

Hello,

My solution to explore all of the design domain consists mostly of the following [four](#) ideas:

- 1) Random sampling: for each result, I randomly sample combinations until I find one that satisfies all constraints. This method is flexible enough to work for any constraints despite not being that efficient.
- 2) Crowding distance: as results are randomly generated, another constraint is applied so that all results are at least a specific distance apart from each other. This way I am using the already known information to explore the design domain and also avoid physically equivalent results. [Distances are non-dimensionalized based on the current known domain, not the \[0,1\] bounds.](#)
- 3) Inflation/deflate: I initially increase (inflate) the number of desired results to increase diversity. [Less diverse solutions are periodically eliminated based on the crowding distance, which is more computationally efficient than eliminating at the end.](#) This second step also guarantees that the distribution is more uniform in the explored areas (look at plot generated)
- 4) Mutation: as an alternative to random sampling, another selecting algorithm is used. Mutation consists of slightly changing a known feasible solution. Inspired by genetic optimization, it helps in problems where most of the design domain is unfeasible. [Dominant selection algorithm when there is an implicit ingredient or an equality constraint.](#)

There are two parameters, `crowding_distance`, `inflation`, [mutation\\_period \(determines how often a mutation is done relative to random sampling\)](#) that could be tweaked to get more uniformly distributed results. [For alloy.txt, I was not able to find all of the feasible domain.](#)

The attached code is also available on [Github](#).

Thanks,

Pedro Leal

Graduate Research Assistant at Texas A&M University