

Benchmarking a Variant of CMAES-APOP on the BBOB Noiseless Testbed

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Outline

- The CMAES-APOP algorithm
- A Variant of CMAES-APOP algorithm
- Numerical Experiments on the BBOB Noiseless Testbed
- Conclusion and Perspectives

The CMAES-APOP algorithm

- Adapting population size seems to be a right way in the CMA-ES to optimize multi-modal functions.
- Some approaches for adapting population size in the CMAES:
 - IPOP-CMA-ES ¹ [AH05, Ros10]: the CMA-ES is restarted with increasing population size by a factor of two whenever one of the stopping criteria is met.
 - BIPOP-CMA-ES ²: define two restart regimes: one with large populations (IPOP part), and another one with small populations. In each restart, BIPOP-CMA-ES selects the restart regime with less function evaluations used so far.

¹[AH05] A. Auger and N. Hansen, A restart cma evolution strategy with increasing population size, 2005 IEEE Congress on Evolutionary Computation, vol. 2, 2005, pp. 1769-1776.

²[Han09] N. Hansen, Benchmarking a bi-population cma-es on the bbo-2009 function testbed, Proceedings of the 11th Annual Conference Companion on Genetic and Evolutionary Computation Conference: Late Breaking Papers, GECCO 09, 2009, pp. 2389-2396.

The CMAES-APOP algorithm

- Ahrari and Shariat-Panahi ³: An adaptation strategy for the CMA-ES which used the oscillation of objective value of x_{mean} to quantify multimodality of the region under exploration.
- Nishida and Akimoto ⁴: An adaptation strategy for the CMA-ES that is based on the estimation accuracy of the natural gradient.

³[ASP15] A. Ahrari and M. Shariat-Panahi, An improved evolution strategy with adaptive population size, Optimization 64 (2015), no. 12, 2567-2586.

⁴[NA16] K. Nishida and Y. Akimoto, Population size adaptation for the cma-es based on the estimation accuracy of the natural gradient, Proceedings of the Genetic and Evolutionary Computation Conference 2016, GECCO 16, 2016, pp. 237-244

The CMAES-APOP Algorithm ⁵

Motivation

- a natural desire when solving any optimization problem
- one prospect when using larger population size to search
"We want to see the decrease of objective function"

Signal?

- We track the non-decrease of objective function (exactly, $f^{\text{med}} := \text{median}(f(\mathbf{x}_{i:\lambda}), i = 1, \dots, \mu)$ - the median of objective function of μ elite solutions in each iteration) in a slot of S successive iterations to adapt the population size in the next S successive iterations
- We do not adapt the population size in each iteration but in each slot of S iterations.
⇒ The variation of population size takes a staircase form in iterations.

⁵[NH17] D. M. Nguyen and N. Hansen, Benchmarking cmaes-apop on the bbob noiseless testbed, Proceedings of the Genetic and Evolutionary Computation Conference Companion (New York, NY, USA), GECCO 17, ACM, 2017, pp. 1756–1763.

A Variant of CMAES-APOP Algorithm

Ideas: $f^{\text{med}} := \text{median}(f(\mathbf{x}_{i:\lambda}), i = 1, \dots, \mu)$ is the 25th percentile of objective function values evaluated on λ candidate points.

⇒ What if we change the 25th percentile to the other percentiles?

Some test functions:

$$f_{\text{Rastrigin}}(\mathbf{x}) = 10n + \sum_{i=1}^n (x_i^2 - 10 \cos(2\pi x_i))$$

$$f_{\text{Schaffer}}(\mathbf{x}) = \sum_{i=1}^{n-1} (x_i^2 + x_{i+1}^2)^{0.25} [\sin^2(50(x_i^2 + x_{i+1}^2)^{0.1}) + 1]$$

$$f_{\text{Ackley}}(\mathbf{x}) = 20 - 20 \cdot \exp \left(-0.2 \sqrt{\frac{1}{n} \sum_{i=1}^n x_i^2} \right) + e - \exp \left(\frac{1}{n} \sum_{i=1}^n \cos(2\pi x_i) \right)$$

$$f_{\text{Bohachevsky}}(\mathbf{x}) = \sum_{i=1}^{n-1} (x_i^2 + 2x_{i+1}^2 - 0.3 \cos(3\pi x_i) - 0.4 \cos(4\pi x_{i+1}) + 0.7)$$

- For each function, 51 runs are conducted.
- $f_{\text{stop}} = 10^{-10}$ ($f_{\text{stop}} = 10^{-8}$ for the Schaffer function).
- the starting point for the functions Rastrigin, Schaffer, Ackley, Bohachevsky is $(5, \dots, 5)$, $(55, \dots, 55)$, $(15, \dots, 15)$, and $(8, \dots, 8)$ respectively; the initial step-size σ for these functions is 2, 20, 5, 3 respectively.

A Variant of CMAES-APOP Algorithm

We run the CMAES-APOP algorithm with the small initial population size $\lambda = \lambda_{\text{default}}$ (i.e, set $k_n = 1$) and without the upper bound for the population size in three dimensions $n = 10, 20, 40$.

Function	n	25-p	1-p	10-p	50-p	75-p	90-p
Rastrigin	10	3.317e+04	4.332e+04	3.527e+04	3.160e+04	3.069e+04	3.250e+04
	20	9.077e+04	1.189e+05	9.254e+04	9.212e+04	9.038e+04	9.286e+04
	40	2.981e+05	3.992e+05	3.163e+05	3.006e+05	3.034e+05	3.133e+05
Schaffer	10	3.098e+04	5.111e+04	3.334e+04	3.051e+04	3.012e+04	3.147e+04
	20	8.175e+04	1.663e+05	8.833e+04	8.024e+04	8.233e+04	8.646e+04
	40	2.255e+05	4.942e+05	2.266e+05	2.224e+05	2.348e+05	2.325e+05
Ackley	10	1.403e+04	2.280e+04	1.481e+04	1.369e+04	1.429e+04	1.498e+04
	20	3.105e+04	6.125e+04	3.263e+04	3.024e+04	3.144e+04	3.326e+04
	40	7.204e+04	1.275e+05	7.379e+04	6.761e+04	7.164e+04	7.617e+04
Bohachevsky	10	1.002e+04	1.494e+04	1.052e+04	1.015e+04	1.064e+04	1.085e+04
	20	2.397e+04	4.261e+04	2.533e+04	2.366e+04	2.378e+04	2.494e+04
	40	5.536e+04	9.881e+04	5.781e+04	5.627e+04	5.810e+04	6.101e+04

Table: The aRT of some variants of CMAES-APOP: the 25-percentile is replaced by the other percentiles (aRT (average Running Time) = number of function evaluations divided by the number of successful trials)

A Variant of CMAES-APOP Algorithm

Some notations:

- P : a set of percentiles.
- $f^P := \text{percentile}(\{f(\mathbf{x}_{i:\lambda}), i = 1, \dots, \lambda\}, p)$: the p -percentile of objective function of λ candidates in each iteration, where p can vary from 0 to 100 (in fact p will be chosen from the set of percentiles P); f_{prev}^P and f_{cur}^P denote the p -percentiles in the previous and current iteration respectively.
- n_{up} : the number of times " $f_{\text{cur}}^P - f_{\text{prev}}^P > 0$ " occurs during a slot of S iterations.
- t_{up} : the history of n_{up} in each slot recorded.
- no_{up} : the number of most recent slots we do not see the non-decrease.
- $\lambda_{\text{max}} := (20n + 30)\lambda_{\text{default}}$: the maximum number of the population, where $\lambda_{\text{default}} = \lfloor 4 + 3 \log(n) \rfloor$.

A Variant of CMAES-APOP Algorithm

```
1 Input:  $\mathbf{m} \in \mathbb{R}^n$ ,  $\sigma \in \mathbb{R}_+$ 
2 Initialize:  $\mathbf{C} = \mathbf{I}$ ,  $\mathbf{p}_c = 0$ ,  $\mathbf{p}_\sigma = 0$ ,  $\lambda = k_n \times \lambda_{\text{default}}$ 
3 Set:  $\mu = \lfloor \lambda/2 \rfloor$ ,  $w_i$ ,  $\mu_w$ ,  $c_c$ ,  $c_\sigma$ ,  $c_1$ ,  $c_\mu$ ,  $d_\sigma$ ,  $\text{iter} = 0$ ,  $S = 5$ ,  $r_{\max} = 30$ ,  $n_{\text{up}} = 0$ ,  $t_{\text{up}} = [ ]$ .
4 While not terminate
5   iter = iter + 1;
6    $\mathbf{x}_i = \mathbf{m} + \sigma \mathbf{y}_i$ ;  $\mathbf{y}_i \sim \mathbf{N}(\mathbf{0}, \mathbf{C})$ , for  $i = 1, \dots, \lambda$ 
7   Take  $p$  randomly from the set of percentiles  $P$ 
8   if iter > 1
9     if  $f_{\text{cur}}^P - f_{\text{prev}}^P > 0$       //Check if  $f^P$  increases
10     $n_{\text{up}} = n_{\text{up}} + 1$ ;
11  end
12 end
13 Update  $\mathbf{m}$ ,  $\mathbf{p}_c$ ,  $\mathbf{p}_\sigma$ ,  $\mathbf{C}$ ,  $\sigma$  as in the CMA-ES
14 if (mod(iter, S) = 1) & (iter > 1)      // Adapting the population size
15    $t_{\text{up}} = [t_{\text{up}}; n_{\text{up}}]$ ;
16   Adapt the population size according to the information of  $n_{\text{up}}$  (... details in the next slide)
17    $n_{\text{up}} \leftarrow 0$       // Reset  $n_{\text{up}}$  back to 0
18 end
```

A Variant of CMAES-APOP Algorithm

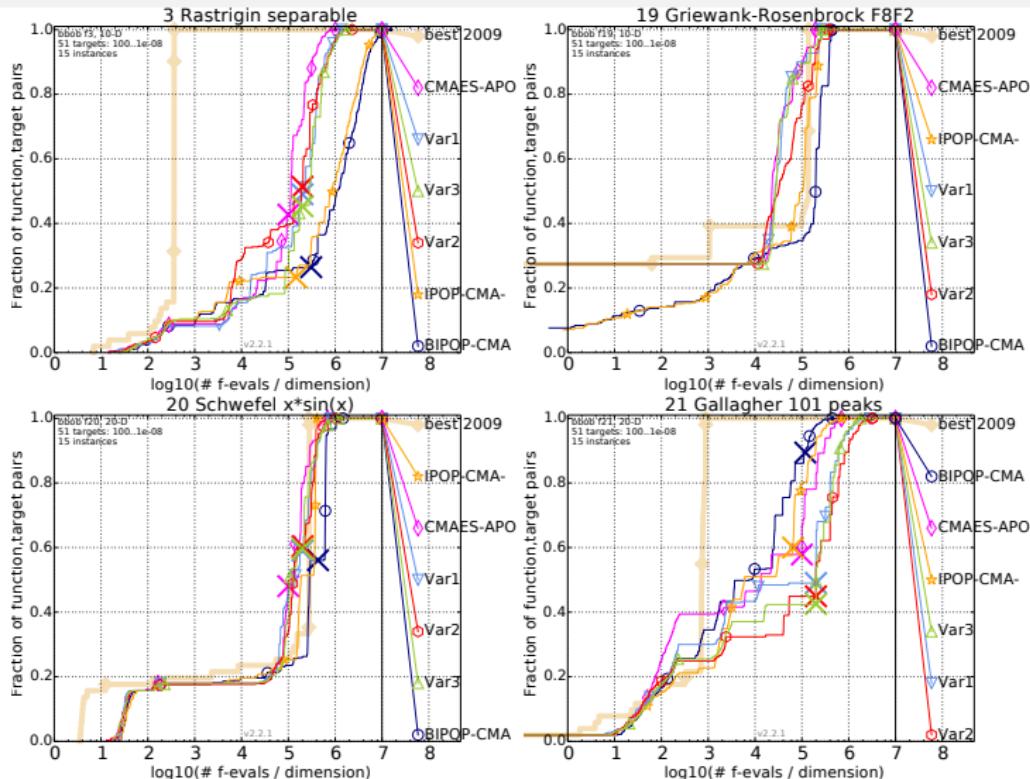
(16) Adapt the population size according to the information of n_{up}

```
16.1      if  $n_{\text{up}} > 1$ 
16.2           $\lambda \leftarrow \left\lfloor \min \left( \exp \left( \frac{n_{\text{up}} \cdot (4 + 3 \log(n))}{S \cdot \sqrt{\lambda - \lambda_{\text{default}} + 1}} \right), r_{\text{max}} \right) \times \lambda \right\rfloor;$ 
16.3           $\lambda \leftarrow \min(\lambda, \lambda_{\text{max}});$ 
16.4           $\sigma \leftarrow \sigma \times \exp \left( \frac{1}{n} \left( \frac{n_{\text{up}}}{S} - \frac{1}{5} \right) \right); // \text{Enlarge } \sigma \text{ a little bit}$ 
16.5      elseif  $n_{\text{up}} = 0$ 
16.6           $\text{no}_{\text{up}} = \text{length}(t_{\text{up}}) - \text{max}(\text{find}(t_{\text{up}} > 0));$ 
16.7          if  $\lambda > 2\lambda_{\text{default}}$ 
16.8               $\lambda \leftarrow \max(\lfloor \lambda \times \exp(-\text{no}_{\text{up}}/10) \rfloor, 2\lambda_{\text{default}});$ 
16.9      end
16.10    end
16.11    if  $\lambda$  is changed // Only when  $n_{\text{up}} > 1$  or  $n_{\text{up}} = 0$ 
16.12        Update  $\mu, w_{i=1 \dots \mu}, \mu_w$  w.r.t the new population size  $\lambda$ 
16.13        Update the parameters  $c_c, c_\sigma, c_1, c_\mu, d_\sigma$ 
16.14    end
```

Numerical Experiments on the BBOB Noiseless Testbed

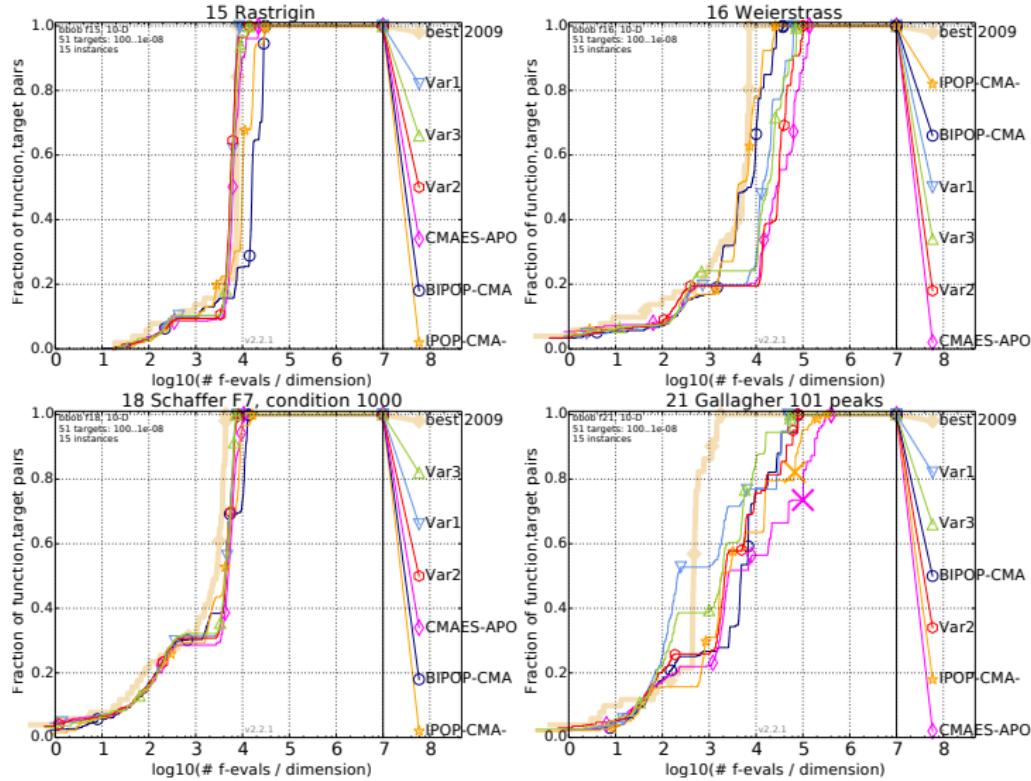
- Test the algorithms with a budget of $2 \times 10^5 \times n$, where n is the problem dimension.
- Denote the variants corresponding to $P_1 = \{1, 25, 50\}$, $P_2 = \{1, 50\}$, and $P_3 = \{1, 50, 75\}$ by Var1, Var2 and Var3 respectively.
- In the first run: the pure CMA-ES with the default population size $\lambda = \lambda_{\text{default}}$. From second run: the pop-size adaptation strategy is applied with the initial population size $\lambda = k_n \times \lambda_{\text{default}}$.
- The parameter k_n is set to 10, 20, 30, 40, 50, 60 for $n = 2, 3, 5, 10, 20, 40$ respectively.
- Take the starting point \mathbf{m}^0 uniformly in $[-4, 4]^n$.
- Set the initial step-size $\sigma_0 = 2$ for all run.

The variants < CMAES-APOP

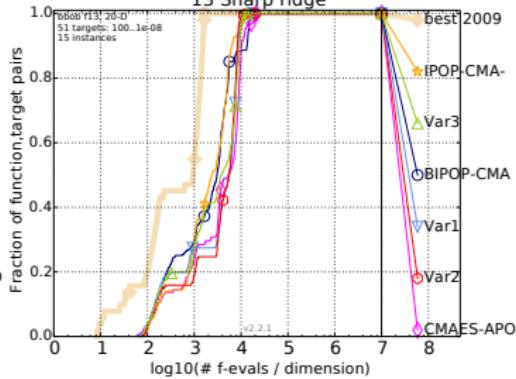
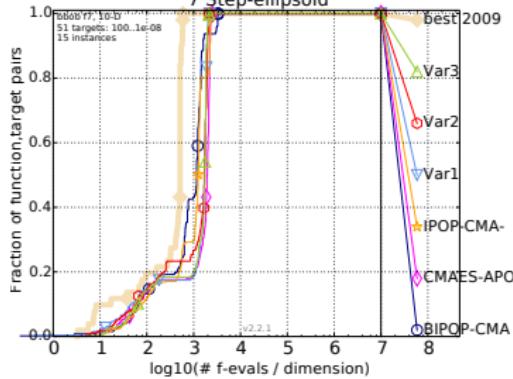
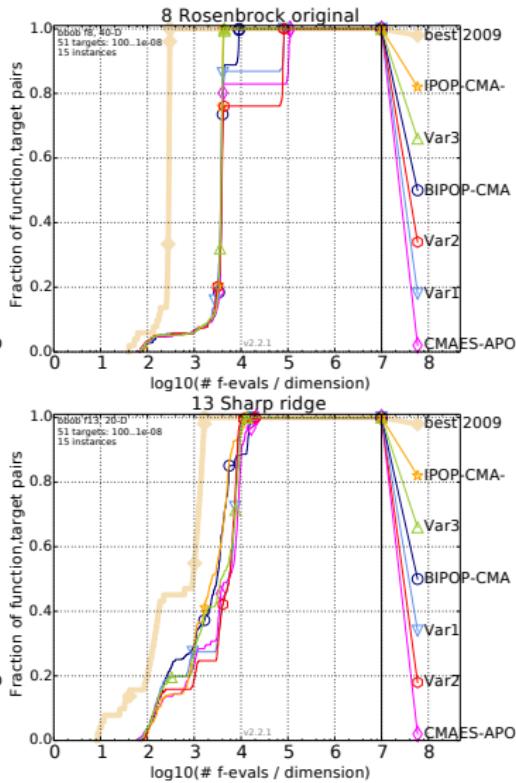
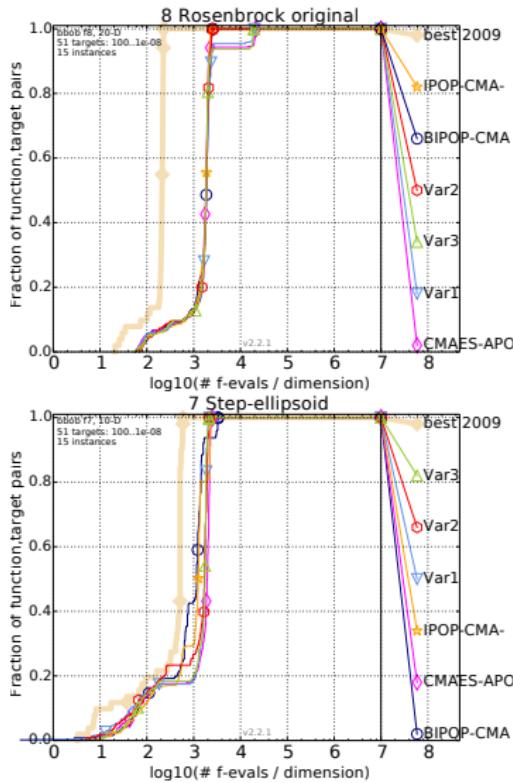


All variants are still better than the IPOP-CMA-ES and BIPOP-CMA-ES on f_3 in 10-D; than the BIPOP-CMA-ES on f_{19} in dimensions 10; and than the BIPOP-CMA-ES on f_{20} in dimensions 20.

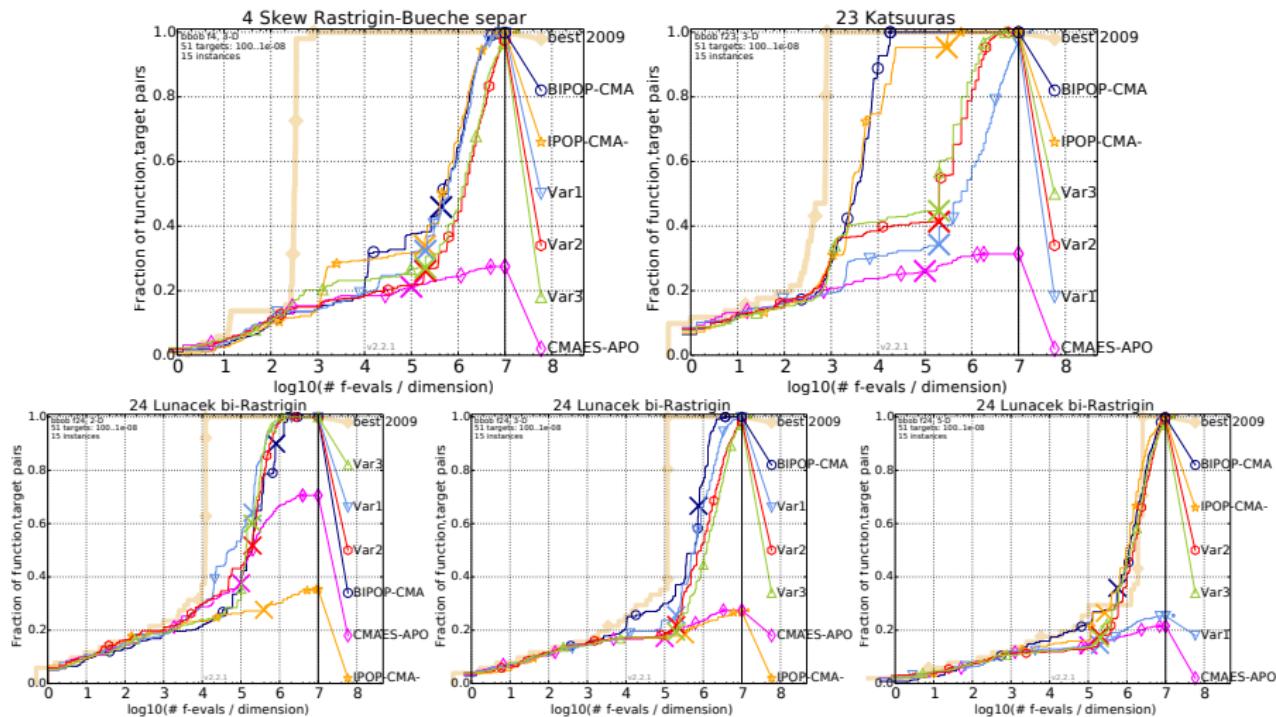
The variants >slightly CMAES-APOP: $f_{15}, f_{16}, f_{18}, f_{21}$ in 10-D



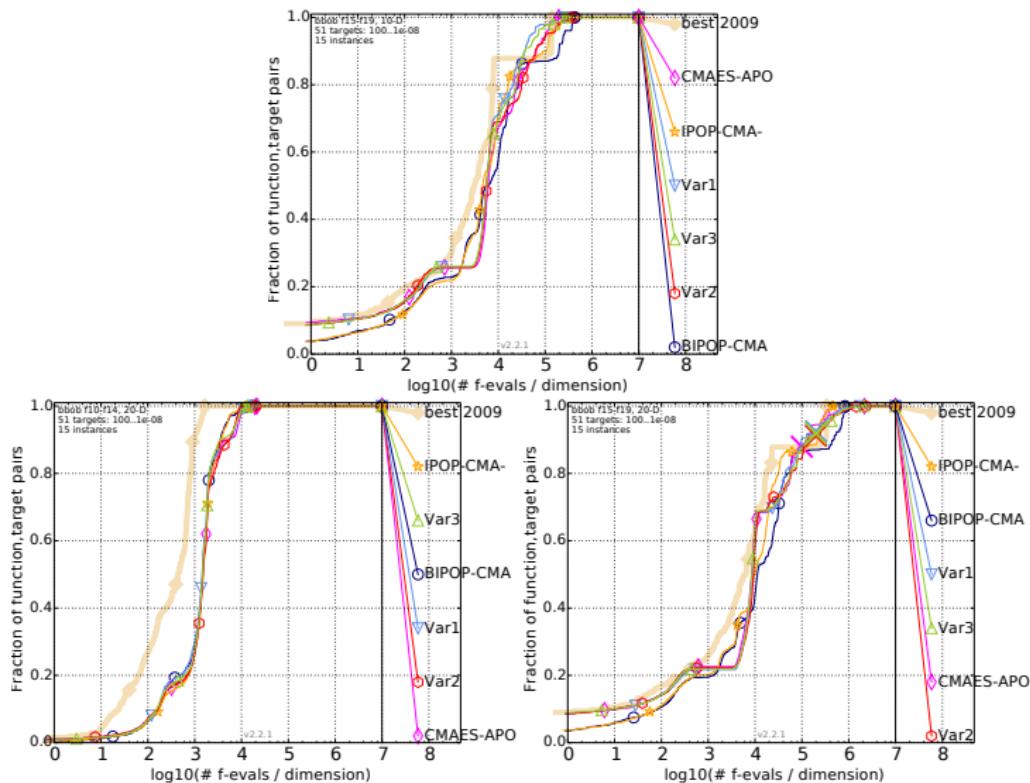
The variants >slightly CMAES-APOP: f_7 , f_8 , f_{13}



The variants > CMAES-APOP: on f_4 , f_{23} , f_{24} in small dimensions

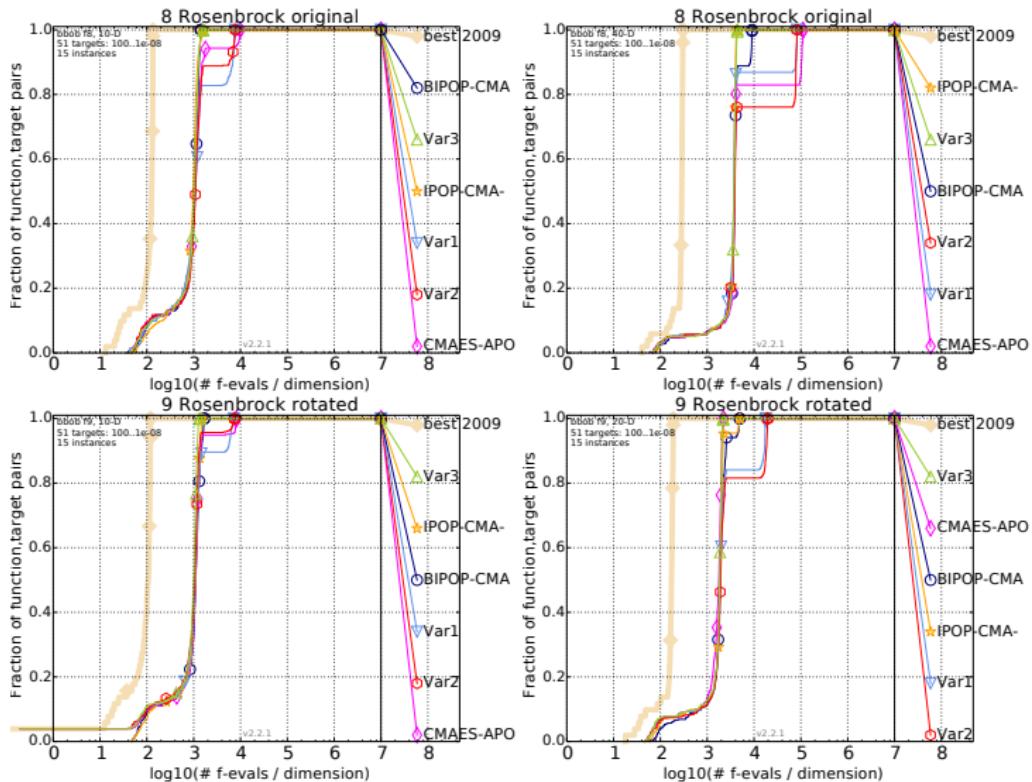


(**Var1** ($\{1, 25, 50\}$) & **Var3** ($\{1, 50, 75\}$)) $>_{\text{slightly}}$ **Var2** ($\{1, 50\}$)

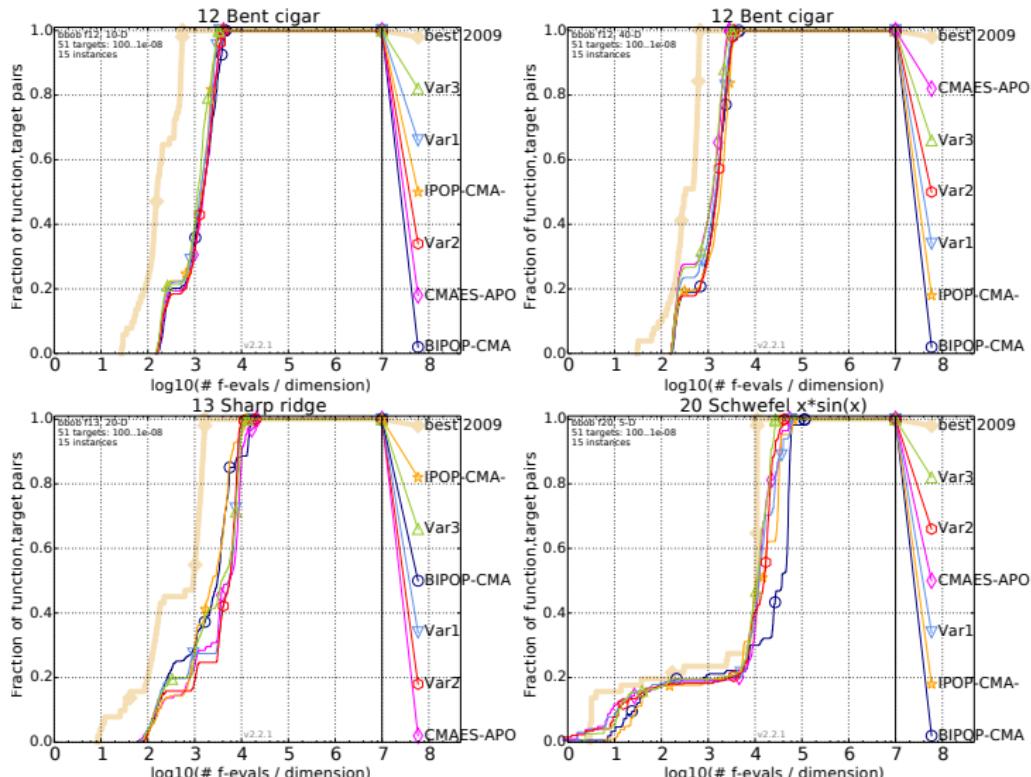


⇒ Tracking more percentiles can help us to make better decisions in adapting population size for the class of conditioned functions, and the class of multi-modal functions with adequate global structure in high dimensions.

Var3 ($\{1, 50, 75\}$) $>_{\text{slightly}}$ ((**Var1** ($\{1, 25, 50\}$) & **Var2** ($\{1, 50\}$))(1/2)



Var3 ($\{1, 50, 75\}$) $>_{\text{slightly}}$ ((**Var1** ($\{1, 25, 50\}$) & **Var2** ($\{1, 50\}$))(2/2)



⇒ The information of non-elite individuals is also useful to adapt the population size.

Conclusion and Perspectives

Conclusion:

- Present a variant of CMAES-APOP: track the change of some percentiles of objective values rather than one percentile; set the upper bound of the population size depending on the problem dimension.
- This approach improves the performance of CMAES-APOP in some cases when the set of percentiles P is chosen appropriately.

Perspectives:

- How to initialize a good set P and how to evaluate the importance of each percentile p in P during the evolution process?
- The information of percentiles could play a deeper role inside the evolution process of the CMA-ES?

References

-  A. Auger and N. Hansen, *A restart cma evolution strategy with increasing population size*, 2005 IEEE Congress on Evolutionary Computation, vol. 2, 2005, pp. 1769–1776 Vol. 2.
-  A. Ahrari and M. Shariat-Panahi, *An improved evolution strategy with adaptive population size*, Optimization **64** (2015), no. 12, 2567–2586.
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Thank you for your attention!