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# Diagonal crossover in genetic algorithms for numerical optimization 

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#### Abstract

In this paper the results of a detailed investigation on a multi-parent recombination operator, diagonal crossover, are reported. Although carlicr publications have indicated the high performance of diagonal crossover on a number of problems, so far it has not been investigated whether high performance is indeed a result of using a high number of parents. Here we formulate three hypotheses to explain why GA performance increases when more parents are used. Based on an extensive study on a test suite containing eight numerical optimization problems we are able to establish that the higher number of parents is indecd one of the sources of higher performance, if and when this occurs. By the diversity of the test functions (unimodal, multimodal, quasi-random landscapes) we can also make obscrvations on the relationship between fitness landscapes and operator performance.

Keywords: genetic algorithms, numerical optimization, recombination, multi-parent crossovers


## 1. Introduction

Multi-parent recombination is a new research arca within evolutionary computation. Although some researchers have incidentally proposed and applied recombination mechanisms using more parents, Bersini and Scront (1996), Bremermann, Rogson and Salaff (1966), Mühlenbcin (1989), the phenomenon of multi-parent recombination has not been given much attention in the past. In this paper we study this phenomenon by investigating the behavior of diagonal crossover (sec the definition in Section 2). Our research goals are two-fold.

1. We try to find connections between the structure of the fitness landscape and the performance of diagonal crossover. In particular we want to establish on what kind of landscapes it is advantageous to increase the number of parents. (Diagonal crossover for 2 parents is identical to the traditional 1-point crossover operator.)
2. We try to disclose the source of increased performance of the diagonal crossover with more parents when it is superior to 2 -parent recombination. Further elaboration of the second rescarch objective requires technical details, and so, we return to this issuc after the exact definition of the operator. The rest of the paper is organized as follows. In the following section we bricfly review multi-parent recombination operators in Evolutionary Algorithms. After the exact definition of diagonal crossover we formulate three hypotheses that can clarify why the performance of the GA increases when the number of parents in diagonal erossover is increased. To this end we design experiments that allow rejection or confirmation of these hypotheses. In Section 4 we present the GA used in the experiments and discuss the performance measures to be used to monitor GA performance. The test suite and the results of the experiments on cach test function are presented in Section 4. Finally, in Section 5 we evaluate the results and draw conclusions.

## 2. Multi-parent recombination

In evolution strategies (ES) global recombination is a multi-parent operator, Bäck (1996), Schwefel (1995). This operator creates a new value in the child chromosome based on two parents, but randomly chooses two parents for cach variable anow. By this particular mechanism the number of parents is mudefined, thus investigations on the effects of different number of recombinants on algorithm performance could not be performed in the traditional ES framework. (Let us note that in Schwefel and Rudolph, 1995, an extension of ES is proposed that allows tuning of the number of recombinants.) So far there are almost no experimental results available on the (dis)advantages of global recombination with respect to usual, two-parent recombination. Schwefel (1995) briefly touches on this issue stating that 'appreciable accoleration' is obtained by changing to bisexual from asexual scheme (i.e. adding recombination using t.wo parents to the mutation-only algorithm), but only 'slight, further increase, is obtained when changing from 'bisexual to multisexual recombination' (i.e. using global recombination instcad of the two-parent variant).

Related work of Beyer (1995), gencralizes the traditional ES recombination operators by introducing the number of parents as an independent parameter $\rho$. The resulting $(\mu / \rho, \lambda)$ evolution strategy is studied for the special case of $\rho=\mu$ and theoretical analysis on the spherical function shows an advantage of using more than two parents.

Global recombination in ES also fertilized Genctic: Algorithms. The genepool recombination of Mühlenbein and Voigt mixes information of possibly more
parents by a similar mechanism as global recombination in ES, Mühlenbein and Voigt (1995), Voigt and Mühlenbein (1995). Hence, the number of parents is not defined here either. Mühlenbein and Voigt report an increase of performance when using genc-pool recombination (GPR) instead of two-parent recombination (TPR). GPR is showed to be approximately $25 \%$ faster than TPR on the ONEMAX problem, and the fuzzyfied GPR outperforms TPR on the spherical function in speed and in realized heritability.

The $N$-parent generalizations of the traditional 1-point crossover and uniform crossover in GAs were introduced in Eiben, Raué and Ruttkay (1994). The resultiing diagonal, respectively, scanning crossover have the number of parents as parameter, and therefore are tumable on the 'extent of sexuality'. This tunability is new feature compared to global recombination and gene pool recombination, where the multi-parent option can only be switched on or off, but it is not, scalable. Several studies, e.g. Eiben, van Kemenade and Kok (1995), Eiben, Rauf and Ruttkay (1994), Eiben and Schippers (1996), van Kemenade, Kok and Eiben (1995), have shown that using more than two parents in either erossover mechanism can increase GA performance, although this does not hold for every problem and the two operators can respond differently to increasing the number of parents.

The main subject of the present investigation, diagonal crossover, generalizes 1-point crossover for $N$ parents by selecting ( $N-1$ ) crossover points and composing $N$ chiddren by taking the resulting $N$ chromosome segments from the parents 'along the diagonals'. The idea is illustrated for $N=3$ in Fig. 1 up.

With respect to our secome rescarch objective let us make the following ohservations. First, the increase in the number of parents automatically leads to an increased number of erossorer points. It can be the case that higher performance for higher $N$ 's is not the result of using more parents, but simply comes from being more disruptive by using more crossover points. This forms our first working hypothesis.
$\mathrm{H}_{1}$ Using more crossover points leads to better performance.
Second, notice that, by application of the diagonal crossover, $N$ parents create $N$ chikdren in one go. Since we use a steady state GA and update the population, i.e. insert offspring, after cach application of crossover (followed by mutation), this means that a (iA using 10 parents diagonal crossover has more information before performing the selection step than a GA using the two-parents version. In wherer words, CiAs with higher operator "arity" have a bigger gencrational gap which might, cause a bias in their favor. Our second working hypothesis is aceordingly the following.
$\mathrm{H}_{2}$ IBigger generational gap leads to better performance.
Finally, we maintain our original conjecture that the advantages of using diagonal crossover with higher "aritics" are not the result of an unintended artifact.


Figure 1. Diagonal crossover with threc parents and three children (up) and with three parents and one child (down).
$\mathrm{H}_{3}$ Using more parents leads to better performance.
Note that the hypotheses $\mathrm{H}_{1}, \mathrm{H}_{2}$ and $\mathrm{H}_{3}$ are not mutually exclusive as there might be more sources of increased GA performance when increasing $N$ in diagonal crossover. The main contribution of the present paper is that these sources are investigated in isolation hence providing a solid ground to check whether higher performance for higher $N$ 's is an artifact $\left(\mathrm{H}_{1}, \mathrm{H}_{2}\right)$, or the higher number of parents is indeed advantagcous.

## 3. Experiment setup and performance measures

All experiments are exccuted using a GA setup as described in Table 1. A nonstandard option is the uniform random parent selection mechanism, whereby no selective pressure is applied when choosing recombinants. The motivation comes from van Kemenade, Kok and Eiben (1995), where we observed that this mechanism is preferable. Note that uniform random parent selection mechanism is standard in Evolution Strategies.

In order to test the working hypotheses presented in Section 2 we run experiments with a set of different crossover operators. For investigating $\mathrm{H}_{1}$ we apply the traditional two-parent two-children $N$-point crossover, De Jong and Spears (1992). This operator is well known from the literature, therefore we ornit a def-

| representation | fixed point binary with Gray coding |
| :--- | :---: |
| GA type | strady state |
| parent selection | uniform random |
| deletion mechanism | worst fitness deletion |
| number of parents | $2-30$ |
| crossover rate $p_{c}$ | 1.0 |
| mutation rate $p_{m}$ | $1 /$ chromosome length |
| population size | 500 |
| termination criterion | population converged OR optimum hit |
| max. nr. of evaluations | 100,000 |
| results averaged over | 50 indcpendent runs |

Table 1. GA setup used in the experiments
inition. If $N$-point crossover exhibits increasing performance when increasing $N$, (the experimental results reported in Eiben, van Kemenade and Kok, 1995, make us expect this) then we accept $\mathrm{H}_{1}$. To test the second hypothesis $\mathrm{H}_{2}$, we will apply a slightly modified version of diagonal crossover that creates only one child. The lower part of Fig. 1 illustrates this operator. When we use the one child version of diagonal crossover the generational gap does not increase with increasing the number of parents. If the original variant outperforms the one child version of diagonal crossover, then we accept the hypothesis $\mathrm{H}_{2}$. Concerning hypothesis $\mathrm{H}_{3}$, note that the number of chromosome segments using $N$-point crossover is $N+1$, which equals the number of chromosome segments obtained by diagonal crossover for $N+1$ parents. This means that the disruptiveness of these operators grows parallelly as $N$ increases. If higher disruptiveness increases GA performance on our test suite, then the performance of both the $N$-point crossover and the diagonal crossover will increase with increasing $N$. This, however, docs not imply that more parents have no additional advantage as the performance of diagonal crossover might grow faster with increasing $N$ than that of the two-parent $N$-point crossover. We accept the hypothesis $\mathrm{H}_{3}$ if diagonal crossover for $N+1$ parents is better than $N$-point crossover.

To evaluate different GA setups, that is, the effect of different number of parents, respectively crossover-points, several performance measures of a run are monitored. The two main performance measures are accuracy and speed. Accuracy is measured by the crror at termination. Since all functions have a minimum of zero, we use the best objective function value at termination as the accuracy measure of one run, and the median of the best objective function values, calculated over the 50 independent runs, as the accuracy belonging to a specific setting. For practical purposes we consider $10^{-10}$ as zero and terminate the run if this value is achieved. Let us remark that using medians
instead of average values has an advantage, namely medians are less sensitive for outliers in the data. On the other hand, if the optimum is found in the majority of the runs, then the median will equal the optimum. Additionally to the medians of the outcomes we also present the $99 \%$ confidence interval bars to the performance curves. The second main performance measure is the speed of the algorithm, measured by the median number of fitness evaluations before termination. If the GA with a certain setup never finds the optimum, this value equals the maximum number of fitness evaluations. A third performance measure is the success rate, i.e. the percentage of rums where the optimal objective function value has been found. We will present figures on success rates and $90 \%$ confidence intervals, whenever the accuracy or the speed curves are (nearly) constant, thus providing (almost) no basis to compare different settings. Finally, for a detailed insight in the behavior of the GA sometimes we also depict the progress curves of the evolution for 18 parents (diagonal crossover), respectively 17 crossover points ( $N$-point crossover). These curves (with a logarithmic $y$-axis) show the population's best objective function value as a function of the number of executed fitness evaluations, averaged over 50 independent runs.

## 4. Experimental results

The experiments have been performed on cight numerical function optimization problems. Each function is to be minimized and is scaled to have an optimal function value of zero. The fitness landscapes defined by these functions have various characteristics, unimodal, multimodal and quasi-random, i.e. very rugged with randomly distributed local optima. Additionally, some of the functions are separable, while others are not. The exact definitions will be given in the corresponding subsections, here we only give a summary on their separability, the dimensions and the representation used in the experiments. As defanlt, we use 20 bits for representing a single variable, but deviate from this value for F1, F2 and F8. For F1 and F2 we use the values originating from de Jong, for F8 30) bits are used, following Bäck (1996). A concise treatment on mumerical optimization problems as test functions can be found in Bäck and Michalewicz (1997).

| Property | F1 | F2 | F3 | F4 | F5 | F6 | F7 | F8 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| separable | y | n | n | n | y | y | y | n |
| dimension | 3 | 2 | 30 | 10 | 10 | 10 | 10 | 30 |
| chrom. length | 30 | 22 | 600 | 200 | 200 | 200 | 200 | 900 |

Table 2. Propertics of the test functions


Figure 2. Speed curves for the spherical function

### 4.1. Spherical function

The first test function is the spherical function:

$$
F 1(\vec{x})=\sum_{i=1}^{n} x_{i}^{2}
$$

where $-5.12 \leq x_{i} \leq 5.12$. This function is one of the most widely used objective functions in Evolutionary Computation, especially for convergence velocity evaluation. It has a unimodal, smooth fitness surface and is separable, making optimization rather casy. We tested the classical version of de Jong with $n=3$. The GA found the optimum with every setting (every operator, for every value of $N$ ). Therefore we omit accuracy and success rate data, only presenting the speed curves in Fig. 2.

From the speed curves it turns out that the two variants of diagonal crossover show almost identical behavior and both are faster than $N$-point crossover. Furthermore, it seems that there is a limit, to increasing $N$ : approximately up to 6 it leads to performance increase, thereafter the performance begins to deteriorate.

### 4.2. Rosenbrock's saddle function

F2 is the saddle function after Rosenbrock:

$$
F 2(\vec{x})=100 \cdot\left(x_{1}^{2}-x_{2}\right)^{2}+\left(1-x_{1}\right)^{2}
$$

where $-2.048 \leq x_{i} \leq 2.048$. The global minimum is zero at $\vec{x}=(1,1)$. The Rosenbrock function is not separable and the unimodal fitness landscape is characterized by an extremely deep valley along the parabola $x_{1}^{2}=x_{2}$.


Figure 3. Accuracy (upper diagram) and speed (lower diagram) for the Rosenbrock function

Recall from Table 2 that we use the classical de Jong setting with chromosome length 22 for F2. Therefore, the maximum number of parents is lowered accordingly in these experiments.

The accuracy and speed curves suggest that increasing $N$ decreases the performance. The success rate curves in Fig. 3 disclose that this is only partly truc. The optimization performance grows with $N$ for $N$-point crossover (up to $N=8$ ), but, deteriorates for the diagonal crossovers. $N$-point crossover outperforms both diagonal crossovers with respect to each performance measure. The two variants of diagonal crossover are practically identical for F2.


Figure 4. Success rates for the Rosenbrock function

### 4.3. Ackley function

Our third test function F3 is the Ackley function:

$$
F 3(\vec{x})=20+e-20 \operatorname{cxp}\left(-0.2 \sqrt{\frac{1}{n} \sum_{i=1}^{n} x_{i}^{2}}\right)-\operatorname{cxp}\left(\frac{1}{n} \sum_{i=1}^{n} \cos \left(2 \pi x_{i}\right)\right),
$$

where $-30 \leq x_{i} \leq 30$. The global minimum of zero is at $\vec{x}=(0,0,0, \ldots)$. This function is not separable and at a low resolution the fitness landscape is unimodal, but the second exponential term covers the landscape with many small peaks and valleys, i.e. many local optima. We tested F3 for $n=30$ and observed that the GA never found the optimum. Accordingly, the speed and the success rate curves are constant, therefore omitted here. We present the accuracy curves in Fig. 5.

The effect of higher $N$ 's is clear from the accuracy curves. Increasing $N$ is advantageous for each operator up to the upper limit we tested. The one child and the $N$-children versions of diagonal crossover perform identically also on this function, and both diagonal crossovers are consistently better than N point crossover.

### 4.4. Griewangk function

F4, the Griewangk function is defined as follows.

$$
F 4(\vec{x})=1+\sum_{i=1}^{n} \frac{x_{i}^{2}}{400 n}-\prod_{i=1}^{n} \cos \left(\frac{x_{i}}{\sqrt{i}}\right)
$$



Figure 5. Accuracy curves for the Ackley function
where $-600 \leq x_{i} \leq 600$. The global minimum of zero is at the point $\vec{x}=$ $(0,0,0, \ldots)$. This function has a product term introducing an interdependency between the variables, thus it is not separable. F4 was tested for 10 dimensions, the results are exhibited in Fig. 6.

On this function the advantages of higher $N$ 's are clear, but the performance increase of accuracy stops at about $N=15$. While the accuracy curves show only modest, differences between the operators, the results on the speed of the algorithm disclose that the diagonal crossovers are significantly faster after $N=$ 5 . The two variants of diagonal erossover do not differ significantly.

### 4.5. Michalewicz function

The fifth test function is taken from Michalewice, Bersini, Dorigo, Langerman, Scront and Gambardella (1996).

$$
F 5(\vec{x})=-\sum_{i=1}^{n} \sin \left(x_{i}\right) \cdot \sin ^{2 n}\left(\frac{i x_{i}^{2}}{\pi}\right)
$$

where $0 \leq x_{i} \leq \pi$. We tested F5 for $n=10$ and observed that the GA found the optimum in the majority of runs. Hence the medians of the accuracy results are equal to the optimal value. Therefore we rather present the sucecess rates instead of the accuracy data.

Increasing $N$ above 2 on the Michalewicz function results in the highest gains so far. The success rates show a spectacular increase from approximately $30 \%$ for 2 parents to $80-90 \%$ for $N$ between $5-20$ and the GA becomes approximately four times faster for $N=5-10$, than for $N=2$. Comparing the operators we see again the superiority of diagonal crossover and no significant difference between the one child and the $N$-child version.


Figure 6. Accuracy (upper diagram) and speed (lower diagram) for the Gricwangk function


Figure 7. Success rates (upper diagram) and speed (lower diagram) for the Michalewicz function


Figure 8. Success rates (upper diagram) and speed (lower diagram) for the Rastrigin function

### 4.6. Rastrigin function

F6 is the Rastrigin function:

$$
F 6(\vec{x})=\alpha n+\sum_{i=1}^{n} x_{i}^{2}-\alpha \cos \left(2 \pi x_{i}\right)
$$

where $-5.12 \leq x_{i} \leq 5.12$. The global minimum of zero is at $\vec{x}=(0,0, \ldots)$. This function is separable and its primary characteristic is the existence of many suboptimal peaks whose valucs increase as the distance from the global optimum point increases. In our tests we used $\alpha=10.0$ and $n=10$.

Since many runs found the optimum, accuracy figures are replaced by success rates curves (notice the $0.65-1.0$ scaling in Fig. 8, upper part). These show that


Figure 9. Speed curves for the Schwefel function
increasing $N$ is advantageous, but the differences between various operators are small. Looking at the speed curves the effect of higher $N$ 's and the differences between operators become clear. We can observe that each operator becomes better for higher $N$ 's and that the two diagonal crossovers (identical again) ontperform N -point crossover.

### 4.7. Schwefel function

F7 is obtained by generalizing Schweffl's 2.26 function (Schwefel, 1995, p. 344):

$$
F 7(\vec{x})=418.9829 n-\sum_{i=1}^{n} x_{i} \sin \left(\sqrt{\left|x_{i}\right|}\right)
$$

where $-512.03 \leq x_{i} \leq 511.97$. The global minimum of zero is at $\vec{x}=(420.9687$, $420.9687, \ldots$ ). Although this function is separable, it is interesting because of the presence of a second-best minimum far away from (in the 'opposite corner' to) the global minimum. This feature, just like two-peaks landscapes, makes the GA sensitive for early commitment with respect to the search direction. F7 was tested for $n=10$ and turned out to be easy. Nearly all runs ended with the global optimum, implying that accuracy and success rates would give no information for comparing the operators. The results on speed, however, show that the GA performance quickly and consequently improves when increasing $N$ from 2 to approximately 10-15, and stagnates thercafter. The algorithm becomes approximately twice as fast for high $N$ 's as for $N=2$. Once again, there is no significant difference between the two diagonal crossovers that both outperform $N$-point crossover.

### 4.8. Fletcher-Powell function

F8, the Fletcher-Powell function is retrieved from Bäck (1996):

$$
\begin{aligned}
F 8(\vec{x}) & =\sum_{i=1}^{n}\left(A_{i}-B_{i}\right)^{2} \\
A_{i} & =\sum_{j=1}^{n}\left(a_{i j} \sin \alpha_{j}+b_{i j} \cos \alpha_{j}\right) \\
B_{i} & =\sum_{j=1}^{n}\left(a_{i j} \sin x_{j}+b_{i j} \cos x_{j}\right),
\end{aligned}
$$

where $n=30, n_{\sigma}=30$, and $-\pi \leq x_{i}^{0} \leq \pi$. The $a_{i j}, b_{i j} \in\{-100, \ldots, 100\}$ are random integers, and $\alpha_{j} \in[-\pi, \pi]$ is the randomly chosen global optimmm position. For the matrices A, B and the vector $\vec{r}$ we used the values given in Bäck (1996) (pp. 265 267).

No runs found the optimmon this function, resulting in constant speed and success rate curves. Accuracy curves reveal differences between performane for different $N$ 's, showing advantageons effects of higher $N$ 's, up to approximately 10. The three operators, however, hardly differ in performance and, as the progress curves for $N=18$ in Fig. 10 indicate, their search behavior is very similar, too.

## 5. Conclusions

Concerning our first research objective, i.e. finding comections betwern the characteristies of the objective functions and the usefuhess of applying more parents, we can observe the following. With the single exerption of Rosenbrock's saddle (F2) it is useful to apply diagonal crossover with "arity" $N>2$. Lookin" for particular features of F2 that may canse this deviance lot us note that it has the lowest dimensionality $(n=2)$ and the shortest chromosomes (of length 22) at opposed to other functions (200)-900) for F3-F8). This makes the dismptiveness of the crossover operators relatively high even for low $N$ 's. The other minodal function in the test set, Fl, is apparently so casy to optimize that the (iA does not suffer from this effect, but on F2 where the optimmu is hidden at the bottom of a long bent valley', see Bäck and Michalewicz (1997), this seems to be disasterous.

Our second researeh objective concerned the identification of the souree(s) of increased performance of diagonal crossover when used with more parents. As for the hypothesis $H_{1}$, i.e. that increasing the mumber of erossover points increases performance, observe that $N$-point crossover did become better for higher $N$ 's on all functions of our test suite. Therefore, we aceept $\mathrm{H}_{1}$ and conchude that higher performance partially comes from a higher number of crossower points. Explanations for this fact are the better mixing of information, see de


Figure 10. Accuracy (upper part) and progress curves (lower part) for the Fletcher-Powell function

Jong and Spears (1992), and perhaps also the increased macro-mutation-like effects of crossover in case of using higher $N$ 's.

Hypothesis $\mathrm{H}_{2}$ about the advantages of a bigger gencrational gap is clearly rejected, since the one child and the $N$ children variants of diagonal crossover exhibited the same behavior on each function. Hence, we can conclude that the advantage of applying diagonal crossover with higher $N$ 's is not the result of a bigger generational gap in the Steady State GA we use (see Section 2 for discussion).

Recall that our working hypotheses are not mutually exclusive. Accepting $\mathrm{H}_{1}$ does not imply rejection of $\mathrm{H}_{3}$, i.e. that better performance for higher $N$ 's would only come from having more crossover points. In fact diagonal crossover was better than $N$-point crossover on all but two functions: on Rosenbrock's saddle (F2) and on the Fletcher-Powell function (F8). On the Fletcher-Powell function diagonal crossover was not significantly better than $N$-point crossover. Such little differences in performance do not clearly justify the acceptence of the hypothesis $\mathrm{H}_{3}$ on the function F8. There is no clear advantage of using more parents for recombination here. Increased performance for higher $N$ 's seems to be the result of the crossovers effect as macro mutation, this effect being intensified by more crossover points. Recall, that F8 spans a very rugged landscape with randomly distributed local optima, which makes it more or less similar to NK-landscapes with relatively high K values. These obscrvations are thus in agreement with carlier conclusions for NK-landscapes, Eiben and Schippers (1996), Hordijk and Manderick (1995), Kauffman (1993), stating that on such surfaces crossover is not useful at all. On Rosenbrock's saddle (F2) diagonal crossover was clearly worse than $N$-point crossover, besides, the performance of diagonal crossover decreased for increasing $N$. This behavior is unique on the test suite we use here and at the moment we do not have a solid clarification for it.

According to the above considerations, hypothesis $\mathrm{H}_{3}$ has to be refined. On quasi-random landscapes, such as F8, increased performance of diagonal crossover for higher $N$ 's may occur, but it seems not to be the result of using more parents, i.e. $\mathrm{H}_{3}$ does not hold. On other types of landscapes (the unimodal F1 and the multimodal, but somewhat regularly shaped F3-F7), diagonal crossover exhibits increased performance when increasing $N$, and it does outperform $N$-point crossover, thus confirming $\mathrm{H}_{3}$. F2 remains an exception, showing that even for unimodal landscapes it is not guaranteed that diagonal crossover will become better when increasing $N$. Yet, with this exception in mind, we can draw the conclusion that if diagonal crossover becomes better for higher $N$ 's then this improvement is not only the consequence of using more crossover points, but also that of using more parents.

Let us close our conclusions with noting that the greatest gain occurred in the speed of the GA, diagonal crossover is usually faster than 2 parent $N$-point crossover (if and when). Clearly, if we had set the maximum number of evaluations lower then this difference in speed would also have resulted in differences
in success rate and accuracy. Thus, although we definitely do not claim that diagonal crossover is a universally superior operator, we have sufficient evidence to say that it is a sound design heuristics to implement it in a GA and set the number of parents above two.

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