

# Synchronous Migration in an Island Model of Differential Evolution

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## Definition

- restricted area of solutions:

$$\Omega = [a_1, b_1] \times [a_2, b_2] \times \dots \times [a_D, b_D], \quad a_j < b_j, \quad j = 1, 2, \dots, D$$

- objective function:

$$f: \Omega \rightarrow \mathbb{R}, \quad D \subseteq \mathbb{R}^D$$

- the global minimum is point  $\mathbf{x}^*$  satisfying condition:

$$\forall \mathbf{x} \in \Omega: f(\mathbf{x}^*) \leq f(\mathbf{x})$$

## Description

- population-based stochastic algorithm heuristically explores the area of possible solutions  $\Omega$
- possible solutions are represented as vectors:  
 $\mathbf{x}_i = \{x_{i,1}, x_{i,2}, \dots, x_{i,D}\}$  and create population of individuals  
 $P = \{\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N\}$
- population of possible solutions is developed in  $\Omega$  by evolutionary operators in order to find solution of task  $\mathbf{x}^*$
- from old population of *parents* a new population of *offsprings* is created and into new generation of population is selected better from couple  $\{\textit{parent}, \textit{offspring}\}$
- evolutionary operators: *mutation, crossover, selection and migration*

## Parameters of DE algorithm

- type of mutation and variant of crossover  $\rightarrow$  *strategy*
  - ▶ eg. *DE/rand/1/bin*
- control parameter of mutation  $F \in (0, 2)$ 
  - ▶ determines length of shift of parent in mutation  $\rightarrow$  *mutant vector*
- control parameter of crossover  $CR \in \langle 0, 1 \rangle$ 
  - ▶ determines probability of selection elements from parent or mutant vector  $\rightarrow$  *offspring*
- population size  $N$ 
  - ▶ bigger population  $\rightarrow$  more detailed exploring of solution area
  - ▶ smaller population  $\rightarrow$  faster searching process

**adaptive DE** is able to adapt values of the control parameters during search process leading to increase efficiency of optimization

## Motivation for using parallelism in optimization

- solving of hard tasks requires large time demands
- distribution time demands of optimization algorithm among several parallel computational units
  - ▶ length of whole parallel algorithm is equivalent to length of the slowest unit
- parallel models distribute computations among several independent processes  $p$ , enable to migrate information,  $p \in \langle 2, N \rangle$
- *master-slave, neighbourhood, **migration**, hybrid, hierarchic*

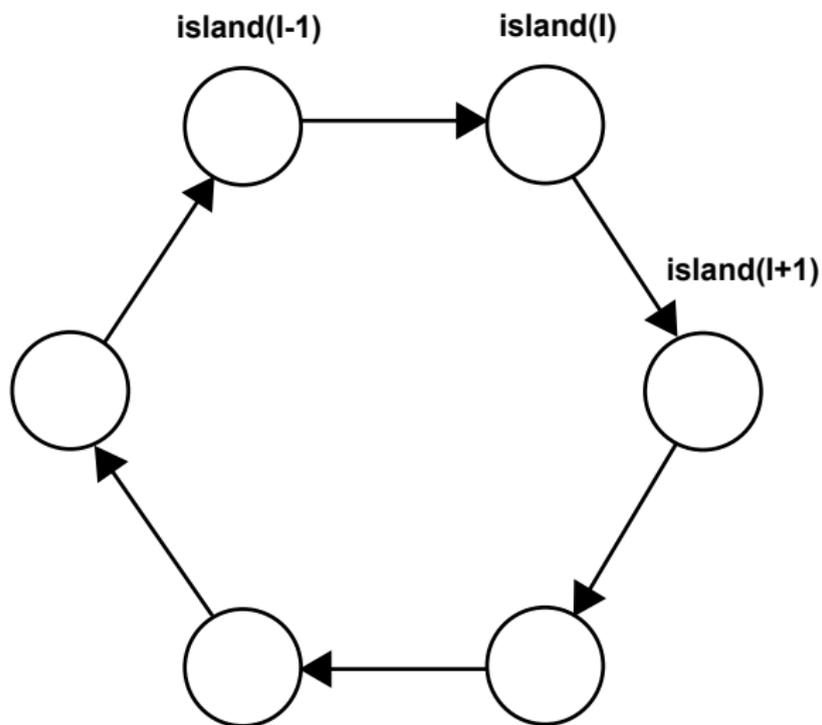
## Model description, operation migration

- distribution population  $P$  onto  $k$  sub-populations of equal sizes and location on isolated islands
- independent development of sub-populations for several generations - epoch
- operation *migration* - moving selected individuals between islands in order to exchange information, after length of epoch
- parameters of migration model - *topology*, *granularity* -  $N_p$ , *migration rate*, *migration policy*, *migration frequency*

## Experiments description

- six state-of-the-art adaptive differential evolution algorithms are paralleled by migration model with **ring** topology
- population of 60 individuals is equidistantly distributed among 6 islands with sub-populations sizes  $N_p = 10$
- all islands are developed by one adaptive DE, after  $nde = 5$  generations synchronous migration occurs (epoch done):
  - 1 1 best and 4 randomly chosen individuals are exchanged between neighbourhood islands
  - 2 better island sends 1 + 5 and less island 1 + 3 individuals
  - 3 one half of individuals are exchanged between islands
- algorithm ends when satisfies condition:
  - ▶  $(f_{max} - f_{min}) \leq 1 \times 10^{-6}$  or  $nfe \geq (D \times 20000)$

# Experiments



## Two main directions of comparison

- all algorithms run on six well known benchmark functions
- reliability rate  $R \in \langle 0, 100 \rangle$  - number of runs satisfying  $|f_{min} - f(\mathbf{x}^*)| < 1 \times 10^{-4}$  in one hundred runs
- parallel percentage speed-up  $p_{su} \in \langle 0, 100 \rangle$ :

$$p_{su} = \frac{nfe_p - nfe_s}{nfe_s} \times 100(\%),$$

where  $nfe_p$  and  $nfe_s$  is number of function evaluation in parallel, respective sequential algorithm

- four categories of classification parallel algorithms based on  $R, p_{su}$ :
  - ▶ 1 - **fast** & **reliable**, 2 - **fast** & low reliable,
  - 3 - slow & **reliable**, 4 - slow & low reliable

# Results

Migration	Adaptive DE	Ack	Dej	Gri	Ras	Ros	Sch
1 + 4	b6e6rl	3	3	3	1	4	1
1 + 4	CoDE0	2	1	2	1	1	1
1 + 4	EPSDE	3	3	3	3	3	3
1 + 4	JADE	1	3	3	3	3	3
1 + 4	jDE	1	1	1	1	4	1
1 + 4	SaDE	4	3	4	4	4	4
1 + 5, 1 + 3	b6e6rl	3	3	3	1	4	1
1 + 5, 1 + 3	CoDE0	2	1	2	1	1	1
1 + 5, 1 + 3	EPSDE	3	3	3	3	3	3
1 + 5, 1 + 3	JADE	2	3	1	3	3	3
1 + 5, 1 + 3	jDE	2	1	1	1	4	1
1 + 5, 1 + 3	SaDE	4	3	4	4	4	3
<i>half exch</i>	b6e6rl	3	3	3	1	4	1
<i>half exch</i>	CoDE0	2	1	2	1	1	1
<i>half exch</i>	EPSDE	3	3	3	3	3	3
<i>half exch</i>	JADE	2	3	3	3	3	3
<i>half exch</i>	jDE	1	1	1	1	4	1
<i>half exch</i>	SaDE	3	3	3	4	4	3

## Brief overview of fast and reliable cases

Migration	Adaptive DE	Problem	Total
1 + 4	4/6	6/6	12×
1 + 5, 1 + 3	4/6	5/6	11×
<i>half exch</i>	3/6	6/6	11×

## Summary

- no significant difference among efficiency of 3 different migrations
- parallel migration model with ring topology increases efficiency compared to sequential algorithm variants
- no improvement of parallel model adaptive variants *SaDE* and *EPSDE*
- the major efficiency of parallel model adaptive variants *CoDE0*, *jDE* and *b6e6rl*

Future research will be focused on asynchronous migration models of adaptive DE variants.

Thank you for attention!