 1) & (y_test_pred == 0))
TP = np.sum((y_test == 1) & (y_test_pred == 1))
```

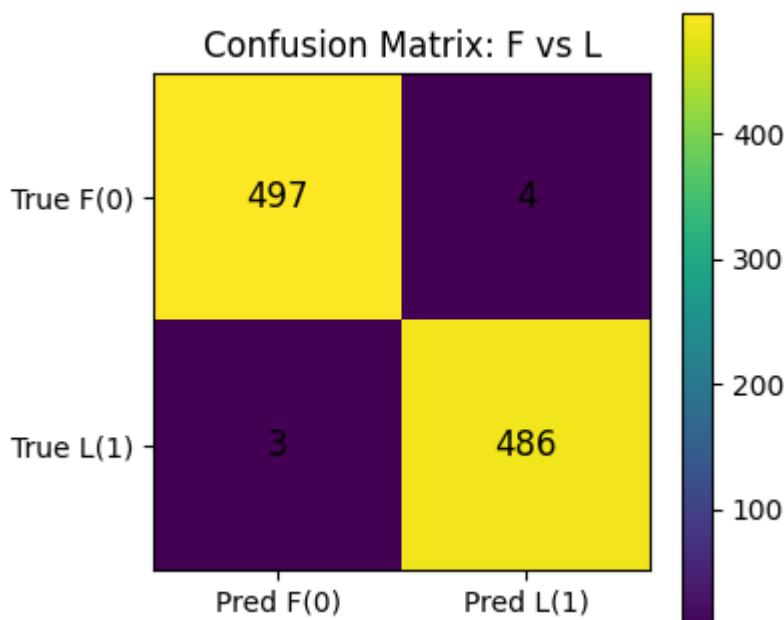
```
Train acc: 1.0
Val acc:  0.9878787878787879
Test acc:  0.9929292929292929
C:\Users\PS3\AppData\Local\Temp\ipykernel_14448\3500915864.py:2: RuntimeWarning:
return 1 / (1 + np.exp(-z))
```

```
cm = np.array([[TN, FP],
              [FN, TP]])

plt.figure(figsize=(4,4))
plt.imshow(cm)
plt.xticks([0,1], ["Pred F(0)", "Pred L(1)"])
plt.yticks([0,1], ["True F(0)", "True L(1)"])

for i in range(2):
    for j in range(2):
        plt.text(j, i, cm[i,j], ha="center", va="center", fontsize=12)

plt.title("Confusion Matrix: F vs L")
plt.colorbar()
plt.show()
```

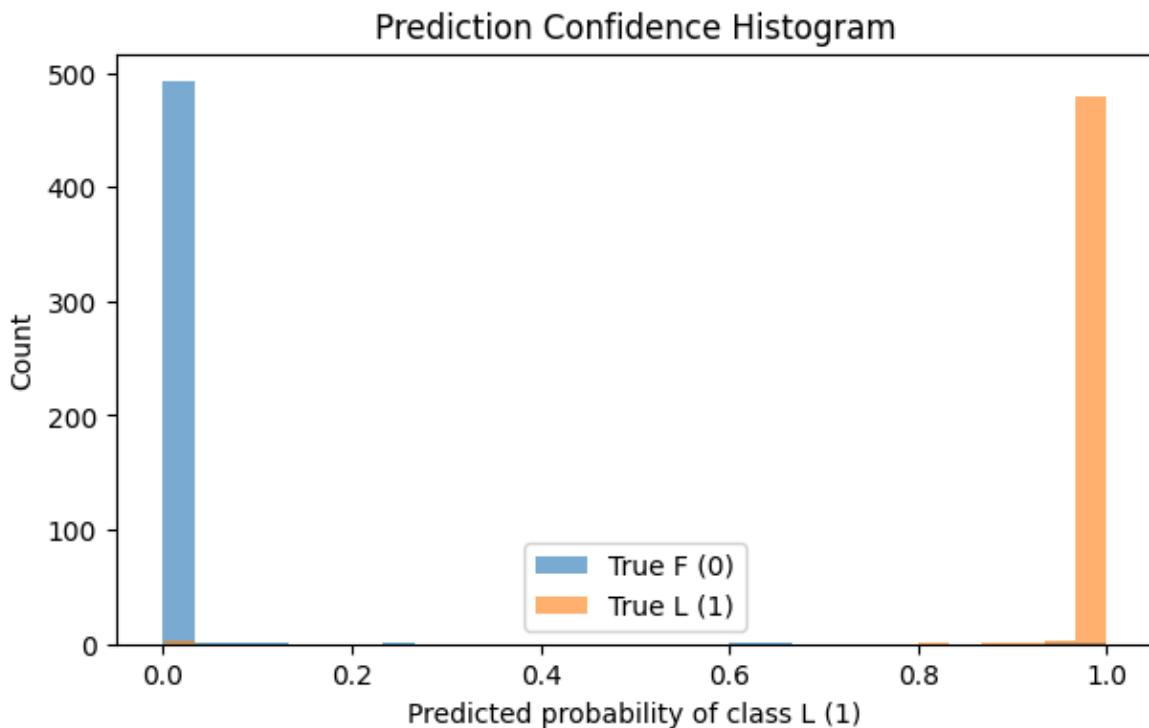


Most predictions are correct (big numbers on the diagonal). Only 4 F's were wrongly called L, and 3 L's were wrongly called F, so the model is getting almost everything right.

```
probs_test = predict_proba_nn(X_test_s, W1b, b1b, W2b, b2b) # (m,)

plt.figure(figsize=(7,4))
plt.hist(probs_test[y_test==0], bins=30, alpha=0.6, label="True F (0)")
plt.hist(probs_test[y_test==1], bins=30, alpha=0.6, label="True L (1)")
plt.xlabel("Predicted probability of class L (1)")
plt.ylabel("Count")
plt.title("Prediction Confidence Histogram")
plt.legend()
plt.show()
```

C:\Users\PS3\AppData\Local\Temp\ipykernel_14448\3500915864.py:2: RuntimeWarning:
return 1 / (1 + np.exp(-z))



- Able to see 2 strong peaks:
 - True F (0) is pushed near probability 0
 - True L (1) is pushed near probability 1 That means the model isn't just correct, it's usually very confident when it predicts.

```
def roc_curve_points(y_true, probs):
    thresholds = np.linspace(0, 1, 200)
    tpr_list, fpr_list = [], []

    for t in thresholds:
```

```
y_pred = (probs >= t).astype(int)
TP = np.sum((y_true==1) & (y_pred==1))
FP = np.sum((y_true==0) & (y_pred==1))
TN = np.sum((y_true==0) & (y_pred==0))
FN = np.sum((y_true==1) & (y_pred==0))

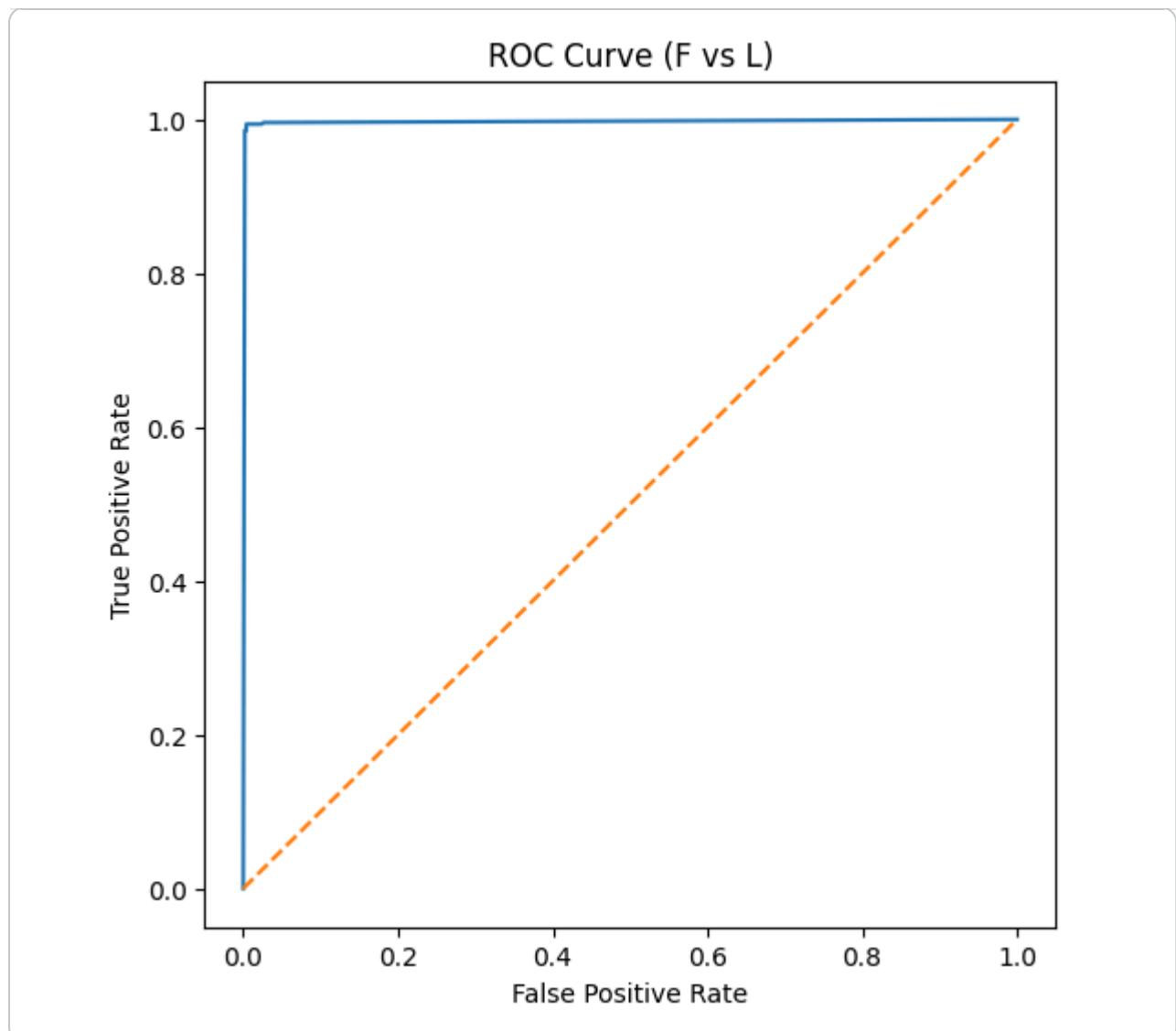
tpr = TP / (TP + FN + 1e-12)
fpr = FP / (FP + TN + 1e-12)

tpr_list.append(tpr)
fpr_list.append(fpr)

return np.array(fpr_list), np.array(tpr_list)

fpr, tpr = roc_curve_points(y_test, probs_test)

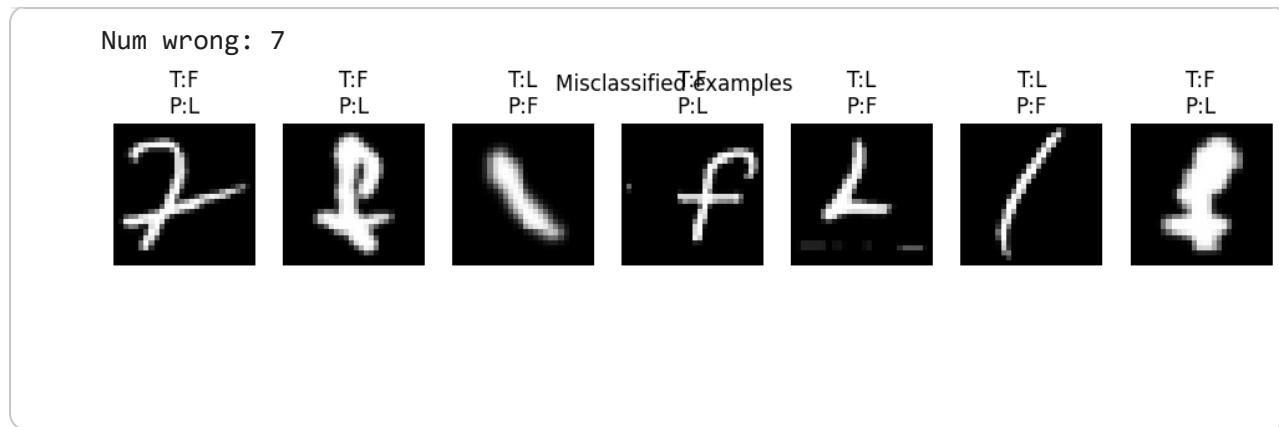
plt.figure(figsize=(6,6))
plt.plot(fpr, tpr)
# rand baseline drawn
plt.plot([0,1], [0,1], "--")
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ROC Curve (F vs L)")
plt.show()
```



The curve hugs the top left corner, far above the diagonal random guessing line. This means the model separates F and L extremely well across different thresholds (very high TPR with very low FPR).

```
wrong = np.where(y_test_pred != y_test)[0]
print("Num wrong:", len(wrong))

n = min(10, len(wrong))
plt.figure(figsize=(12,2))
for i in range(n):
    idx = wrong[i]
    plt.subplot(1, n, i+1)
    plt.imshow(X_test[idx].reshape(28,28), cmap="gray")
    plt.title(f"T:{'L' if y_test[idx]==1 else 'F'}\nP:{'L' if y_test_pred[id
    plt.axis("off")
plt.suptitle("Misclassified examples")
plt.show()
```



The few wrong images look like messy handwriting cases where F and L shapes can resemble each other (missing strokes, extra curves/ very faint writing). These are the natural hard cases in real handwriting.

CONCLUSION

- Using the same one-hidden-layer neural network from Task 3 (with 784 inputs, $h=128$, $\alpha=0.1$), the model achieved:
 - Train accuracy = 1.000
 - Validation accuracy ~ 0.988
 - Test accuracy ~ 0.993
- The confusion matrix ($TN=497$, $FP=4$, $FN=3$, $TP=486$) shows only 7 errors in the whole test set.
- The training curve and confidence histogram also confirm that the model converged and makes highly confident predictions for most samples.
- Overall, a single hidden layer is sufficient to distinguish F vs L very accurately, with remaining errors mainly coming from ambiguous handwriting.

▼ TASK 5

- In this task I extend my 1-hidden-layer neural network to support an arbitrary number of layers, where each layer can have a different number of nodes (given by a list like [784,128,64,1]).
- I also add a simple method to reduce overfitting on a small dataset: L2 regularization (weight decay).
- Finally, I evaluate the deep network on the Task 4 dataset (F vs L) and compare performance against the previous 1-hidden-layer baseline, including results with and without regularization.

Parameter initialization for an arbitrary number of layers

Here I initialize weights and biases for the architecture defined by a list: [n0,n1..nL] where n0 is input size and nL = 1 is the output node. For each layer l, weights and biases are:

$$W^l \rightarrow R^{(n_l * n_{l-1})}, b^l \rightarrow R^{(n_l * 1)}$$

They are stored in a dictionary like W1, b1, W2, b2,..

```
# parameter initialization for an arbitrary number of layers
def init_params_deep(layers, seed=0):
    rng = np.random.default_rng(seed)
    params = {}
    for l in range(1, len(layers)):
        n_in = layers[l-1]
        n_out = layers[l]
        params[f"W{l}"] = rng.normal(0, 0.1, size=(n_out, n_in))
        params[f"b{l}"] = np.zeros((n_out, 1))
    return params
```

Compute the output probability by applying the same pattern repeatedly across layers:

$$\text{For layer } l: Z^l = W^l \cdot a^{l-1} + b^l, a^l = \sigma(z^l)$$

with $a^0 = x$ (1 / many samples)

The final activation: a^L is the predicted probability: $y' = a^L$

I also store intermediate values (Z^l, A^l) in a cache so backpropagation can reuse them.

```
def forward_deep(x_or_X, params, return_cache=False):
    A = np.array(x_or_X, dtype=float)

    # many samples: X is (m,n)
    if A.ndim == 2 and A.shape[0] > 1:
        X = A
        A_prev = X.T # (n, m)
        cache = {"A0": A_prev}

        L = len([k for k in params.keys() if k.startswith("W")])
        for l in range(1, L+1):
            W = params[f"W{l}"]
            b = params[f"b{l}"]
            Z = W @ A_prev + b # (n_l, m)
            A_prev = sigmoid(Z) # (n_l, m)
            cache[f"Z{l}"] = Z
            cache[f"A{l}"] = A_prev

    y_prob = A_prev.T # (m,1)
```

```

        if return_cache:
            return y_prob, cache
        return y_prob

        # SINGLE sample
        x = to_col(A) # (n,1)
        A_prev = x
        cache = {"A0": A_prev}

        L = len([k for k in params.keys() if k.startswith("W")])
        for l in range(1, L+1):
            W = params[f"W{l}"]
            b = params[f"b{l}"]
            Z = W @ A_prev + b
            A_prev = sigmoid(Z)
            cache[f"Z{l}"] = Z
            cache[f"A{l}"] = A_prev

        y_prob = float(A_prev) #scalar
        if return_cache:
            return y_prob, cache
        return y_prob
    
```

✓ Loss function: Binary Cross Entropy (BCE) + L2 penalty

- The main loss for binary classification is binary cross entropy.
- To reduce overfitting, I add L2 regularization (weight decay) $J = J(\text{BCE}) + \lambda / 2m \cdot \sum_{l=1}^L \|W^l\|^2$
- When $\lambda = 0$, there is no regularization.
- When $\lambda > 0$, large weights are penalized, encouraging simpler models.

REF: [L2 Penalty – Ridge loss](#)

```

def bce_loss(y, y_prob, eps=1e-12):
    y = to_1d_labels(y)

    if np.isscalar(y_prob):
        p = float(np.clip(y_prob, eps, 1-eps))
        yy = int(y[0])
        return float(-(yy*np.log(p) + (1-yy)*np.log(1-p)))

    p = np.array(y_prob, dtype=float).reshape(-1)
    p = np.clip(p, eps, 1-eps)
    m = y.shape[0]
    return float(-(1/m)*np.sum(y*np.log(p) + (1-y)*np.log(1-p)))

def l2_penalty(params):
    s = 0.0
    L = len([k for k in params.keys() if k.startswith("W")])
    for l in range(1, L+1):
        W = params[f"W{l}"]
        s += np.sum(W**2)
    return s
    
```

```

        s += np.sum(W*W)
        return float(s)

def total_loss(y, y_prob, params, lam=0.0):
    J = bce_loss(y, y_prob)
    if lam > 0:
        m = len(to_1d_labels(y))
        J = J + (lam/(2*m)) * l2_penalty(params)
    return float(J)

```

Backpropagation for one training example (SGD)

Computing gradients for a single sample (because training uses SGD).

For the output layer with sigmoid + BCE, the delta simplifies to: $\delta^{L-1} = y' - y$ Then for each layer going backwards:

```

diffJ/diff W^L = del^L(a^(L-1))^T,
diffJ/diff b^L = del^L

```

For hidden layers: $\delta^{L-1} = (W^L)^T \cdot \delta^L * \sigma'(z^{L-1})$

With L2 regularization (weight decay), the weight gradient becomes:

```
diffJ/diff W^L <- diffJ/diff W^L + lambda.W^L
```

```

def backprop_one_deep(cache, y, params, lam=0.0):
    """
    cache contains A0..AL and Z1..ZL for single sample (all are (n,1))
    y is 0/1 scalar
    """
    y = int(to_1d_labels(y)[0])

    L = len([k for k in params.keys() if k.startswith("W")])
    grads = {}

    # Output layer
    AL = cache[f"A{L}"]           # (1,1)
    dZ = (AL - y)                # (1,1) delta at output (sigmoid + BCE)

    for l in reversed(range(1, L+1)):
        A_prev = cache[f"A{l-1}"]   # (n_{l-1},1)
        W = params[f"W{l}"]         # (n_l, n_{l-1})

        dW = dZ @ A_prev.T         # (n_l, n_{l-1})
        db = dZ                     # (n_l,1)

        # L2 term on weights (weight decay)
        if lam > 0:

```

```

dW = dW + (lam * W)

grads[f"dw{l}"] = dW
grads[f"db{l}"] = db

if l > 1:
    dA_prev = W.T @ dZ
    A_lm1 = cache[f"A{l-1}"]
    dZ = dA_prev * sigmoid_grad_from_a(A_lm1)

return grads

```

▼ Training loop: Stochastic Gradient Descent (SGD)

- Pick one random training sample (x_i, y_i)
- Forward pass $\rightarrow y'$
- Compute loss
- Backprop \rightarrow gradients
- Update parameters: $W^l \leftarrow W^l - \alpha * \frac{\partial J}{\partial W^l}$ $b^l \leftarrow b^l - \alpha * \frac{\partial J}{\partial b^l}$

I track a running cost every N iterations (roughly one pass over the dataset) to plot a training curve and monitor convergence.

```

def fit_deep_sgd(X, y, layers, alpha=0.1, lam=0.0, max_iterations=50000, thr=1e-05):
    X = np.array(X, dtype=float)
    y = to_1d_labels(y)
    m, n = X.shape
    assert layers[0] == n and layers[-1] == 1

    if N is None:
        N = m

    params = init_params_deep(layers, seed=seed)
    rng = np.random.default_rng(seed)

    J_running = 0.0
    J_prev = 0.0
    hist = []
    it = 0

    while True:
        i = rng.integers(0, m)
        x_i = X[i]
        y_i = y[i]

        y_prob, cache = forward_deep(x_i, params, return_cache=True)
        J = total_loss(y_i, y_prob, params, lam=lam)

        grads = backprop_one_deep(cache, y_i, params, lam=lam)

        for l in range(len(layers) - 1, 0, -1):
            dW = grads[f"dw{l}"]
            db = grads[f"db{l}"]

            dA_prev = W.T @ dZ
            A_lm1 = cache[f"A{l-1}"]
            dZ = dA_prev * sigmoid_grad_from_a(A_lm1)

            dW = dW + (lam * W)
            db = db + (lam * b)

            grads[f"dw{l}"] = dW
            grads[f"db{l}"] = db

        W = params[f"W{l}"]
        b = params[f"b{l}"]
        W -= alpha * dW
        b -= alpha * db

        J_running += J
        if it % N == 0:
            J_running /= N
            hist.append(J_running)
            J_running = 0.0
            print(f"iteration {it}, J: {J_running:.6f}")

        if abs(J - J_prev) < thr:
            break
        J_prev = J
        it += 1

```

```

L = len([k for k in params.keys() if k.startswith("W")])
for l in range(1, L+1):
    params[f"W{l}"] -= alpha * grads[f"dW{l}"]
    params[f"b{l}"] -= alpha * grads[f"db{l}"]

it += 1
J_running += J

if it >= max_iterations:
    if debug_every_N:
        print("Stopped: reached max_iterations.")
    break

if it % N == 0:
    change = abs(J_running - J_prev)
    hist.append((it, J_running, change))
    if debug_every_N:
        print(f"After {it} iters: J_running={J_running:.4f}, change={change:.4f}")
    if change < threshold:
        if debug_every_N:
            print("Stopped: convergence reached.")
        break
    J_prev = J_running
    J_running = 0.0

return params, hist

```

```

def predict_proba_deep(A_in, params):
    """
    Returns:
        - single sample -> scalar probability
        - many samples -> (m,) probabilities
    """
    y_prob = forward_deep(A_in, params) # scalar or (m,1)

    if np.isscalar(y_prob):
        return float(y_prob)

    return np.array(y_prob, dtype=float).reshape(-1)

def predict_class_deep(A_in, params, threshold=0.5):
    """
    Returns:
        - single sample -> scalar class 0/1
        - many samples -> (m,) int classes
    """
    probs = predict_proba_deep(A_in, params)

    if np.isscalar(probs):
        return int(probs >= threshold)

```