GP TEST ONE 2010-08-25

Objective: Find program whose output matches x^2+x+1 over the range $-1 \le x \le 1$.

Function Set: +, -, *, % (protected division).

Terminal Set: X, and 1

Fitness: Sum for absolute errors of x (-1.0, -0.9, ...0.9, 1.0)

Selection: Fitness proportionate non elitist.

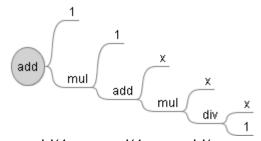
Initial Ramped

Population:

Parameters: Population size: 100

Termination: 100 generations.

Best Individual: Run One:



 $gp_add(1, gp_mul(1, gp_add(x, gp_mul(x, gp_div(x, 1)))))$

Run Two:

$$gp_add(gp_mul(gp_add(x, 1), x), 1)$$

(((x + 1) * x) + 1)

Run Three:

/**Same as Run One**/

Run Four:

Run Five:

Run Six:

$$gp_add(gp_mul(x, gp_add(gp_mul(x, 1), 1)), 1)$$

((x * ((x * 1) + 1)) + 1)

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Code: from pyevolve import *
import math
error accum = Util.ErrorAccumulator()
def gp add(a, b):
   return a+b
def qp sub(a,b):
   return a-b
def gp mul(a, b):
   return a*b
def gp div(a,b):
   "Safe" division, if divide by 0, return 1.
   if b == 0:
      return 1.0
   else:
      return a/(b*1.0)
def rangef(min, max, step):
   result = []
   while 1:
      result.append(min)
      min = min + step
      if min>=max:
         break
   return result
def eval func (chromosome):
   global error accum
   error accum.reset()
   code_comp = chromosome.getCompiledCode()
   for x in rangef(-1, 1, .1):
      evaluated = eval(code comp)
      target = x**2 + x + 1
      error accum += (target, evaluated)
   return error accum.getRMSE()
def main run():
   genome = GTree.GTreeGP()
   genome.setParams(max depth=5, method="ramped")
   genome.evaluator.set(eval func)
   ga = GSimpleGA.GSimpleGA(genome)
   ga.setParams(gp_terminals = ['x', '1'],
                     gp_function_prefix = "gp")
   ga.setMinimax(Consts.minimaxType["minimize"])
   ga.setGenerations(100)
   ga.setMutationRate(0.08)
   ga.setCrossoverRate(1.0)
   ga.setPopulationSize(100)
   ga.evolve(freq stats=5)
   print ga.bestIndividual()
if name == " main ":
   main run()
```

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