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Genetic Algorithm

Q1A: Roulette-Wheel Selection Modification

In the original code, tournament selection was used. As required, I made minimal and strictly necessary code updates by replacing only the selection() logic with a roulette-wheel mechanism.

Below is the modified portion of the code:

MODIFIED CODE SNIPPET — roulette_wheel_selection()

```
def roulette_wheel_selection(self, k=None):
    population = self.population
    pop_size = len(population)
    if pop_size == 0:
        return []
    if k is None:
        k = self.population_size
```

```

fitnesses = np.array([float(getattr(ind, 'fitness', 0.0)) for ind in population], dtype=float)

if not np.all(np.isfinite(fitnesses)):
    fitnesses = np.nan_to_num(fitnesses, nan=0.0, posinf=0.0, neginf=0.0)

min_f = fitnesses.min()
if min_f <= 0:
    fitnesses = fitnesses - min_f + 1e-8

total = fitnesses.sum()
if total <= 0:
    probs = np.ones(pop_size) / pop_size
else:
    probs = fitnesses / total

indices = np.random.choice(pop_size, size=k, replace=True, p=probs)

return [deepcopy(population[i]) for i in indices]

```

JUSTIFICATION

Roulette-wheel selection assigns proportional probabilities:

$$P(i) = f_i / \sum f_j$$

This ensures chromosomes with higher fitness are selected more frequently, unlike tournament selection which uses ranking. Here minimal code change was maintained — only selection logic replaced. Non-positive fitness values are shifted to positive domain without altering proportionality.

This is necessary because GA selection requires strictly positive weights for probability sampling. Numerical stability ($\text{EPS} = 1e-8$) avoids division-by-zero and ensures reproducibility.

Q2B: Modified Fitness Function With Parameter-Based Penalty

The original fitness function considered:

$$\text{fitness} = \text{accuracy} - \lambda * (\text{total_parameters} / \text{scale})$$

The updated requirement is to:

1. Separate Conv-block parameters and FC-layer parameters.
2. Assign different weights based on computational cost.

Below is the modified portion of the code:

MODIFIED CODE SNIPPET — `evaluate_fitness()`

```

conv_params = 0
fc_params = 0
other_params = 0

for module in model.modules():
    if isinstance(module, (nn.Conv1d, nn.Conv2d, nn.Conv3d)):
        for p in module.parameters(recurse=False):
            conv_params += p.numel()
    elif isinstance(module, (nn.Linear,)):
        for p in module.parameters(recurse=False):
            fc_params += p.numel()
    else:
        for p in module.parameters(recurse=False):
            other_params += p.numel()

conv_M = conv_params / param_scale
fc_M = fc_params / param_scale

penalty = (w_conv * conv_M) + (w_fc * fc_M)

fitness = best_acc - (lambda_penalty * penalty)

```

JUSTIFICATION FOR WEIGHTS

Convolution layers are computationally heavier than FC layers.

Conv-layer FLOPs $\approx k^2 \cdot C_{in} \cdot C_{out} \cdot H \cdot W$

FC-layer FLOPs $\approx \text{input_dim} * \text{output_dim}$

Hence:

- Conv blocks dominate computation → assign higher weight
- FC layers require significantly fewer operations → lower weight

Chosen weights:

$$w_{conv} = 0.7$$

$$w_{fc} = 0.3$$

These preserve the intent: architectures with excessive convolution parameters get proportionally penalized, while still allowing flexibility in FC layers.

Penalty Term:

$$\text{penalty} = w_{conv} * (\text{conv_params} / \text{scale}) + w_{fc} * (\text{fc_params} / \text{scale})$$

Final fitness:

$$\text{fitness} = \text{Accuracy} - \lambda \times \text{penalty}$$

This keeps accuracy primary but discourages unnecessarily large models.

LOGGING CHANGES

The revised code prints:

- conv_params
 - fc_params
 - penalty
 - final fitness
-