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class EvoAmp:
         def init (self, population tot, crossover rate, mutation rate, generations, N, d, target):
             self.population tot = population tot
             self.crossover_rate = crossover_rate
             self.mutation_rate = mutation_rate
             self.generations = generations
             self.N = N
             self.target = target
             self.pop_dis = np.ndarray(shape=(self.population_tot, self.N-1))
             self.pop_pha = np.random.uniform(size = (self.population_tot, self.N), low = 0, high = 2*np.pi*0) # randians
             self.pop_amp = np.random.uniform(size=(self.population_tot, self.N), low = 0, high=1)
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             # Population with same spacing and phase -> Only optimizing amplitude
             for i in range(self.population tot):
                 self.pop_dis[i, :] = d
             self.pop_fit = np.zeros(self.population_tot, dtype = np.float)
             self.evolve()
         def results(self):
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             # Return the best solution
             q = np.argmin(self.pop_fit)
             return self.pop amp[q]
         def evolve(self):
             # probabilities for cross over. To avoid calculationg random numbers every itteration.
             crossovers = np.random.uniform(size=(self.generations), low = 0, high=1)
             mutations = np.random.uniform(size=(self.generations), low = 0, high=1)
             for i in range(self.generations):
                 for dna in range(self.population tot):
                     score = fitness(self.pop dis[dna], self.pop pha[dna], self.pop amp[dna], self.target)
                     self.pop fit[dna] = score;
                 index = np.argmin(self.pop fit)
                 self.pop amp[0] = self.pop amp[index]
                 self.pop amp[-1] = self.pop amp[index]
                 # Stop simulating when the generation matches the maximum no of generations.
                 if i == self.generations-1:
                     print('DONE')
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                 for _ in range(1):
                     if crossovers[i] < self.crossover rate:</pre>
                         s 1 = selection(self.pop fit)
                         s 2 = selection(self.pop fit)
                         self.pop\_amp[s\_1] = (self.pop\_amp[s\_1] + self.pop\_amp[s\_2])/2
                     if mutations[i] < self.mutation rate:</pre>
                         # mutation occurs
                         s 1 = selection(self.pop fit)
                         self.pop_amp[s_1][np.random.randint(low=0, high=self.N)] = np.random.uniform(low = 0, high=1)
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import numpy as np
from functions import *