Both Nature and SB use Simulated Annealing (SA) [5] as a baseline for comparison. Table 2 of [2] gives a concise comparison of hyperparameters used by the two works. As in [3], we implement and run SA based on the description given in the SB manuscript. Our implementation differs from that described in Nature in its use of move and shuffle in addition to swap, shift and mirror actions. We also use two initial macro placement schemes, i.e., "spiral macro placement" whereby macros are sequentially placed around the boundary of the chip canvas in a counterclockwise spiral manner, and "greedy packer" whereby macros are packed in sequence from the lower-left corner to the top-right corner of the chip canvas [2]. Force-directed (FD) placement is used to update the locations of standard-cell clusters every $\{2n, 3n, 4n, 5n\}$ macro actions, where n is the number of hard macros; FD is not itself an action. The SA cost function is the proxy cost, which consists of wirelength, density and congestion.

As described in [3], Google's implementations of FD and proxy cost calculation are not open-sourced, but are only available via the plc_client in [7]. For speed and transparency, our SA experiments use our own C++ reimplementations of FD and proxy cost calculation; however, the Google plc_client is used for final FD soft macro placement and proxy cost evaluation at the end of the SA run. (Section 3.2 in [3] provides details of force-directed placement and proxy cost congestion.) Our SA codes are open-sourced in MacroPlacement [8].

A Go-With-the-Winners Enhancement. Similar to *Nature* and SB, our previous work in [3] runs 320 SA workers in parallel for 12.5 hours. Each worker, with its own hyperparameter setting, operates independently and does not communicate with other workers. Then, the macro placement solution with minimum proxy cost (as calculated by our C++ code) is used as the final SA solution. However, running 320 SA workers in parallel requires multiple servers, which may be impractical for users with limited computing resources.

To obtain a stronger SA baseline, we adopt the "go-with-the-winners" (GWTW) scheme [1] in a multi-threading implementation. In essence, GWTW allows a set of solution threads to proceed independently, but periodically executes a 'syncup' whereby (i) the best threads are identified, (ii) their solutions are cloned to fill up the entire set of threads, and (iii) the threads then independently continue the solution process until the next 'sync-up'. This approach has seen previous adoption in physical design, e.g., for gate sizing [4].

The detailed algorithm is shown in Algorithm 1. The algorithm can be divided into following steps:

- Lines 1-10: We initialize all the SA workers in parallel. Each SA worker uses a unique random seed to shuffle same-size macros. Placement is initialized via spiral initialization for workers with odd IDs and greedy packing for those with even IDs.
- Lines 14-17: Each SA worker is run for sync_iter iterations in parallel. sync_iter is set based on sync_freq. We use sync_freq = 0.1, meaning there are 9 synchronizations among the workers.
- Lines 18-21: The algorithm stops when *Iter* iterations are performed; otherwise, *syncWorkers* selects the top *k* workers based on proxy cost and replicates their macro locations and orientations to the remaining workers evenly.

• Line 22: writes out the best macro placement solution in terms of proxy cost for each worker.

```
Algorithm 1: Simulated Annealing
```

```
Input: Random seeds: seed = 1,
             Number of iterations: Iters,
             N \times \# macro moves per iteration (N = 20),
             Initial temperature: T_0 = 0.005,
             Minimum temperature: T_{min} = 1 \times 10^{-8},
             Cooling rate: \alpha = \exp\left(\frac{\ln(T_{min}/T_0)}{Iters}\right),
             Number of workers: W = 80,
             Replicated top k = 8 workers,
             Synchronization frequency: sync\_freq = 0.1
   Output: Macro placement solutions.
   // SA wrapped Go-With-the-Winners
 1 workers \leftarrow create W workers;
   for i \leftarrow 0 to W - 1 in parallel do
       workers[i].seed \leftarrow seed + i;
       workers[i].N \leftarrow N;
 4
       workers[i].T \leftarrow T_0;
 5
       workers[i].\alpha \leftarrow \alpha;
 6
       if (i \mod 2) = 0 then
 7
           workers[i].macro_placement \leftarrow "spiral macro placement";
 8
 9
          workers[i].macro_placement \leftarrow "greedy packer";
11 iter_count \leftarrow 0;
12 sync_iter \leftarrow Iters \times sync\_freq;
13 while true do
       end\_iter \leftarrow min(Iters, iter\_count + sync\_iter);
14
       for i \leftarrow 0 to W - 1 in parallel do
15
           Each worker performs (end_iter – iter_count) SA iterations;
16
             applying N \times \#macro moves per iteration and updating
             temperature;
       iter\_count \leftarrow end\_iter;
17
       \mathbf{if} \ \mathit{iter\_count} = \mathit{Iters} \ \mathbf{then}
18
          break;
19
       candidate_solutions \leftarrow extractTopK(workers, k);
20
       Evenly distribute these top-k solutions across all the workers;
```

22 Write out the best solution of each worker.

As noted above, after placing soft macros (standard-cell clusters) with GWTW SA, we use CT's plc_client to evaluate the proxy cost of the best macro place-

ment solutions for each worker, and then return the best solution in terms of proxy cost for P&R evaluation. In our runs, the probabilities for five solution move operators (swap, shift, move, shuffle and flip) are respectively set to 0.24, 0.24, 0.24 and 0.04. The number of iterations is set to ensure that overall runtime for each testcase is less than 12 hours on our slowest CPU server.¹

Relative to the SA implementation in [3], our present SA implementation achieves similar or better results while using only one-fourth of the CPU resources: 80 threads instead of 320 threads, enabling execution on a single CPU server. Further, to ensure exact reproducibility across different platforms, (i) we use lookup tables for exponent computation and provide a binarized version of the lookup table in our repository; and (ii) we provide scripts to generate Docker and Singularity images that reproduce the same environment. Our testing across a range of Intel Xeon Gold and AMD EPYC CPUs confirms exact matching of SA solutions obtained by all 80 workers using the same Docker or Singularity image.

References

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 $^{^1}For$ Ariane, BlackParrot, MemPoolGroup, Ariane-X2 and Ariane-X4, we set the number of iterations to 18K, 9K, 4.5K, 9K and 4K, respectively. This corresponds, e.g., to $\sim\!\!11$ hours on an Intel Xeon Gold 6148 CPU, or $\sim\!\!3$ hours on an AMD EPYC 9684X CPU.