

parameters	value	explanation
λ	1000	Cardinality of down-sampled \mathcal{S}_c
N	50	Max. number of GGGP generations.
M	32	Maximum number of individuals.
d	7	Maximum tree depth of genotypes.
(c_c, c_m)	(0.5, 0.5)	Crossover and mutation chance.
σ_0	1	Initial standard deviation.
N_{ES}	10	Max. number of CMA-ES generations.