Project Homepage » DEAP 1.2.2 documentation » Library Reference » previous | next | modules | index **Table of Contents** Algorithms Algorithms Complete Algorithms The algorithms module is intended to contain some specific algorithms in order to execute very Variations common evolutionary algorithms. The method used here are more for convenience than reference as Covariance Matrix Adaptation **Evolution Strategy** the implementation of every evolutionary algorithm may vary infinitely. Most of the algorithms in this module use operators registered in the toolbox. Generally, the keyword used are mate() for crossover, Previous topic mutate() for mutation, select() for selection and evaluate() for evaluation. **Evolutionary Tools** You are encouraged to write your own algorithms in order to make them do what you really want them Next topic to do. Genetic Programming Complete Algorithms This Page **Show Source** These are complete boxed algorithms that are somewhat limited to the very basic evolutionary computation concepts. All algorithms accept, in addition to their arguments, an initialized statistics Quick search object to maintain stats of the evolution, an initialized HallofFame to hold the best individual(s) to Go appear in the population, and a boolean verbose to specify whether to log what is happening during the evolution or not. deap.algorithms.eaSimple(population, toolbox, cxpb, mutpb, ngen[, stats, halloffame, verbose]) This algorithm reproduce the simplest evolutionary algorithm as presented in chapter 7 of [Back2000]. **Parameters:** • population - A list of individuals. • toolbox - A Toolbox that contains the evolution operators. • cxpb - The probability of mating two individuals. • mutpb - The probability of mutating an individual. • **ngen** - The number of generation. • stats - A statistics object that is updated inplace, optional. • halloffame - A Halloffame object that will contain the best individuals, optional. • verbose - Whether or not to log the statistics. The final population **Returns:** A class:~deap.tools.Logbook with the statistics of the evolution **Returns:** The algorithm takes in a population and evolves it in place using the varAnd() method. It returns the optimized population and a Logbook with the statistics of the evolution. The logbook will contain the generation number, the number of evalutions for each generation and the statistics if a statistics is given as argument. The cxpb and mutpb arguments are passed to the varAnd() function. The pseudocode goes as follow evaluate(population) for g in range(ngen): population = select(population, len(population)) offspring = varAnd(population, toolbox, cxpb, mutpb) evaluate(offspring) population = offspring As stated in the pseudocode above, the algorithm goes as follow. First, it evaluates the individuals with an invalid fitness. Second, it enters the generational loop where the selection procedure is applied to entirely replace the parental population. The 1:1 replacement ratio of this algorithm requires the selection procedure to be stochastic and to select multiple times the same individual, for example, selTournament() and selRoulette(). Third, it applies the varAnd() function to produce the next generation population. Fourth, it evaluates the new individuals and compute the statistics on this population. Finally, when ngen generations are done, the algorithm returns a tuple with the final population and a Logbook of the evolution. **Note:** Using a non-stochastic selection method will result in no selection as the operator selects *n* individuals from a pool of *n*. This function expects the toolbox.mate(), toolbox.mutate(), toolbox.select() and toolbox.evaluate() aliases to be registered in the toolbox. [Back2000] Back, Fogel and Michalewicz, "Evolutionary Computation 1: Basic Algorithms and Operators", 2000. deap.algorithms.eaMuPlusLambda(population, toolbox, mu, lambda\_, cxpb, mutpb, ngen[, stats, halloffame, verbose]) This is the  $(\mu + \lambda)$  evolutionary algorithm. **Parameters:** • population - A list of individuals. • toolbox - A Toolbox that contains the evolution operators. • mu - The number of individuals to select for the next generation. • lambda\_ - The number of children to produce at each generation. • **cxpb** - The probability that an offspring is produced by crossover. • mutpb - The probability that an offspring is produced by mutation. • **ngen** – The number of generation. • stats - A statistics object that is updated inplace, optional. • halloffame - A на110ffame object that will contain the best individuals, optional • verbose - Whether or not to log the statistics. The final population Returns: A class:~deap.tools.Logbook with the statistics of the evolution. **Returns:** The algorithm takes in a population and evolves it in place using the varor() function. It returns the optimized population and a Logbook with the statistics of the evolution. The logbook will contain the generation number, the number of evalutions for each generation and the statistics if a statistics is given as argument. The cxpb and mutpb arguments are passed to the varor() function. The pseudocode goes as follow evaluate(population) for g in range(ngen): offspring = varOr(population, toolbox, lambda\_, cxpb, mutpb) evaluate(offspring) population = select(population + offspring, mu) First, the individuals having an invalid fitness are evaluated. Second, the evolutionary loop begins by producing lambda\_ offspring from the population, the offspring are generated by the varor() function. The offspring are then evaluated and the next generation population is selected from both the offspring and the population. Finally, when ngen generations are done, the algorithm returns a tuple with the final population and a Logbook of the evolution. This function expects toolbox.mate(), toolbox.mutate(), toolbox.select() and toolbox.evaluate() aliases to be registered in the toolbox. This algorithm uses the varor() variation. deap.algorithms.eaMuCommaLambda(population, toolbox, mu, lambda\_, cxpb, mutpb, ngen[, stats, halloffame, verbose]) This is the  $(\mu, \lambda)$  evolutionary algorithm. **Parameters:** • population - A list of individuals. • toolbox - A Toolbox that contains the evolution operators. • mu - The number of individuals to select for the next generation. • lambda\_ - The number of children to produce at each generation. • **cxpb** - The probability that an offspring is produced by crossover. • mutpb - The probability that an offspring is produced by mutation. • **ngen** – The number of generation. • stats - A statistics object that is updated inplace, optional. • halloffame - A Halloffame object that will contain the best individuals, optional. • verbose - Whether or not to log the statistics. The final population **Returns:** A class:~deap.tools.Logbook with the statistics of the evolution The algorithm takes in a population and evolves it in place using the varor() function. It returns the optimized population and a Logbook with the statistics of the evolution. The logbook will contain the generation number, the number of evalutions for each generation and the statistics if a statistics is given as argument. The cxpb and mutpb arguments are passed to the varor() function. The pseudocode goes as follow evaluate(population) for g in range(ngen): offspring = varOr(population, toolbox, lambda\_, cxpb, mutpb) evaluate(offspring) population = select(offspring, mu) First, the individuals having an invalid fitness are evaluated. Second, the evolutionary loop begins by producing lambda\_ offspring from the population, the offspring are generated by the varor() function. The offspring are then evaluated and the next generation population is selected from only the offspring. Finally, when ngen generations are done, the algorithm returns a tuple with the final population and a Logbook of the evolution. Note: Care must be taken when the lambda:mu ratio is 1 to 1 as a non-stochastic selection will result in no selection at all as the operator selects *lambda* individuals from a pool of *mu*. This function expects toolbox.mate(), toolbox.mutate(), toolbox.select() and toolbox.evaluate() aliases to be registered in the toolbox. This algorithm uses the varor() variation. deap.algorithms.eaGenerateUpdate(toolbox, ngen[, stats, halloffame, verbose]) This is algorithm implements the ask-tell model proposed in [Colette2010], where ask is called generate and tell is called update. **Parameters:** • toolbox - A Toolbox that contains the evolution operators. • **ngen** – The number of generation. • stats - A statistics object that is updated inplace, optional. • halloffame - A Halloffame object that will contain the best individuals, optional. • **verbose** – Whether or not to log the statistics. The final population **Returns:** A class:~deap.tools.Logbook with the statistics of the evolution **Returns:** The algorithm generates the individuals using the toolbox.generate() function and updates the generation method with the toolbox.update() function. It returns the optimized population and a Logbook with the statistics of the evolution. The logbook will contain the generation number, the number of evalutions for each generation and the statistics if a statistics is given as argument. The pseudocode goes as follow for g in range(ngen): population = toolbox.generate() evaluate(population) toolbox.update(population) [Colette2010] Collette, Y., N. Hansen, G. Pujol, D. Salazar Aponte and R. Le Riche (2010). On Object-Oriented Programming of Optimizers - Examples in Scilab. In P. Breitkopf and R. F. Coelho, eds.: Multidisciplinary Design Optimization in Computational Mechanics, Wiley, pp. 527-565; **Variations** Variations are smaller parts of the algorithms that can be used separately to build more complex algorithms. deap.algorithms.varAnd(population, toolbox, cxpb, mutpb) Part of an evolutionary algorithm applying only the variation part (crossover and mutation). The modified individuals have their fitness invalidated. The individuals are cloned so returned population is independent of the input population. **Parameters:** • population - A list of individuals to vary. • toolbox - A Toolbox that contains the evolution operators. • **cxpb** - The probability of mating two individuals. • mutpb - The probability of mutating an individual. A list of varied individuals that are independent of their parents. **Returns:** The variation goes as follow. First, the parental population  $P_{
m p}$  is duplicated using the toolbox.clone() method and the result is put into the offspring population  $P_{
m o}$ . A first loop over  $P_{
m o}$  is executed to mate pairs of consecutive individuals. According to the crossover probability *cxpb*, the individuals  $\mathbf{x}_i$  and  $\mathbf{x}_{i+1}$  are mated using the toolbox.mate() method. The resulting children  $\mathbf{y}_i$  and  $\mathbf{y}_{i+1}$  replace their respective parents in  $P_{\mathrm{o}}$ . A second loop over the resulting  $P_{\mathrm{o}}$  is executed to mutate every individual with a probability mutpb. When an individual is mutated it replaces its not mutated version in  $P_{
m o}$ . The resulting  $P_{
m o}$  is returned. This variation is named And beceause of its propention to apply both crossover and mutation on the individuals. Note that both operators are not applied systematicaly, the resulting individuals can be generated from crossover only, mutation only, crossover and mutation, and reproduction according to the given probabilities. Both probabilities should be in [0,1]. deap.algorithms.varOr(population, toolbox, lambda\_, cxpb, mutpb) Part of an evolutionary algorithm applying only the variation part (crossover, mutation or reproduction). The modified individuals have their fitness invalidated. The individuals are cloned so returned population is independent of the input population. **Parameters:** • population - A list of individuals to vary. • toolbox - A Toolbox that contains the evolution operators. • lambda\_ - The number of children to produce • **cxpb** - The probability of mating two individuals. • mutpb - The probability of mutating an individual. The final population. **Returns:** The variation goes as follow. On each of the lambda\_ iteration, it selects one of the three operations; crossover, mutation or reproduction. In the case of a crossover, two individuals are selected at random from the parental population  $P_{
m p}$ , those individuals are cloned using the toolbox.clone() method and then mated using the toolbox.mate() method. Only the first child is appended to the offspring population  $P_{
m o}$ , the second child is discarded. In the case of a mutation, one individual is selected at random from  $P_{
m p}$ , it is cloned and then mutated using using the toolbox.mutate() method. The resulting mutant is appended to  $P_{
m o}$ . In the case of a reproduction, one individual is selected at random from  $P_{
m p}$ , cloned and appended to  $P_{
m o}$ . This variation is named Or beceause an offspring will never result from both operations crossover and mutation. The sum of both probabilities shall be in [0,1], the reproduction probability is 1 cxpb - mutpb. Covariance Matrix Adaptation Evolution Strategy A module that provides support for the Covariance Matrix Adaptation Evolution Strategy. class deap.cma. Strategy(centroid, sigma[, \*\*kargs]) A strategy that will keep track of the basic parameters of the CMA-ES algorithm ([Hansen2001]). **Parameters:** • centroid - An iterable object that indicates where to start the evolution. • **sigma** – The initial standard deviation of the distribution. • parameter - One or more parameter to pass to the strategy as described in the following table, optional. **Details** Default **Parameter** Number of children to produce at int(4 + 3 \* log(N))lambda\_ each generation, N is the individual's size (integer). The number of parents to keep int(lambda\_ / 2) mu from the lambda children (integer). The initial covariance matrix of the cmatrix identity(N) distribution that will be sampled. Decrease speed, can be weights "superlinear" "superlinear", "linear" Or "equal". Cumulation constant for step-size. CS (mueff + 2) / (N + mueff + 3)Damping for step-size. 1 + 2 \* max(0, sqrt((mueff - 1) / (N +damps 1)) - 1) + csCumulation constant for covariance 4 / (N + 4)ccum matrix. Learning rate for rank-one update.  $2 / ((N + 1.3)^2 + mueff)$ ccov1 Learning rate for rank-mu update.  $2 * (mueff - 2 + 1 / mueff) / ((N + 2)^2$ ccovmu + mueff) Hansen and Ostermeier, 2001. Completely Derandomized Self-Adaptation in [Hansen2001] Evolution Strategies. Evolutionary Computation computeParams(params) Computes the parameters depending on  $\lambda$ . It needs to be called again if  $\lambda$  changes during evolution. **Parameters:** params - A dictionary of the manually set parameters. generate(ind\_init) Generate a population of  $\lambda$  individuals of type *ind\_init* from the current strategy. Parameters: ind\_init - A function object that is able to initialize an individual from a list. A list of individuals. **Returns:** update(population) Update the current covariance matrix strategy from the population. **Parameters:** population - A list of individuals from which to update the parameters. class deap.cma. StrategyOnePlusLambda(parent, sigma[, \*\*kargs]) A CMA-ES strategy that uses the  $1+\lambda$  paradigm ([Igel2007]). **Parameters:** • parent - An iterable object that indicates where to start the evolution. The parent requires a fitness attribute. • sigma - The initial standard deviation of the distribution. • lambda - Number of offspring to produce from the parent. (optional, defaults • parameter - One or more parameter to pass to the strategy as described in the following table. (optional) Other parameters can be provided as described in the next table Default Details **Parameter** 1.0 + N / (2.0 \* lambda)Damping for step-size. 1.0 / (5 + sqrt(lambda\_) / 2.0) Taget success rate. ptarg ptarg \* lambda / (2.0 + ptarg \* Step size learning rate. lambda\_) Cumulation time horizon. CC 2.0 / (N + 2.0)Covariance matrix learning rate. 2.0 / (N\*\*2 + 6.0)ccov Threshold success rate. 0.44 pthresh [Igel2007] | Igel, Hansen, Roth, 2007. Covariance matrix adaptation for multi-objective optimization. *Evolutionary Computation* Spring;15(1):1-28 computeParams(params) Computes the parameters depending on  $\lambda$ . It needs to be called again if  $\lambda$  changes during evolution. **Parameters:** params - A dictionary of the manually set parameters. generate(ind\_init) Generate a population of  $\lambda$  individuals of type *ind\_init* from the current strategy. **Parameters:** ind\_init - A function object that is able to initialize an individual from a list. A list of individuals. **Returns:** update(population) Update the current covariance matrix strategy from the *population*. **Parameters:** population - A list of individuals from which to update the parameters. class deap.cma. StrategyMultiObjective(population, sigma[, \*\*kargs]) Multiobjective CMA-ES strategy based on the paper [Voss2010]. It is used similarly as the standard CMA-ES strategy with a generate-update scheme. **Parameters:** • population - An initial population of individual. • sigma - The initial step size of the complete system. • mu - The number of parents to use in the evolution. When not provided it defaults to the length of *population*. (optional) • lambda - The number of offspring to produce at each generation. (optional, defaults to 1) • indicator - The indicator function to use. (optional, default to hypervolume()) Other parameters can be provided as described in the next table **Default** Details **Parameter** 1.0 + N / 2.0Damping for step-size. 1.0 / (5 + 1.0 / 2.0)Taget success rate. ptarg Step size learning rate. ptarg / (2.0 + ptarg)ср Cumulation time horizon. CC 2.0 / (N + 2.0)2.0 / (N\*\*2 + 6.0)Covariance matrix learning ccov rate. Threshold success rate. pthresh 0.44 [Voss2010] Voss, Hansen, Igel, "Improved Step Size Adaptation for the MO-CMA-ES", 2010. generate(ind\_init) Generate a population of  $\lambda$  individuals of type *ind\_init* from the current strategy. **Parameters:** ind\_init - A function object that is able to initialize an individual from a list. A list of individuals with a private attribute \_ps. This last attribute is essential **Returns:** to the update function, it indicates that the individual is an offspring and the index of its parent. update(population) Update the current covariance matrix strategies from the *population*. **Parameters:** population - A list of individuals from which to update the parameters. Project Homepage » DEAP 1.2.2 documentation » Library Reference » previous | next | modules | index © Copyright 2009-2019, DEAP Project. Built on Jun 05, 2019. Found a bug? Created using Sphinx 1.8.5.