Experiment results log.

Planned order of scenarios

| Scenarios | Parameters | Week |
|-----------|----------------|---------------------|
| A1 | N | March 23 - March 29 |
| A2 | В | March 30 - April 5 |
| A3 | Datasets | March 30 - April 5 |
| B1 | Chain Strength | April 6 - April 12 |
| B2 | Embedding | April 13 - April 19 |
| В3 | Shots | April 20 - April 26 |
| B4 | Annealing | April 27 - May 3 |

Actual order of scenarios

| Scenarios | Parameters | Week |
|-----------|----------------|---------------------|
| A1 | N | March 23 - March 29 |
| B1 | Chain Strength | March 30 - April 5 |
| A2 | В | April 6 - April 12 |
| В3 | Shots | April 13 - April 19 |
| A2B3 | B and Shots | April 20 - April 26 |
| B2 | Embedding | April 20 - April 26 |
| B4 | Annealing | April 27 - May 3 |
| A3 | Datasets | April 27 - May 3 |

Sidenotes to research about

• Find what is the maximum N value that is supported by dwave

Scenario A1 - N

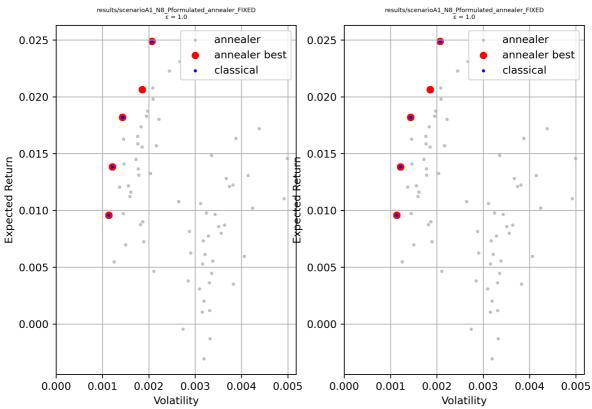
We started by experimenting several values of N, in order to find the maximum possible value of N that could be solved in a reasonable time by the classical solver.

The N values are: 8, 16, 32, and 64. P was calculated as $P = -q * min_sigma + max_mu$

For this scenario, we used the "diversified" dataset and 1000 shots per execution. The q_values are listed in the following table:

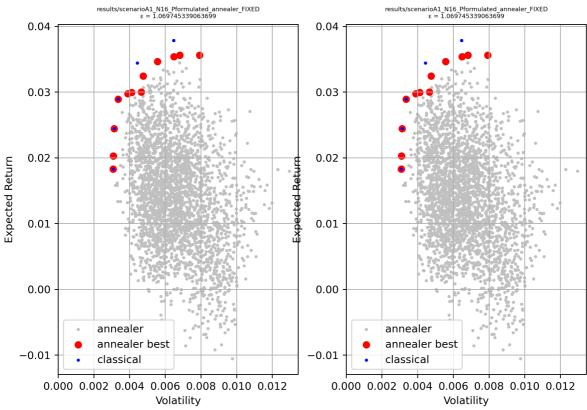
| N | q values | Epsilon Indicator |
|----|--|-------------------|
| 8 | 0, 11, 20, 54 | 1.0 |
| 16 | 0, 2, 6, 100, 500 | 1.070 |
| 32 | 0, 0.4, 0.9, 2, 3, 9, 100 | 1.967 |
| 64 | 0, 0.2, 0.4, 0.6, 1.1, 1.3, 1.5, 2, 5, 6, 7, 8, 10, 100, 500 | 2.022 |

Epsilon Indicator - scenario1Y2021M04D18h23m07s48



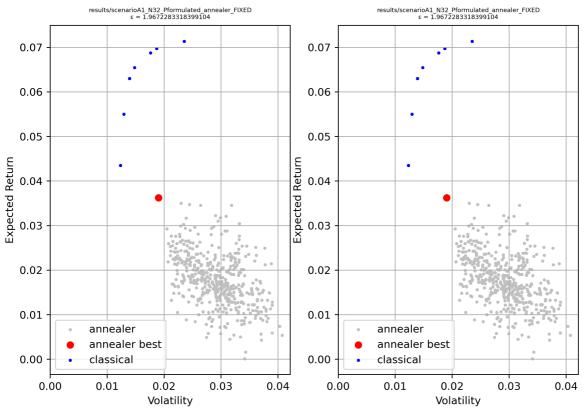
How to interpret: Blue markers are part of the efficient frontier. The epsilon indicator is the minimum factor by which the red set has to be multiplied in the objective so as to weakly dominate Hence, the closer to 1 is the epsilon indicator, the better the red set.

Epsilon Indicator - scenario1Y2021M04D18h23m08s05



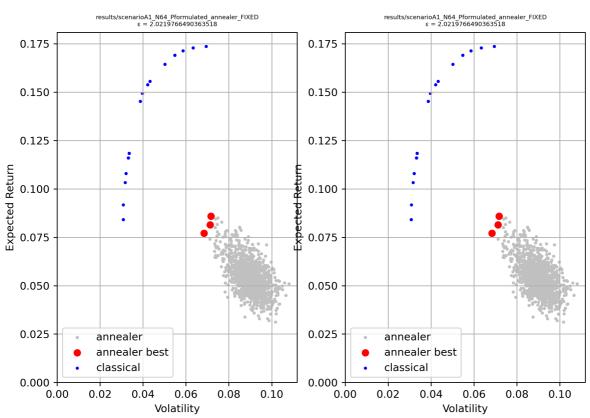
How to interpret: Blue markers are part of the efficient frontier. The epsilon indicator is the minimum factor by which the red set has to be multiplied in the objective so as to weakly dominate Hence, the closer to 1 is the epsilon indicator, the better the red set.

Epsilon Indicator - scenario1Y2021M04D18h23m08s14



How to interpret: Blue markers are part of the efficient frontier. The epsilon indicator is the minimum factor by which the red set has to be multiplied in the objective so as to weakly dominate Hence, the closer to 1 is the epsilon indicator, the better the red set.

Epsilon Indicator - scenario1Y2021M04D18h23m08s20



How to interpret: Blue markers are part of the efficient frontier. The epsilon indicator is the minimum factor by which the red set has to be multiplied in the objective so as to weakly dominate Hence, the closer to 1 is the epsilon indicator, the better the red set.

Key Takeaways:

As expected, the epsilon indicator increases with the N value. However, during those executions, dwave's problem inspector warned that the chains were too weak, and that, in the case of N=64, all samples had

broken chains. Based on this warning, we decided to immediately execute scenario B1, changing the original order of scenarios.

Scenario B1 - Chain Strength

Looking at the fraction of chain breaks in Scenario A1, we know that on average each sample had almost a third (0.31) of its chains broken when N=32. This fraction increases to over half (0.54) when N=64! Those values are very high and are another clue that the chain strength needs to be adjusted, especially for those values of N.

A good starting value for the chain strength is the maximum absolute value (maxAbs) of the QUBO matrix. However, this is not always the most optimal value. We need to test several values based on this initial value. By testing those values, we can find a value near the sweet spot between the probability that the chains are intact and the probability of finding optimal values. Refer to:

https://www.dwavesys.com/sites/default/files/2_Wed_Am_PerfTips.pdf

We have three tables, one for the epsilon indicator, one for the fractions of valid solutions, and one for the average fractions of chain breaks.

Starting with the average fractions of chain breaks (Lower is better):

| Chain strength | N8 | N16 | N32 | N64 |
|----------------|---------|---------|---------|---------|
| default value | 0.00081 | 0.01153 | 0.31350 | 0.54426 |
| 0.125 * maxAbs | 0.00397 | 0.02741 | 0.31014 | 0.38301 |
| 0.250 * maxAbs | 0.00034 | 0.00106 | 0.00170 | 0.00683 |
| 0.375 * maxAbs | 0.00006 | 0.00032 | 0.00111 | 0.00453 |
| 0.500 * maxAbs | 0.00006 | 0.00026 | 0.00149 | 0.00475 |
| 0.625 * maxAbs | 0.00006 | 0.00031 | 0.00112 | 0.00453 |
| 0.750 * maxAbs | 0.00006 | 0.00029 | 0.00130 | 0.00454 |
| 0.875 * maxAbs | 0.00006 | 0.00017 | 0.00102 | 0.00461 |
| 1.000 * maxAbs | 0.00003 | 0.00034 | 0.00100 | 0.00439 |
| 1.125 * maxAbs | 0.00000 | 0.00030 | 0.00119 | 0.00401 |
| 1.250 * maxAbs | 0.00000 | 0.00042 | 0.00125 | 0.00419 |
| 1.375 * maxAbs | 0.00006 | 0.00028 | 0.00108 | 0.00424 |
| 1.500 * maxAbs | 0.00009 | 0.00025 | 0.00201 | 0.00430 |

Next, we obtained the following fractions of valid solutions (Higher is better):

| Chain strength | N8 | N16 | N32 | N64 |
|----------------|-------|-------|-------|-------|
| default value | 0.877 | 0.688 | 0.121 | 0.094 |

| Chain strength | N8 | N16 | N32 | N64 |
|----------------|-------|-------|-------|-------|
| 0.125 * maxAbs | 0.001 | 0.002 | 0.076 | 0.205 |
| 0.250 * maxAbs | 0.934 | 0.622 | 0.395 | 0.243 |
| 0.375 * maxAbs | 0.848 | 0.543 | 0.325 | 0.220 |
| 0.500 * maxAbs | 0.781 | 0.485 | 0.299 | 0.186 |
| 0.625 * maxAbs | 0.703 | 0.444 | 0.261 | 0.172 |
| 0.750 * maxAbs | 0.665 | 0.388 | 0.252 | 0.170 |
| 0.875 * maxAbs | 0.630 | 0.406 | 0.242 | 0.163 |
| 1.000 * maxAbs | 0.598 | 0.366 | 0.235 | 0.151 |
| 1.125 * maxAbs | 0.594 | 0.370 | 0.219 | 0.148 |
| 1.250 * maxAbs | 0.556 | 0.342 | 0.223 | 0.129 |
| 1.375 * maxAbs | 0.540 | 0.330 | 0.212 | 0.136 |
| 1.500 * maxAbs | 0.512 | 0.310 | 0.198 | 0.138 |

Finally, we obtained the following epsilon indicators (Lower is better):

| Chain strength | N8 | N16 | N32 | N64 |
|----------------|-------|--------|-------|-------|
| default value | 1,000 | 1,070 | 1,967 | 2,022 |
| 0,125 * maxAbs | 1,368 | 18,844 | 1,767 | 1,977 |
| 0,250 * maxAbs | 1,000 | 1,075 | 1,178 | 1,474 |
| 0,375 * maxAbs | 1,000 | 1,057 | 1,203 | 1,580 |
| 0,500 * maxAbs | 1,000 | 1,099 | 1,331 | 1,500 |
| 0,625 * maxAbs | 1,000 | 1,098 | 1,269 | 1,410 |
| 0,750 * maxAbs | 1,000 | 1,120 | 1,429 | 1,523 |
| 0,875 * maxAbs | 1,000 | 1,123 | 1,430 | 1,587 |
| 1,000 * maxAbs | 1,000 | 1,119 | 1,250 | 1,526 |
| 1,125 * maxAbs | 1,000 | 1,099 | 1,142 | 1,539 |
| 1,250 * maxAbs | 1,000 | 1,092 | 1,355 | 1,610 |
| 1,375 * maxAbs | 1,000 | 1,110 | 1,352 | 1,465 |
| 1,500 * maxAbs | 1,000 | 1,101 | 1,345 | 1,423 |

To validate such results, this scenario has been repeated for N=16, N=32, and N=64.

| Chain strength | N16 | N32 | N64 |
|----------------|-----|-----|-----|
| | | | |

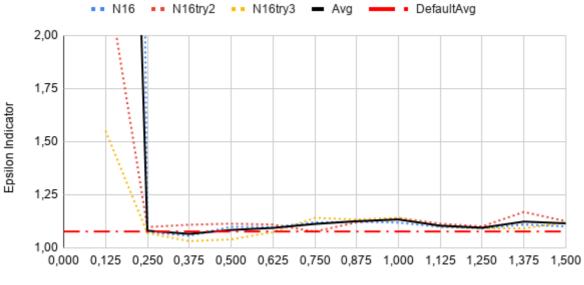
| Chain strength | N16 | N32 | N64 |
|----------------|-------|-------|-------|
| default value | 1,092 | 1,739 | 1,981 |
| 0,125 * maxAbs | 2,314 | 1,813 | 2,185 |
| 0,250 * maxAbs | 1,098 | 1,256 | 1,550 |
| 0,375 * maxAbs | 1,109 | 1,336 | 1,492 |
| 0,500 * maxAbs | 1,114 | 1,257 | 1,502 |
| 0,625 * maxAbs | 1,110 | 1,322 | 1,503 |
| 0,750 * maxAbs | 1,077 | 1,299 | 1,516 |
| 0,875 * maxAbs | 1,120 | 1,307 | 1,489 |
| 1,000 * maxAbs | 1,141 | 1,350 | 1,485 |
| 1,125 * maxAbs | 1,114 | 1,327 | 1,430 |
| 1,250 * maxAbs | 1,101 | 1,266 | 1,549 |
| 1,375 * maxAbs | 1,169 | 1,198 | 1,508 |
| 1,500 * maxAbs | 1,126 | 1,325 | 1,597 |

And one more time:

| Chain strength | N16 | N32 | N64 |
|----------------|-------|-------|-------|
| default value | 1,070 | 1,728 | 1,988 |
| 0,125 * maxAbs | 1,551 | 1,760 | 1,906 |
| 0,250 * maxAbs | 1,070 | 1,266 | 1,462 |
| 0,375 * maxAbs | 1,032 | 1,235 | 1,583 |
| 0,500 * maxAbs | 1,040 | 1,325 | 1,514 |
| 0,625 * maxAbs | 1,074 | 1,332 | 1,551 |
| 0,750 * maxAbs | 1,141 | 1,270 | 1,458 |
| 0,875 * maxAbs | 1,134 | 1,229 | 1,515 |
| 1,000 * maxAbs | 1,141 | 1,252 | 1,536 |
| 1,125 * maxAbs | 1,101 | 1,248 | 1,547 |
| 1,250 * maxAbs | 1,092 | 1,303 | 1,523 |
| 1,375 * maxAbs | 1,092 | 1,247 | 1,560 |
| 1,500 * maxAbs | 1,120 | 1,391 | 1,519 |
| 5,000 * maxAbs | 1.177 | 1,297 | 1,627 |

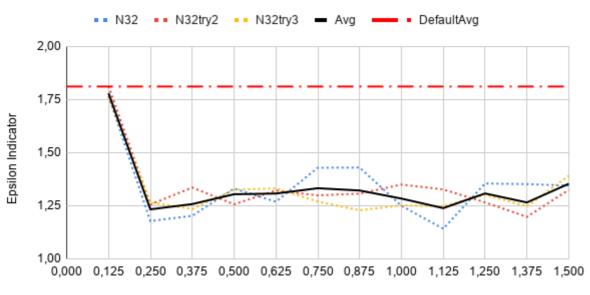
The results are summarized in the following charts.



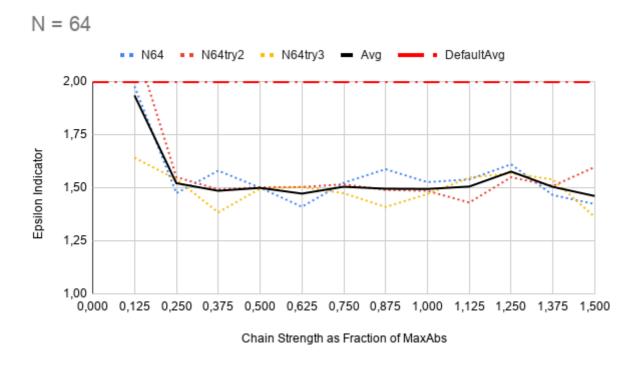








Chain Strength as Fraction of MaxAbs



Key Takeaways:

Looking at the results, we notice that the impact of any change to the chain strength is higher for higher values of N.

It also becomes clear that, especially for higher values of N, the default chain strength is far from being the best value. It seems that for higher values of N, the farther is the default chain strength value from the best value.

Another thing that also becomes clear is that the fractions of chain breaks and valid solutions are not directly synonymous with the quality of the solutions.

For the case of N=8, every try gave a perfect score of 1.000.

For the case N=16, the epsilon values are so similar that they fall under the margin of variation. Thus we cannot place conclusions based on these results. (Note: in this case, the default strength is always the best!)

There is an exception for both cases of N=8 and N=16. When chain_strength = 0.125 * maxAbs there is a high fraction of chain breaks and almost no samples are valid solutions. Thus, for this value of chain strength, the results are very bad.

This behavior is also noticeable for N=32 and N=64, that present a relatively high epsilon indicator with this chain strength.

It seems that, after this very weak chain strength, the following values of chain strength rapidly attain the lowest epsilon indicators registered, with a very slow climb afterwards.

In the end, the results suggest that it is okay to choose any value that is part of the slow climb. However, from theory, we know that we should avoid any value over 1.000 * maxAbs, since it scales down the problem.

Therefore, for all N values, a safe range seems to be between 0.250 * maxAbs and 1.000 * maxAbs.

Based on those findings, the case N=8 will not be tested in the remaining scenarios, since the annealer already achieved optimality.

Scenario A2 - B OLD

For this scenario, we will be looking at how different budgets affect the performance of the annealer. Therefore, different fractions of B are going to be tested: 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, and 0.9.

I will be using chain_strength = 1.000 * maxAbs, for the reasons explained in the previous scenario.

Reminder: the fraction used in previous scenarios was B=0.5!

Obviously, for each value of B, we first need to solve it classically. Then, from the results, we get the sequences of q_values to be used in the annealer.

| N | q values | Budget fraction |
|----|--|-----------------|
| 16 | 0, 20, 500 | 0.1 (1) |
| 32 | 0, 7, 20, 40 | 0.1 (3) |
| 64 | 0, 0.6, 2, 4, 6, 8, 20, 40, 80, 500 | 0.1 (6) |
| 16 | 0, 8, 10, 40 | 0.2 (3) |
| 32 | 0, 5, 8, 20, 30, 80 | 0.2 (6) |
| 64 | 0, 0.3, 0.8, 2, 4, 5, 7, 9, 20, 30, 500 | 0.2 (12) |
| 16 | 0, 2, 6, 20, 60 | 0.3 (4) |
| 32 | 0, 3, 4, 10, 20, 50 | 0.3 (9) |
| 64 | 0, 0.2, 2, 3, 4, 5, 7, 9, 20, 30, 100 | 0.3 (19) |
| 16 | 0, 2, 5, 10, 30 | 0.4 (6) |
| 32 | 0, 0.2, 0.9, 2, 4, 20, 30, 70, 500 | 0.4 (12) |
| 64 | 0, 0.3, 0.6, 1, 2, 3, 4, 6, 8, 20, 30, 90 | 0.4 (25) |
| 16 | 0, 2, 6, 100, 500 | 0.5 (8) |
| 32 | 0, 0.4, 0.9, 2, 3, 9, 100 | 0.5 (16) |
| 64 | 0, 0.2, 0.4, 0.6, 1.1, 1.3, 1.5, 2, 5, 6, 7, 8, 10, 100, 500 | 0.5 (32) |
| 16 | 0, 0.1, 0.8, 3, 20, 30 | 0.6 (9) |
| 32 | 0, 0.1, 0.5, 1, 2, 3, 7, 8, 20, 30 | 0.6 (19) |
| 64 | 0, 0.1, 0.2, 0.3, 0.4, 0.6, 0.7, 2, 3, 7, 9, 20 | 0.6 (38) |
| 16 | 0, 0.7, 20 | 0.7 (11) |
| 32 | 0, 0.4, 2 | 0.7 (22) |
| 64 | 0, 0.1, 0.2, 0.3, 0.7, 1, 2, 3, 4, 6, 20 | 0.7 (44) |
| | | |

| N | q values | Budget fraction |
|----|--|------------------------|
| 16 | 0, 4 | 0.8 (12) |
| 32 | 0, 0.8, 7, 9 | 0.8 (25) |
| 64 | 0, 0.1, 0.2, 0.4, 0.5, 0.6, 1, 2, 3, 6, 20 | 0.8 (51) |
| 16 | 0, 50 | 0.9 (14) |
| 32 | 0, 0.8, 3 | 0.9 (28) |
| 64 | 0, 0.6, 1, 2, 5, 500 | 0.9 (57) |

Question: Is it bad to have different number of samples between cases?

With those results, we obtained the following epsilon indicators:

| Budget fraction | N16 (AvgChainBreak) | N32 (AvgChainBreak) | N64 (AvgChainBreak) |
|-----------------|---------------------|---------------------|---------------------|
| 0,1 | 1,000 (0,00406) | 1,142 (0,00979) | 1,846 (0,01572) |
| 0,2 | 1,000 (0,00048) | 1,334 (0,00151) | 2,929 (0,00493) |
| 0,3 | 1,026 (0,00044) | 1,373 (0,00152) | 1,750 (0,00473) |
| 0,4 | 1,113 (0,00024) | 1,281 (0,00134) | 1,640 (0,00452) |
| 0,5 | 1,075 (0,00031) | 1,311 (0,00127) | 1,513 (0,00484) |
| 0,6 | 1,146 (0,00033) | 1,293 (0,00141) | 1,441 (0,00470) |
| 0,7 | 1,162 (0,00038) | 1,421 (0,00108) | 1,907 (0,00464) |
| 0,8 | 1,005 (0,00031) | 1,408 (0,00144) | inf (0,00462) |
| 0,9 | 1,103 (0,00034) | inf (0,00129) | inf (0,00447) |

Gráficos com os resultados deste cenário estão no cenário seguinte

Key Takeaways:

The first thing I notice is that there is a high chain break fraction when the budget is B=0.1. Afterwards, it attains a consistently low fraction, with small variation.

For N=32 and N=64, as expected from theory, the epsilon indicator is lower when the budget is or is close to B=0.5, since this value has the highest number of admissible solutions.

Behavior for all N values is hard to grasp. Nonetheless, budget fraction is a parameter that is particular to each practitioner.

Scenario B3 - Shots OLD

The previous scenario, A2, made us wonder about the number of samples. That is, there is a possibility that the cases where B is farthest from B=0.5 have worse performance because of having less values of q and thus less samples taken.

Therefore, we pose a question: Is it better to increase the number of shots per value of q or to add more values of q to be executed?

Since the results so far seem to have a good coverage of the efficient frontier, but still far from it, we believe that the issue is related to the number of samples per value of q. Hence, we are going to repeat the previous scenario with a new methodology to define the number of samples per value of q. This methodology is called Allocated.

Initially, each value of q had 1000 shots, i.e., 1000 samples taken. This time, each case will have a total allocated number of shots for every value of q. For example, if we have a case with three values of q and another case with five values of q, then, with a total allocation of 5000 shots per case, then the first case will have 1666 shots per value, while the second case will have 1000 shots per value.

Based on this methodology, we will start with a total allocation of 15000 shots, such that each of the 15 values of q from case B=0.5 have 1000 shots.

| N | q values | Budget fraction | Shots per value of q |
|----|--|------------------------|----------------------|
| 16 | 0, 20, 500 | 0.1 (1) | 5000 |
| 32 | 0, 7, 20, 40 | 0.1 (3) | 3750 |
| 64 | 0, 0.6, 2, 4, 6, 8, 20, 40, 80, 500 | 0.1 (6) | 1500 |
| 16 | 0, 8, 10, 40 | 0.2 (3) | 3750 |
| 32 | 0, 5, 8, 20, 30, 80 | 0.2 (6) | 2500 |
| 64 | 0, 0.3, 0.8, 2, 4, 5, 7, 9, 20, 30, 500 | 0.2 (12) | 1363 |
| 16 | 0, 2, 6, 20, 60 | 0.3 (4) | 3000 |
| 32 | 0, 3, 4, 10, 20, 50 | 0.3 (9) | 2500 |
| 64 | 0, 0.2, 2, 3, 4, 5, 7, 9, 20, 30, 100 | 0.3 (19) | 1363 |
| 16 | 0, 2, 5, 10, 30 | 0.4 (6) | 3000 |
| 32 | 0, 0.2, 0.9, 2, 4, 20, 30, 70, 500 | 0.4 (12) | 1666 |
| 64 | 0, 0.3, 0.6, 1, 2, 3, 4, 6, 8, 20, 30, 90 | 0.4 (25) | 1250 |
| 16 | 0, 2, 6, 100, 500 | 0.5 (8) | 3000 |
| 32 | 0, 0.4, 0.9, 2, 3, 9, 100 | 0.5 (16) | 2142 |
| 64 | 0, 0.2, 0.4, 0.6, 1.1, 1.3, 1.5, 2, 5, 6, 7, 8, 10, 100, 500 | 0.5 (32) | 1000 |
| 16 | 0, 0.1, 0.8, 3, 20, 30 | 0.6 (9) | 2500 |
| 32 | 0, 0.1, 0.5, 1, 2, 3, 7, 8, 20, 30 | 0.6 (19) | 1500 |
| 64 | 0, 0.1, 0.2, 0.3, 0.4, 0.6, 0.7, 2, 3, 7, 9, 20 | 0.6 (38) | 1250 |
| 16 | 0, 0.7, 20 | 0.7 (11) | 5000 |
| 32 | 0, 0.4, 2 | 0.7 (22) | 5000 |
| | | | |

| N | q values | Budget fraction | Shots per value of q |
|----|--|------------------------|----------------------|
| 64 | 0, 0.1, 0.2, 0.3, 0.7, 1, 2, 3, 4, 6, 20 | 0.7 (44) | 1363 |
| 16 | 0, 4 | 0.8 (12) | 7500 |
| 32 | 0, 0.8, 7, 9 | 0.8 (25) | 3750 |
| 64 | 0, 0.1, 0.2, 0.4, 0.5, 0.6, 1, 2, 3, 6, 20 | 0.8 (51) | 1363 |
| 16 | 0, 50 | 0.9 (14) | 7500 |
| 32 | 0, 0.8, 3 | 0.9 (28) | 5000 |
| 64 | 0, 0.6, 1, 2, 5, 500 | 0.9 (57) | 2500 |

We obtained the following epsilon indicators:

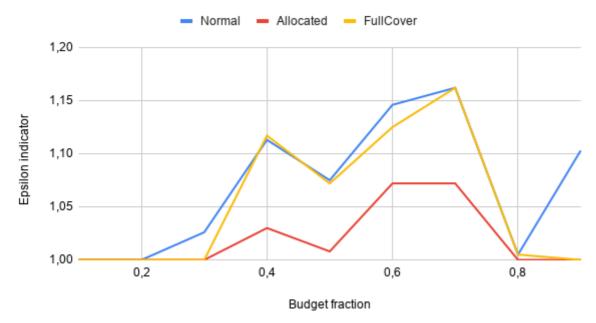
| Budget fraction | N16 | N32 | N64 |
|-----------------|-------|-------|-------|
| 0,1 | 1,000 | 1,069 | 1,667 |
| 0,2 | 1,000 | 1,295 | 2,434 |
| 0,3 | 1,000 | 1,293 | 1,651 |
| 0,4 | 1,030 | 1,277 | 1,632 |
| 0,5 | 1,008 | 1,271 | 1,507 |
| 0,6 | 1,072 | 1,298 | 1,521 |
| 0,7 | 1,072 | 1,333 | 1,926 |
| 0,8 | 1,000 | 1,413 | inf |
| 0,9 | 1,000 | inf | inf |

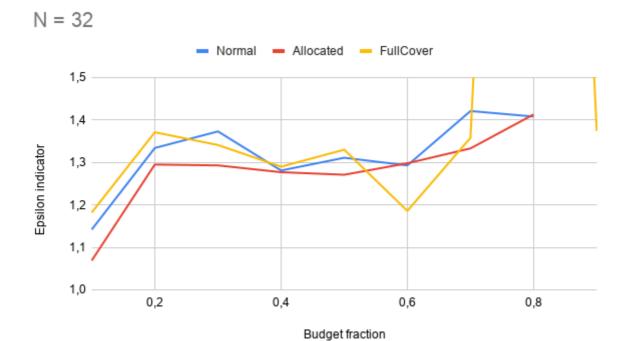
However, we need to take into account that in real case scenarios, we won't be able to have these carefully chosen values of q. In fact, they were discovered because it was feasible to classically solve these scenarios! For this reason, we introduce another methodology, called FullCoverage. This methodology will execute the same values of q for every scenario. The list of values of q is based on guesswork and gained experience with the given scenarios: 0, 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 1000. As with Allocated methodology, this list is allocated to a total of 15000 samples (500 per value of q).

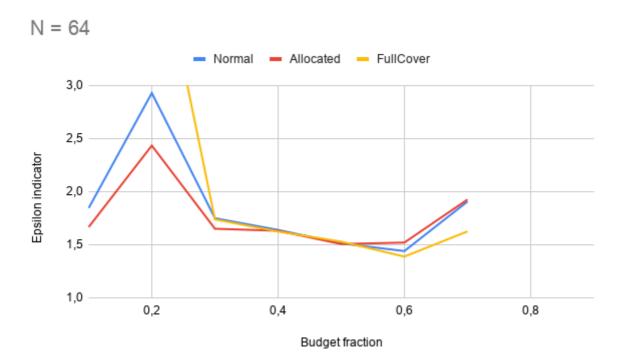
| Budget fraction | N16 | N32 | N64 |
|------------------------|-------|-------|-------|
| 0,1 | 1,000 | 1,182 | 5,000 |
| 0,2 | 1,000 | 1,371 | 4,625 |
| 0,3 | 1,000 | 1,341 | 1,739 |
| 0,4 | 1,117 | 1,290 | 1,625 |
| 0,5 | 1,072 | 1,330 | 1,530 |

| Budget fraction | N16 | N32 | N64 |
|------------------------|-------|-------|-------|
| 0,6 | 1,125 | 1,186 | 1,389 |
| 0,7 | 1,162 | 1,358 | 1,626 |
| 0,8 | 1,005 | 4,442 | inf |
| 0,9 | 1,000 | 1,374 | inf |









Key Takeaways:

Compared to the previous methodology, called Simple, the Allocated methodology brings improvements in almost every case. This is expected, since all the cases had their number of samples increased, minus the case N=64 B=0.5, which keeps the same number of samples (and also has the same performance in both methodologies).

When looking at the more "realistic" FullCover methodology, the results are not the best, but don't fall shortly compared to Allocated.

For the next scenarios, we are going to use the Allocated methodology, as well as B=0.5.

Scenario A2 and B3 - B and Shots

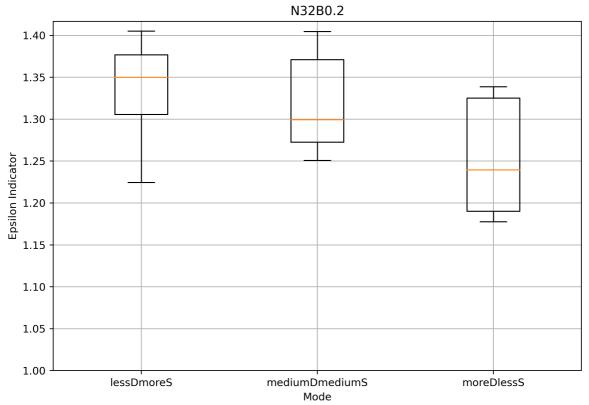
2 factors: B and Shots

"B" factor has three levels: Small Budget, Medium Budget and Large Budget (fractions 0.2, 0.5, and 0.8, respectively).

"Shots" factor has three levels: Less directions and More shots per direction, Medium directions and Medium shots per direction, More directions and Less shots per direction (codenamed lessDmoreS, mediumDmediumS, and moreDlessS, respectively).

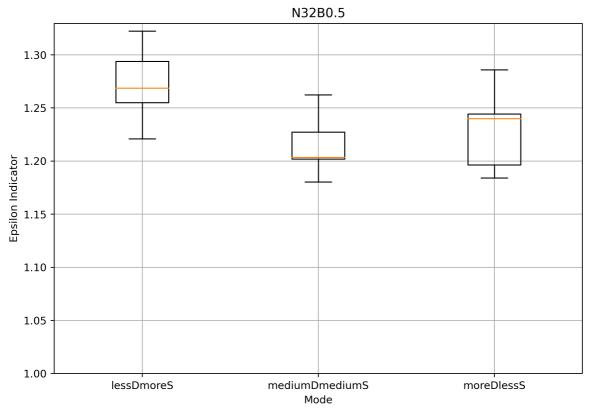
Starting with N=32:

Boxplots - scenario1Y2021M04D27h19m08s33



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set. Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

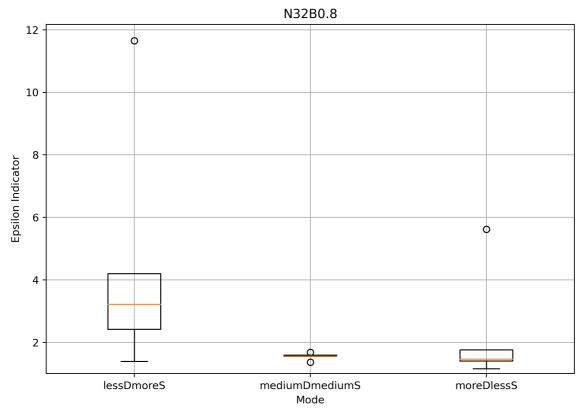
Boxplots - scenario1Y2021M04D27h19m09s04



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set.

Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

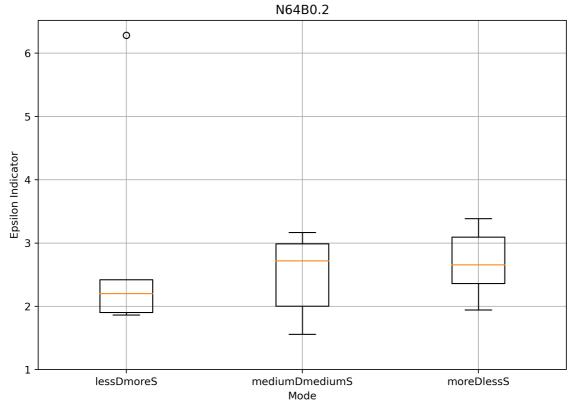
Boxplots - scenario1Y2021M04D27h19m05s21



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set. Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

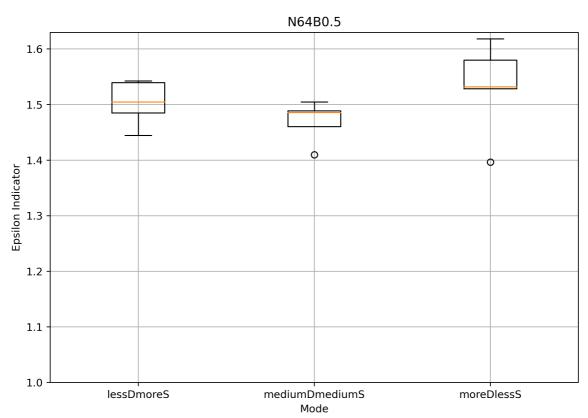
And for N=64:

Boxplots - scenario1Y2021M04D27h19m04s33



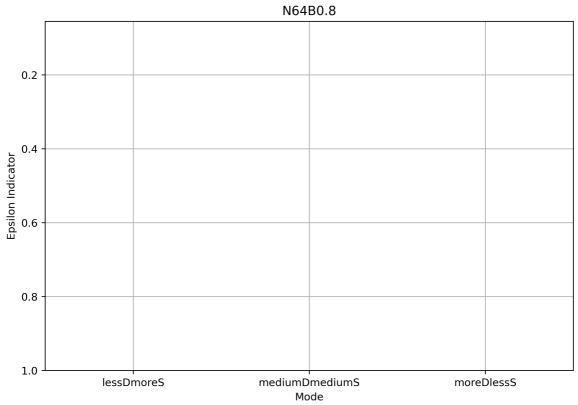
How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set. Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

Boxplots - scenario1Y2021M04D27h19m09s24



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set. Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

Boxplots - scenario1Y2021M04D27h19m04s58



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set.

Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

Key Takeaways:

Looking at N=32, moreDlessS is better when B=0.2. When B=0.5, mediumDmediumS is better. Finally, when B=0.8, moreDlessS is again the best, but closely followed by mediumDmediumS.

Looking at N=64, lessDmoreS is better when B=0.2. When B=0.5, mediumDmediumS is better, followed by lessDmoreS. Finally, when B=0.8, there is nothing displayed, but moreDlessS was the only one to provide a valid answer, with an epsilon indicator of 1.988.

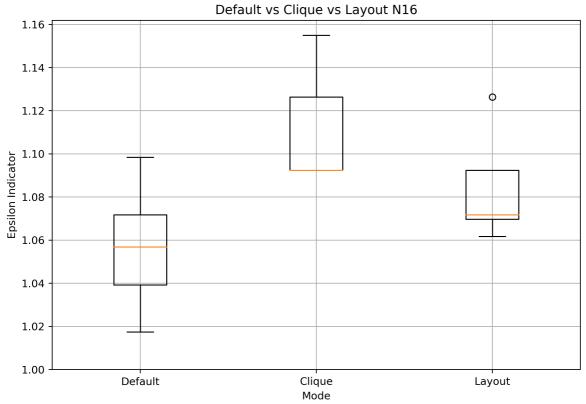
Scenario B2 - Embedding

So far, we used the general embedding. D-Wave offers another two embedding options, clique and layout embeddings. The three options are going to be compared.

| Embedding | N16 | N32 | N64 |
|--------------|-------|-------|-------|
| general try1 | 1,057 | 1,165 | 1,593 |
| general try2 | 1,039 | 1,275 | 1,548 |
| general try3 | 1,072 | 1,275 | 1,568 |
| general try4 | 1,017 | 1,290 | 1,487 |
| general try5 | 1,098 | 1,252 | 1,335 |
| clique try1 | 1,092 | 1,316 | 1,546 |
| clique try2 | 1,092 | 1,320 | 1,510 |
| | | | |

| Embedding | N16 | N32 | N64 |
|-------------|-------|-------|-------|
| clique try3 | 1,155 | 1,381 | 1,428 |
| clique try4 | 1,126 | 1,316 | 1,577 |
| clique try5 | 1,092 | 1,254 | 1,518 |
| layout try1 | 1.070 | 1.250 | 1.454 |
| layout try2 | 1.072 | 1.326 | 1.389 |
| layout try3 | 1.062 | 1.301 | 1.464 |
| layout try4 | 1.092 | 1.248 | 1.472 |
| layout try5 | 1.126 | 1.275 | 1.459 |

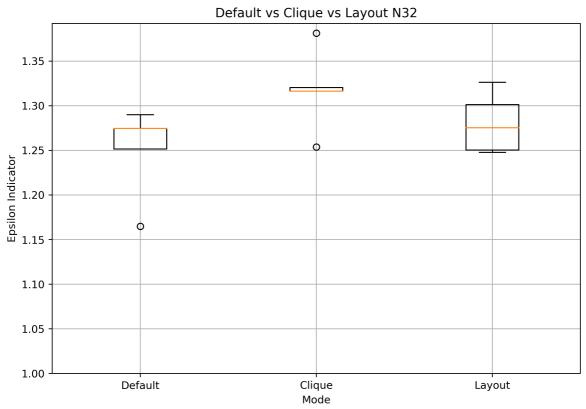
Boxplots - scenario1Y2021M04D30h16m33s35



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set.

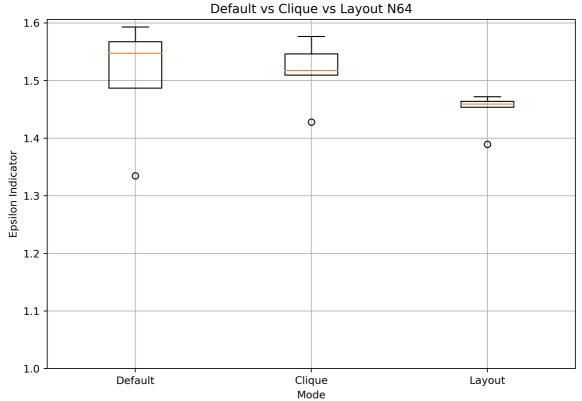
Hence, the closer to 1 is the epsilon indicator, the better the annealer set,

Boxplots - scenario1Y2021M04D30h16m33s10



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set. Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

Boxplots - scenario1Y2021M04D30h16m32s38



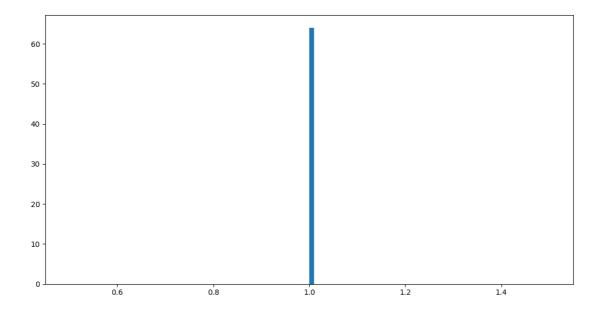
How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set.

Key Takeaways:

It seems that the higher the value of N, the better is the layout embedding compared to the general. Concretely, for N=16 the general embedding shows the best performance, closely followed by layout embedding. For N=32, the gap between those general and layout embeddings gets narrower. Finally, for N=64, general embedding falls short of the other two options, with layout embedding being clearly better. This suggests that, for N>=64, we should choose layout embedding.

In conversations with Jose Pinilla, a Ph.D. student that authored an implementation of a layout-aware embedding, layout embedding is much more suited for *sparse* graphs, which is not the case of the POP. In fact, POP usually generates fully connected graphs. However, Jose Pinilla said "if there are clusters of high connectivity, you'll immediately be rewarded with faster results, or a higher chance of at least finding an embedding". I noticed that, in fact, layout embedding was much faster than the other two options.

It is interesting that those faster results were also accompanied by better performance. Again, in conversations with Jose Pinilla, he provided me with some code to plot a histogram that let us confirm that the graph is in fact fully connected.



The graph is fully connected, which means that there are no clusters of high connectivity and no speed boost should be expected. So, why did it have better performance? This is an interesting question that I pose for further research.

For the remaining scenarios, we will use the layout embedding.

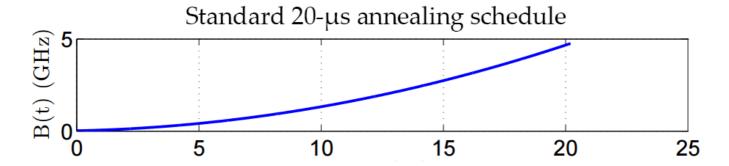
Scenario B4 - Annealing

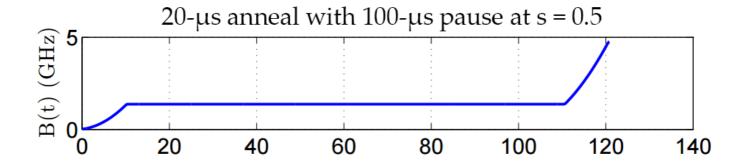
It is time to study the impact of Annealing, if it has any!

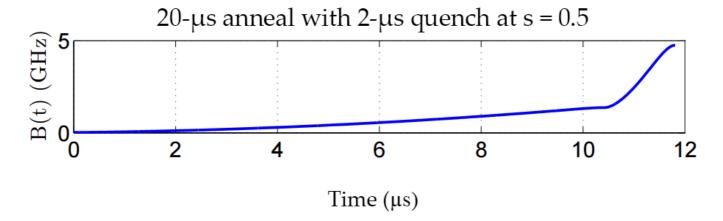
So far, we used the default annealing strategy. We will study another three common strategies that may provide significant improvements to the annealer performance.

- default Standard 20µs annealing schedule
- long Standard 100µs annealing schedule
- pause 20μs anneal with 100μs pause at s=0.5
- quench 20μs anneal with 2μs quench at s=0.5

default, pause, and quench are illustrated in the following image:

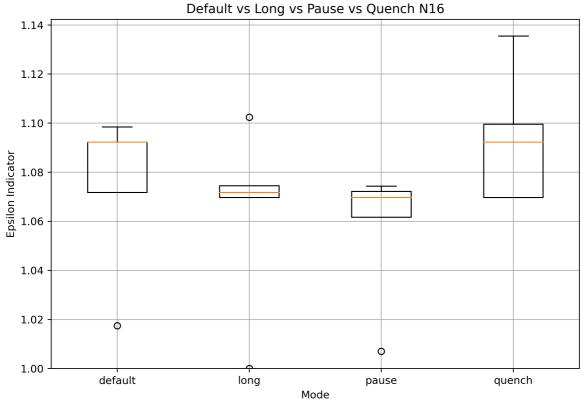






The experiments were run five times:

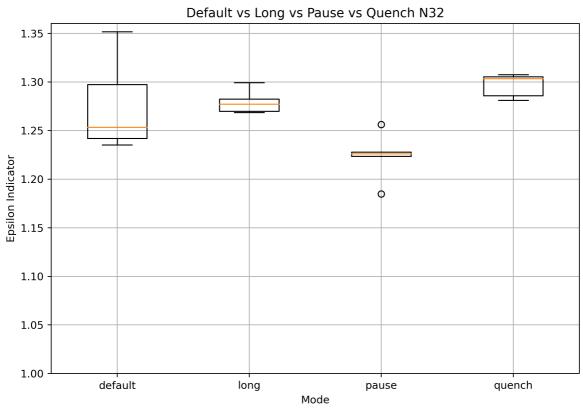
Boxplots - scenario1Y2021M05D01h02m05s12



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set.

Hence, the closer to 1 is the epsilon indicator, the better the annealer set,

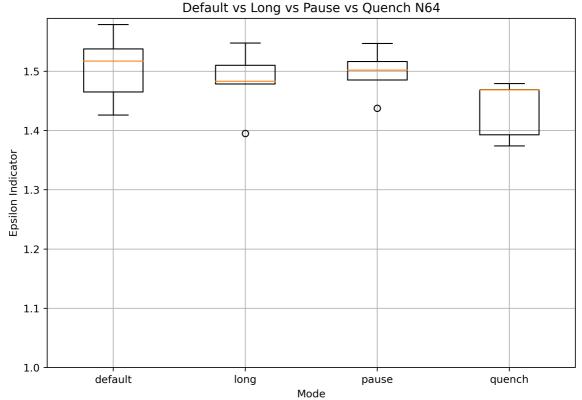
Boxplots - scenario1Y2021M05D01h02m06s15



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set.

Hence, the closer to 1 is the epsilon indicator, the better the annealer set.

Boxplots - scenario1Y2021M05D01h02m07s15



How to interpret: The epsilon indicator is the minimum factor by which the annealer set has to be multiplied in the objective so as to weakly dominate the classical set.

As expected, long and pause are consistently better than default, since they have at least as much anneal time as default. This is, however, at the cost of more machine time budget. In fact, long achieved a perfect score at N=16 in one of the runs.

quench is interesting, since it falls short when N=16 and N=32, but outperforms when N=64.

We noticed that for N=32, pause clearly outperformed the other schedules.

Scenario A3 - Dataset UNFINISHED AND NEEDS CSV FIX!

For this scenario, we will study the influence from the dataset. Previous scenarios used a diversified dataset, with assets as uncorrelated as possible. Therefore, we are going to introduce another dataset, called strongly_correlated, from the same source, however, with strongly correlated assets. That is, with assets from the same sub-industry.

The results are executed for sizes N=32 and N=64, with parameters chain_strength = 1.000 * maxAbs and B=0.5. Since this scenario is small, the results have been repeated two more times, for a total of three tries.

| N and Dataset | q values | |
|-------------------------|--|--|
| N32_diversified | 0, 0.4, 0.9, 2, 3, 9, 100 | |
| N32_strongly_correlated | 0, 1, 6, 10, 70, 90 | |
| N64_diversified | 0, 0.2, 0.4, 0.6, 1.1, 1.3, 1.5, 2, 5, 6, 7, 8, 10, 100, 500 | |
| N64_strongly_correlated | 0, 0.1, 0.2, 0.3, 0.6, 1, 2, 3, 4, 6, 10, 20, 80 | |
| Dataset | N32 (AvgChainBreak) N64 (AvgChainBreak) | |
| | | |

| Dataset | N32 (AvgChainBreak) | N64 (AvgChainBreak) |
|--------------------------|---------------------|---------------------|
| diversified try1 | 1.433 (0.00075) | 1.516 (0.00409) |
| diversified try2 | 1.505 (0.00092) | 1.435 (0.00413) |
| diversified try3 | 1.500 (0.00074) | 1.501 (0.00384) |
| strongly_correlated try1 | 1.300 (0.00093) | 1.701 (0.00370) |
| strongly_correlated try2 | 1.352 (0.00103) | 1.641 (0.00363) |
| strongly_correlated try3 | 1.482 (0.00076) | 1.668 (0.00360) |

Key Takeaways:

The dataset choice does make a significant difference in the performance of the annealer. However, it does not seem to be caused by whether it is diversified or not.