NEAT

Evolving Neural Networks through Augmenting Topologies

Fall 2018

Slides by: Aref Moqadam Mehr

Intuition

- Easy to understand
- Easy to trace
- GPU friendly



Image From: http://medium.com/

Intuition

- More Optimized
- Fewer calculations
- Reinforcement Learning application



Basic Idea

- 1. Select an Empty Network
- 2. Randomly add Connections
- 3. Randomly mutate Connections
- 4. Optimize via Genetics Algorithms



By the way, Why?

- Can evolving topologies along with weights provide an advantage over evolving weights on a fixed-topology?
- A fully connected network can in principle approximate any continuous function.
- So why waste valuable effort permuting over different topologies?

Encoding

- TWEANNs Encoding
- Binary Encoding
- Graph Encoding
- Indirect Encoding

Problems

- Mating between different genes.
- Initial Populations
- Protecting Speciation





Genetic Encoding

| Genome (Genotype) | | | | | | | | | | | |
|---|---|-------------|---|------------------|---|--------------|---|---|---|--|--|
| Node Genes | Node 1 Sensor | Nod Sen: | e 2 sor | Node 3 Sensor | Node 4 Output | Node Hidd | e 5 len | | | | |
| Connect. Genes | In 1 Out 4 Weight 0.7 Enabled Innov 1 | | In 2 Out 4 Weight-0.5 DISABLED Innov 2 | | In 3 Out 4 Weight 0.5 Enabled Innov 3 | | In 2 Out 5 Weight 0.2 Enabled Innov 4 | In 5 Out 4 Weight 0.4 Enabled Innov 5 | In 1 Out 5 Weight 0.6 Enabled Innov 6 | In 4 Out 5 Weight 0.6 Enabled Innov 11 | |
| Network (Phenotype) 4 1 2 3 3 | | | | | | | | | | | |

Mutation

- Add Connection
- Add Node



Tracking Genes through Historical Markings

- When Structural Mutation Happens
- Global Innovation Number incrementally increases.
- Crossover within same GIN
- Crossover with a Gene with different GIN



Speciation

- δ: distance of different structures
- E: the number of excess genes
- D: the number of disjoint genes
- W: the average weight matching genes (including disabled genes)

$$\delta = \frac{c_1 E}{N} + \frac{c_2 D}{N} + c_3 \cdot \overline{W}.$$

Explicit Fitness Sharing

- Organism in same niches share same fitness.
- a species cannot afford to become too big even if many of its organisms perform well.
- sh = 0 : if $\delta(i, j) < \delta t$ sh = 1 : otherwise

$$f'_i = \frac{f_i}{\sum_{j=1}^n \operatorname{sh}(\delta(i,j))}.$$

Minimal Solution

- Minimizing Dimensionality through Incremental Growth from Minimal Structure
- only those structures survive that are found to be useful through fitness evaluations

Performance Evaluations

- Can NEAT evolve the necessary structures?
- Can NEAT find solutions more efficiently than other Neuro-Evolution systems?

Performance Evaluations

- Evolving XORs
- Pole balancing



Bench-marks

- Pole Balancing as a Benchmark Task
- Pole Balancing Comparisons
- Double Pole Balancing with Velocities

| Method | Evaluations | Generations | No. Nets |
|-----------------|-------------|-------------|----------|
| Ev. Programming | 307,200 | 150 | 2048 |
| Conventional NE | 80,000 | 800 | 100 |
| SANE | 12,600 | 63 | 200 |
| ESP | 3,800 | 19 | 200 |
| NEAT | 3,600 | 24 | 150 |

Analysis of NEAT

| Method | Evaluations | Failure Rate |
|-----------------------------------|-------------|--------------|
| No-Growth NEAT (Fixed-Topologies) | 30,239 | 80% |
| Nonspeciated NEAT | 25,600 | 25% |
| Initial Random NEAT | 23,033 | 5% |
| Nonmating NEAT | 5,557 | 0 |
| Full NEAT | 3,600 | 0 |

Conclusion

 Evolving topology along with weights



Any Question?

- Paper: Evolving Neural Networks through Augmenting Topologies, K. O. Stanley, et.al., 2006, MIT Press Journal
- Find this presentation online: <u>https://arefmq.github.io/</u> <u>downloads/NEAT-Presntation.pdf</u>

Link of Video: <u>https://www.youtube.com/watch?</u> <u>v=qv6UVOQ0F44&t=31s</u>

Find me on web: <u>https://arefmq.github.io/</u>

Read More: <u>https://en.wikipedia.org/wiki/</u> <u>Neuroevolution of augmenting topologies</u>