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SUPPLEMENTAL MATERIAL for Software for data-based stochastic programming using bootstrap estimation

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Appendix A: Experiments

We demonstrate experiments on the algorithms over a few examples as detailed in the following subsections. For simulation purpose, we grant ourselves the access to the distribution of the entire population F_Ω , and approximate the theoretical optimal function value z^* by drawing an extremely large number of samples from the distribution F_Ω , then compute the corresponding optimal function value, which we use as z^* for the simulations.

We replicate the execution of each algorithm a large number (e.g., 500) times with a fixed candidate solution \hat{x} . At each replication, a new set D is drawn independently from F_Ω , and the algorithm is executed to return a $(1 - \alpha)$ confidence interval for the optimal function value z^* . The coverage rate is then reported as the percentages of the confidence intervals that contains z^* over the replications. We also report the average length of the confidence intervals, denoted as “avg len”, over the repeated experiments for each algorithm. The average length of the CIs to some extent represents the sharpness of the intervals computed by the algorithms.

A.1. Small Schultz Examples

A.1.1. Unique Solution This example is from [Schultz et al. \(1998\)](#) used in ([Eichhorn and Römisch 2007](#), p. 129). Results are shown in Table 1.

A.1.2. Nonunique Solution This example is a modified version of the previous problem that used in ([Eichhorn and Römisch 2007](#), p. 131). Results are shown in Table 2. This problem has multiple optima.

| method | N | B | k | avg len | coverage |
|-----------------------------|----|-----|----|---------|----------|
| Classical_gaussian | 40 | 100 | - | 8.04 | 0.92 |
| Classical_gaussian | 80 | 100 | - | 5.58 | 0.90 |
| Classical_gaussian | 40 | 500 | - | 8.01 | 0.92 |
| Classical_gaussian | 80 | 500 | - | 5.73 | 0.93 |
| Classical_quantile | 40 | 100 | - | 7.82 | 0.88 |
| Classical_quantile | 80 | 100 | - | 5.41 | 0.87 |
| Classical_quantile | 40 | 500 | - | 7.97 | 0.90 |
| Classical_quantile | 80 | 500 | - | 5.69 | 0.89 |
| Bagging_with_replacement | 40 | 100 | 16 | 9.36 | 0.93 |
| Bagging_with_replacement | 40 | 100 | 24 | 9.38 | 0.95 |
| Bagging_with_replacement | 40 | 100 | 32 | 9.42 | 0.96 |
| Bagging_with_replacement | 80 | 100 | 32 | 7.62 | 0.97 |
| Bagging_with_replacement | 80 | 100 | 48 | 7.39 | 0.97 |
| Bagging_with_replacement | 80 | 100 | 64 | 7.49 | 0.98 |
| Bagging_with_replacement | 40 | 500 | 16 | 8.21 | 0.90 |
| Bagging_with_replacement | 40 | 500 | 24 | 8.15 | 0.89 |
| Bagging_with_replacement | 40 | 500 | 32 | 8.26 | 0.91 |
| Bagging_with_replacement | 80 | 500 | 32 | 6.08 | 0.93 |
| Bagging_with_replacement | 80 | 500 | 48 | 6.05 | 0.95 |
| Bagging_with_replacement | 80 | 500 | 64 | 6.07 | 0.95 |
| Bagging_without_replacement | 40 | 100 | 16 | 9.32 | 0.95 |
| Bagging_without_replacement | 40 | 100 | 24 | 9.59 | 0.93 |
| Bagging_without_replacement | 40 | 100 | 32 | 9.43 | 0.94 |
| Bagging_without_replacement | 80 | 100 | 32 | 7.54 | 0.98 |
| Bagging_without_replacement | 80 | 100 | 48 | 7.83 | 0.98 |
| Bagging_without_replacement | 80 | 100 | 64 | 7.68 | 0.95 |
| Bagging_without_replacement | 40 | 500 | 16 | 8.42 | 0.91 |
| Bagging_without_replacement | 40 | 500 | 24 | 8.58 | 0.94 |
| Bagging_without_replacement | 40 | 500 | 32 | 8.60 | 0.93 |
| Bagging_without_replacement | 80 | 500 | 32 | 6.16 | 0.93 |
| Bagging_without_replacement | 80 | 500 | 48 | 6.29 | 0.95 |
| Bagging_without_replacement | 80 | 500 | 64 | 6.26 | 0.95 |
| Subsampling | 40 | 100 | 16 | 6.15 | 0.79 |
| Subsampling | 40 | 100 | 24 | 5.08 | 0.73 |
| Subsampling | 40 | 100 | 32 | 3.52 | 0.58 |
| Subsampling | 80 | 100 | 32 | 4.36 | 0.80 |
| Subsampling | 80 | 100 | 48 | 3.62 | 0.72 |
| Subsampling | 80 | 100 | 64 | 2.54 | 0.55 |
| Subsampling | 40 | 500 | 16 | 6.30 | 0.81 |
| Subsampling | 40 | 500 | 24 | 5.18 | 0.74 |
| Subsampling | 40 | 500 | 32 | 3.65 | 0.56 |
| Subsampling | 80 | 500 | 32 | 4.45 | 0.82 |
| Subsampling | 80 | 500 | 48 | 3.68 | 0.74 |
| Subsampling | 80 | 500 | 64 | 2.58 | 0.56 |
| Extended | 40 | 100 | - | 8.29 | 0.88 |
| Extended | 80 | 100 | - | 5.37 | 0.81 |
| Extended | 40 | 500 | - | 8.22 | 0.90 |
| Extended | 80 | 500 | - | 5.60 | 0.86 |

Table 1 Results for unique_schultz ($z^* \approx -62.29$) with $\alpha=0.1$ based on 100 replications.

| method | N | B | k | avg len | coverage |
|-----------------------------|----|-----|----|---------|----------|
| Classical_gaussian | 40 | 100 | - | 7.92 | 0.90 |
| Classical_gaussian | 80 | 100 | - | 5.46 | 0.91 |
| Classical_gaussian | 40 | 500 | - | 7.89 | 0.89 |
| Classical_gaussian | 80 | 500 | - | 5.60 | 0.93 |
| Classical_quantile | 40 | 100 | - | 7.72 | 0.89 |
| Classical_quantile | 80 | 100 | - | 5.30 | 0.86 |
| Classical_quantile | 40 | 500 | - | 7.84 | 0.90 |
| Classical_quantile | 80 | 500 | - | 5.57 | 0.92 |
| Bagging_with_replacement | 40 | 100 | 16 | 9.23 | 0.93 |
| Bagging_with_replacement | 40 | 100 | 24 | 9.24 | 0.94 |
| Bagging_with_replacement | 40 | 100 | 32 | 9.28 | 0.96 |
| Bagging_with_replacement | 80 | 100 | 32 | 7.45 | 0.96 |
| Bagging_with_replacement | 80 | 100 | 48 | 7.23 | 0.97 |
| Bagging_with_replacement | 80 | 100 | 64 | 7.32 | 0.97 |
| Bagging_with_replacement | 40 | 500 | 16 | 8.10 | 0.89 |
| Bagging_with_replacement | 40 | 500 | 24 | 8.02 | 0.89 |
| Bagging_with_replacement | 40 | 500 | 32 | 8.13 | 0.91 |
| Bagging_with_replacement | 80 | 500 | 32 | 5.95 | 0.92 |
| Bagging_with_replacement | 80 | 500 | 48 | 5.92 | 0.94 |
| Bagging_with_replacement | 80 | 500 | 64 | 5.94 | 0.94 |
| Bagging_without_replacement | 40 | 100 | 16 | 9.19 | 0.95 |
| Bagging_without_replacement | 40 | 100 | 24 | 9.43 | 0.93 |
| Bagging_without_replacement | 40 | 100 | 32 | 9.27 | 0.92 |
| Bagging_without_replacement | 80 | 100 | 32 | 7.38 | 0.98 |
| Bagging_without_replacement | 80 | 100 | 48 | 7.63 | 0.98 |
| Bagging_without_replacement | 80 | 100 | 64 | 7.50 | 0.96 |
| Bagging_without_replacement | 40 | 500 | 16 | 8.29 | 0.90 |
| Bagging_without_replacement | 40 | 500 | 24 | 8.44 | 0.91 |
| Bagging_without_replacement | 40 | 500 | 32 | 8.46 | 0.93 |
| Bagging_without_replacement | 80 | 500 | 32 | 6.03 | 0.93 |
| Bagging_without_replacement | 80 | 500 | 48 | 6.14 | 0.96 |
| Bagging_without_replacement | 80 | 500 | 64 | 6.11 | 0.94 |
| Subsampling | 40 | 100 | 16 | 6.04 | 0.82 |
| Subsampling | 40 | 100 | 24 | 4.99 | 0.72 |
| Subsampling | 40 | 100 | 32 | 3.45 | 0.61 |
| Subsampling | 80 | 100 | 32 | 4.28 | 0.81 |
| Subsampling | 80 | 100 | 48 | 3.52 | 0.76 |
| Subsampling | 80 | 100 | 64 | 2.48 | 0.57 |
| Subsampling | 40 | 500 | 16 | 6.21 | 0.83 |
| Subsampling | 40 | 500 | 24 | 5.10 | 0.78 |
| Subsampling | 40 | 500 | 32 | 3.59 | 0.60 |
| Subsampling | 80 | 500 | 32 | 4.36 | 0.83 |
| Subsampling | 80 | 500 | 48 | 3.60 | 0.78 |
| Subsampling | 80 | 500 | 64 | 2.52 | 0.56 |
| Extended | 40 | 100 | - | 8.16 | 0.86 |
| Extended | 80 | 100 | - | 5.16 | 0.81 |
| Extended | 40 | 500 | - | 8.08 | 0.88 |
| Extended | 80 | 500 | - | 5.45 | 0.89 |

Table 2 Results for nonunique_schultz ($z^* \approx -61.36$) with $\alpha=0.1$ based on 100 replications.

A.2. CVaR

A one-stage CVaR problem is used by Lam and Qian (2018):

$$\min_x \left\{ x + \frac{1}{a} E[(\xi - x)_+] \right\}$$

where $(\cdot)_+$ is defined as $\max\{\cdot, 0\}$, $a = 0.1$ and ξ is a drawn from a standard normal distribution. Results are shown in Table 3.

A.3. Scalable Farmer

This example is based on the well-known farmer example from Birge and Louveaux (1997) as modified for stress-testing various pieces of software such as Knueven et al. (2020, updated 2022). To make it scalable, two instance creation parameters *cropsmult* and *numscens* are added. The original problem has three crops and three scenarios. The scalable instances have *cropsmult* sets of the original three crops with the characteristics as in the original problem and yields that depend on the scenario. Scenarios are in groups of three with a uniformly distributed pseudo-random number added to the yield values of the original three scenarios.

For the results shown in Tables 4 and 5, we used the original three crops and 1000 scenarios to get an assumed value for z^* .

A.4. Discussion of Results

These experiments are intended mainly to illustrate that the software can be used for such experiments. They do illustrate the unsurprising result that if the samples are too small, the confidence intervals will not be very good. They also suggest that the method we call Extended, which is sort of an afterthought in Eichhorn and Römisich (2007), does not seem to work all that well.

The results are mostly reasonable, but mixed and depend on the availability of enough data as well as the choice of method and parameters. Detailed conclusions are beyond the scope of this small study. A preliminary indication is that bagging might be the best thing to try first.

One other thing to note, though, concerns CVaR. Since CVaR considers the tail, getting good confidence intervals requires a larger value of N . Perhaps for similar reasons, quantile-based intervals do not seem to be as good as Gaussian.

References

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- Lam H, Qian H (2018) Assessing solution quality in stochastic optimization via bootstrap aggregating. *2018 Winter Simulation Conference (WSC)*, 2061–2071 (IEEE).
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| method | N | B | k | avg len | coverage |
|-----------------------------|-----|------|-----|---------|----------|
| Classical_gaussian | 300 | 100 | - | 0.36 | 0.90 |
| Classical_gaussian | 600 | 100 | - | 0.26 | 0.82 |
| Classical_gaussian | 300 | 1000 | - | 0.36 | 0.94 |
| Classical_gaussian | 600 | 1000 | - | 0.26 | 0.80 |
| Classical_quantile | 300 | 100 | - | 0.35 | 0.50 |
| Classical_quantile | 600 | 100 | - | 0.25 | 0.57 |
| Classical_quantile | 300 | 1000 | - | 0.36 | 0.69 |
| Classical_quantile | 600 | 1000 | - | 0.26 | 0.58 |
| Bagging_with_replacement | 300 | 100 | 120 | 0.63 | 1.00 |
| Bagging_with_replacement | 300 | 100 | 180 | 0.62 | 1.00 |
| Bagging_with_replacement | 300 | 100 | 240 | 0.62 | 1.00 |
| Bagging_with_replacement | 600 | 100 | 240 | 0.62 | 1.00 |
| Bagging_with_replacement | 600 | 100 | 360 | 0.63 | 1.00 |
| Bagging_with_replacement | 600 | 100 | 480 | 0.62 | 1.00 |
| Bagging_with_replacement | 300 | 1000 | 120 | 0.20 | 1.00 |
| Bagging_with_replacement | 300 | 1000 | 180 | 0.20 | 1.00 |
| Bagging_with_replacement | 300 | 1000 | 240 | 0.20 | 1.00 |
| Bagging_with_replacement | 600 | 1000 | 240 | 0.20 | 1.00 |
| Bagging_with_replacement | 600 | 1000 | 360 | 0.20 | 1.00 |
| Bagging_with_replacement | 600 | 1000 | 480 | 0.20 | 1.00 |
| Bagging_without_replacement | 300 | 100 | 120 | 0.81 | 1.00 |
| Bagging_without_replacement | 300 | 100 | 180 | 0.99 | 1.00 |
| Bagging_without_replacement | 300 | 100 | 240 | 1.39 | 1.00 |
| Bagging_without_replacement | 600 | 100 | 240 | 0.80 | 1.00 |
| Bagging_without_replacement | 600 | 100 | 360 | 0.99 | 1.00 |
| Bagging_without_replacement | 600 | 100 | 480 | 1.39 | 1.00 |
| Bagging_without_replacement | 300 | 1000 | 120 | 0.26 | 1.00 |
| Bagging_without_replacement | 300 | 1000 | 180 | 0.31 | 1.00 |
| Bagging_without_replacement | 300 | 1000 | 240 | 0.45 | 1.00 |
| Bagging_without_replacement | 600 | 1000 | 240 | 0.26 | 1.00 |
| Bagging_without_replacement | 600 | 1000 | 360 | 0.32 | 1.00 |
| Bagging_without_replacement | 600 | 1000 | 480 | 0.45 | 1.00 |
| Subsampling | 300 | 100 | 120 | 0.36 | 0.65 |
| Subsampling | 300 | 100 | 180 | 0.35 | 0.61 |
| Subsampling | 300 | 100 | 240 | 0.35 | 0.60 |
| Subsampling | 600 | 100 | 240 | 0.26 | 0.64 |
| Subsampling | 600 | 100 | 360 | 0.26 | 0.60 |
| Subsampling | 600 | 100 | 480 | 0.25 | 0.50 |
| Subsampling | 300 | 1000 | 120 | 0.36 | 0.67 |
| Subsampling | 300 | 1000 | 180 | 0.36 | 0.65 |
| Subsampling | 300 | 1000 | 240 | 0.36 | 0.62 |
| Subsampling | 600 | 1000 | 240 | 0.26 | 0.63 |
| Subsampling | 600 | 1000 | 360 | 0.26 | 0.62 |
| Subsampling | 600 | 1000 | 480 | 0.26 | 0.67 |
| Extended | 300 | 100 | - | 0.50 | 0.79 |
| Extended | 600 | 100 | - | 0.36 | 0.71 |
| Extended | 300 | 1000 | - | 0.51 | 0.85 |
| Extended | 600 | 1000 | - | 0.36 | 0.81 |

Table 3 Results for $\text{cvar}(z^* \approx 1.79)$ with $\alpha=0.1$ based on 100 replications.

| method | N | B | k | avg len | coverage |
|-----------------------------|----|------|----|----------|----------|
| Classical_gaussian | 30 | 100 | - | 28259.46 | 0.887 |
| Classical_gaussian | 60 | 100 | - | 20181.52 | 0.892 |
| Classical_gaussian | 30 | 1000 | - | 28294.08 | 0.885 |
| Classical_gaussian | 60 | 1000 | - | 20272.41 | 0.907 |
| Classical_quantile | 30 | 100 | - | 27343.29 | 0.870 |
| Classical_quantile | 60 | 100 | - | 19645.95 | 0.877 |
| Classical_quantile | 30 | 1000 | - | 28236.30 | 0.880 |
| Classical_quantile | 60 | 1000 | - | 20235.36 | 0.905 |
| Bagging_with_replacement | 30 | 100 | 12 | 31422.27 | 0.932 |
| Bagging_with_replacement | 30 | 100 | 18 | 31257.34 | 0.938 |
| Bagging_with_replacement | 30 | 100 | 24 | 32061.31 | 0.932 |
| Bagging_with_replacement | 60 | 100 | 24 | 25164.00 | 0.955 |
| Bagging_with_replacement | 60 | 100 | 36 | 25298.68 | 0.953 |
| Bagging_with_replacement | 60 | 100 | 48 | 25285.28 | 0.950 |
| Bagging_with_replacement | 30 | 1000 | 12 | 28323.03 | 0.895 |
| Bagging_with_replacement | 30 | 1000 | 18 | 28489.61 | 0.902 |
| Bagging_with_replacement | 30 | 1000 | 24 | 28554.10 | 0.900 |
| Bagging_with_replacement | 60 | 1000 | 24 | 20614.13 | 0.907 |
| Bagging_with_replacement | 60 | 1000 | 36 | 20688.71 | 0.907 |
| Bagging_with_replacement | 60 | 1000 | 48 | 20751.51 | 0.905 |
| Bagging_without_replacement | 30 | 100 | 12 | 32638.99 | 0.935 |
| Bagging_without_replacement | 30 | 100 | 18 | 32640.54 | 0.920 |
| Bagging_without_replacement | 30 | 100 | 24 | 32558.19 | 0.930 |
| Bagging_without_replacement | 60 | 100 | 24 | 25546.72 | 0.965 |
| Bagging_without_replacement | 60 | 100 | 36 | 25369.97 | 0.963 |
| Bagging_without_replacement | 60 | 100 | 48 | 25711.17 | 0.970 |
| Bagging_without_replacement | 30 | 1000 | 12 | 29438.81 | 0.902 |
| Bagging_without_replacement | 30 | 1000 | 18 | 29629.62 | 0.905 |
| Bagging_without_replacement | 30 | 1000 | 24 | 29664.24 | 0.895 |
| Bagging_without_replacement | 60 | 1000 | 24 | 21069.04 | 0.915 |
| Bagging_without_replacement | 60 | 1000 | 36 | 21175.03 | 0.917 |
| Bagging_without_replacement | 60 | 1000 | 48 | 21287.06 | 0.922 |
| Subsampling | 30 | 100 | 12 | 21560.28 | 0.777 |
| Subsampling | 30 | 100 | 18 | 17703.34 | 0.680 |
| Subsampling | 30 | 100 | 24 | 12593.48 | 0.502 |
| Subsampling | 60 | 100 | 24 | 15389.09 | 0.785 |
| Subsampling | 60 | 100 | 36 | 12457.75 | 0.680 |
| Subsampling | 60 | 100 | 48 | 8943.31 | 0.500 |
| Subsampling | 30 | 1000 | 12 | 22124.33 | 0.797 |
| Subsampling | 30 | 1000 | 18 | 18209.22 | 0.690 |
| Subsampling | 30 | 1000 | 24 | 12877.64 | 0.510 |
| Subsampling | 60 | 1000 | 24 | 15739.58 | 0.790 |
| Subsampling | 60 | 1000 | 36 | 12903.63 | 0.700 |
| Subsampling | 60 | 1000 | 48 | 9166.89 | 0.525 |
| Extended | 30 | 100 | - | 26890.50 | 0.848 |
| Extended | 60 | 100 | - | 19620.82 | 0.863 |
| Extended | 30 | 1000 | - | 28080.65 | 0.870 |
| Extended | 60 | 1000 | - | 20435.31 | 0.875 |

Table 4 Results for farmer ($z^* \approx -132750.32$) with $\alpha=0.1$ based on 400 replications.

| method | N | B | k | avg len | coverage |
|-----------------------------|----|------|----|----------|----------|
| Classical_gaussian | 30 | 100 | - | 21809.65 | 0.750 |
| Classical_gaussian | 60 | 100 | - | 15622.85 | 0.740 |
| Classical_gaussian | 30 | 1000 | - | 21955.60 | 0.770 |
| Classical_gaussian | 60 | 1000 | - | 15648.47 | 0.730 |
| Classical_quantile | 30 | 100 | - | 21348.92 | 0.740 |
| Classical_quantile | 60 | 100 | - | 15337.17 | 0.740 |
| Classical_quantile | 30 | 1000 | - | 22030.79 | 0.740 |
| Classical_quantile | 60 | 1000 | - | 15641.96 | 0.740 |
| Bagging_with_replacement | 30 | 100 | 12 | 24013.20 | 0.800 |
| Bagging_with_replacement | 30 | 100 | 18 | 24468.50 | 0.790 |
| Bagging_with_replacement | 30 | 100 | 24 | 25117.26 | 0.830 |
| Bagging_with_replacement | 60 | 100 | 24 | 19617.59 | 0.830 |
| Bagging_with_replacement | 60 | 100 | 36 | 19544.47 | 0.820 |
| Bagging_with_replacement | 60 | 100 | 48 | 19788.37 | 0.860 |
| Bagging_with_replacement | 30 | 1000 | 12 | 21923.88 | 0.730 |
| Bagging_with_replacement | 30 | 1000 | 18 | 22105.87 | 0.760 |
| Bagging_with_replacement | 30 | 1000 | 24 | 22159.09 | 0.770 |
| Bagging_with_replacement | 60 | 1000 | 24 | 16040.28 | 0.750 |
| Bagging_with_replacement | 60 | 1000 | 36 | 16049.88 | 0.760 |
| Bagging_with_replacement | 60 | 1000 | 48 | 16071.97 | 0.760 |
| Bagging_without_replacement | 30 | 100 | 12 | 25178.82 | 0.830 |
| Bagging_without_replacement | 30 | 100 | 18 | 25316.79 | 0.780 |
| Bagging_without_replacement | 30 | 100 | 24 | 25361.81 | 0.790 |
| Bagging_without_replacement | 60 | 100 | 24 | 19903.04 | 0.890 |
| Bagging_without_replacement | 60 | 100 | 36 | 19887.07 | 0.860 |
| Bagging_without_replacement | 60 | 100 | 48 | 19734.46 | 0.880 |
| Bagging_without_replacement | 30 | 1000 | 12 | 22641.34 | 0.760 |
| Bagging_without_replacement | 30 | 1000 | 18 | 23047.99 | 0.780 |
| Bagging_without_replacement | 30 | 1000 | 24 | 23065.88 | 0.770 |
| Bagging_without_replacement | 60 | 1000 | 24 | 16416.91 | 0.750 |
| Bagging_without_replacement | 60 | 1000 | 36 | 16428.75 | 0.780 |
| Bagging_without_replacement | 60 | 1000 | 48 | 16453.14 | 0.760 |
| Subsampling | 30 | 100 | 12 | 17165.90 | 0.700 |
| Subsampling | 30 | 100 | 18 | 13965.11 | 0.550 |
| Subsampling | 30 | 100 | 24 | 9980.19 | 0.400 |
| Subsampling | 60 | 100 | 24 | 12253.03 | 0.640 |
| Subsampling | 60 | 100 | 36 | 9822.92 | 0.520 |
| Subsampling | 60 | 100 | 48 | 6949.98 | 0.370 |
| Subsampling | 30 | 1000 | 12 | 17245.27 | 0.670 |
| Subsampling | 30 | 1000 | 18 | 14230.53 | 0.550 |
| Subsampling | 30 | 1000 | 24 | 10112.13 | 0.410 |
| Subsampling | 60 | 1000 | 24 | 12336.30 | 0.630 |
| Subsampling | 60 | 1000 | 36 | 10034.51 | 0.510 |
| Subsampling | 60 | 1000 | 48 | 7141.78 | 0.350 |
| Extended | 30 | 100 | - | 21392.68 | 0.760 |
| Extended | 60 | 100 | - | 16125.62 | 0.730 |
| Extended | 30 | 1000 | - | 21603.71 | 0.770 |
| Extended | 60 | 1000 | - | 16038.97 | 0.780 |

Table 5 Results for farmer ($z^* \approx -132750.32$) with $\alpha=0.2$ based on 100 replications.