Genetic Algorithms for Federated Learning: Two Distinct Approaches

GenFed vs. FedCSGA - Comprehensive Analysis

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Federated Learning: The Core Problem

Given N clients with local datasets \mathcal{D}_i , the goal is to solve:

$$\min_{w \in \mathbb{R}^d} f(w) = \sum_{i=1}^N \frac{n_i}{n} F_i(w)$$
 (1)

where $F_i(w) = \frac{1}{n_i} \sum_{(x,y) \in \mathcal{D}_i} \ell(w; x, y)$ is the local loss.

Key Challenges Addressed by these Papers:

- Data Heterogeneity (Non-IID): Clients' data distributions \mathcal{D}_i vary significantly, causing local models to drift and slowing convergence.
- System Heterogeneity: Clients have different computation, memory, and network speeds.
- **Communication Bottlenecks:** Uploading models is slow; total training time is often constrained by a deadline.
- Participant Reliability: Some clients may be slow, drop out, or be malicious (Byzantine).

The Landscape of FL Client/Model Optimization

Our papers tackle the challenges of heterogeneity and efficiency. They fit into a broader landscape of FL optimization research:

Optimization-Based

- Goal: Maximize clients or minimize time.
- **FedCS:** Greedily selects clients who can meet a deadline.
- Hybrid-FL: Heuristic to select clients based on time and data distribution.
- FedCSGA Uses a full Genetic Algorithm to solve this NP-hard selection problem, outperforming simple greedy heuristics.

Utility-Function-Based

- **Goal:** Select clients with the highest "utility".
- Oort: Defines utility by training loss and time, prioritizing clients with high impact and speed.
- **PyramidFL:** A fine-grained utility approach.
- GenFed Uses a GA-inspired "fitness" (validation accuracy) to select *models* post-training, not *clients* pre-training.

Two GA Approaches: When to Apply?

FedCSGA (Wu et al.) Genetic Client Selection

- When: Before training round.
- **What:** Selects the optimal *set* of clients to participate.
- Goal: Maximize number of clients that can finish within a time deadline T.
- GA Use: Full GA (crossover, mutation) to solve the complex, NP-hard scheduling problem.

GenFed (Zheng et al.) Genetic Model Aggregation

- When: After clients train.
- **What:** Selects the optimal *set* of models to aggregate.
- Goal: Maximize global model quality by filtering out low-performing or malicious models.
- GA Use: GA-inspired (fitness, selection) but only uses the "Selection" step.

GenFed: Genetic Mechanism Mapping

Idea: FL as Genetic Evolution

| Genetic Mechanism | Federated Learning Component |
|--------------------|--|
| Population | Set of client models $\{w_i\}$ |
| Crossover | Server aggregation $(\sum w_i)$ |
| Mutation | Local client training (Gradient Descent) |
| Fitness Evaluation | Model performance on a test set |
| Selection | Missing in traditional FL! |

GenFed's Contribution: Add the missing **Selection** step.

- How: Server maintains a small, public, and privacy-safe global validation set \mathcal{D}_g^{ν} .
- Before aggregation, the server evaluates all k received models:

$$fitness_i = Accuracy(w_i^t, \mathcal{D}_g^v)$$

- It selects only the top- ρ_t models with the highest fitness.
- It aggregates only these elite models to form w_{t+1} .

GenFed: Pseudocode (Algorithm 2)

```
1: Server-Side Execution:
 2: Initialize w_{\sigma}^{0}, Global validation set \mathcal{D}_{\sigma}^{v}
      Initialization Phase: (Optional) Send supplemental data to clients with little data.
      Training Phase:
      for each round t = 1, 2, ..., T do
 6:
           S_t \leftarrow \text{Select a subset of clients (e.g., 10 random clients)}
 7:
           for each client k \in S_t in parallel do
 8:
                Send w_{\pi}^{t} to client k
                w_{\iota}^{t} \leftarrow \mathring{\mathsf{C}} \mathsf{lientUpdate}(w_{\varrho}^{t}, \mathcal{D}_{k}) \; \{\mathsf{Local \; training} = \mathsf{Mutation}\}
 9:
10:
                Send w_k^t back to server
11:
           end for
12:
           — GenFed Step —
13: for each received model w_k^t do
14:
                \mathsf{fitness}_k \leftarrow \mathsf{Evaluate}(w_k^t, \mathcal{D}_{g}^{\mathsf{v}}) \; \{\mathsf{Evaluate} \; \mathsf{fitness}\}
15:
           end for
16: \rho_t \leftarrow \text{GetAggregationQuantity}(t, \rho_{\text{max}}, \text{strategy})
17:
      C_t \leftarrow \text{Select top-}\rho_t \text{ models from } S_t \text{ based on fitness } \{\text{Selection}\}
18:
           w_{g}^{t+1} \leftarrow \sum_{i \in C_{t}} p_{i} w_{i}^{t} \{ Crossover (of elite models) \}
19: end for
20: Return w_{\sigma}^{T}
```

FedCSGA: System Model & Problem

Goal: Maximize clients selected, given a hard deadline T.

Time Model:

- Computation: $t_i^{\text{comp}} = (e \cdot d_i)/c_i$ (epochs × data / capacity)
- Communication: $t_i^{\text{comm}} = S/B_i$ (model size / bandwidth)
- Clients upload **sequentially** (one-by-one).

Key Insight: Computation can overlap with the *previous* client's communication. The total time depends on the **sequence q** of clients. **Optimization Problem:**

$$\max_{\boldsymbol{q}} |\boldsymbol{q}| \quad \text{s.t.} \quad \boldsymbol{\Theta}^{\boldsymbol{q}}_{|\boldsymbol{q}|} \leq \mathcal{T}$$

- $\mathbf{q} = \langle q_1, q_2, \ldots \rangle$ is the client upload order.
- $\Theta_{|\mathbf{q}|}^{\mathbf{q}}$ is the total time for sequence \mathbf{q} (a complex function of overlaps).
- This is a variant of BIN PACKING, which is **NP-hard**.
- A simple greedy approach will fail; a GA is well-suited.

FedCSGA: Fitness Function

Chromosome: A sequence of clients, $\mathbf{q}_i = \langle q_{i1}, q_{i2}, \ldots \rangle$. **Fitness Function (with Adaptive Penalty):**

$$F(\mathbf{q}_i) = h(\mathbf{q}_i) - \lambda g(\mathbf{q}_i)$$

- **Objective** $h(\mathbf{q}_i)$: How "good" is the sequence?
 - IID: $h(\mathbf{q}_i) = |\mathbf{q}_i|$ (Just maximize client count)
 - Non-IID: $h(\mathbf{q}_i) = \sum_{j=1}^{|\mathbf{q}_i|} (1 \alpha \cdot A(q_{ij}))$ (Balances count with accuracy A, prioritizing low-accuracy clients)
- **Penalty** $g(\mathbf{q}_i)$: How "infeasible" is it?
 - $g(\mathbf{q}_i) = e^{\max(0,(\Theta-T)/T)} 1$ (Exponential penalty for exceeding deadline T)
- Penalty Factor λ :
 - $\lambda = \lambda_0 e^{\sqrt{r}}$ (where r = generation)
 - **Intuition:** Start with low penalty (explore), end with high penalty (enforce deadline).

FedCSGA: Pseudocode (Algorithm 1, Part 1/2)

```
    Input: Client set M, deadline T, population size n, generations R
    Output: Optimal client sequence q*
    Initialization:
    P<sub>r</sub> ← ∅ (population)
    for i = 1 to n do
    k ← random client from M
    q<sub>i</sub> ← ⟨k⟩
    Greedily add cheapest clients to q<sub>i</sub> until Θ<sup>q<sub>i</sub></sup> > T
    Add q<sub>i</sub> to P<sub>r</sub>
    end for
```

FedCSGA: Pseudocode (Algorithm 1, Part 2/2)

```
12: Evolution:
13: while r < R do
14:
           P_{r+1} \leftarrow \emptyset
15:
           Crossover:
16:
       for c = 1 to n/2 do
17:
               Select \mathbf{q}_i, \mathbf{q}_i from P_r (e.g., tournament)
18:
               \mathbf{q}'_i, \mathbf{q}'_i \leftarrow \mathsf{UniformCrossover}(\mathbf{q}_i, \mathbf{q}_i) with adaptive p_c
19:
           end for
20:
           Mutation:
21:
           for c = 1 to n do
22:
               \mathbf{q}'_i \leftarrow \mathsf{AdjacentSwapMutation}(\mathbf{q}_i) with adaptive p_m
23:
           end for
24:
           Selection:
25:
         P_{r+1} \leftarrow \mathsf{SelectBest}(P_r \cup P', n) \text{ using fitness } F(\mathbf{q})
26:
           r \leftarrow r + 1
27: end while
28: \mathbf{q}^* \leftarrow \text{Best } feasible \text{ chromosome } (\Theta < T) \text{ from } P_R
29: Return q*
```

Theoretical Analysis

GenFed (Zheng et al.):

- Contribution is conceptual (mapping FL to GA), lacking a formal convergence proof.
- **Implied Guarantee:** Selecting models by validation accuracy biases the global model toward a better solution .
- This filtering also provides inherent robustness by discarding low-quality or Byzantine models

FedCSGA (Wu et al.):

- NP-Hardness: The paper formally states that the deadline-constrained, order-dependent client selection problem is NP-hard.
- This justifies using a meta-heuristic (like a GA) over a simple greedy algorithm (like FedCS), which is likely to get stuck in a local optimum.
- The GA is a tool to find a high-quality approximate solution to this NP-hard problem in polynomial time.

Base Training Configuration (Our Re-Implementation)

Model Architecture:

- MLP: 784 (input) ightarrow 128 ightarrow 64 ightarrow 10 (output)
- Loss: CrossEntropyLoss
- Optimizer: SGD with momentum

Training Hyperparameters:

- Learning rate: $\eta = 0.01$
- Local epochs: E = 5
- Batch size: 32
- Device: CUDA/CPU auto-detect

Federated Setup:

- Datasets: MNIST
- Total Clients N: 10 (for our test) or 100 (in papers)
- Clients per Round k: 5 (our test) or 10 (in papers)
- Non-IID: Dirichlet distribution with α (lower $\alpha =$ more heterogeneous)

Key Hyperparameters Explained

GenFed (Zheng et al.):

- $\mathcal{D}_{\sigma}^{\mathsf{v}}$: **Global Validation Set**. A public, balanced dataset (e.g., 10% of MNIST test set) held by the server to evaluate model fitness.
- ρ_t : Aggregation Quantity. The number of "elite" models to select for aggregation.
- ρ_{max} : Max Aggregation. The upper bound for ρ_t .
- Strategy (Eq 5-9): The function (Linear, Sinusoidal, etc.) that governs how ρ_t changes over time t.

FedCSGA (Wu et al.):

- T: **Time Deadline**. The hard constraint (in seconds) that the entire round (computation + sequential upload) must fit into.
- n: Population Size. The number of chromosomes (sequences) in the GA (e.g., 90).
- R: **Generations**. The number of evolution steps the GA takes.
- p_c, p_m : Crossover/Mutation Probabilities. Dynamically adapted based on chromosome fitness to balance exploration/exploitation.
- α: Non-IID Factor. A value in [0,1] that balances maximizing client Team 12 (CS6007 Project)

Paper Results: GenFed (Zheng et al.)

Setup: 100 clients, 10 selected/round, Dirichlet $\alpha = 0.1$

Performance (vs. FedAvg)

• **MNIST:** 15× faster, +1.17% Acc

• SVHN: $34 \times$ faster, +1.62%Acc

• Fashion: $36 \times$ faster, +7.71% Acc

• CIFAR-10: $16 \times$ faster, +3.98%Acc

Byzantine Robustness (CIFAR-10)

| Attack | FedAvg Drop | GenFed Drop |
|------------|-------------|-------------|
| Label Flip | -5.90% | -3.01% |
| IPM . | -2.93% | -0.77% |
| Mimic | -3.96% | -1.38% |

Insight: The validation filtering naturally discards malicious models as "low fitness", reducing attack impact by > 50%.

Scalability

- FedAvg performance degrades as client count ↑
- GenFed performance improves

Paper Results: FedCSGA (Wu et al.)

Setup: 100 clients, GA (n=90, R=10), Non-IID ($\alpha = 0.7$)

IID Setting (T = 3min)

Non-IID Setting (T = 5min)

| Method | Clients/Round | Accuracy | | |
|----------|---------------|----------|--|--|
| | Fashion-MNIST | | | |
| FedAvg | 2.3 | 88.7% | | |
| FedCS | 7.8 | 89.1% | | |
| FedCSGA | 8.5 (+9%) | 89.9% | | |
| CIFAR-10 | | | | |
| FedAvg | 1.6 | 70.6% | | |
| FedCS | 6.5 | 72.3% | | |
| FedCSGA | 7.0 (+8%) | 76.0% | | |

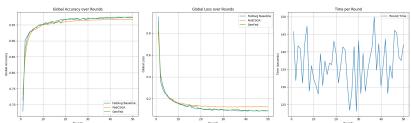
| Method | ${\sf Clients}/{\sf Round}$ | Accuracy | | |
|-----------|-----------------------------|----------|--|--|
| | Fashion-MNIST | | | |
| FedAvg | 3.0 | 57.8% | | |
| PyramidFL | 9.0 | 66.1% | | |
| FedCSGA | 13.5 (+50%) | 66.5% | | |
| | CIFAR-10 | | | |
| FedAvg | 2.9 | 38.8% | | |
| PyramidFL | 7.5 | 53.3% | | |
| FedCSGA | 11.2 (+49%) | 54.0% | | |
| | | | | |

Key Insights:

- GA (FedCSGA) consistently finds better solutions (more clients) than the greedy (FedCS) or random (FedAvg) approaches.
- The Non-IID fitness function ($\alpha=0.7$) is critical, dramatically improving accuracy over FedAvg (54.0% vs 38.8%) by balancing client count and data diversity.

Our Results: Accuracy & Loss

Setup: N = 10 clients, k = 5 selected, MNIST, $\alpha = 0.5$

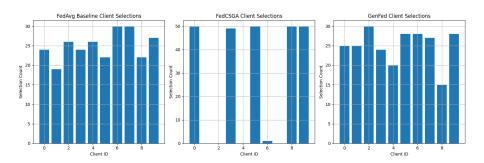


Analysis:

| Method | Avg Accuracy |
|------------------------------|-----------------|
| FedAvg (Baseline) | 96.61% |
| GenFed ($\rho = 3$) | 97.02% (+0.41%) |
| FedCSGA | 96.18% (-0.43%) |

- GenFed (Green): Achieved the highest accuracy, confirming the paper's claim that filtering models improves quality.
- FedCSGA (Orange): Underperformed.
- FedAvg (Blue): Baseline → ✓ へ ∼

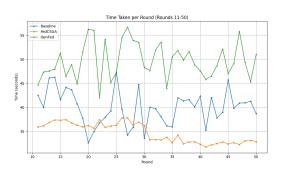
Our Results: Client Selection Analysis



Analysis:

- FedAvg (Left): Selection is random, as expected. All 10 clients were selected roughly 20-30 times over 50 rounds.
- FedCSGA (Center): The GA was *highly* selective. It almost exclusively picked clients 0, 2, 4, 6, 8.
- GenFed (Right): Selection is random (like FedAvg) because GenFed selects *models* after training, it doesn't change the *client selection* policy.

Our Results: Timing Analysis (Extrapolated)



- Time per Round (Left): FedCSGA is stable; GenFed varies due to validation, FedAvg fluctuates moderately.
- Time Ratios (Center): GenFed slows with complexity; FedCSGA stays consistently fast.
- Cumulative Time (Right): FedCSGA is fastest overall; GenFed is slowest from validation overhead.

Lessons Learned: FedCSGA Failure Analysis

Root Cause: Stale Fitness Proxy Problem in Our Implementation:

- FedCSGA's Non-IID fitness $h(\mathbf{q}) = \sum (1 \alpha \cdot A(q_j))$ requires knowing client accuracy $A(q_j)$.
- We computed this accuracy **once** at round $0 (w_0)$.
- The GA used this stale accuracy as a proxy for all 50 rounds.
- As the global model w_t evolved, the round 0 accuracy became a meaningless proxy.
- The GA "overfit" to the stale proxy, picking clients (0,2,4,6,8) that were good at round 0, but not necessarily at round 40.

Proposed Fixes:

- **QUALIFY Re-evaluate Fitness:** Re-run all clients on a validation set every \sim 5 rounds to get fresh accuracy data for the GA.
- **Scale Up:** The GA's advantage over greedy is minimal at N = 10. The papers used N = 100, where the search space is vast and a GA is necessary.

GenFed vs. FedCSGA: Key Differences

| Aspect | GenFed | FedCSGA |
|------------|---|---|
| Stage | Post-training (Model Aggregation) | Pre-training (Client Selection) |
| Decision | Which models to aggregate | Which clients to select |
| Objective | Maximize model quality/fitness | Maximize clients under a time deadline |
| GA Use | Minimal (Fitness + Selection only) | Full (Crossover, Mutation, Selection) |
| Chromosome | N/A (simple ranking) | A sequence of clients $\langle q_1, \ldots \rangle$ |
| Fitness | Validation accuracy on \mathcal{D}^{v}_{σ} | Function of $ \mathbf{q} $, time Θ , and accuracy $A(\mathbf{q})$ |
| Key Tool | Global validation set | Time model + Adaptive GA operators |
| Solves | Data heterogeneity, Byzantine attacks | System heterogeneity, Time constraints |

Verdict

Use GenFed when...

- √ You have a public, representative validation dataset.
- ✓ **Participant reliability** is a major concern (e.g., malicious clients, stragglers with buggy code).
- √ Your main goal is fastest convergence (fewer rounds) and highest final accuracy.
- imes You have no hard per-round time deadlines.

Use FedCSGA when...

- √ You have strict time deadlines per round (e.g., real-time applications).
- ✓ **System heterogeneity** (client speeds) is the main bottleneck.
- ✓ You can accurately profile client computation (c_i) and bandwidth (B_i) .
- × You do not have a public validation set to filter models.

Hybrid Approach: Use **FedCSGA** to select the max clients for a constraint T, then use **GenFed** to filter the received models.

Conclusion & Future Work

Summary

- **GenFed:** A simple, powerful model selection filter.
 - 10-35× faster convergence (in rounds)
 - Inherent Byzantine robustness
 - Validated in our re-implementation
- **FedCSGA**: A powerful *client* selection scheduler.
 - Solves NP-hard time-constrained selection
 - Needs scale ($N \gg 10$) and fresh fitness data

Future Work

- Our Project:
 - Fix FedCSGA fitness staleness
 - Scale to N = 100 clients
 - Implement and test the Hybrid approach
- Research:
 - Multi-objective GA for FedCSGA (time, accuracy, and fairness)

Code

https://github.com/sithtsar/FedGA

References

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Thank You Questions?

Backup: GenFed ρ_t Scheduling

Goal: Find the best strategy for varying ρ_t (number of models to aggregate).

- Strategy (1): Constant $\rho_t = \rho_{\text{max}}$. (Stable, but slow start)
- Strategy (3): Linear $\rho_t = \min(\rho_{\text{max}} \cdot t/c + 1, \rho_{\text{max}})$
- Strategy (4): Sinusoidal $(\pi/2)$ $\rho_t = \min(\rho_{\sf max} \cdot \sin(t\pi/2c) + 1, \rho_{\sf max})$
- Strategy (5): Sinusoidal (π) (Unstable)

Result: Strategies (3) and (4) performed best.

- Intuition: Start with small ρ_t (e.g., $\rho_t = 1$) to aggregate only the *very* best model, allowing for rapid exploration.
- ullet As training progresses, *increase* ho_t to include more models, which provides stability and fine-tuning.

Backup: FedCSGA α Parameter Impact

Non-IID Fitness Function: $h(\mathbf{q}) = \sum_{j=1}^{|\mathbf{q}|} (1 - \alpha \cdot A(q_j))$

Table: Final Accuracy vs. α (Non-IID, T = 5min)

| α | 0.0 | 0.4 | 0.7 | 1.0 |
|---------------------------|----------------|----------------|-----|-----|
| Fashion-MNIST CIFAR-10 | 63.5% 49.1% | 65.2% 52.2% | | |

Analysis:

- $\alpha = 0$: Maximize clients only (same as IID). Ignores data heterogeneity.
- $\alpha=1$: Only cares about accuracy. Selects too few clients, hurting diversification.
- $\alpha = 0.7$: **Optimal trade-off.** Balances selecting *many* clients (for diversity) with prioritizing clients that *need* training (low-accuracy clients).