

Jayant Parmar
Postgraduate Diploma in Artificial Intelligence
Artificial Intelligence
Indian Institute of Technology Jodhpur, Jodhpur, India
Email: g25ait1072@iitj.ac.in

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 ($\text{EPS} = 1\text{e-}8$) 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 $\approx k*k*C_{in}*C_{out}*H*W$

FC-layer FLOPs $\approx input_dim * output_dim$

Hence:

- Conv blocks dominate computation \rightarrow assign higher weight
- FC layers require significantly fewer operations \rightarrow 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:

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

Final fitness:

$$fitness = Accuracy - \lambda \times penalty$$

This keeps accuracy primary but discourages unnecessarily large models.

LOGGING CHANGES

The revised code prints:

- conv_params
- fc_params
- penalty
- final fitness
