EVOLUTIONARY AND FUZZY SYSTEMS

COURSEWORK 2

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TASK 1

GENETIC IMPLEMENTATION OF THE INPUT MEMBERSHIP FUNCTION:

Genetic algorithm (GA) is a search technique that takes inspiration from natural selection and genetics. It updates a population of solutions to acquire a global optimum. The new solutions generated by the GA, also referred to as offspring, from the parental solutions by applying the crossover and mutation operation. Said operations should be applied in such a way that offspring inherit important factors of parents.

In this report, the GA is applied on the initial Fuzzy System design of the Ambient Assisted Living. The temperature and humidity system has been taken into consideration; the genotypes are implemented from its membership functions. Please observe the below-mentioned implementation.

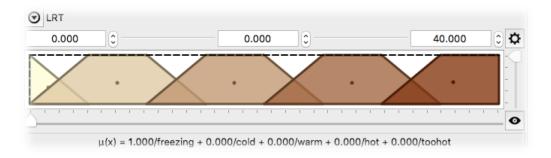


Figure 1: Living Room Temperature

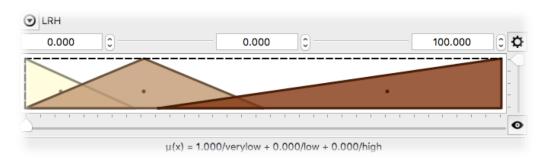


Figure 2: Living Room Humidity

5	5	10	15	10	15	20	25	20	25	30	35	30	35

Table 1: Genotype of The Temperature Membership Functions (pop1)

6	8	11	17	13	17	22	27	22	28	33	37	34	38
	Table 2: Population Increment (pop2)												
3	9	9	18	15	16	21	26	23	27	31	36	35	39
				Ta	ible 3: Po	opulatio	n Increm	ent (pop	3)				
2	7	8	19	13	18	23	28	23	26	30	34	35	37

Table 4: Population Increment (pop4)

Algorithm Structure:

As the selection operation, the Minimal Generation Gap (MGG). The GA incorporating the MCG. The operation is as follows:

1. Create the initial population, and set $g \leftarrow 1$. (g = generation).

Created the population of 4 members (genotypes), pop1, pop2, pop3, and pop4.

The first gene in the genotype (pop1) represents the first initiation point (crisp value) of the first membership function unless it's as same as the primary initiator (the starting point of the universe of discourse) of the particular MF. However, if it is the same then the next distinct turning point of that MF is considered to be the first gene, we move on to the next MF and store its initiator, the turning points, and the end of the membership function.

The same implementation can be applied to the other inputs, however, there will be very small changes. Since we are using trapezoidal MFs in the current implementation, there are 4 genes representing each point of the trapezoid but the triangular MFs would probably generate 3 genes and similarly, Gaussian MFs would produce 2 genes (representing its height and the curve's width).

2. Chose 2 solutions, say par1 and par2 randomly from the created population as parents.

Picked pop1 and pop3 as par1 and par2, respectively.

3. Create two further solutions par3 and par4 using the crossover operation. This operation is applied to every generation. Picked non-similar (distinct) genes from the par1 and par2 and generated the following genotypes.

3	25	15	9	30	9	30	18	20	16	25	21	35	26

Table 5: par3

	5	5	10	15	10	15	21	21	26	23	27	31	36	39
L														

Table 6: par4

- 4. Derive a solution *par5* from *par1* using the mutation operation. Do the same to generate solution *par6* from *par2*. This operation is also applied to every generation with the probability of *P*.
- 5. From the set of generated solutions: {par1, par2, ···, par6} select 2 best possible solutions. Remove par1 and par2 (parents) from the population, and replace with the best possible selected solutions in the population.

As mentioned in the task description, there are n examples of the mapping (Xi, Yi), to expect output Yi for each Xi. Mean Square Error (MSE) was employed as a fitness function to evaluate the accuracy between the given defuzzified output (Y) and the genetically generated output (Y') of each population member.

$$MSE = 1/n \sum_{k=1}^{n} (Yik - Y'ik)^{2}$$

- 6. Given If g=G, end the algorithm. (Where G =final generation; and its value is initially defined).
- 7. Given that $g \neq G$, increment g ($g \leftarrow g+1$) and continue from step 2.

GENETIC IMPLEMENTATION OF THE OUTPUT MEMBERSHIP FUNCTION:

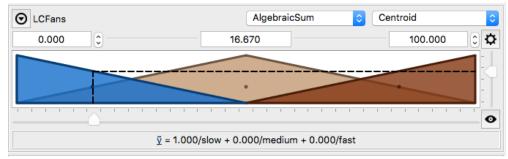


Figure 3: Cooling Fans (fuzzy output)

H/T - Cooling	freezing	cold	warm	hot	Too hot
Fans					
Very-low	No change	No change	slow	medium	fast
low	No change	slow	medium	fast	fast
high	No change	slow	fast	fast	fast

Table 7: Matrix- Temperature and Humidity output

Every output of each input membership function's combination represents each gene in the genotype. Please observe the below-mentioned genetic value assignation and the genotype.

Output Membership Function	Genetic value
No change	0
slow	1
Medium	2
fast	3

Table 7: Genetic values of each MF

0	0	1	2	3	0	1	2	3	3	0	1	3	3	3

Table 8: Genotype of the cooling fan output (pop1)

The above-mentioned genotype is generated by taking the first combination's rule base output, from the matrix table, as the first gene in genotype, similarly, the second gene is generated by the second the next input combination (row-wise), and so on. The same algorithm structure that is employed by the input MFs genetic population, would be applied to the output genotype. There are n examples of the combination and their mapped rules (Xi, Yi), representing the combination (Xi) and its consecutive rule (Yi). The same fitness function (MSE) would be applied to assess the accuracy of the current rule base and the predicted rule base.

HYBRID IMPLEMENTATION:

The above-mentioned Fuzzy Logic Control System is implemented by the hybrid functionalities of the evolutionary computing. The MFs are represented by the classes in hybrid systems, which consecutively produces a probable output of the distinct generated combinations with the respect of Q, the rule-base is developed by the continues reductions and increments of the size (space on the universe of discourse) of MFs.

The same fitness function, that is used in the first question, is employed to extract the changes in the accuracy of the probable rule base, in each iteration of the increment and decrement with the mutation probability of P_u . The probability policy is updated by the count of the combination classes representing membership functions as sets, the update in the probability is represented by u.

Task 2

The employed optimization techniques on the used benchmark functions, Rastrigin and Sphere, are Particle Swarm Optimization (PSO), Genetic Algorithm (GA), and Differential Evolution (DE). The optimization experiment is implemented in Python and certain libraries, such as PyEvolve and SwarmPackagePy, were adopted.

However, not all the optimizers and functions were available as pre-defined methods, for which the methods had to be created from scratch for some of the optimizations' implementations.

RASTRIGIN FUNCTION:

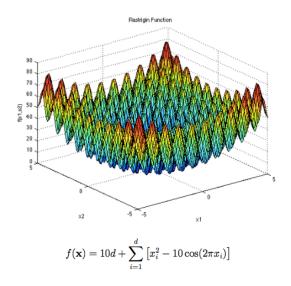


Figure 4: Rastrigin Function

$$f(\mathbf{x}) = An + \sum_{i=1}^n \left[x_i^2 - A\cos(2\pi x_i)
ight]$$

Dimensions: d; Global minimum at: x = 0; Function of x: f(x) = 0

In other way:

$$f(\mathbf{x}^*) = 0$$
, at $\mathbf{x}^* = (0, \dots, 0)$

The Rastrigin function (non-convex function) is exercised for optimization algorithms as a performance analysis problem. It contains various local minima, and it is a multimodal function with non-linearity. It was first introduced as a 2-dimensional problem, however, later upgraded to be used as a multi-dimensional performance scale. It is quite to difficult to optimize this function because of it having an extensive search space.

SPHERE FUNCTION:

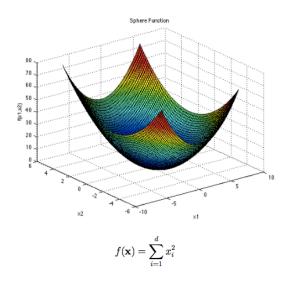


Figure 5: Sphere Function

Dimensions: d; Global minimum: $f(\mathbf{x}^*) = 0$, at $\mathbf{x}^* = (0, ..., 0)$

The Sphere function contains local minima in multi-dimensional, however, its global minima lacks multi-dimensionality. The sphere function is convex, unimodal, continuous, and it produces twofold flat areas for the problem.

OPTIMIZATION:

Swarm Particle Optimization on Rastrigin Function-2-Dimensions:

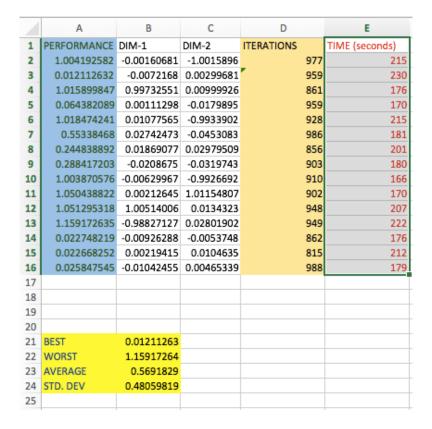


Figure 6: SPO on Rastrigin-2-D

Swarm Particle Optimization on Rastrigin Function-10-Dimensions:

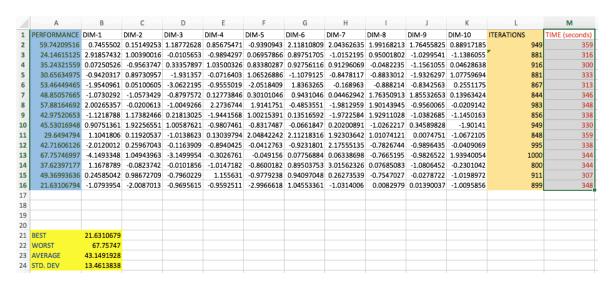


Figure 7: SPO on Rastrigin-10-D

Differential Evolution on Rastrigin Function-2-Dimensions:

	Α	В	С	D	E
1	PERFORMANCE	DIM-1	DIM-2	ITERATIONS	TIME (seconds)
2	8.847069706	-2.60742214	-1.4312301	932	155
3	8.847069706	-4.80395713	2.73234694	961	199
4	5.140096766	-1.60313704	-1.603137	858	197
5	0.127745934	-0.08740604	-0.3465633	961	209
6	8.847069706	4.220687413	-2.5837963	826	195
7	5.140096766	4.050964728	3.42059323	879	199
8	0.127745934	-5.90068888	-5.8062172	813	189
9	8.847069706	-3.98088785	1.02487408	926	224
10	5.140096766	-3.11981156	2.75717971	915	154
11	0.127745934	-3.11981156	2.75717971	907	189
12	6.585744794	-2.52660504	-1.1182457	817	201
13	8.847069706	-1.05323074	-2.3401816	885	222
14	5.140096766	3.063906155	3.41642919	992	186
15	0.127745934	1.101649384	-5.7585351	887	169
16	4.770919102	0.503390167	-2.55975	902	220
17					
18					
19					
20					
21					
22	BEST	0.127745934			
23	WORST	8.847069706			
24	AVERAGE	5.110892215			
25	STD. DEV	3.469207256			

Figure 8: DE on Rastrigin-2-D

Differential Evolution on Rastrigin Function-10-Dimensions:

	Α	В	C	D	Е	F	G	н		J	K	L	M
1	PERFORMANCE	DIM-1	DIM-2	DIM-3	DIM-4	DIM-5	DIM-6	DIM-7	DIM-8	DIM-9	DIM-10	ITERATIONS	TIME (seconds)
2	130.8344353	1.97469293	2.0397964	3.36103054	-4.8185669	-1.6290748	2.57606268	-4.5063734	-5.5876368	-3.8422993	-3.5604505	923	338
3	101.9256436	0.22789955	-1.1760037	-4.5542214	5.80835256	1.10131726	1.10131726	4.21670953	1.99005756	4.65811644	-0.3847234	815	312
4	109.4465424	-3.4342784	-0.5129366	-5.4725421	-3.9720492	-1.5890964	0.44586041	3.37646259	1.68963424	5.80318616	-1.0034571	817	321
5	112.3744508	-1.9554116	-4.3061299	2.64818168	-3.7598188	-1.73354	-2.5013733	-4.1918966	-4.8544498	-2.0552954	-3.7729312	815	313
6	112.1907942	1.27100072	-1.8990895	-5.1199023	-1.6871479	-3.0677096	-1.5732467	4.96470558	4.74099447	3.32295095	2.80279553	954	306
7	94.44079632	3.37206109	3.37206109	2.46248568	5.31675122	-3.9404475	-1.0185174	3.11143624	1.15518141	1.1785364	2.89803271	941	349
8	130.8344353	4.10253867	-2.4998395	-5.6687808	5.83712566	4.98511816	0.34291741	4.46527617	3.78291626	-4.9026733	5.72241283	818	351
9	101.9256436	-4.7354331	-0.8214756	3.52166839	-5.8630522	1.72743795	1.41879715	-2.7980326	-4.2450934	-3.102583	3.77576147	957	320
10	107.4454036	-2.4823708	0.56184674	-2.0525742	1.8224971	-0.9166007	-5.8209749	5.78163547	-1.6916259	4.63775984	0.95466279	880	312
11	109.4465424	-4.3605712	1.38079341	-1.8337172	3.0288714	-2.8832798	0.11438546	-0.9307129	-4.4015638	-5.2363862	4.69753351	844	302
12	112.3744508	-4.9090954	-0.5735374	2.74605799	-2.1078902	-1.4751279	-2.3068021	5.48732259	-5.7457803	-5.6627132	0.41749508	963	339
13	112.1907942	5.74672586	-4.4040517	3.43792366	-1.5139241	4.27410306	5.80474567	-0.9999863	-0.7075368	4.93092844	3.47115989	943	317
14	117.1305335	-0.7410481	-4.6596414	-4.4845706	-4.0425095	1.74806926	3.76725207	5.85682069	5.80637023	5.2676639	-0.9655178	951	359
15	111.0248688	-0.3068435	-4.953961	5.71093444	-4.7997597	1.6536106	-2.36435	0.77890405	-0.932229	1.60546853	4.28470182	991	311
16	93.94481935	0.36107743	4.27146367	-3.3209099	-2.2836741	3.86185999	-0.9707687	-0.9707687	0.76725157	4.33051527	4.81504381	953	333
17													
18													
19													
20													
21													
22	BEST	93.9448194											
23	WORST	130.834435											
24	AVERAGE	110.50201											
25	STD. DEV	9.32980055											

Figure 9: DE on Rastrigin-10-D

Genetic Algorithm on Rastrigin Function-2-Dimensions:

	Α	В	С	D	E
1	PERFORMANCE	DIM-1	DIM-2	ITERATIONS	TIME (seconds)
2	1.958874	0.00176782	-0.0012622	816	197
3	2.486346	-0.0030262	-0.0037378	890	218
4	1.129793	-0.0032606	-0.0037211	955	217
5	2.777357	-0.001082	-0.0010592	890	185
6	4.13996	0.00053875	2.64E-05	822	170
7	1.769212	0.00236801	0.00289143	919	195
8	3.46181	-0.0022134	0.00337133	851	202
9	3.313474	-0.0008425	0.00399817	939	213
10	2.623246	-0.0011595	0.00277111	834	213
11	2.62E+00	4.93E-05	0.00116189	911	224
12	3.23E+00	-2.40E-05	-0.0015999	823	156
13	2.32E+00	0.00093021	0.00076735	843	200
14	4.61E+00	0.00074259	-0.0022746	997	226
15	5.52E+00	0.00251189	0.00249516	913	157
16	4.49E+00	0.00307271	-9.81E-05	876	219
17					
18					
19					
20					
21					
22	BEST	1.129793			
23	WORST	5.519475			
24	AVERAGE	3.0955934			
25	STD. DEV	1.19132217			

Figure 10: GA on Rastrigin-2-D

Genetic Algorithm on Rastrigin Function-10-Dimensions:

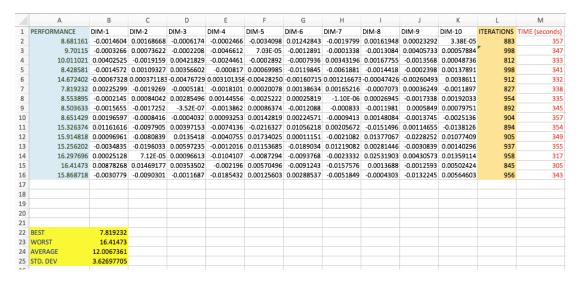


Figure 11: GA on Rastrigin-10-D

Particle Swarm Optimization on Sphere Function-2-Dimensions:

	Α	В	С	D	E
1	PERFORMAN	DIM-1	DIM-2	ITERATIONS	TIME (seconds)
2	0.01033317	-0.0141447	0.0031242	951	204
3	0.01033317	-0.0141447	0.0031242	816	227
4	0.00453538	0.00626378	0.00125962	819	226
5	0.00453538	0.00626378	0.00125962	816	170
6	0.00453538	0.00626378	0.00125962	859	189
7	0.01033317	-0.0141447	0.0031242	951	189
8	0.00453538	0.00626378	0.00125962	910	177
9	0.00141418	-0.0019975	-6.32E-06	874	160
10	0.00453538	0.00626378	0.00125962	974	185
11	0.00141418	-0.0019975	-6.32E-06	859	174
12	0.00141418	-0.0019975	-6.32E-06	835	180
13	0.00453538	0.00626378	0.00125962	928	229
14	0.00141418	-0.0019975	-6.32E-06	969	169
15	0.00453538	0.00626378	0.00125962	980	186
16	0.00141418	-0.0019975	-6.32E-06	915	176
17					
18					
19					
20					
21					
22	BEST	0.00141418			
23	WORST	0.01033317			
24	AVERAGE	0.00465454			
25	STD. DEV	0.00297152			
20					

Figure 12: PSO on Sphere-2-D

Particle Swarm Optimization on Sphere Function-10-Dimensions:

	A	В	C	D	Е	F	G	Н	1	J	K	L		M
1	PERFORMANCE	DIM-1	DIM-2	DIM-3	DIM-4	DIM-5	DIM-6	DIM-7	DIM-8	DIM-9	DIM-10	ITERATIONS		TIME (seconds)
2	-52.3261573	-0.2886796	-0.4110168	-1.2807931	-0.8328798	-0.0760306	-0.2470847	0.28790082	-1.280172	-0.0636256	-0.0015219		893	314
3	-77.03102941	-0.3522932	-0.5212038	0.40919157	-0.204329	1.13048629	-0.054802	-0.1213445	-0.7217496	-0.4226325	-0.1240379		935	325
4	-77.03102941	-0.3522932	-0.5212038	0.40919157	-0.204329	1.13048629	-0.054802	-0.1213445	-0.7217496	-0.4226325	-0.1240379		947	354
5	-119.0253656	0.1920824	0.1681016	-0.0322747	0.12873423	0.01382064	0.35666254	0.30758713	-0.1228919	-0.6629622	-0.1894011		935	300
6	-119.0253656	0.1920824	0.1681016	-0.0322747	0.12873423	0.01382064	0.35666254	0.30758713	-0.1228919	-0.6629622	-0.1894011		858	303
7	-77.03102941	-0.3522932	-0.5212038	0.40919157	-0.204329	1.13048629	-0.054802	-0.1213445	-0.7217496	-0.4226325	-0.1240379		951	319
8	-119.0253656	0.1920824	0.1681016	-0.0322747	0.12873423	0.01382064	0.35666254	0.30758713	-0.1228919	-0.6629622	-0.1894011		949	331
9	-119.0253656	0.1920824	0.1681016	-0.0322747	0.12873423	0.01382064	0.35666254	0.30758713	-0.1228919	-0.6629622	-0.1894011		810	307
10	-52.3261573	-0.2886796	-0.4110168	-1.2807931	-0.8328798	-0.0760306	-0.2470847	0.28790082	-1.280172	-0.0636256	-0.0015219		880	351
11	-77.03102941	-0.3522932	-0.5212038	0.40919157	-0.204329	1.13048629	-0.054802	-0.1213445	-0.7217496	-0.4226325	-0.1240379		998	321
12	-119.0253656	0.1920824	0.1681016	-0.0322747	0.12873423	0.01382064	0.35666254	0.30758713	-0.1228919	-0.6629622	-0.1894011		845	351
13	-52.3261573	-0.2886796	-0.4110168	-1.2807931	-0.8328798	-0.0760306	-0.2470847	0.28790082	-1.280172	-0.0636256	-0.0015219		861	334
14	-77.03102941	-0.3522932	-0.5212038	0.40919157	-0.204329	1.13048629	-0.054802	-0.1213445	-0.7217496	-0.4226325	-0.1240379		853	315
15	-119.0253656	0.1920824	0.1681016	-0.0322747	0.12873423	0.01382064	0.35666254	0.30758713	-0.1228919	-0.6629622	-0.1894011		894	344
16	-77.03102941	-0.3522932	-0.5212038	0.40919157	-0.204329	1.13048629	-0.054802	-0.1213445	-0.7217496	-0.4226325	-0.1240379		877	319
17														
18														
19														
20														
21														
22	BEST	-119.02537												
23	WORST	-52.326157												
24	AVERAGE	-88.887789												
25	STD. DEV	27.1285253												

Figure 13: PSO on Sphere-10-D

Differential Evolution on Sphere Function-2-Dimensions:

	Α	В	С	D	E
1	PERFORMANCE	DIM-1	DIM-2	ITERATIONS	TIME (seconds)
2	0.007636838	0.07418719	-0.0461855	958	176
3	0.004316778	-0.0467298	-0.0461855	991	220
4	0.004316778	-0.0467298	-0.0461855	970	150
5	0.004316778	-0.0467298	-0.0461855	991	171
6	0.004316778	-0.0467298	-0.0461855	854	215
7	0.001533246	0.02590879	-0.0293595	997	223
8	0.001533246	0.02590879	-0.0293595	913	220
9	0.001533246	0.02590879	-2.94E-02	976	223
10	0.001533246	0.02590879	-0.0293595	896	183
11	8.58E-05	0.00191364	-9.07E-03	890	175
12	8.58E-05	0.00191364	-9.07E-03	883	193
13	8.58E-05	0.00191364	-0.0090656	987	225
14	8.58E-05	0.00191364	-9.07E-03	998	177
15	5.04E-05	0.00315999	-0.0063579	898	209
16	5.90E-06	0.00224791	-9.21E-04	915	161
17					
18					
19					
20					
21					
22	BEST	5.9015E-06			
23	WORST	0.00763684			
24	AVERAGE	0.00209578			
25	STD. DEV	0.00182955			

Figure 14: DE on Sphere-2-D

Differential Evolution on Sphere Function-10-Dimensions:

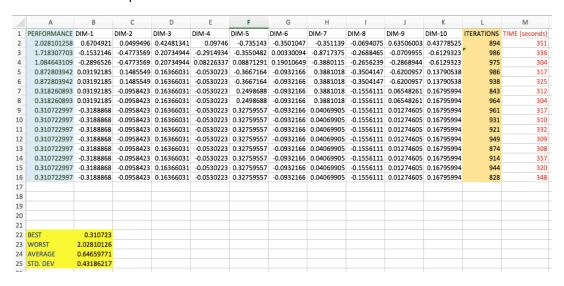


Figure 15: DE on Sphere-10-D

Genetic Algorithm on Sphere Function-2-Dimensions:

	Α	В	С	D	E
1	PERFORMAN	DIM-1	DIM-2	ITERATIONS	TIME (seconds)
2	0.006031	0.02731402	0.00314857	918	186
3	0.098876	0.00325733	-0.0086354	872	166
4	0.040693	0.00094705	0.00090878	812	180
5	0.171818	0.00090562	0.00032712	872	167
6	0.19003	-0.0009501	-1.77E-03	981	208
7	0.138657	0.00278058	0.00156539	813	174
8	0.080647	-0.0010343	4.64E-05	873	190
9	0.188584	0.00054731	-0.0006641	1000	194
10	0.143247	0.00028613	3.81E-06	856	219
11	2.68E-01	-3.13E-04	-4.40E-05	974	154
12	2.13E-01	-2.75E-04	-0.0001502	926	176
13	1.60E-01	-0.0002437	-5.12E-05	956	207
14	4.08E-01	-0.0005358	0.00121683	974	151
15	3.50E-01	-0.0001549	-0.0003624	804	221
16	2.52E-01	6.41E-05	5.45E-04	863	221
17					
18					
19					
20					
21					
22	BEST	0.006031			
23	WORST	0.407679			
24	AVERAGE	0.180567			
25	STD. DEV	0.10068446			

Figure 16: GA on Sphere-2-D

Genetic Algorithm on Sphere Function-10-Dimensions:

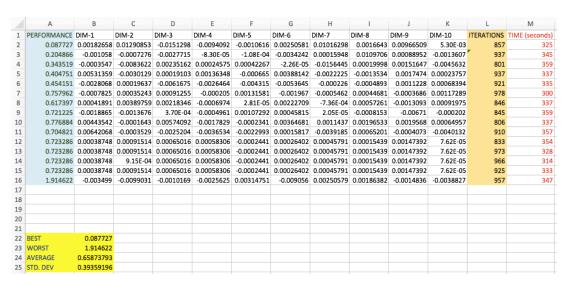


Figure 17: GA on Sphere-10-D

RESULT ANALYSIS:

	Α	В	С	D	Е	F	G	Н	1
1	perf/opt	PSO-rast-2D	DE-rast-2D	GA-rast-2D		perf/opt	PSO-sph-2D	DE-sph-2D	GA-sph-2D
2	BEST	0.01211263	0.12774593	1.129793		BEST	0.00141418	5.9015E-06	0.006031
3	WORST	1.15917264	8.84706971	5.519475		WORST	0.01033317	0.00763684	0.407679
4	AVERAGE	0.5691829	5.11089222	3.0955934		AVERAGE	0.00465454	0.00209578	0.180567
5	STD. DEV	0.48059819	3.46920726	1.19132217		STD. DEV	0.00297152	0.00182955	0.10068446
6		BEST	WORST					BEST	WORST
7									
8	perf/opt	PSO-rast-100	DE-rast-10D	GA-rast-10D		perf/opt	PSO-sph-10D	DE-sph-10D	GA-sph-10D
9	BEST	21.6310679	93.9448194	7.819232		BEST	-119.02537	0.310723	0.087727
10	WORST	67.75747	130.834435	16.41473		WORST	-52.326157	2.02810126	1.914622
11	AVERAGE	43.1491928	110.50201	12.0067361		AVERAGE	-88.887789	0.64659771	0.65873793
12	STD. DEV	13.4613838	9.32980055	3.62697705		STD. DEV	27.1285253	0.43186217	0.39359196
13			WORST	BEST			BEST	WORST	
14									

Figure 18: Result Analysis

The code and methods' implementation is provided in *Appendix-A (SPO-Rastrigin)*, *Appendix-B (DE-Rastrigin)*, *Appendix-C (GA-Rastrigin)*, *Appendix-D (SPO-Sphere)*, *Appendix-E (DE-Sphere)*, and *Appendix-F (GA-Sphere)*. After executing each optimization algorithm for *30-50* generations, on Rastrigin and Sphere function, the below-mentioned analysis was composed.

RASTRIGIN FUNCTION:

Best Performance on the Rastrigin Function-2-D: Particle Swarm Optimization performed really well with the fitness of 0.01211263 (Standard Dev: 0.48059819).

Best Performance on the Rastrigin Function-10-D: Genetic Algorithm performed really well with the fitness of 7.819232 (Standard Dev: 3.62697705).

Worst Performance on the Rastrigin Function-2-D: Differential Evolution performed really bad with the fitness of 8.84706971 (Standard Dev: 3.46920726).

Worst Performance on the Rastrigin Function-10-D: Differential Evolution performed really bad with the fitness of 130.834435 (Standard Dev: 9.32980055).

SPHERE FUNCTION:

Best Performance on the Sphere Function-2-D: Differential Evolution performed really well with the fitness of 5.9015E-06 (Standard Dev: 0.00182955).

Best Performance on the Sphere Function-10-D: Particle Swarm Optimization performed really well with the fitness of -119.02537 (Standard Dev: 27.1285253).

Worst Performance on the Sphere Function-2-D: Genetic Algorithm performed really bad with the fitness of 0.407679 (Standard Dev: 0.10068446).

Worst Performance on the Sphere Function-10-D: Differential Evolution performed really bad with the fitness of 2.02810126 (Standard Dev: 0.43186217).

Task 3

BIN PACKING-1-DIMENSIONAL:

Let bin capacity be C, N is the number of bins, M is the set of items, where items = $\{i1, i2, iM\}$, and $\{S1, S2, \dots, SM\}$ is the integer sizes.

Definition of the problem:

The bin packing problem (BPP) is defined as follows:

Given a non-negative number C and a set of M items of integer size S1, the bin packing problem refers to packing given items into the bins such that the total size of items in any given bin doesn't exceed the capacity C and the number of bins N used is kept to a minimum.

Complexity Class Membership:

This problem resides in NP-Hard complexity class.

Existing Solutions:

The existing solutions to the BPP include the first-fit decreasing method (FFD) and best-fit decreasing method (BFD) which are popular heuristic methods. The computation of these methods works in such a way that the items are sorted in descending order, and then the methods applied for the items in turn.

Implementation:

The implemented Genetic Algorithm (GA) the set of item numbers (brick sizes) in a bin was made to act as a "gene". The genotype is expressed by sequencing the item sets for all the bins. For instance, a situation where In = 15, an example genotype g1 would look as follows:

$$g_1: (1,3,10)(2,9,11)(5,7,13,15)(4,6,14)(8,12)$$

The above *g1* means that (In1, In3, In10), (In2, In9, In11), (In5, In7, In13, In15), (In4, In6, In14) and (In8, In12) are assigned to B1, B2, B3, B4, and B5, respectively. The number of genes is variable.

Even if the order of the genes is changed, the genotype decodes to the same solution. For example, a genotype

$$g_2: (8,12)(2,9,11)(5,7,13,15)(4,6,14)(1,3,10)$$

The above genotype shows the exact same genotype with the same set of genes as in g1 but in a variable order, despite this, they produce the same solution.

The solution is independent of the order of the genes like in the case of sets.

Solution Proposal:

Existing solutions in GA are random generations specifically aimed at optimization problems. To obtain better initial solutions, the FF (First-Fit) was incorporated into the GA. The items are first sequenced, and then the FF applied for each item. The sequencing is done at random in this generation process to generate unprobable, unexpected solutions.

Crossover:

This operation is designed in a way that offspring inherit important factors of the parents. An imperative aspect of BPP is the set of items. To incur a better solution, local optimization and heuristic rules such as FF (First-Fit) and MBS (Minimum Bin Slack)' should be introduced to the crossover operation of the GA.

In this crossover operation, offspring c1 and c2 are generated from parents P1 and P2. Initially, a few bins (genes) are randomly chosen from P1 and the items inside them are replicated in c1, then from P2, only those bins are picked which don't contain the same items as P1's bins that are replicated in c1. The remaining sets of P2 are replicated to St (temporary set), excluding the items similar to the items of the P1's sets that were copied to c1. The sets in St are not replicated in c1 because the quantity of the bricks (items) in St is quite less. The St sets are divided into two sets, St_A and St_B, St_A contains all the sets which just have one item in them and rest of the sets are stored in St_B.

Mutation:

This procedure works in such a way that 2 to 3 bins (gene equivalents) in the parent are randomly chosen. The items in the chosen bins are copied to *Stm* individually, that is, as item-sets containing a single item.

The remaining bins are copied to the offspring. Then, the replacement procedure between the offspring and *Stm* is executed such that is uses in the crossover operation. Lastly, a two-phase procedure is executed to get the items remaining in the *Stm* to be used in the crossover operation.

Loss Function:

$$\sum_{i=0}^{n} Ri + S$$

R: Remaining size of the ith bin

S: Total size of the bin (new bin)

The solution should be considered good if the output of above-mentioned function gets smaller with each iteration. To calculate the Ri, the below-mentioned formula is used.

$$Ri = B - \sum_{i=0}^{n} Oi$$

B: Current size of the particular bin

O: ith Brick's (item) size

Note: As long as $O \ge B \ge 0$, new bin won't be added

The proposed fitness function should work in each situation since the number of bins doesn't affect the solution, it recommences with each bin-packing iteration. For each bin, the remaining size is calculated after a new bin is created and added to the size of the new bin, which, as a result, should give an approximate sum of total remaining space.

BIN PACKING-2-DIMENSIONAL:

The same implementation can be applied to the 2-D BPP, however, some changes need to be made with the current genotypes. The purpose is to pack 2D bins (rectangles) with random 2D shapes while avoiding extensions amongst the shapes and large unused spaces on rectangular bounds.

Instead of a simple item number (brick sizes), the sets should contain the coordinates of the bricks with the consideration of (0,0) bottom-left-corner, so the figure is definable with respect to three parameters, coordinates (x, y) and orientation angle (Q). Furthermore, the mutation should be enhanced in such a way that it considers the distance between shapes and their corresponding sizes.

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APPENDICES

APPENDIX-A

```
# Created By Huzefa Shaikh
           import sys
import random
import math
import numpy as np
9 dim = 2
10 perfRes = []
11 #Particle-count-in-Swarm
 # Bounds-on-positions-velocities
v_max = 20
 16 v_min = -20
17 p_min = -32
18 p_max = 32
# updates_no
cmax = 1000

# amt-of-dampen-velocity-on-updates
dampener = 1
dampen_rate = 1

# Leaving original variables distinct
orig_dampen_rate = dampen_rate
orig_dampener = dampener

# function-to-optimize (minimize)
           def F(x):
   global dim
   D = dim
               summation = 0
               i = 0
summation = D*10
```

```
for x in range(15):
 global cmax, dampener, dampen_rate, dim
 dampen_rate = orig_dampen_rate
 dampener = orig_dampener
 swarm = []
 while i < particle_no:</pre>
    swarm.append(Particle())
 best_pos = []
 worst_pos = []
 all_pos = []
 AvgX = 0

AvgXc = []
 best_velocity = []
 best_err = -1
 worst_err = -1
nerr = []
 # Run-updates-swarm-and-output-best-position/error
 while i <= cmax:</pre>
     err = swarm[j].Evaluate()
      if err < best_err or best_err == -1:</pre>
       best_pos = []
        best_velocity = []
        while k < dim:
          best_pos.append(swarm[j].pos[len(swarm[j].pos)-1][k])
          best_velocity.append(swarm[j].velocity[len(swarm[j].velocity)-1][k])
        best_err = err
```

```
swarm[j].UpdateVelocity(best_pos)
        swarm[j].UpdatePosition()
     dampener = dampener * dampen_rate # Dampen the velocity
  # Output-stats
  perfRes.append(best_err)
  print best_err, best_pos
velocity: The list of particle velocities
nRess = np.array(perfRes)
print '\n0ver all best performance: ', min(perfRes)
print '\n0ver all worst performance: ', max(perfRes)
print '\nAverage performance: ', np.mean(nRess)
def __init__(self):
    global dim
```

```
self.best_pos = []
self.best_err = -1 # this is set to -1 so we update after the first step
self.pos = []
temp_pos = []
temp_velocity = []
  temp_pos.append(self.InitPosition())
   temp_velocity.append(self.InitVelocity())
   self.best_pos.append(0) # initialize the best position array
self.velocity.append(temp_velocity)
global dim
# The F function that we are trying to minimize self.err = F(self.pos[len(self.pos)-1]) if self.best_err == -1 or self.err < self.best_err: self.first_update = False self.best_err = self.err
   self.best_pos = []
     self.best_pos.append(self.pos[len(self.pos)-1][j])
if temp > p_max:
  return p_min
return temp
```

```
## Initialize the velocity of the particle between 1 and -1
## for each dimension

def Initvelocity(self):
    if random.random()
    return random.random()
    return random.random()

## A function that is used to randomize the cognitive term

def RandomizeCognitive(self):
    return random.random()

## A function that is used to randomize the social term

def RandomizeSocial(self):
    return random.random()

## A function that is used to update the velocity

## A function that is used to update the velocity

## of the particle the particle's past and the global best position seen

def UpdateVelocity(self, global_best_pos):
    global v_max, dampener, dia

## us a control parameter that tells the particle

## how much to discount the previous velocity

## c1 is a control parameter that tells the particle

## how much to weight its own previous positions

c1 = 2

## c2 is a control parameter that tells the particle

## how much to veight the swarms best best position

c2 = 2

## r1 and r2 are random numbers that weight the

## cognitive and social terms

r1 = self.RandomizeSocial()

t = len(self.velocity)

## Construct the new velocity for the particle

new_velocity_arr = []
    j = 0

while j < din:

## Apply the control parameters to the particle's previous velocity

## in the direction that we are working on

v_term = demonraces.

"t-rem = demonraces.

## Create the cognitive and social terms

own_term = c1 = r1 * (self.best_pos[]) = self.pos[t-1][j])

## A function that is used to random terms

own_term = c2 = r2 * (global_best_pos[]) = self.pos[t-1][j])

## Add the velocities together to make the new velocity

## A function that we remove the particle that the previous velocity

## A function that we are working on

v_term = cdmparer=wself.velocity[t-1][j]

## Add the velocities together to make the new velocity

## A function that is used to randomize the cognitive and social terms

own_term = c1 = r1 * (self.best_pos[j] - self.pos[t-1][j])

## Add the velocities together to make the new velocity
```

```
if new_velocity > v_max:
      new_velocity = v_max
    elif new_velocity < v_min:</pre>
     new_velocity = v_min
    new_velocity_arr.append(new_velocity)
  self.velocity.append(new_velocity_arr)
def UpdatePosition(self):
  global p_max, p_min, dim
  t1 = len(self.velocity)
  t2 = len(self.pos)
  new_position_arr = []
    new_position = self.pos[t2-1][j] + self.velocity[t1-1][j]
# If the position is smaller or larger than the bounds, change them
    if new_position > p_max:
      new_position = p_max
    elif new_position < p_min:</pre>
     new_position = p_min
    new_position_arr.append(new_position)
  self.pos.append(new_position_arr)
```

Appendix-B

```
import math
import numpy as np
i
```

```
max_gen = input("Enter the number of runs::
dim=input("Enter the dimension of the problem:: ")
    for i in xrange(0,dim): print "Enter the lower and upper bound of %d th variable" %i
l = input()
u = input()
for i in range(0,dim):
    Xl.insert(i,l)
# Open the file to store the best individual of every generation
f_best=open("best-population.out","w")
#f_worst=open("worst-population.out","w")
global pop,fvals,num_fe
pop=[]
     i in range
X=[]
for j in range(0,dim):
    #fill-up-X-add-population
    X.insert(j,(Xl[j] + (Xu[j]-Xl[j])*rand_n()))
       #bounds-check
for j in range(0,dim):
    while X[j] < Xl[j] or X[j] > Xu[j]:
        if X[j]-Xl[j]:
            X[j]=2*Xl[j]-X[j]
        if X[j]>Xu[j]:
            X[j]=2*Xu[j]-X[j]
```

```
#BOUNDS-CHECK
for k in range(0,dim):

while U[k] < X1[k] or U[k] > Xu[k]:

if U[k]<X1[k]:

U[k]=2*X1[k]-U[k]

if U[k]>Xu[k]:

U[k]=2*Xu[k]-U[k]

U.insert(dim,func(U)) #the function value (the-last-value)

#SELECTION
#Comparing the trial vector and past individual
if U[dim] <= fvals[j]:

for k in range(0,dim):
 (pop[j])[k]=U[k]

fvals.insert(j,func(pop[j]))
```

```
#Find the best objective func. value and write it to the file
    best_val=fvals[0]
    worst_val=fvals[0]
    best_index=0
for i in range(0,NP):
        if fvals[i] < best_val:</pre>
             best_index=i
            best_val=fvals[i]
            best_index=i
             worst_val=fvals[i]
    f_best.write(str(best_val))
    f_best.write('\n')
#Report the best pop and save the population
    for i in range(0,NP):
        for j in range(0,dim):
    f.write(str((pop[i])[j]) + '\t')
    best_val=fvals[0]
    best_index=0
    worst_index=0
    popss = []
    for i in range(0,NP):
        popss.append(pop[i])
        results.append(fvals[i])
        if fvals[i] < best_val:</pre>
             best_index=i
             best_val=fvals[i]
```

```
for wi in range(0,NP):
    if fvals[wi] > best_val:
        worst_index=wi
        best_val=fvals[wi]

nres = np.array(results)

print results,',',popss
    print '\nBest : ',fvals[best_index],'--:--',pop[best_index]
    print '\nworst : ',fvals[worst_index],'--:--',pop[worst_index]
    print '\naverage : ',np.mean(nres)

print 'function-evaluations-total-number : ', num_fe

print("Differential-Evolution->>>")
    print("Differential-Evolution->>>")
    print("Evolution-process-running..")
    evolve_de_rand_1()
    print("\nSUMMARY")
    print("\nSUMMARY")
    print("\nSUMMARY")
    print("\nSUMMARY")
    print("\nSUMMARY")
    print("\nSUMMARY")
    report()
```

Appendix-C

```
Created by Huzefa Shaikh
from pyevolve import Mutators, Initializators from pyevolve import Selectors import math from pyevolve import GSimpleGA from pyevolve import GDList from pyevolve import Consts import numpy as np #Rastrigin function intitiation def rast(pen):
     total_ret = 0
len_genome = len(gen)
     for i in range(len_genome):
   total_ret += gen[i]**2 - 10*math.cos(2*math.pi*gen[i])
return (10*len_genome) + total_ret
   results = []
for x in range(15):
     gen.setParams(rangemin=-5.2, rangemax=5.30, bestrawscore=0.00, rounddecimal=2) gen.initializator.set(Initializators.G1DListInitializatorReal)
     gen_Algo = GSimpleGA.GSimpleGA(gen)
     gen_Algo.terminationCriteria.set(GSimpleGA.RawScoreCriteria)
     gen_Algo.setMinimax(Consts.minimaxType["minimize"])
     gen_Algo.setCrossoverRate(0.8)
gen_Algo.setPopulationSize(50)
     gen_Algo.setMutationRate(lr)
lr += 0.01
     gen_Algo.evolve(freq_stats=40)
     #print 'Run: ',x,'\n'
exec_best = gen_Algo.bestIndividual()
print exec_best
     results.append(exec_best)
   minn = min(results)
   print 'Best Performacne: ', minn
print 'Worst Performacne: ', maxx
print 'Average of all the performance', meann
     __name__ == "__main__":
algorith_run()
```

Appendix-D

Appendix-E

Appendix-F

```
# created by Huzefa Shaikh
from pyevolve import G1DList
from pyevolve import Mutators, Initializators
from pyevolve import G5impleGA, Consts
import numpy as np
     genome = G1DList.G1DList(2)
     genome.setParams(rangemin=-5.12, rangemax=5.13)
genome.initializator.set(Initializators.G1DListInitializatorReal)
     ga = GSimpleGA.GSimpleGA(genome, seed=666)
ga.setMinimax(Consts.minimaxType["minimize"])
      lr += 0.01
     ga.evolve(freq_stats=100)
     #print 'Run: ',x,'\n'
best = ga.bestIndividual()
      results.append(best)
   meann = np.mean(nresult)
```

```
print 'Best Performacne: ', minn
print 'Worst Performacne: ', maxx
print 'Average of all the performance', meann
return
if __name__ == "__main__":
    run_main()
```