

# MultiGA: Leveraging Multi-Source Seeding in Genetic Algorithms



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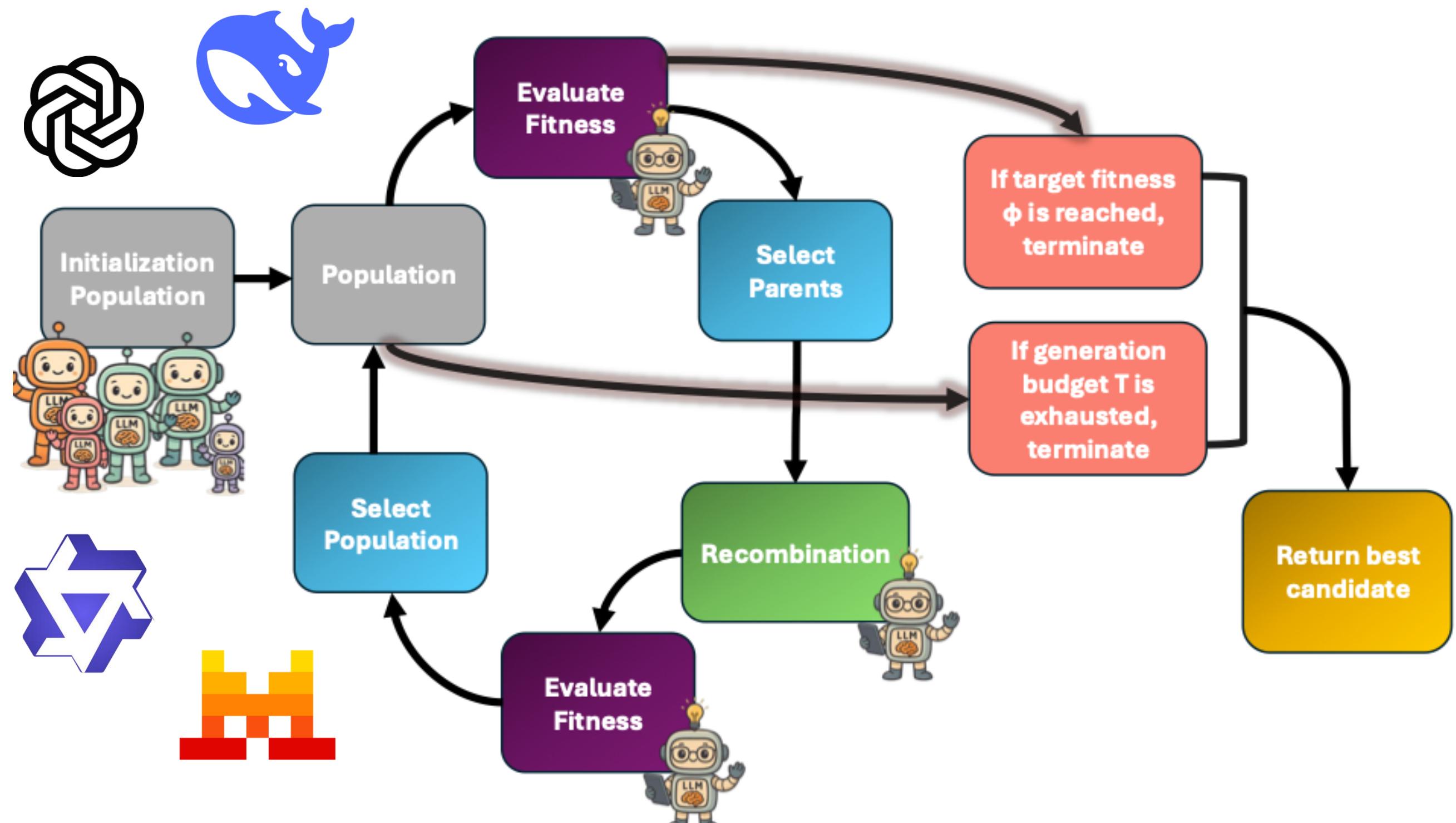


1

Can heterogeneous LLMs, each with different strengths, jointly improve robustness on code generation and complex reasoning tasks?

2

Idea: Initialize the genetic algorithm cycle with candidates produced by multiple LLMs, creating a diverse starting population.



Overview of the MultiGA framework. Populations are initialized with multiple LLMs, while an independent LLM  $E$  handles fitness evaluation (scoring candidates) and recombination (combining two parent solutions). The process terminates once target fitness  $\phi$  or maximum number  $T$  generations is reached. Then, the top candidate solution is returned.

## Algorithm 1: MultiGA: Multi-Source Genetic Algorithm

**Input:** Task spec  $Q$ ; Generator LLMs  $\mathcal{G} = \{g_1, \dots, g_m\}$ ; Evaluator/crossover LLM  $E$  with fitness  $f : \mathcal{X} \rightarrow [0, 1]$ ; Population size  $n$ ; Top- $k$ ; Threshold  $\tau \in [0, 1]$ ; Max generations  $T$ ; Target fitness  $\phi \in [0, 1]$ .

**Output:** Best solution  $\hat{x} \in \mathcal{X}$ .

```
1  $P_0 \leftarrow \text{initialize\_population}(\mathcal{G}, Q)$ ;  
2  $t \leftarrow 0$ ;  
3 while  $t < T - 1$  do  
4    $scores \leftarrow \text{evaluate\_fitness}(P_t, f)$ ;  
5   if  $\max(scores) \geq \phi$  then  
6     break  
7    $S_t \leftarrow \text{select\_parents}(P_t, scores, k)$ ;  
8    $C_t \leftarrow \text{recombination}(S_t, P_t, E, Q)$  # Create next generation (children solutions);  
9    $P_{t+1} \leftarrow \text{select\_population}(P_t, C_t, \tau)$  # Retire unfit candidates;  
10   $t \leftarrow t + 1$ ;  
11   $scores \leftarrow \text{evaluate\_fitness}(P_t, f)$ ;  
12   $\hat{x} \leftarrow \arg \max_{x \in P_t} f(x)$ ;  
13 return  $\hat{x}$ ;
```

3

This diversity gives crossover and mutation far more variation to work with, enabling hybrid reasoning and code patterns that no single model would generate.

4

MultiGA produces accuracy that consistently approaches the performance of the strongest single model baseline and reduces reliance on any single model through iterative refinement and recombination.

Example of recombination on a text-to-SQL task (“Which publisher published the slowest superhero?”). Parent 1 (gpt-4o, score 0.95) was paired with a randomly selected parent (deepseek-r1, score 0.1). The resulting child achieved a perfect score of 1.0 by preserving gpt-4o’s overall structure while incorporating the MIN aggregation from r1.

Dataset	Test Rows	Total Items	Label Choices
$\mathcal{D}_{SQL}$ (BIRD Mini-Dev)	100	500	$\infty$
$\mathcal{D}_{NATPLAN}$ (Meeting Planning)	100	1000	$\infty$
$\mathcal{D}_{GPQA}$ (Grad-Level Science Questions)	198	198	4
$\mathcal{D}_{BBQ}$ (BBQ Bias Data)	104	6879	3

Datasets used in our experiments, with test set size, total available items, and number of label choices.

Seed Model	$\mathcal{D}_{SQL}$	$\mathcal{D}_{NATPLAN}$	$\mathcal{D}_{GPQA}$	$\mathcal{D}_{BBQ}$
gpt-4o	<b>0.56</b>	0.39	<b>0.44</b>	<b>1.00</b>
qwen2.5-coder-32b-instruct	0.45	0.28	0.30	0.87
deepseek-r1-distill-qwen-32b	0.46	0.23	0.41	0.97
openai/gpt-oss-20b	0.48	0.19	0.41	0.93
mistral-7b-instruct-v0.3	0.25	0.10	0.22	0.87
$\mathcal{G}$ (MultiGA)	0.55	<b>0.40</b>	0.40	<b>1.00</b>

Accuracy across tasks when seeding with each individual model versus seeding with all five simultaneously ( $G$ ).

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I am applying to CS PhD programs for the Fall 2026 cycle & seeking potential advisors!

