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scipy.optimize.linprog(c, A_ub=None, b_ub=None, A_eq=None, b_eq=None, bounds=None, method='highs', callback=None, options=None, x0=None, integrality=None) [source]

Linear programming: minimize a linear objective function subject to linear equality and inequality constraints.

Linear programming solves problems of the following form:

$$egin{aligned} \min_{x} \ c^T x \ & ext{such that} \ A_{ub} x \leq b_{ub}, \ & A_{eq} x = b_{eq}, \ & l \leq x \leq u, \end{aligned}$$

where x is a vector of decision variables; c, b_{ub} , b_{eq} , l, and u are vectors; and A_{ub} and A_{eq} are matrices.

Alternatively, that's:

• minimize

c @ x

such that

```
A_ub @ x <= b_ub
A_eq @ x == b_eq
1b <= x <= ub
```

Note that by default 1b = 0 and ub = None. Other bounds can be specified with bounds.

Parameters: c: 1-D array

The coefficients of the linear objective function to be minimized.

A_ub : 2-D array, optional

The inequality constraint matrix. Each row of A_ub specifies the coefficients of a linear inequality constraint on x.

b_ub : 1-D array, optional

The inequality constraint vector. Each element represents an upper bound on the corresponding value of A_ub @ x.

A_eq: 2-D array, optional

The equality constraint matrix. Each row of A_eq specifies the coefficients of a linear equality constraint on x.

b_eq: 1-D array, optional

The equality constraint vector. Each element of $A_{eq} @ \times \text{must}$ equal the corresponding element of b_{eq} .

bounds: sequence, optional

A sequence of (min, max) pairs for each element in x, defining the minimum and maximum values of that decision variable. If a single tuple (min, max) is provided, then min and max will serve as bounds for all decision variables. Use None to indicate that there is no bound. For instance, the default bound (0, None) means that all decision variables are non-negative, and the pair (None, None) means no bounds at all, i.e. all variables are allowed to be any real.

method: str, optional

The algorithm used to solve the standard form problem. <u>'highs'</u> (default), <u>'highs-ds'</u>, <u>'highs-ipm'</u>, <u>'interior-point'</u> (legacy), <u>'revised simplex'</u> (legacy), and <u>'simplex'</u> (legacy) are supported. The legacy methods are deprecated and will be removed in SciPy 1.11.0.

callback : callable, optional

If a callback function is provided, it will be called at least once per iteration of the algorithm. The callback function must accept a single scipy.optimize.OptimizeResult consisting of the following fields:

x: 1-D array

The current solution vector.

fun : float

The current value of the objective function $c \otimes x$.

success: bool

True when the algorithm has completed successfully.

slack: 1-D array

The (nominally positive) values of the slack, b_ub - A_ub @ x.

con: 1-D array

The (nominally zero) residuals of the equality constraints, b_eq - A_eq @ x.

phase : int

The phase of the algorithm being executed.

status : int

An integer representing the status of the algorithm.

- 0: Optimization proceeding nominally.
- 1: Iteration limit reached.
- 2 : Problem appears to be infeasible.
- 3: Problem appears to be unbounded.
- 4: Numerical difficulties encountered.

nit: int

The current iteration number.

message : str

A string descriptor of the algorithm status.

Callback functions are not currently supported by the HiGHS methods.

options: dict, optional

A dictionary of solver options. All methods accept the following options:

maxiter : int

Maximum number of iterations to perform. Default: see method-specific documentation.

disp: bool

Set to True to print convergence messages. Default: False.

presolve : bool

Set to False to disable automatic presolve. Default: True.

All methods except the HiGHS solvers also accept:

tol: float

A tolerance which determines when a residual is "close enough" to zero to be considered exactly zero.

autoscale : bool

Set to True to automatically perform equilibration. Consider using this option if the numerical values in the constraints are separated by several orders of magnitude. Default: False.

rr: bool

Set to False to disable automatic redundancy removal. Default: True.

rr_method : string

Method used to identify and remove redundant rows from the equality constraint matrix after presolve. For problems with dense input, the available methods for redundancy removal are:

"SVD":

Repeatedly performs singular value decomposition on the matrix, detecting redundant rows based on nonzeros in the left singular vectors that correspond with zero singular values. May be fast when the matrix is nearly full rank.

"pivot":

Uses the algorithm presented in [5] to identify redundant rows.

"ID":

Uses a randomized interpolative decomposition. Identifies columns of the matrix transpose not used in a full-rank interpolative decomposition of the matrix.

None:

Uses "svd" if the matrix is nearly full rank, that is, the difference between the matrix rank and the number of rows is less than five. If not, uses "pivot". The behavior of this default is subject to change without prior notice.

Default: None. For problems with sparse input, this option is ignored, and the pivot-based algorithm presented in [5] is used.

For method-specific options, see show_options('linprog').

x0: 1-D array, optional

Guess values of the decision variables, which will be refined by the optimization algorithm. This argument is currently used only by the 'revised simplex' method, and can only be used if *x0* represents a basic feasible solution.

integrality: 1-D array or int, optional

Indicates the type of integrality constraint on each decision variable.

- 0: Continuous variable; no integrality constraint.
- 1: Integer variable; decision variable must be an integer within bounds.
- 2 : Semi-continuous variable; decision variable must be within bounds or take value 0.
- 3 : Semi-integer variable; decision variable must be an integer within *bounds* or take value 0.

By default, all variables are continuous.

For mixed integrality constraints, supply an array of shape *c.shape*. To infer a constraint on each decision variable from shorter inputs, the argument will be broadcasted to *c.shape* using *np.broadcast_to*.

This argument is currently used only by the 'highs' method and ignored otherwise.

Returns: **res** : OptimizeResult

A <u>scipy.optimize.OptimizeResult</u> consisting of the fields below. Note that the return types of the fields may depend on whether the optimization was successful, therefore it is recommended to check *OptimizeResult.status* before relying on the other fields:

x: 1-D array

The values of the decision variables that minimizes the objective function while satisfying the constraints.

fun: float

The optimal value of the objective function c @ x.

slack: 1-D array

The (nominally positive) values of the slack variables, b_ub - A_ub @ x.

con: 1-D array

The (nominally zero) residuals of the equality constraints, b_eq - A_eq @ x.

success: bool

True when the algorithm succeeds in finding an optimal solution.

status : int

An integer representing the exit status of the algorithm.

- Optimization terminated successfully.
- 1 : Iteration limit reached.
- 2 : Problem appears to be infeasible.
- 3 : Problem appears to be unbounded.
- 4 : Numerical difficulties encountered.

nit : int

The total number of iterations performed in all phases.

message : str

A string descriptor of the exit status of the algorithm.

See also

show options

Additional options accepted by the solvers.

Notes

This section describes the available solvers that can be selected by the 'method' parameter.

'highs-ds' and 'highs-ipm' are interfaces to the HiGHS simplex and interior-point method solvers [13], respectively. 'highs' (default) chooses between the two automatically. These are the fastest linear programming solvers in SciPy, especially for large, sparse problems; which of these two is faster is problemdependent. The other solvers ('interior-point', 'revised simplex', and 'simplex') are legacy methods and will be removed in SciPy 1.11.0.

Method highs-ds is a wrapper of the C++ high performance dual revised simplex implementation (HSOL) [13], [14]. Method highs-ipm is a wrapper of a C++ implementation of an interior-point method [13]; it features a crossover routine, so it is as accurate as a simplex solver. Method highs chooses between the two automatically. For new code involving linprog, we recommend explicitly choosing one of these three method values.

New in version 1.6.0.

Method interior-point uses the primal-dual path following algorithm as outlined in [4]. This algorithm supports sparse constraint matrices and is typically faster than the simplex methods, especially for large, sparse problems. Note, however, that the solution returned may be slightly less accurate than those of the simplex methods and will not, in general, correspond with a vertex of the polytope defined by the constraints.

New in version 1.0.0.

Method revised simplex uses the revised simplex method as described in [9], except that a factorization [11] of the basis matrix, rather than its inverse, is efficiently maintained and used to solve the linear systems at each iteration of the algorithm.

New in version 1.3.0.

Method simplex uses a traditional, full-tableau implementation of Dantzig's simplex algorithm [1], [2] (not the Nelder-Mead simplex). This algorithm is included for backwards compatibility and educational purposes.

• New in version 0.15.0.

Before applying interior-point, revised simplex, or simplex, a presolve procedure based on [8] attempts to identify trivial infeasibilities, trivial unboundedness, and potential problem simplifications. Specifically, it checks for:

- rows of zeros in A_eq or A_ub, representing trivial constraints;
- columns of zeros in A_eq and A_ub, representing unconstrained variables;
- column singletons in A_eq, representing fixed variables; and
- column singletons in A_ub, representing simple bounds.

If presolve reveals that the problem is unbounded (e.g. an unconstrained and unbounded variable has negative cost) or infeasible (e.g., a row of zeros in A_{eq} corresponds with a nonzero in b_{eq}), the solver terminates with the appropriate status code. Note that presolve terminates as soon as any sign of

unboundedness is detected; consequently, a problem may be reported as unbounded when in reality the problem is infeasible (but infeasibility has not been detected yet). Therefore, if it is important to know whether the problem is actually infeasible, solve the problem again with option presolve=False.

If neither infeasibility nor unboundedness are detected in a single pass of the presolve, bounds are tightened where possible and fixed variables are removed from the problem. Then, linearly dependent rows of the A_eq matrix are removed, (unless they represent an infeasibility) to avoid numerical difficulties in the primary solve routine. Note that rows that are nearly linearly dependent (within a prescribed tolerance) may also be removed, which can change the optimal solution in rare cases. If this is a concern, eliminate redundancy from your problem formulation and run with option rr=False or presolve=False.

Several potential improvements can be made here: additional presolve checks outlined in [8] should be implemented, the presolve routine should be run multiple times (until no further simplifications can be made), and more of the efficiency improvements from [5] should be implemented in the redundancy removal routines.

After presolve, the problem is transformed to standard form by converting the (tightened) simple bounds to upper bound constraints, introducing non-negative slack variables for inequality constraints, and expressing unbounded variables as the difference between two non-negative variables. Optionally, the problem is automatically scaled via equilibration [12]. The selected algorithm solves the standard form problem, and a postprocessing routine converts the result to a solution to the original problem.

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Examples

Consider the following problem:

$$egin{array}{ll} \min_{x_0,x_1} & -x_0 + 4x_1 \ \mathrm{such\ that} & -3x_0 + x_1 \leq 6, \ & -x_0 - 2x_1 \geq -4, \ & x_1 \geq -3. \end{array}$$

The problem is not presented in the form accepted by <u>linprog</u>. This is easily remedied by converting the "greater than" inequality constraint to a "less than" inequality constraint by multiplying both sides by a factor of -1. Note also that the last constraint is really the simple bound $-3 \le x_1 \le \infty$. Finally, since there are no bounds on x_0 , we must explicitly specify the bounds $-\infty \le x_0 \le \infty$, as the default is for variables to be non-negative. After collecting coeffecients into arrays and tuples, the input for this problem is:

```
>>> from scipy.optimize import linprog
>>> c = [-1, 4]
>>> A = [[-3, 1], [1, 2]]
>>> b = [6, 4]
>>> x0_bounds = (None, None)
>>> x1_bounds = (-3, None)
>>> res = linprog(c, A_ub=A, b_ub=b, bounds=[x0_bounds, x1_bounds])
>>> res.fun
-22.0
>>> res.x
array([10., -3.])
>>> res.message
'Optimization terminated successfully. (HiGHS Status 7: Optimal)'
```

The marginals (AKA dual values / shadow prices / Lagrange multipliers) and residuals (slacks) are also available.

```
>>> res.ineqlin
    residual: [ 3.900e+01      0.000e+00]
    marginals: [-0.000e+00 -1.000e+00]
```

For example, because the marginal associated with the second inequality constraint is -1, we expect the optimal value of the objective function to decrease by eps if we add a small amount eps to the right hand side of the second inequality constraint:

```
>>> eps = 0.05
>>> b[1] += eps
>>> linprog(c, A_ub=A, b_ub=b, bounds=[x0_bounds, x1_bounds]).fun
-22.05
```

Also, because the residual on the first inequality constraint is 39, we can decrease the right hand side of the first constraint by 39 without affecting the optimal solution.

```
>>> b = [6, 4] # reset to original values
>>> b[0] -= 39
>>> linprog(c, A_ub=A, b_ub=b, bounds=[x0_bounds, x1_bounds]).fun
-22.0
```

Previous
scipy.optimize.milp

Next linprog(method='simplex')

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