

# Distributor2 2.4

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## 1 Distributor2: block distribution module

### 1.1 Preamble

This module provides functions to distribute blocks on a given number of processors. At the end of the process, each block will have a number corresponding to the processor it must be affected to for a balanced computation, depending on given criterias. This module doesn't perform splitting (see the Transform module for that).

This module is part of Cassiopee, a free open-source pre- and post-processor for CFD simulations.

To use the module with the Converter array interface, you must import it as:

```
import Distributor2 as D2
```

To use the module with the pyTree interface, you must import it as:

```
import Distributor2.PyTree as D2
```

### 1.2 Automatic load balance

**D2.distribute:** distribute automatically the blocks amongst N processors.

With the array interface, where A is a list of blocks:

- prescribed is a list of blocks that are forced to be on a given processor. `prescribed[2] = 0` means that block 2 MUST be affected to processor 0.
  - perfo is a tuple or a tuple list for each processor. Each tuple describes the relative weight of solver CPU time regarding the communication speed and latence (solverWeight, latenceWeight, comSpeedWeight).
  - weight is a list of weight for each block indicating the relative cost for solving each block.
  - com is a  $i \times j$  matrix describing the volume of points exchanged between bloc  $i$  and bloc  $j$ .
- Algorithm can be chosen in: 'gradient', 'genetic', 'fast'.

The function output is a stats dictionary. `stats['distrib']` is a vector describing the attributed processor for each block, `stats['meanPtsPerProc']` is the mean number of points per proc, `stats['varMin']` is the minimum variation of number of points, `stats['varMax']` is the maximum variation of number of points, `stats['varRMS']` is the mean variation of number of points, `stats['nptsCom']` is the number of points exchanged between processors for communication, `stats['comRatio']` is the ratio of points exchanged between processors in this configuration divided by the total number of points needed in communications, `stats['adaptation']` is the value of the optimized function:

```
stats = D2.distribute(A, N, prescribed=[], perfo=[], weight=[], com=[], algorithm='gradient',
nghost=0)
```

With the pyTree interface, the user-defined node `.Solver#Param/proc` is updated with the attributed processor number.

If `useCom=0`, only the grid number of points is taken into account.

If `useCom='all'`, matching and overlap communications are taken into account.

If `useCom='match'`, only match connectivity are taken into account.

if `useCom='overlap'`, only overlap connectivity are taken into account.

if `useCom='bbox'`, overlap between zone bbox is taken into account.

When using distributed trees, `prescribed` must be a dictionary containing the zones names as key, and the prescribed proc as value. `weight` is also a dictionary where the keys are the zone names and the weight as the value. It is not mandatory to assign a weight to all the zones of the pyTree. Default value is assumed 1, only different weight values can be assigned to zones. `t` can be either a skeleton or a loaded skeleton pyTree for `useCom=0` or `useCom='match'`, but must be a loaded skeleton tree only for the other settings:

```
t, stats = D2.distribute(t, N, prescribed=, perfo=[], weight=, useCom='all', algorithm='gradient')
```

(See: [distribute.py](#)) (See: [distributePT.py](#))

### 1.3 Various operations

**D2.addProcNode:** add a "proc" node to all zones of A with given value:

```
B = D2.addProcNode(A, 12)
```

(See: [addProcNodePT.py](#))

**D2.getProc:** get the proc node of a zone or a list of zones:

```
proc = D2.getProc(a) .or: [proc1,proc2,...] = D2.getProc(A)
```

(See: [getProcPT.py](#))

**D2.getProcDict:** return a dictionary `proc['blocName']` identifying the attributed processor for a given bloc name. If `prefixByBase` is True, then the dictionary is `proc['baseName/blocName']`. A must be an already distributed tree:

```
procDict = D2.getProcDict(A, prefixByBase=False)
```

(See: [getProcDictPT.py](#))

**D2.copyDistribution:** copy the distribution of B to A matching zones by their name:

```
A = D2.copyDistribution(A, B)
```

(See: [copyDistributionPT.py](#))

**D2.redispach:** redispach a tree where a new distribution is defined in the node 'proc':

```
B = D2.redispach(A)
```

(See: [redispachPT.py](#))

## 1.4 Example files

Example file: [distribute.py](#)

```
# - distribute (array) -
import Generator as G
import Distributor2 as D2
import numpy

# Distribution sans communication entre blocs
N = 11
arrays = []
for i in xrange(N):
    a = G.cart( (0,0,0), (1,1,1), (10+i, 10, 10) )
    arrays.append(a)
out = D2.distribute(arrays, NProc=5); print out

# Distribution avec des perfos differentes pour chaque proc
out = D2.distribute(arrays, NProc=3, perfo=[(1,0,0), (1.2,0,0), (0.2,0,0)]); print out

# Distribution avec forçage du bloc 0 sur le proc 1, du bloc 2 sur le proc 3
# -1 signifie que le bloc est a equilibrer
prescribed = [-1 for x in xrange(N)]
prescribed[0] = 1; prescribed[2] = 3
out = D2.distribute(arrays, NProc=5, prescribed=prescribed); print out

# Distribution avec communications entre blocs, perfos identique pour tous
# les procs
volCom = numpy.zeros( (N, N), numpy.int32 )
volCom[0,1] = 100; # Le bloc 0 echange 100 pts avec le bloc 1
out = D2.distribute(arrays, NProc=5, com=volCom, perfo=(1,0.,0.1)); print out

# Distribution avec des solveurs differents pour les blocs (le solveur est 2
# fois plus couteux pour les bloc 2 et 4)
out = D2.distribute(arrays, weight=[1,2,1,2,1,1,1,1,1,1,1], NProc=3); print out
```

Example file: [distributePT.py](#)

```
# - distribute (pyTree) -
import Generator.PyTree as G
import Distributor2.PyTree as D2
import Converter.PyTree as C
import Connector.PyTree as X

N = 11
t = C.newPyTree(['Base'])
pos = 0
for i in xrange(N):
```

```

    a = G.cart((pos,0,0), (1,1,1), (10+i, 10, 10))
    pos += 10 + i - 1
    t[2][1][2].append(a)
t = X.connectMatch(t)

# Distribute on 3 processors
t, stats = D2.distribute(t, 3)
C.convertPyTree2File(t, 'out.cgns')

```

Example file: [addProcNodePT.py](#)

```

# - addProcNode (pyTree) -
import Converter.PyTree as C
import Generator.PyTree as G
import Distributor2.PyTree as D2

a = G.cart((0,0,0), (1,1,1), (10,10,10))
a = D2.addProcNode(a, 12)
C.convertPyTree2File(a, 'out.cgns')

```

Example file: [getProcPT.py](#)

```

# - getProc (pyTree) -
import Converter.PyTree as C
import Generator.PyTree as G
import Distributor2.PyTree as D2

a = G.cart((0,0,0), (1,1,1), (10,10,10))
a = D2.addProcNode(a, 12)
proc = D2.getProc(a); print proc

```

Example file: [getProcDictPT.py](#)

```

# - getProcDict (pyTree) -
import Generator.PyTree as G
import Distributor2.PyTree as D2
import Converter.PyTree as C
import Connector.PyTree as X

N = 11
t = C.newPyTree(['Base'])
pos = 0
for i in xrange(N):
    a = G.cart((pos,0,0), (1,1,1), (10+i, 10, 10))
    pos += 10 + i - 1
    t[2][1][2].append(a)

t = X.connectMatch(t)
t, stats = D2.distribute(t, 3)

proc = D2.getProcDict(t)
zoneNames = C.getZoneNames(t, prefixByBase=False)
for z in zoneNames: print z, proc[z]

# - or with base prefix -
proc = D2.getProcDict(t, prefixByBase=True)
zoneNames = C.getZoneNames(t, prefixByBase=True)
for z in zoneNames: print z, proc[z]

```

Example file: [copyDistributionPT.py](#)

```

# - copyDistribution (pyTree) -
import Converter.PyTree as C
import Distributor2.PyTree as D2
import Converter.Internal as Internal
import Generator.PyTree as G

# Case
N = 11
t = C.newPyTree(['Base'])
pos = 0
for i in xrange(N):
    a = G.cart((pos,0,0), (1,1,1), (10+i, 10, 10))
    a[0] = 'cart%d'%i
    pos += 10 + i - 1
    D2._addProcNode(a, i)
    t[2][1][2].append(a)

t2 = C.newPyTree(['Base'])
for i in xrange(N):
    a = G.cart((pos,0,0), (1,1,1), (10+i, 10, 10))
    a[0] = 'cart%d'%i
    pos += 10 + i - 1
    t2[2][1][2].append(a)
t2 = D2.copyDistribution(t2, t)
C.convertPyTree2File(t2, 'out.cgns')

```

### Example file: [redispatchPT.py](#)

```

# - redispatch (pyTree) -
import Converter.PyTree as C
import Distributor2.PyTree as D2
import Distributor2.Mpi as D2mpi
import Converter.Mpi as Cmpi
import Transform.PyTree as T
import Connector.PyTree as X
import Converter.Internal as Internal
import Generator.PyTree as G

# Case
N = 11
t = C.newPyTree(['Base'])
pos = 0
for i in xrange(N):
    a = G.cart((pos,0,0), (1,1,1), (10+i, 10, 10))
    pos += 10 + i - 1
    t[2][1][2].append(a)
t = X.connectMatch(t)
if (Cmpi.rank == 0): C.convertPyTree2File(t, 'in.cgns')
Cmpi.barrier()

# lecture du squelette
a = Cmpi.convertFile2SkeletonTree('in.cgns')

# equilibrage 1
(a, dic) = D2.distribute(a, NProc=Cmpi.size, algorithm='fast', useCom=0)

# load des zones locales dans le squelette
a = Cmpi.readZones(a, 'in.cgns', proc=Cmpi.rank)

# equilibrage 2 (a partir d'un squelette charge)
(a, dic) = D2.distribute(a, NProc=Cmpi.size, algorithm='gradient1',

```

```

        useCom='match')

a = D2mpi.redispatch(a)

# force toutes les zones sur 0
zones = Internal.getNodesFromType(a, 'Zone_t')
for z in zones:
    nodes = Internal.getNodesFromName(z, 'proc')
    Internal.setValue(nodes[0], 0)

a = D2mpi.redispatch(a)

# Reconstitue l'arbre complet a l'écriture
Cmpi.convertPyTree2File(a, 'out.cgns')

```