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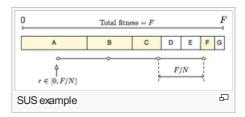
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Stochastic universal sampling

From Wikipedia, the free encyclopedia

Stochastic universal sampling (SUS) is a technique used in genetic algorithms for selecting potentially useful solutions for recombination. It was introduced by James Baker.^[1]

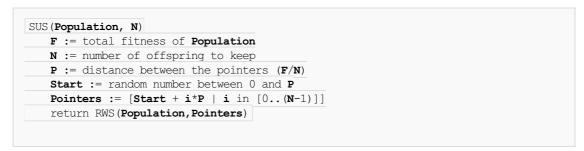
SUS is a development of fitness proportionate selection (FPS) which exhibits no bias and minimal spread. Where FPS chooses several solutions from the population by repeated random sampling, SUS uses a single random value to sample all of the



solutions by choosing them at **evenly spaced intervals**. This gives weaker members of the population (according to their fitness) a chance to be chosen and thus reduces the unfair nature of fitness-proportional selection methods.

Other methods like roulette wheel can have bad performance when a member of the population has a really large fitness in comparison with other members. Using a comb-like ruler, SUS starts from a small random number, and chooses the next candidates from the rest of population remaining, not allowing the fittest members to saturate the candidate space.

Described as an algorithm, pseudocode for SUS looks like:



```
RWS(Population, Points)
Keep = []
i := 0
for P in Points
    while fitness sum of Population[0..i] < P
        i++
        add Population[i] to Keep
return Keep</pre>
```

Where Population[0..i] is the set of individuals with array-index 0 to (and including) i.

Here RWS() describes the bulk of fitness proportionate selection (also known as "roulette wheel selection") - in true fitness proportional selection the parameter **Points** is always a (sorted) list of random numbers from 0 to **F**. The algorithm above is intended to be illustrative rather than canonical.

See also [edit]

- Fitness proportionate selection
- Reward-based selection

References [edit]

 A Baker, James E. (1987). "Reducing Bias and Inefficiency in the Selection Algorithm". Proceedings of the Second International Conference on Genetic Algorithms and their Application (Hillsdale, New Jersey: L. Erlbaum Associates): 14–21.

Categories: Genetic algorithms | Stochastic algorithms

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