

# Package: n1qn1 (via r-universe)

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**Title** Port of the 'Scilab' 'n1qn1' Module for Unconstrained BFGS Optimization

**Version** 6.0.1-12

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**Description** Provides 'Scilab' 'n1qn1'. This takes more memory than traditional L-BFGS. The n1qn1 routine is useful since it allows prespecification of a Hessian. If the Hessian is near enough the truth in optimization it can speed up the optimization problem. The algorithm is described in the 'Scilab' optimization documentation located at <[https://www.scilab.org/sites/default/files/optimization\\_in\\_scilab.pdf](https://www.scilab.org/sites/default/files/optimization_in_scilab.pdf)>. This version uses manually modified code from 'f2c' to make this a C only binary.

**URL** <https://github.com/nlmixr2/n1qn1c>

**BugReports** <https://github.com/nlmixr2/n1qn1c/issues>

**Depends** R (>= 3.2)

**Imports** Rcpp (>= 0.12.3)

**Suggests** testthat, covr

**License** CeCILL-2

**Biarch** true

**NeedsCompilation** yes

**LinkingTo** RcppArmadillo (>= 0.5.600.2.0), Rcpp (>= 0.12.3)

**Encoding** UTF-8

**RoxygenNote** 7.3.2

**Repository** <https://nlmixr2.r-universe.dev>

**RemoteUrl** <https://github.com/nlmixr2/n1qn1c>

**RemoteRef** HEAD

**RemoteSha** 1a6bbf0634a16b660d5e336b8a646bddealad16d

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|           |  |
|-----------|--|
| .n1qn1ptr | <i>This gives the function pointers in the n1qn1 library</i> |
|-----------|--|

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### Description

Using this will allow C-level linking by function pointers instead of abi.

### Usage

```
.n1qn1ptr()
```

### Value

list of pointers to the n1qn1 functions

### Author(s)

Matthew L. Fidler

### Examples

```
.n1qn1ptr()
```

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|       |                           |
|-------|---------------------------|
| n1qn1 | <i>n1qn1 optimization</i> |
|-------|---------------------------|

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### Description

This is an R port of the n1qn1 optimization procedure in scilab.

### Usage

```
n1qn1(
  call_eval,
  call_grad,
  vars,
  environment = parent.frame(1),
  ...,
  epsilon = .Machine$double.eps,
  max_iterations = 100,
```

```

nsim = 100,
imp = 0,
invisible = NULL,
zm = NULL,
restart = FALSE,
assign = FALSE,
print.functions = FALSE
)

```

## Arguments

|                 |  |
|-----------------|--|
| call_eval       | Objective function   |
| call_grad       | Gradient Function  |
| vars            | Initial starting point for line search   |
| environment     | Environment where call_eval/call_grad are evaluated.   |
| ...             | Ignored additional parameters.   |
| epsilon         | Precision of estimate  |
| max_iterations  | Number of iterations   |
| nsim            | Number of function evaluations   |
| imp             | Verbosity of messages.   |
| invisible       | boolean to control if the output of the minimizer is suppressed.   |
| zm              | Prior Hessian (in compressed format; This format is output in c.hess).                                       |
| restart         | Is this an estimation restart?   |
| assign          | Assign hessian to c.hess in environment environment? (Default FALSE)   |
| print.functions | Boolean to control if the function value and parameter estimates are echoed every time a function is called. |

## Value

The return value is a list with the following elements:

- **value** The value at the minimized function.
- **par** The parameter value that minimized the function.
- **H** The estimated Hessian at the final parameter estimate.
- **c.hess** Compressed Hessian for saving curvature.
- **n.fn** Number of function evaluations
- **n.gr** Number of gradient evaluations

## Author(s)

C. Lemarechal, Stephen L. Campbell, Jean-Philippe Chancelier, Ramine Nikoukhah, Wenping Wang & Matthew L. Fidler

## Examples

```

## Rosenbrock's banana function
n=3; p=100

fr = function(x)
{
  f=1.0
  for(i in 2:n) {
    f=f+p*(x[i]-x[i-1]**2)**2+(1.0-x[i])**2
  }
  f
}

grr = function(x)
{
  g = double(n)
  g[1]=-4.0*p*(x[2]-x[1]**2)*x[1]
  if(n>2) {
    for(i in 2:(n-1)) {
      g[i]=2.0*p*(x[i]-x[i-1]**2)-4.0*p*(x[i+1]-x[i]**2)*x[i]-2.0*(1.0-x[i])
    }
  }
  g[n]=2.0*p*(x[n]-x[n-1]**2)-2.0*(1.0-x[n])
  g
}

x = c(1.02,1.02,1.02)
eps=1e-3
n=length(x); niter=100L; nsim=100L; imp=3L;
nzm=as.integer(n*(n+1L)/2L)
zm=double(nzm)

(op1 <- n1qn1(fr, grr, x, imp=3))

## Note there are 40 function calls and 40 gradient calls in the above optimization

## Now assume we know something about the Hessian:
c.hess <- c(797.861115,
           -393.801473,
           -2.795134,
           991.271179,
           -395.382900,
           200.024349)
c.hess <- c(c.hess, rep(0, 24 - length(c.hess)))

(op2 <- n1qn1(fr, grr, x, imp=3, zm=c.hess))

## Note with this knowledge, there were only 29 function/gradient calls

(op3 <- n1qn1(fr, grr, x, imp=3, zm=op1$c.hess))

## The number of function evaluations is still reduced because the Hessian

```

```
## is closer to what it should be than the initial guess.  
## With certain optimization procedures this can be helpful in reducing the  
## Optimization time.
```

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