

## [fictional Work]

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# Dynamic Pruning Meets Gene Networks: A Biologically Inspired AI Model for Emotion Recognition and Cultural Adaptation

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### Abstract

Emotion recognition and cultural adaptation are critical challenges in advancing artificial intelligence (AI) systems toward greater human-like understanding and interaction. Inspired by the dynamic control mechanisms of gene regulatory networks, we propose a novel AI framework that integrates *dynamic pruning* with biologically inspired gene network dynamics. By embedding “gene-like regulatory behaviors,” our model adapts its neural architecture dynamically, mirroring the biological processes of gene expression control. This novel approach bridges biology and AI, demonstrating how biological dynamic control principles can enhance AI adaptability and efficiency.

We validate our framework through simulations on standard emotion recognition datasets (e.g., FER-2013, AffectNet) and achieve state-of-the-art accuracy of up to 92% while reducing computational costs by 50%. The proposed model further demonstrates enhanced adaptability to multi-cultural datasets, providing context-sensitive emotional insights. This work highlights the potential of integrating biological principles into AI for creating more efficient, adaptive, and culturally aware systems.

## 1 Introduction

Artificial intelligence systems that can recognize human emotions and adapt to cultural contexts are critical for advancing human-machine interactions. These systems find applications in healthcare, education, customer service, and beyond. However, existing emotion recognition models face several limitations:

1. **Static Architectures:** Current models are often fixed in structure and struggle to adapt to dynamic changes in tasks or environments.
2. **Lack of Cultural Awareness:** Many models fail to generalize across diverse cultural datasets, leading to biased outcomes.
3. **Computational Inefficiency:** As models grow in size, their computational and memory requirements become impractical for real-world scenarios.

To address these challenges, this study draws inspiration from the dynamic regulatory processes of gene networks. Previous pruning approaches, such as gradient-based pruning [1] and deep compression [2], primarily rely on static criteria and thus lack the adaptability needed for dynamically changing tasks like emotion recognition. In contrast, biological gene networks exhibit remarkable adaptability, controlling gene expression through context-sensitive interactions [3]. Based on these insights, our work integrates gene-like regulatory behaviors into AI models, aiming to enhance adaptability, efficiency, and cultural awareness.

This paper introduces a novel framework that combines gene network-inspired dynamics with a dynamic pruning algorithm. The proposed method dynamically optimizes the network structure during training, maintaining high performance while significantly reducing computational overhead. We demonstrate the efficacy of our approach through experiments on standard and multicultural emotion recognition datasets.

## 2 Related Work

### 2.1 Emotion Recognition

Deep learning has revolutionized emotion recognition with models analyzing visual, auditory, and textual inputs. Public datasets such as FER-2013 and AffectNet have enabled significant advancements, yet they still suffer from cultural biases due to limited diversity in training data [4]. This motivates research into more adaptive architectures.

### 2.2 Dynamic Pruning

Pruning techniques reduce network complexity by removing redundant parameters. For example, gradient-based pruning [1] calculates per-weight importance during training, and deep compression [2] combines pruning, quantization, and Huffman coding. However, these methods typically use static thresholds, and do not account for dynamic changes in input distributions—particularly crucial for tasks like emotion recognition, which may involve culturally diverse or evolving data.

## 2.3 Gene Network Dynamics

In biological systems, gene regulatory networks control gene expression through dynamic, context-dependent interactions. These networks can be modeled using Boolean networks for discrete on/off states or systems of differential equations to capture continuous changes [5]. Such models highlight how dynamic feedback and adaptive thresholds can yield robust, context-aware behavior. Our approach translates these principles to neural network training, allowing the model to prune and rewire connections adaptively.

## 3 Proposed Method

### 3.1 Overview

Our dynamic pruning framework is inspired by gene regulatory networks, combining:

1. **Gene-like Regulatory Behaviors:** Adaptive thresholding and rewiring, mirroring gene expression control.
2. **Dynamic Pruning & Reconstruction:** Periodically removing and re-adding network connections based on real-time importance scores and rewiring probabilities.

By integrating these components, our model aims to maintain high accuracy while significantly reducing computational overhead, even under culturally diverse or rapidly changing inputs.

### 3.2 Mathematical Model

#### 3.2.1 Gene Network Dynamics

We draw an analogy to biological gene expression, where mRNA ( $m_i$ ) and protein ( $p_i$ ) levels evolve over time:

$$\frac{dm_i}{dt} = f_i(m, p) - \gamma_m m_i, \quad \frac{dp_i}{dt} = g_i(m) - \gamma_p p_i, \quad (1)$$

where  $f_i$  and  $g_i$  represent regulatory interactions and  $\gamma_m, \gamma_p$  are decay rates. These dynamics inspire how we update connection importance and decide to prune or restore weights.

#### 3.2.2 Dynamic Pruning

We define an importance score  $S_{ij}$  for each weight  $w_{ij}$  as:

$$S_{ij} = |w_{ij}| + \alpha \cdot I_{ij} + \beta \cdot \text{Time}_i, \quad (2)$$

where  $I_{ij}$  may represent a gradient-based saliency metric and  $\text{Time}_i$  accounts for training progress. Connections with  $S_{ij} < \tau$  are pruned, where the threshold  $\tau$  is dynamically adjusted during training.

### 3.2.3 Reconstruction (Rewiring)

Pruned connections may be restored based on a rewiring probability  $P_{ij}$ :

$$P_{ij} = \exp\left(-\frac{\Delta_{ij}}{\sigma}\right), \quad (3)$$

where  $\Delta_{ij}$  represents the "importance difference" for potential connections and  $\sigma$  controls the scale. Over time,  $\sigma$  is gradually decreased to reduce new connection formation, mirroring biological homeostasis.

## 3.3 Enhancing Reproducibility: Dynamic Threshold $\tau$ and Rewiring Probability $P_{ij}$

### Dynamic Threshold $\tau$ Setting:

- **Initial Value:** Set as  $\tau = \mu + 0.5\sigma$ , where  $\mu$  and  $\sigma$  are the mean and standard deviation of initial weight importance scores.
- **Adaptive Updates:** At each epoch's end, recalculate the distribution of remaining weights and adjust  $\tau$  to the median score if necessary.
- **Convergence:** Once accuracy stabilizes over several epochs,  $\tau$  remains fixed.

### Rewiring Probability $P_{ij}$ Adjustment:

- **Initial Setting:** For a pruned connection candidate  $(i, j)$ ,  $P_{ij}$  is computed as in Eq. (3).
- **Normalization:** Normalize probabilities so that  $\sum_{i,j} P_{ij} = 1$ .
- **Time Decay:** Gradually decrease  $\sigma$  across epochs to limit excessive rewiring during later stages.

## 4 Algorithm

The overall procedure is as follows:

- Step 1. Initialize** network weights  $\mathbf{W}$  and gene-inspired parameters ( $\gamma, \sigma, \alpha, \beta$ , etc.).
- Step 2. Train** the model in mini-batches, updating weights via standard backpropagation (e.g., Adam).
- Step 3. Compute** importance scores  $\{S_{ij}\}$  each epoch.
- Step 4. Prune** weights for which  $S_{ij} < \tau$ .

**Step 5. Rewire** pruned connections based on probability  $P_{ij}$ .

**Step 6. Update**  $\tau$  and  $\sigma$  as described.

**Step 7. Repeat** Steps 2–6 until convergence or maximum epochs.

## 5 Experiments

### 5.1 Datasets

We evaluate our model on several emotion recognition datasets:

- **FER-2013:** Labeled facial emotion data, achieving 92% accuracy (4% improvement over a static baseline of  $\sim 88\%$ ).
- **AffectNet:** A culturally diverse dataset, with average accuracy of 91.5% (improving over the baseline by 4–5%).
- **RAVDESS:** An audio-based emotion dataset, with accuracy around 90% (baseline  $\sim 85\text{--}86\%$ ).

### 5.2 Implementation Details

- **Learning Rate:** 0.001
- **Batch Size:** 64
- **Epochs:** 50
- **Optimizer:** Adam
- **Data Preprocessing:** Normalization to  $[0, 1]$ , data augmentation (random rotations  $\pm 15^\circ$ , flips, brightness adjustments  $\pm 20\%$ ), and ROI cropping using OpenCV/Dlib.

### 5.3 Evaluation Metrics

We consider:

1. **Accuracy:** Classification accuracy across test sets.
2. **Efficiency:** Parameter reduction, inference time, and memory footprint.
3. **Adaptability:** Consistency of performance across culturally diverse subsets and dynamic input conditions.

## 6 Results and Discussion

### 6.1 Performance

- **FER-2013:** 92% accuracy (+4% vs. baseline).
- **AffectNet:** 91.5% accuracy ( +5% vs. baseline), highlighting adaptability to cultural diversity.
- **RAVDESS:** 90% accuracy ( +4–5% vs. baseline) on audio-based emotions.

### 6.2 Efficiency

- Model size reduced by 50% without sacrificing accuracy.
- Inference time decreased by 35%, making the approach suitable for real-time or resource-limited environments.

### 6.3 Cultural Adaptation

- Average performance improvement of about +5% for underrepresented cultural subsets.
- Experiments on AffectNet subsets (e.g., Asian vs. African facial expressions) consistently yield > 90% accuracy, demonstrating robust generalization.

### 6.4 Illustrative Case Study

- **Microexpressions:** Dynamic pruning and rewiring retain subtle features (e.g., East Asian microexpressions), enabling fine-grained recognition.
- **Reduced Bias:** The adaptive rewiring mechanism helps mitigate biases typically observed in static architectures.

### 6.5 Limitations

- **Dependence on High-Quality Data:** Performance may degrade with noisy or unrepresentative datasets.
- **Computational Complexity:** While pruning reduces parameters, the rewiring mechanism introduces additional computational overhead, which might require further optimization for large-scale deployments.

## 7 Additional Insights on Cultural Adaptation

To assess cross-cultural adaptability, we generated synthetic subsets from AffectNet focusing on culturally salient features:

- **Consistent Accuracy:** The model maintains  $> 90\%$  accuracy across diverse subsets.
- **Dynamic Feature Adjustment:** Importance-based pruning selectively preserves culturally significant features, reducing bias.
- **Future Integration:** Incorporating audio, text, and physiological signals may further enhance cultural adaptability.

## 8 Application to Non-Emotion Tasks

Our gene-inspired dynamic pruning approach is applicable to other domains:

### 1. Medical AI:

- *Imaging Diagnostics:* Efficient pruning for high-dimensional medical images.
- *Genomic Analysis:* Modeling gene interactions using the gene-network perspective.

### 2. Time-Series Data Analysis:

- *Financial Forecasting:* Adapting quickly to volatile market patterns.
- *IoT Sensor Streams:* Lightweight models for continuous sensor data.

### 3. Robotics and Control Systems:

- *Adaptive Control:* Real-time reconfiguration of robotic movements.
- *Fault Detection:* Rewiring to focus on critical sensor inputs.

### 4. Education Technology:

- *Personalized Learning:* Adjusting to student performance patterns.
- *Language Processing:* Pruning for domain-specific vocabularies in real-time transcription.

## 9 Conclusion

We proposed a novel AI model that integrates dynamic pruning with gene network-inspired dynamics to address the limitations of static architectures, cultural biases, and computational

inefficiencies in emotion recognition. By mirroring the adaptability of biological gene regulation, our model achieves high accuracy while significantly reducing model size and inference time. This interdisciplinary fusion of biology and AI opens avenues for more efficient, adaptive, and culturally aware intelligent systems with potential applications across healthcare, robotics, education, and beyond.

## 10 Research Highlights

### 1. Biological Inspiration:

The dynamic pruning process, inspired by gene regulatory mechanisms, offers a context-sensitive alternative to static pruning methods.

### 2. Interdisciplinary Innovation:

By integrating concepts from biology and AI, the framework provides a blueprint for constructing adaptive, efficient neural architectures.

## 11 Future Directions

### 1. Extended Applications:

Investigate domains such as time-series analysis, autonomous robotics, and genomics.

### 2. Experimental Refinement:

Conduct longitudinal and multi-modal experiments (combining audio, text, and physiological signals) to further validate cross-cultural emotional understanding.

### 3. Ethical and Societal Considerations:

Ensure that dynamic, adaptive emotion recognition systems address privacy, bias, and ethical implications when deployed at scale.

---

## Appendix: Simplified Dynamic Pruning Implementation

```
1 # -----  
2 # FICTIONAL CODE FOR DEMONSTRATION PURPOSES ONLY  
3 # This simplified implementation demonstrates the core concepts of  
4 # dynamic pruning inspired by gene networks as described in the paper  
5 # -----  
6  
7 import torch
```



```

8 import torch.nn as nn
9 import torch.optim as optim
10 import numpy as np
11 import matplotlib.pyplot as plt
12 from torchvision import datasets, transforms
13 from torch.utils.data import DataLoader
14
15 # -----
16 # 1. Model Definition with Gene-inspired Dynamics
17 # -----
18 class GeneInspiredNet(nn.Module):
19     def __init__(self, input_size, hidden_sizes, output_size):
20         super(GeneInspiredNet, self).__init__()
21         self.layers = nn.ModuleList()
22
23         # Create layers based on sizes
24         layer_sizes = [input_size] + hidden_sizes + [output_size]
25         for i in range(len(layer_sizes) - 1):
26             self.layers.append(nn.Linear(layer_sizes[i], layer_sizes[
27                 i+1]))
28
29         # Gene-inspired parameters
30         self.importance_scores = {} # Store importance scores for
31             each weight
32         self.pruning_mask = {} # Binary masks for pruning
33         self.rewiring_candidates = {} # Potential connections to
34             restore
35
36         # Initialize importance tracking
37         for i, layer in enumerate(self.layers):
38             self.importance_scores[i] = torch.ones_like(layer.weight)
39             self.pruning_mask[i] = torch.ones_like(layer.weight)
40             self.rewiring_candidates[i] = set() # Empty set
41                 initially
42
43     def forward(self, x):
44         for i, layer in enumerate(self.layers):
45             # Apply the pruning mask during forward pass
46             masked_weight = layer.weight * self.pruning_mask[i]
47             x = F.linear(x, masked_weight, layer.bias)
48
49             # Apply activation function (except for output layer)

```

```

46         if i < len(self.layers) - 1:
47             x = torch.relu(x)
48         return x
49
50     # -----
51     # 2. Dynamic Pruning Functions
52     # -----
53     def compute_importance_scores(model, alpha=0.5, beta=0.2, time_factor
54                                   =0.1):
55         """Update importance scores using gene-inspired formula"""
56         for i, layer in enumerate(model.layers):
57             weight_magnitude = torch.abs(layer.weight.data)
58             gradient_info = torch.abs(layer.weight.grad) if layer.weight.
59                 grad is not None else torch.zeros_like(layer.weight)
60
61             # Gene-inspired importance score from Eq. 2
62             model.importance_scores[i] = (
63                 weight_magnitude +
64                 alpha * gradient_info +
65                 beta * time_factor
66             )
67
68     def dynamic_pruning(model, current_epoch, prune_rate=0.3, sigma=1.0):
69         """Prune connections with low importance scores"""
70         for i, layer in enumerate(model.layers):
71             # Calculate dynamic threshold (simplified from paper)
72             scores = model.importance_scores[i]
73             mu = torch.mean(scores)
74             std = torch.std(scores)
75             threshold = mu + 0.5 * std
76
77             # Create pruning mask based on threshold
78             new_mask = (scores >= threshold).float()
79
80             # Store pruned connections as rewiring candidates
81             pruned_indices = torch.nonzero((model.pruning_mask[i] -
82                 new_mask) > 0)
83             for idx in pruned_indices:
84                 model.rewiring_candidates[i].add((idx[0].item(), idx[1].
85                     item()))
86
87             # Update the pruning mask

```

```

84     model.pruning_mask[i] = new_mask
85
86 def rewiring(model, sigma=1.0, max_reconnections=100):
87     """Restore some pruned connections based on rewiring probability
88     """
89     for i, layer in enumerate(model.layers):
90         if not model.rewiring_candidates[i]:
91             continue
92
93     # For each candidate, calculate rewiring probability (Eq. 3)
94     candidates = list(model.rewiring_candidates[i])
95     probabilities = []
96
97     for idx in candidates:
98         row, col = idx
99         # Calculate delta (simplified version)
100         importance = model.importance_scores[i][row, col].item()
101         mu = torch.mean(model.importance_scores[i]).item()
102         delta = abs(importance - mu)
103
104         # Eq. 3: Probability of rewiring
105         prob = np.exp(-delta / sigma)
106         probabilities.append(prob)
107
108     # Normalize probabilities
109     probabilities = np.array(probabilities) / sum(probabilities)
110
111     # Select connections to rewire
112     num_reconnect = min(max_reconnections, len(candidates))
113     if num_reconnect > 0:
114         reconnect_idx = np.random.choice(
115             len(candidates),
116             size=num_reconnect,
117             replace=False,
118             p=probabilities
119         )
120
121     # Rewire selected connections
122     for idx in reconnect_idx:
123         row, col = candidates[idx]
124         model.pruning_mask[i][row, col] = 1.0
125         model.rewiring_candidates[i].remove((row, col))

```

```

125
126 # -----
127 # 3. Training Loop with Dynamic Pruning
128 # -----
129 def train_with_dynamic_pruning(model, train_loader, valid_loader,
130                                epochs=50, lr=0.001, device="cpu"):
131     criterion = nn.CrossEntropyLoss()
132     optimizer = optim.Adam(model.parameters(), lr=lr)
133
134     # Tracking metrics
135     train_losses = []
136     valid_accs = []
137     pruning_rates = []
138
139     for epoch in range(epochs):
140         model.train()
141         running_loss = 0.0
142
143         for inputs, labels in train_loader:
144             inputs, labels = inputs.to(device), labels.to(device)
145             inputs = inputs.view(inputs.size(0), -1) # Flatten for
146                                                         simplicity
147
148             # Forward pass
149             optimizer.zero_grad()
150             outputs = model(inputs)
151             loss = criterion(outputs, labels)
152
153             # Backward pass
154             loss.backward()
155             optimizer.step()
156
157             running_loss += loss.item()
158
159         # Update importance scores based on current weights and
160         # gradients
161         compute_importance_scores(model, time_factor=epoch/epochs)
162
163         # Apply dynamic pruning and rewiring
164         if epoch > 5: # Start pruning after initial training
165             dynamic_pruning(model, epoch)

```

```

165         # Gradually decrease sigma to reduce rewiring over time
166         sigma = max(0.5, 2.0 * (1.0 - epoch/epochs))
167         rewiring(model, sigma=sigma)
168
169     # Validation
170     model.eval()
171     correct = 0
172     total = 0
173     with torch.no_grad():
174         for inputs, labels in valid_loader:
175             inputs, labels = inputs.to(device), labels.to(device)
176             inputs = inputs.view(inputs.size(0), -1)
177             outputs = model(inputs)
178             _, predicted = torch.max(outputs.data, 1)
179             total += labels.size(0)
180             correct += (predicted == labels).sum().item()
181
182     # Calculate metrics
183     train_loss = running_loss / len(train_loader)
184     valid_acc = 100 * correct / total
185
186     # Calculate pruning rate (percentage of zeros)
187     total_weights = 0
188     pruned_weights = 0
189     for i, layer in enumerate(model.layers):
190         total_weights += model.pruning_mask[i].numel()
191         pruned_weights += (model.pruning_mask[i] == 0).sum().item()
192     pruning_rate = 100 * pruned_weights / total_weights
193
194     # Store metrics
195     train_losses.append(train_loss)
196     valid_accs.append(valid_acc)
197     pruning_rates.append(pruning_rate)
198
199     print(f"Epoch {epoch+1}/{epochs}, Loss: {train_loss:.4f}, "
200           f"Validation Acc: {valid_acc:.2f}%, Pruning Rate: {pruning_rate:.2f}%")
201
202     return train_losses, valid_accs, pruning_rates
203
204     # -----

```

```

205 # 4. Demonstration with a Simple Emotion Dataset
206 # -----
207 def demo_emotion_recognition():
208     """Simplified demo with MNIST (as a stand-in for emotion data)"""
209     # Load MNIST (as a simplified proxy for emotion recognition)
210     transform = transforms.Compose([
211         transforms.ToTensor(),
212         transforms.Normalize((0.1307,), (0.3081,))
213     ])
214
215     train_dataset = datasets.MNIST('./data', train=True, download=
        True, transform=transform)
216     test_dataset = datasets.MNIST('./data', train=False, transform=
        transform)
217
218     train_loader = DataLoader(train_dataset, batch_size=64, shuffle=
        True)
219     test_loader = DataLoader(test_dataset, batch_size=64, shuffle=
        False)
220
221     # Create and train model
222     input_size = 28 * 28 # MNIST image size
223     hidden_sizes = [128, 64]
224     output_size = 10 # 10 digits (replace with emotion classes for
        real implementation)
225
226     device = torch.device("cuda" if torch.cuda.is_available() else "
        cpu")
227     model = GeneInspiredNet(input_size, hidden_sizes, output_size).to
        (device)
228
229     # Train with dynamic pruning
230     train_losses, valid_accs, pruning_rates =
        train_with_dynamic_pruning(
231         model, train_loader, test_loader, epochs=20, device=device
232     )
233
234     # Plot results
235     plt.figure(figsize=(15, 5))
236
237     plt.subplot(1, 3, 1)
238     plt.plot(train_losses)

```

```

239 plt.title('Training Loss')
240 plt.xlabel('Epoch')
241 plt.ylabel('Loss')
242
243 plt.subplot(1, 3, 2)
244 plt.plot(valid_accs)
245 plt.title('Validation Accuracy')
246 plt.xlabel('Epoch')
247 plt.ylabel('Accuracy (%)')
248
249 plt.subplot(1, 3, 3)
250 plt.plot(pruning_rates)
251 plt.title('Pruning Rate')
252 plt.xlabel('Epoch')
253 plt.ylabel('Pruned Weights (%)')
254
255 plt.tight_layout()
256 plt.show()
257
258 return model, (train_losses, valid_accs, pruning_rates)
259
260 # -----
261 # Run the demonstration if called directly
262 # -----
263 if __name__ == "__main__":
264     import torch.nn.functional as F
265     print("Starting Gene-Inspired Dynamic Pruning Demo")
266     model, metrics = demo_emotion_recognition()
267     print("Demo completed")

```

Listing 1: Dynamic Pruning Implementation (Simplified for Demonstration)

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