

# Package ‘pbdBASE’

December 8, 2018

**Type** Package

**Title** Programming with Big Data -- Base Wrappers for Distributed Matrices

**Version** 0.5-0

**Description** An interface to and extensions for the 'PBLAS' and 'ScaLAPACK' numerical libraries. This enables R to utilize distributed linear algebra for codes written in the 'SPMD' fashion. This interface is deliberately low-level and mimics the style of the native libraries it wraps. For a much higher level way of managing distributed matrices, see the 'pbdDMAT' package.

**License** Mozilla Public License 2.0

**Depends** R (>= 3.0.0), methods

**Imports** utils, pbdMPI (>= 0.3-8), pbdSLAP(>= 0.2-4)

**SystemRequirements** OpenMPI (>= 1.5.4) on Solaris, Linux, Mac, and FreeBSD. MS-MPI (Microsoft HPC Pack 2012) or MPICH2 (>= 1.4.1p1) on Windows.

**LazyLoad** yes

**LazyData** yes

**ByteCompile** yes

**NeedsCompilation** yes

**URL** <http://r-pbd.org/>

**BugReports** <http://group.r-pbd.org/>

**MailingList** Please send questions and comments regarding pbdR to RBigData@gmail.com

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**RoxygenNote** 6.0.1

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**Repository** CRAN

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**Description**

A package contains the basic methods for dealing with distributed data types, as well as the data types themselves.

**Details**

Package: pbdBASE  
Type: Package  
License: MPL  
LazyLoad: yes

This package requires an MPI library (OpenMPI, MPICH2, or LAM/MPI).

### Author(s)

Drew Schmidt <wrathematics AT gmail.com>, Wei-Chen Chen, George Ostrouchov, and Pragneshkumar Patel.

### References

Programming with Big Data in R Website: <http://r-pbd.org/>

---

BASE Global Environment

*Global Environment for the pbdBASE Package*

---

### Description

The environment for the pbdBASE package where "global" variables are stored.

### Usage

`.pbdBASEEnv`

### Format

An object of class environment of length 0.

### Details

The `._blacs_gridinfo_` and `._blacs_initialized` objects are stored in this environment.

---

base.blacs\_gridinit     *Creating Grid From A System Context*

---

### Description

Creates a grid from a System Context obtained from a call to 'sys2blacs\_handle'.

### Usage

```
base.blacs_gridinit(SYSCTX, NPROW, NPCOL, nprocs = pbdMPI::comm.size(comm),
  comm = .pbd_env$SPMD.CT$comm)
```

### Arguments

SYSCTX	System context obtained from a call to 'sys2blacs_handle'
NPROW	Number of rows in the process grid
NPCOL	Number of columns in the process grid
nprocs	Number of processors in the communicator
comm	An MPI (not BLACS) communicator.

### Value

A blacs context number

---

base.crossprod     *crossprod*

---

### Description

Crossproduct.

### Usage

```
base.crossprod(uplo, trans, x, descx, descc)
```

### Arguments

uplo	Triangle whose values to use.
trans	tcrossprod or crossprod.
x	Matrix to crossprod.
descx	ScaLAPACK descriptor array.
descc	ScaLAPACK descriptor array of output.

### Details

For advanced users only.

---

base.dallreduce	<i>dallreduce</i>
-----------------	-------------------

---

**Description**

Allreduce

**Usage**

```
base.dallreduce(x, descx, op = "sum", scope = "All")
```

**Arguments**

x	Matrix.
descx	ScaLAPACK descriptor array.
op	Operation.
scope	Rows, columns, or both.

**Details**

For advanced users only.

---

base.descinit	<i>descinit</i>
---------------	-----------------

---

**Description**

Creates ScaLAPACK descriptor array.

**Usage**

```
base.descinit(dim, bldim, ldim, ICTXT = 0)
```

**Arguments**

dim	Global dim.
bldim	Blocking dim.
ldim	Local dim.
ICTXT	BLACS context.

**Details**

For advanced users only.

---

 base.dgamx2d

*BLACS Min*


---

### Description

Min value across a process grid.

### Usage

base.dgamx2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)

base.igamn2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)

base.dgamn2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)

### Arguments

ICTXT	BLACS ICTXT.
SCOPE	Rows, cols, or both.
m, n	Problem size.
x	Local values.
lda	Leading dimension.
RDEST	Row destination.
CDEST	Col destination.

### Details

For advanced users only.

---

 base.dgesd2d

*BLACS Point to Poin*


---

### Description

Sent value across a process grid.

### Usage

base.dgesd2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)

base.dgerv2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)



**Arguments**

ICTXT	BLACS ICTXT.
SCOPE	Rows, cols, or both.
m, n	Problem size.
x	Local values.
lda	Leading dimension.
RDEST	Row destination.
CDEST	Col destination.

**Details**

For advanced users only.

---

<i>base.dhilbmk</i>	<i>dhilbmk</i>
---------------------	----------------

---

**Description**

Create Hilbert matrix.

**Usage**

`base.dhilbmk(n)`

**Arguments**

n	Size.
---	-------

**Details**

For advanced users only.

---

base.dim0	<i>maxdim</i>
-----------	---------------

---

**Description**

Compute dimensions on process MYROW=MYCOL=0

**Usage**

```
base.dim0(dim, ICTXT = 0)
```

**Arguments**

dim	Global dim.
ICTXT	BLACS context.

**Details**

For advanced users only.

---

base.free_blacs_system_handle
<i>Free Blacs System Handle</i>

---

**Description**

Free Blacs System Handle

**Usage**

```
base.free_blacs_system_handle(SHANDLE)
```

**Arguments**

SHANDLE	A system handle. Obtained via a call to 'sys2blacs.handle'
---------	--

---

base.igamx2d	<i>BLACS Max</i>
--------------	------------------

---

**Description**

Max value across a process grid.

**Usage**

```
base.igamx2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)
```

**Arguments**

ICTXT	BLACS ICTXT.
SCOPE	Rows, cols, or both.
m, n	Problem size.
x	Local values.
lda	Leading dimension.
RDEST	Row destination.
CDEST	Col destination.

**Details**

For advanced users only.

---

base.igsum2d	<i>BLACS Sums</i>
--------------	-------------------

---

**Description**

Sum across a process grid.

**Usage**

```
base.igsum2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)
```

```
base.dgsum2d(ICTXT, SCOPE, m, n, x, lda, RDEST, CDEST)
```

**Arguments**

ICTXT	BLACS ICTXT.
SCOPE	Rows, cols, or both.
m, n	Problem size.
x	Local values.
lda	Leading dimension.
RDEST	Row destination.
CDEST	Col destination.

**Details**

For advanced users only.

---

base.indxg2p	<i>indxg2p</i>
--------------	----------------

---

**Description**

Computes the process coordinate which contains the entry of a distributed matrix specified by a global index INDXGLOB. Simplified reimplemention of the ScaLAPACK aux INDXG2P function.

**Usage**

```
base.indxg2p(INDXGLOB, NB, NPROCS)
```

**Arguments**

INDXGLOB	Global index.
NB	Block size.
NPROCS	Total number of processors over which matrix is distributed.

**Details**

For advanced users only.

---

base.matexp	<i>matexp</i>
-------------	---------------

---

**Description**

Serial matrix exponentiation.

**Usage**

```
base.matexp(A, p = 6, t = 1)
```

**Arguments**

A	Matrix to exponentiate.
p	Pade' expansion size.
t	Scaling factor.

**Details**

For advanced users only.

---

base.maxdim	<i>maxdim</i>
-------------	---------------

---

**Description**

Compute maximum dimension across all nodes

**Usage**

```
base.maxdim(dim)
```

**Arguments**

dim	Global dim.
-----	-------------

**Details**

For advanced users only.

---

base.minctxt	<i>Get BLACS Context Grid Information</i>
--------------	---

---

**Description**

Finds the smallest integers for creating a new BLACS context.

**Usage**

```
base.minctxt(after = 0)
```

**Arguments**

after                    ignores all values below this integer as possibilities

**Details**

For advanced users only.

Returns the smallest integer which could become a new BLACS context value.

For example, if contexts 0, 1, and 2 are taken, and after=0, then the function returns 3. If 0, 1, 2, and 5 are taken, the function returns 3 if after=0, but returns 6 if after=4.

The function is useful when a transitory grid is needed, such as for reading in data onto a subset of processors before distributing out to the full grid.

**Value**

Returns the minimum value.

---

base.mksubmat	<i>(Un)Distribute</i>
---------------	-----------------------

---

**Description**

(Un)Distribute matrix.

**Usage**

```
base.mksubmat(x, descx)
```

```
base.mkgblmat(x, descx, rsrc, csrc)
```

**Arguments**

x                        Matrix.  
descx                    ScaLAPACK descriptor array.  
rsrc, csrc                Row/column source.

**Details**

For advanced users only.

---

base.nbd	<i>Next Best Divisor</i>
----------	--------------------------

---

**Description**

Given integers  $n$  and  $d$ , with  $n > d$ , this function finds the "next best divisor" of  $n$  which is greater than or equal to  $d$ .

**Usage**

```
base.nbd(n, d)
```

**Arguments**

$n$	The dividend (number divided into).
$d$	The candidate divisor.

**Details**

Surprisingly useful for thinking about processor grid shapes.

**Examples**

```
spmd.code = "
  pbdBASE::base.nbd(100, 10) # 10 divides 100, so 10 is returned
  pbdBASE::base.nbd(100, 11) # 11 does not, so the 'next best' divisor, 20, is returned
"

pbdMPI::execmpi(spmd.code = spmd.code, nrank = 1L)
```

---

base.numroc	<i>numroc</i>
-------------	---------------

---

**Description**

NUMBER of Rows Or Columns

**Usage**

```
base.numroc(dim, bldim, ICTXT = 0, fixme = TRUE)
```

**Arguments**

dim	Global dim.
bldim	Blocking dim.
ICTXT	BLACS context.
fixme	Should ldims be "rounded" to 0 or not.

**Details**

For advanced users only.

---

base.ownany

*Determining Local Ownership of a Distributed Matrix*

---

**Description**

For advanced users only.

**Usage**

```
base.ownany(dim, bldim, ICTXT = 0)
```

**Arguments**

dim	global dimension
bldim	blocking dimension
ICTXT	BLACS context

**Details**

A simple wrapper of numroc. The return is the answer to the question 'do I own any of the global matrix?'. Passing a distributed matrix is allowed, but often it is convenient to determine that information without even having a distributed matrix on hand. In this case, explicitly passing the appropriate information to the arguments dim=, bldim= (and CTXT= as necessary, since it defaults to 0) while leaving x missing will produce the desired result. See the examples below for more clarity.

The return for each function is local.

**Examples**

```
smpd.code = "
  suppressMessages(library(pbdBASE))
  init.grid()

  iown <- ownany(dim=c(4, 4), bldim=c(2, 2), CTXT=0)
  comm.print(iown, all.rank=T)

  finalize()
```



```
"
pbdMPI::execmpi(spmc.code = spmd.code, nrank = 2L)
```

---

base.pdchtri	<i>pdchtri</i>
--------------	----------------

---

### Description

Inverse of cholesky.

### Usage

```
base.pdchtri(uplo, x, descx, desc)
```

### Arguments

uplo	Triangle whose values to use.
x	Matrix to crossprod.
descx	ScaLAPACK descriptor array.
desc	ScaLAPACK descriptor array of output.

### Details

For advanced users only.

---

base.pdclvar	<i>Column Variances</i>
--------------	-------------------------

---

### Description

Computes the variances of a ScaLAPACK-like distributed matrix. Significantly faster than using `apply()`, even in compared to the performance differences you would find comparing these two approaches using just base R.

### Usage

```
base.pdclvar(x, descx)
```

### Arguments

x	The matrix.
descx	ScaLAPACK descriptor array.

---

base.pdhilbmk	<i>pdhilbmk</i>
---------------	-----------------

---

**Description**

Create Hilbert matrix.

**Usage**

base.pdhilbmk(descx)

**Arguments**

descx            ScaLAPACK descriptor matrix.

**Details**

For advanced users only.

---

base.pdmkcpn1	<i>pdmkcpn1</i>
---------------	-----------------

---

**Description**

Create Companion Matrix

**Usage**

base.pdmkcpn1(coef, descx)

**Arguments**

coef            Coefficients vector.  
descx           ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.pdmvsum	<i>R-like Matrix-Vector Sum</i>
--------------	---------------------------------

---

**Description**

For advanced users only.

**Usage**

```
base.pdmvsum(x, descx, y, descy)
```

**Arguments**

x	Matrix.
descx, descy	ScaLAPACK descriptor array.
y	Vector.

---

base.pdsweep	<i>pdsweep</i>
--------------	----------------

---

**Description**

Matrix-Vector Sweep

**Usage**

```
base.pdsweep(x, descx, vec, MARGIN, FUN)
```

**Arguments**

x	Matrix.
descx	ScaLAPACK descriptor array.
vec	Vector
MARGIN	Rows or columns.
FUN	Function.

**Details**

For advanced users only.

base.procgrid      *procgrid*

---

**Description**

"Optimal" process grid when nrow and ncol are empty

**Usage**

base.procgrid(nprocs)

**Arguments**

nprocs      Number of processors.

**Details**

For advanced users only.

---

base.p\_matexp\_pade\_wrap  
*p\_matexp\_pade\_wrap*

---

**Description**

Pade' expansion.

**Usage**

base.p\_matexp\_pade\_wrap(A, desca, p = 6)

**Arguments**

A      Matrix.  
desca      ScaLAPACK descriptor array.  
p      Order of the Pade' approximation.

**Details**

For advanced users only.

---

base.p\_matpow\_by\_squaring\_wrap  
*p\_matpow\_by\_squaring\_wrap*

---

### Description

Matrix power by squaring.

### Usage

```
base.p_matpow_by_squaring_wrap(A, desca, b = 1)
```

### Arguments

A	Matrix.
desca	ScaLAPACK descriptor array.
b	Power.

### Details

For advanced users only.

---

base.rcolcpy            *R Column Copy*

---

### Description

For advanced users only.

### Usage

```
base.rcolcpy(x, descx, y, descy, xcol, ycol)
```

### Arguments

x, y	Matrix.
descx, descy	ScaLAPACK descriptor array.
xcol, ycol	Columns.

---

base.rcolcpy2	<i>R Column Copy-2</i>
---------------	------------------------

---

**Description**

For advanced users only.

**Usage**

```
base.rcolcpy2(x, descx, y, descy, xcol, ycol)
```

**Arguments**

x, y	Matrix.
descx, descy	ScaLAPACK descriptor array.
xcol, ycol	Columns.

---

base.redist	<i>base.redist</i>
-------------	--------------------

---

**Description**

Redistribute a matrix from rank 0 to all ranks in block cyclic fashion.

**Usage**

```
base.redist(desc, A)
```

**Arguments**

desc	ScaLAPACK descriptor array.
A	Matrix.

---

base.rl2blas	<i>Level 2 R-like BLAS</i>
--------------	----------------------------

---

**Description**

For advanced users only.

**Usage**

```
base.rl2blas(x, descx, vec, FUN)
```

**Arguments**

x	Matrix.
descx	ScaLAPACK descriptor array.
vec	Global vector.
FUN	Function.

---

base.rl2insert	<i>R-like Matrix-Vector Insertion</i>
----------------	---------------------------------------

---

**Description**

For advanced users only.

**Usage**

```
base.rl2insert(x, descx, vec, i, j)
```

**Arguments**

x	Matrix.
descx	ScaLAPACK descriptor array.
vec	Global vector.
i, j	Indices.

---

base.rpdgecon	<i>rpdgecon</i>
---------------	-----------------

---

**Description**

Inverse condition number of a general matrix.

**Usage**

```
base.rpdgecon(norm, m, n, a, desca)
```

**Arguments**

norm	Type of norm.
m, n	Problem size
a	Matrix.
desca	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdgelqf	<i>rpdgelqf</i>
---------------	-----------------

---

**Description**

LQ.

**Usage**

```
base.rpdgelqf(m, n, x, descx)
```

**Arguments**

m, n	Problem size.
x	Matrix.
descx	ScaLAPACK descriptor array.

**Details**

For advanced users only.



---

base.rpdgels	<i>rpdgels</i>
--------------	----------------

---

**Description**

Linear model fitter via rank-revealing QR (with pivoting).

**Usage**

```
base.rpdgels(tol, m, n, nrhs, a, desca, b, descb)
```

**Arguments**

tol	Numerical tolerance for the QR.
m, n	Problem size.
nrhs	Number of right hand sides.
a	Left hand side.
desca	ScaLAPACK descriptor array.
b	Right hand side.
descb	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdgemm	<i>rpdgemm</i>
--------------	----------------

---

**Description**

Matrix-Matrix Multiply.

**Usage**

```
base.rpdgemm(transx, transy, x, descx, y, descy, descc)
```

**Arguments**

transx, transy	'T' or 'N' for transpose or not.
x, y	Matrix.
descx, descy, descc	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdgemr2d	<i>rdgemr2d</i>
----------------	-----------------

---

**Description**

General 2d block cyclic redistribution function.

**Usage**

```
base.rpdgemr2d(x, descx, descy)
```

**Arguments**

x	Matrix.
descx, descy	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdgeqpf	<i>rdgeqpf</i>
---------------	----------------

---

**Description**

QR.

**Usage**

```
base.rpdgeqpf(tol, m, n, x, descx, comm = .pbd_env$SPMD.CT$comm)
```

**Arguments**

tol	Numerical tolerance for the QR.
m, n	Problem size.
x	Matrix.
descx	ScaLAPACK descriptor array.
comm	An MPI (not BLACS) communicator.

**Details**

For advanced users only.

---

base.rpdgesv	<i>rpdgesv</i>
--------------	----------------

---

**Description**

Solving a (square) system of equations.

**Usage**

```
base.rpdgesv(n, nrhs, a, desca, b, descb)
```

**Arguments**

n	Problem size.
nrhs	Number of right hand sides.
a, b	Matrix.
desca, descb	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdgesvd	<i>rpdgesvd</i>
---------------	-----------------

---

**Description**

SVD.

**Usage**

```
base.rpdgesvd(jobu, jobvt, m, n, a, desca, descu, descvt, ...,
  inplace = FALSE, comm = .pbd_env$SPMD.CT$comm)
```

**Arguments**

jobu, jobvt	Control for u/vt return.
m, n	Problem size.
a	Matrix.
desca, descu, descvt	ScaLAPACK descriptor array.
...	Ignored
inplace	Should the computation be done in-place or not. For REALLY advanced users only.
comm	An MPI (not BLACS) communicator.

**Details**

For advanced users only.

---

base.rpdgetrf	<i>rpdgetrf</i>
---------------	-----------------

---

**Description**

LU factorization.

**Usage**

base.rpdgetrf(a, desca)

**Arguments**

a	Matrix.
desca	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdgetri	<i>rpdgetri</i>
---------------	-----------------

---

**Description**

Matrix inversion.

**Usage**

base.rpdgetri(n, a, desca)

**Arguments**

n	Problem size.
a	Matrix.
desca	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdlange	<i>rpdlange</i>
---------------	-----------------

---

**Description**

Matrix norms.

**Usage**

```
base.rpdlange(norm, m, n, a, desca)
```

**Arguments**

norm	Type of norm.
m, n	Problem size
a	Matrix.
desca	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdlaprnt	<i>rpdlaprnt</i>
----------------	------------------

---

**Description**

Matrix printer.

**Usage**

```
base.rpdlaprnt(m, n, a, desca)
```

**Arguments**

m, n	Number rows/cols.
a	Matrix.
desca	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdorglq	<i>rpdorglq</i>
---------------	-----------------

---

**Description**

Recover Q.

**Usage**

base.rpdorglq(m, n, k, lq, desc, tau)

**Arguments**

m, n	Problem size.
k	Number of elementary reflectors.
lq	QR decomposition.
desc	ScaLAPACK descriptor array.
tau	Elementary reflectors.

**Details**

For advanced users only.

---

base.rpdorgqr	<i>rpdorgqr</i>
---------------	-----------------

---

**Description**

Recover Q.

**Usage**

base.rpdorgqr(m, n, k, qr, descqr, tau)

**Arguments**

m, n	Problem size.
k	Number of elementary reflectors.
qr	QR decomposition.
descqr	ScaLAPACK descriptor array.
tau	Elementary reflectors.

**Details**

For advanced users only.

---

base.rpdormqr	<i>rpdormqr</i>
---------------	-----------------

---

**Description**

$op(Q) * y$ .

**Usage**

base.rpdormqr(side, trans, m, n, k, qr, descqr, tau, c, desc)

**Arguments**

side	'L' or 'R', for left or righth application of Q matrix.
trans	Q or Q <sup>T</sup> .
m, n	Problem size.
k	Number of elementary reflectors.
qr	QR decomposition.
descqr	ScaLAPACK descriptor array.
tau	Elementary reflectors.
c	Vector.
desc	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdpotrf	<i>rpdpotrf</i>
---------------	-----------------

---

**Description**

Cholesky factorization.

**Usage**

base.rpdpotrf(uplo, n, a, desca)

**Arguments**

uplo	Triangle where the information is stored (in the symmetric matrix).
n	Problem size.
a	Matrix.
desca	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdsyevr	<i>rpdsyevr</i>
---------------	-----------------

---

**Description**

Symmetric eigenvalue decomposition.

**Usage**

```
base.rpdsyevr(jobz, uplo, n, a, desca, descz)
```

**Arguments**

jobz	Control for if vectors/values/both are returned.
uplo	Triangle where the information is stored (in the symmetric matrix).
n	Problem size.
a	Matrix.
desca, descz	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdsyevx	<i>rpdsyevx</i>
---------------	-----------------

---

**Description**

Generalized eigenvalue problem.

**Usage**

```
base.rpdsyevx(jobz, range, n, a, desca, vl, vu, il, iu, abstol = 1e-08,
  orfac = 0.001)
```



**Arguments**

jobz	Control for if vectors/values/both are returned.
range	Parameter to determine the search criteria for eigenvalues.
n	Problem size.
a	Matrix.
desca	ScaLAPACK descriptor array.
v1, vu	Endpoints of the interval subset of the real line in which to search for eigenvalues, if specified by range.
il, iu	Eigenvalues with indices il, ..., iu will be found, if specified by range.
abstol	Absolute error tolerance for the eigenvalues.
orfac	Eigenvectors with eigenvalues below orfac*norm(a) of each other are reorthogonalized.

**Details**

For advanced users only.

---

base.rpdtran	<i>rpdtran</i>
--------------	----------------

---

**Description**

Transpose.

**Usage**

base.rpdtran(a, desca, descc)

**Arguments**

a	Matrix.
desca, descc	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rpdtrcon	<i>rpdrcon</i>
---------------	----------------

---

**Description**

Inverse condition number of a triangular matrix.

**Usage**

```
base.rpdtrcon(norm, uplo, diag, n, a, desca)
```

**Arguments**

norm	Type of norm.
uplo	Triangle where information is stored.
diag	Specifies if the matrix is unit triangular or not.
n	Problem size
a	Matrix.
desca	ScaLAPACK descriptor array.

**Details**

For advanced users only.

---

base.rrowcpy	<i>R Row Copy</i>
--------------	-------------------

---

**Description**

For advanced users only.

**Usage**

```
base.rrowcpy(x, descx, y, descy, xrow, yrow)
```

**Arguments**

x, y	Matrix.
descx, descy	ScaLAPACK descriptor array.
xrow, yrow	Rows.

---

base.rrowcpy2	<i>R Row Copy-2</i>
---------------	---------------------

---

**Description**

For advanced users only.

**Usage**

```
base.rrowcpy2(x, descx, y, descy, xrow, yrow)
```

**Arguments**

x, y	Matrix.
descx, descy	ScaLAPACK descriptor array.
xrow, yrow	Rows.

---

base.tri2zero	<i>tri2zero</i>
---------------	-----------------

---

**Description**

Zero Triangle

**Usage**

```
base.tri2zero(x, descx, uplo = "L", diag = "N")
```

**Arguments**

x	Matrix.
descx	ScaLAPACK descriptor array.
uplo	Triangle.
diag	Zero diagonal as well.

**Details**

For advanced users only.

---

base.valid\_context      *BLACS Context Validation*

---

### Description

Checks if a supplied ICTXT is valid.

### Usage

```
base.valid_context(ICTXT, ..., override = FALSE)
```

### Arguments

ICTXT	BLACS context number.
...	Not used.
override	If override=FALSE, the context number will produce an error if it is any of the reserved contexts (0, 1, or 2).

---

blacsexit                      *BLACS Exit*

---

### Description

Shuts down all BLACS communicators.

### Usage

```
base.blacsexit(CONT = TRUE)
```

```
blacsexit(CONT = TRUE)
```

### Arguments

CONT	logical; determines whether or not to shut down <i>all</i> MPI communicators
------	--

### Details

If the user wishes to shut down BLACS communicators but still have access to MPI, then call this function with CONT=TRUE. Calling blacsexit(CONT=FALSE) will shut down all MPI communicators, equivalent to calling

```
> blacsexit(CONT=TRUE) > finalize(mpi.finalize=TRUE)
```

This function is automatically invoked if BLACS communicators are running and finalize() is called.

**Value**

Has an invisible return of 0 when successful.

**Examples**

```

spmd.code = "
  suppressMessages(library(pbdBASE))
  init.grid()

  blacsexit()

  # finalize() # This should be off since blacexit().
"

pbdMPI::execmpi(spmd.code = spmd.code, nrank = 2L)

```

---

blacs\_apt

*Functions to set and get BLACS\_APTS*


---

**Description**

To set and get BLACS array/object/whatever pointers needed in and from R. Because other packages has it's own memory stack vision that may not be visiable by this package or vice versa.

**Usage**

```

set.blacs.apt()

get.blacs.apt()

```

**Details**

The 'set.blacs.apt()' is for advanced users. This one is needed to be called within R from 'pbd-BASE' package to set the pointers to the memory where BLACS had initialized so that the pointers are set to the right address of the memory stack.

The 'get.blacs.apt()' is for debugging only. The advanced user mainly calls the C version 'get\_BLACS\_APTS\_from\_R()' in 'src/export\_blacs/pkg\_oos.c'.

I am lazy to use .C(), but should not hurt performance here. Eventually, .pbdBASEEnv should pass to .C() and set/get pointers from it instead of .GlobalEnv.

---

coords *Local to Global/Global to Local Indexing*

---

### Description

Get the local index given global information.

### Usage

```
indxg2l(INDXGLOB, NB, IPROC, ISRCPROC, NPROCS)
```

```
indxl2g(INDXLOC, NB, IPROC, ISRCPROC, NPROCS)
```

### Arguments

INDXGLOB	Global index.
NB	Block size.
IPROC	Coordinate of the process whose local info is to be determined.
ISRCPROC	The coordinate of the process that possesses the first row/column of the distributed matrix. That's always 0 pbdDMAT.
NPROCS	Total number of processors over which matrix is distributed.
INDXLOC	Local index.

### Details

For advanced users only.

---

coordspair *Global to Local/Local to Global Pair Indexing*

---

### Description

Get the local index-pair given global information.

### Usage

```
g2lpair(gi, gj, bldim, ICTXT)
```

```
l2gpair(i, j, bldim, ICTXT)
```

**Arguments**

<code>gi, gj</code>	Global indices.
<code>bldim</code>	Blocking dimension
<code>ICTXT</code>	BLACS context.
<code>i, j</code>	Local indices.

**Details**

For advanced users only.

---

<code>diag</code>	<i>diag</i>
-------------------	-------------

---

**Description**

Grab diagonal or create distributed diagonal matrix.

**Usage**

```
base.ddiagtk(x, descx, proc.dest = "all")
```

```
base.ddiagmk(diag, descx)
```

**Arguments**

<code>x</code>	Matrix.
<code>descx</code>	ScaLAPACK descriptor array.
<code>proc.dest</code>	Who owns the result.
<code>diag</code>	Diagonal.

**Details**

For advanced users only.

---

finalizer	<i>Finalizer</i>
-----------	------------------

---

**Description**

A replacement for `pbdMPI::finalize()` that automatically shuts BLACS communicators down.

**Usage**

```
base.finalize(mpi.finalize = .pbd_env$SPMD.CT$mpi.finalize)
```

```
finalize(mpi.finalize = .pbd_env$SPMD.CT$mpi.finalize)
```

**Arguments**

`mpi.finalize` If MPI should be shut down.

---

g2lcoord	<i>g2lcoord</i>
----------	-----------------

---

**Description**

Global to local coordinates with explicit ownership given.

**Usage**

```
g2lcoord(dim, bldim, gi, gj, gridinfo)
```

**Arguments**

`dim` Global dimension.

`bldim` Blocking dimension.

`gi, gj` Global row and column indices, respectively.

`gridinfo` The return of `base.blacs(ICTXT(x))`. See the Details section for more information.

**Value**

For the process that owns the desired local data at global indices (`gi, gj`), the return is the local index. Otherwise, NA is returned.



---

g2l_coord	<i>g2l_coord</i>
-----------	------------------

---

**Description**

Global to local coords.

**Usage**

```
base.g2l_coord(ind, bldim, ICTXT = 0, dim = NULL)
```

```
g2l_coord(ind, bldim, ICTXT = 0, dim = NULL)
```

**Arguments**

ind	Matrix indices.
bldim	Blocking dimension.
ICTXT	BLACS context.
dim	Ignored; will be removed in a future version.

**Details**

For advanced users only.

---

get.comm.from.ICTXT	<i>Getting Communicator From BLACS Context</i>
---------------------	--

---

**Description**

Blacs context are associated with a certain communicator. It can be useful to retrieve this communicator to manipulate the matrix accordingly.

**Usage**

```
get.comm.from.ICTXT(ICTXT)
```

**Arguments**

ICTXT	a BLACS context
-------	-----------------

---

gridexit	<i>gridexit</i>
----------	-----------------

---

### Description

Frees a BLACS context.

### Usage

```
base.gridexit(ICTXT, override = FALSE)
```

```
gridexit(ICTXT, override = FALSE)
```

### Arguments

ICTXT	BLACS context number.
override	logical; if TRUE, ignores normal check preventing the closing of ICTXT values of 0, 1, and 2. This could cause things to go crazy and I do not recommend it.

### Details

For advanced users only.

The function frees the requested BLACS context. It is a trivial wrapper for the BLACS routine BLACS\_GRIDEXIT. Also removes the object `._blacs_gridinfo_ICTXT`.

Contexts 0, 1, and 2 can not be freed in this way unless the argument `override=FALSE`. This will probably break something and I do not recommend it.

### Value

Silently returns 0 when successful. Silently returns 1 when requested ICTXT does not exist.

---

gridinfo	<i>Get BLACS Context Grid Information</i>
----------	---

---

### Description

Grabs the existing BLACS context grid information.

### Usage

```
base.blacs(ICTXT = 0)
```

```
blacs(ICTXT = 0)
```

**Arguments**

ICTXT            BLACS context number.

**Details**

BLACS contexts have important internal use, and advanced users familiar with ScaLAPACK might find some advantage in directly manipulating these process grids. Most users should not need to directly manage BLACS contexts, in this function or elsewhere.

The function effectively serves as a shorthand for

```
eval(parse(text=paste("__blacs_gridinfo_", ICTXT, sep="")))
```

**Value**

Returns a list with 5 elements: NPROW and NPCOL, the number of process rows and columns respectively; ICTXT, the associated BLACS context number; MYROW and MYCOL, the current process' row and column position in the process grid.

**Examples**

```
spmd.code = "
  suppressMessages(library(pbdBASE))
  init.grid()

  mygrid <- blacs(0)

  pbdMPI::comm.print(mygrid)

  finalize()
"

pbdMPI::execmpi(spmd.code = spmd.code, nrank = 2L)
```

---

gridinit

*blacs\_init*

---

**Description**

BLACS grid initialization.

**Usage**

```
base.blacs_init(ICTXT, NPROW, NPCOL, ..., quiet = FALSE)
```

```
blacs_init(ICTXT, NPROW, NPCOL, ..., quiet = FALSE)
```

```
blacs_gridinit(ICTXT, NPROW, NPCOL, ..., quiet = FALSE)
```

**Arguments**

ICTXT	BLACS context.
NPROW, NPCOL	Number of process rows/cols.
...	Additional arguments.
quiet	Verbose initialization or not.

**Details**

For advanced users only.

---

InitGrid	<i>Initialize Process Grid</i>
----------	--------------------------------

---

**Description**

Manages the creation of BLACS context grids.

**Usage**

```
init.grid(NPROW, NPCOL, ICTXT, quiet = FALSE)
```

**Arguments**

NPROW	number of process rows. Can be missing; see details.
NPCOL	number of process columns. Can be missing; see details.
ICTXT	BLACS context number.
quiet	logical; controls whether or not information about grid size should be printed.

**Details**

`blacs_init()` is for experienced users only. It is a shallow wrapper of the BLACS routine `BLACS_INIT`, with the addition of creating the `.__blacs_gridinfo_``ICTXT` objects, as described below.

The remainder of this section applies only to `init.grid()`.

If `ICTXT` is missing, three variables will be created in the `.pbdBASEEnv` environment:

```
.__blacs_gridinfo_0
.__blacs_gridinfo_1
.__blacs_gridinfo_2
```

These variables store the BLACS process grid information for the BLACS context corresponding to the trailing digit of the variable. Most users should invoke `init.grid()` in this fashion, namely with `ICTXT` missing, and only do so once.

Contexts 0, 1, and 2 are reserved. Additional custom contexts are possible to create, but they must be integers  $\geq 3$ .

Context 0 is the “full” process grid of NPROW by NPCOL processes; contexts 1 is the process grid consisting of 1 process row and NPROW\*NPCOL processes columns; context 2 is the process grid consisting of NPROW\*NPCOL processes rows and 1 process column. These contexts can be redundant depending on the number of processes available.

BLACS contexts have important internal use, and advanced users familiar with ScaLAPACK might find some advantage in directly manipulating these process grids. Most users should not need to directly manage BLACS contexts, in this function or elsewhere.

If the NPROW and NPCOL values are missing, then a best process grid will be chosen for the user based on the total available number of processes. Here “best” means as close to a square grid as possible.

The variables `._blacs_gridinfo_ICTXT` are just storage mechanisms to avoid needing to directly invoke the BLACS routine `BLACS_GRIDINFO`.

Additionally, another variable is created in the `.pbdBASEEnv` environment, namely `._blacs_initialized`. Its existence is to alert `finalize()` to shut down BLACS communicators, if necessary, to prevent memory leaks.

### Value

Silently returns 0 when successful. Additionally, several variables are created in the `.pbdBASEEnv` environment. See Details section.

### Examples

```
spmd.code = "
  suppressMessages(library(pbdBASE))
  init.grid()

  finalize()
"

pbdMPI::execmpi(spmd.code = spmd.code, nrank = 2L)
```

---

l2g\_coord

*l2g\_coord*


---

### Description

Local to global coords.

### Usage

```
base.l2g_coord(ind, bldim, ICTXT = 0, dim = NULL)
```

```
l2g_coord(ind, bldim, ICTXT = 0, dim = NULL)
```

**Arguments**

ind	Matrix indices.
bldim	Blocking dimension.
ICTXT	BLACS context.
dim	Ignored; will be removed in a future version.

**Details**

For advanced users only.

---

numroc2	<i>numroc2</i>
---------	----------------

---

**Description**

A better version of NUMROC (NUMBER Rows Or Columns). Returns the local dimension given global matrix + distribution parameters.

**Usage**

```
numroc2(N, NB, IPROC, NPROCS)
```

**Arguments**

N	Global number of rows/cols.
NB	Block size.
IPROC	Coordinate of the process whose local info is to be determined.
NPROCS	Total number of processors over which matrix is distributed.

**Details**

For advanced users only.

pcoords

*Interchange Between Process Number and BLACS Coordinates***Description**

Grabs the existing BLACS context grid information.

**Usage**

```
base.pnum(ICTXT, PROW, PCOL)
```

```
base.pcoord(ICTXT, PNUM)
```

**Arguments**

ICTXT	BLACS context number.
PROW, PCOL	BLACS grid location row/column
PNUM	process rank

**Details**

For advanced users only. These functions are simple recreations of the BLACS routines BLACS\_PNUM and BLACS\_PCOORD. The former gets the process number associated with the BLACS process grid location c(MYPROW, MYPCOL), while the latter does the reverse.

**Value**

pnum returns an integer; pcoord returns a list containing elements PROW and PCOL.

**Examples**

```
spmd.code = "
  suppressMessages(library(pbdBASE))
  init.grid()

  blacs_ <- blacs(ICTXT = 0)

  # get the ICTXT = 0 BLACS coordinates for process 0
  myCoords <- base.pcoord(ICTXT = 0, PNUM = 0)

  comm.print(myCoords)

  finalize()
"
```

```
pbdMPI::execmpi(spmd.code = spmd.code, nranks = 2L)
```

---

sys2blacs.handle      *Context Within a Given Communicator*

---

**Description**

Creates a context that will be valid for a given communicator

**Usage**

```
sys2blacs.handle(comm)
```

**Arguments**

comm                      Communicator for which you want to set the BLACS context

**Value**

A system handle, i.e. the system context number. System contexts can be used to have ScalaPACK methods run in different communicators.

**See Also**

base.free\_blacs\_system\_handle, base.blacs\_gridinit



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