

## The LMDZORINCA configurations

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The LMDZORINCA configurations allow you to couple the LMDz atmospheric circulation model, the ORCHIDEE land model, and the INCA atmospheric chemistry model close to the CMIP6 coupled model.

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## 1. Description of the configuration

### 1.1. LMDZORINCA\_v6.1.11

LMDZORINCA\_v6.1.11 is the **default configuration** for Inca forced model. It is compatible with the IPSLCM6.1.11\_LR, LMDZOR\_v6.1.11 models. It is constructed with the following models:

- LMDZ branches/IPSLCM6.0.15 rev 3643 for the gcm and the create\_etat0\_limit
- ORCHIDEE tag tags/ORCHIDEE\_2\_0/ORCHIDEE rev 6592
- INCA tags/INCA5\_CMIP rev 928

This configuration allows you to work with the INCA version tuned for 39 levels and old LMDZ physics scheme for NMHC, NMHC\_AER, NMHC\_AER\_S, AER, GES and DUSS, and on 79 levels and new LMDZ physics scheme for AER.

The same version of INCA is used in IPSLCM5A2CHT.1 configuration.

### 1.2. LMDZORINCA\_v6.2\_work

LMDZORINCA\_v6.2\_work is a working configuration close to IPSLCM6\_v6.2\_work using last testing rev for each component

- LMDZ6 trunk for the gcm and the create\_etat0\_limit
- ORCHIDEE branches/ORCHIDEE\_2\_2/ORCHIDEE
- INCA trunk/INCA6

This configuration use the new way of compilation (with a script). Read [here](#) the documentation on it.

### 1.3. LMDZORINCA\_v6.2.2

LMDZORINCA\_v6.2.2 is the 6.2 configuration for production, close to IPSLCM6.2.2 and LMDZOR\_v6.2.2

- LMDZ6/branches/IPSL-CM6A-MR rev 3855 for the gcm and the create\_etat0\_limit
- branches/ORCHIDEE\_Quest/ORCHIDEE rev 7086
- tags/INCA6.2 rev 1082

### 1.4. ICOLMDZORINCA\_v7\_work

ICOLMDZORINCA\_v7\_work is a working configuration close to LMDZORINCA\_v6.2\_work configuration with the possibility to use Dynamico's dynamic instead of Lmdz's one. This configuration use last testing rev for each component :

- LMDZ6 trunk for the physic
- DYNAMICO master or LMDZ6 trunk for the dynamic
- ORCHIDEE branches/ORCHIDEE\_2\_2/ORCHIDEE
- INCA trunk/INCA6

With this configuration you can choose between running with a regular grid as in v6 configurations, or with an unstructured grid (with Dynamico)

### 1.5. ICOLMDZORINCA\_v7.2

ICOLMDZORINCA\_v7.2 is a configuration for production using dynamico's dynamic.

- LMDZ6 branches/LMDZ-INCA-Dyn rev 3965 for the physic
- DYNAMICO rebase/trunk/nudging for the dynamic

- ORCHIDEE branches/ORCHIDEE\_2\_2/ORCHIDEE rev 6594
- INCA branches/INCA\_DYNAMICOXIOS

## 2. Technical details

### 2.1. How to use it

#### 2.1.1. Compilation LMDZORINCA\_v6.2\_work

To compile v6.2, and v7 configuration you need to use the script `compile_lmdzorinca.sh`. Options to use this script are available with the command :

```
./compile_lmdzorinca.sh -h
```

or

```
./compile_icolmdzorinca.sh -h
```

If you compile in debug or in dev mode you need to modify the parameter `OptMode` in `config.card`. If you change the resolution at the compilation you need to modify the parameter `ResolAtm` in `config.card`

#### 2.1.2. Compilation of LMDZORINCA\_v6.1.10

To compile LMDZORINCA\_v6.1.10 configuration you need to use the Makefile store in `config/LMDZORINCA/` directory. You can choose between several target :

- VLR and Old LMDZ Physics scheme
  - NMHC\_AERxLMD9695-L39 (default one)
  - NMHC\_AER\_SxLMD9695-L39
  - NMHCxLMD9695-L39
  - AERxLMD9695-L39
  - DUSSxLMD9695-L39
  - GESxLMD9695-L39
- LR and New LMDZ Physics scheme
  - AERxLMD144142-L79
- no maintained anymore - ask to Anne COZIC if you want to use it
  - DUSSxLMD144142-L39

You can add the compilation of another resolution by copying the syntax of an existing resolution. If you do so, you will need to regrid all input files for the chosen chemistry model. [documentation To regrid the inca files](#)

### 2.2. Computing performances

- for VLR resolution (96x95x39) we advice to work with 48MPI and 12OMP for the `gcm` executable (`config.card`)
- for LR resolution (144x142x79) we advice to work with 71MPI and 8 OMP for the `gcm` executable (`config.card`)

```
ATM= (gcm_${ResolAtm}_${OptMode}_${ConfChem}).e, lmdz.x, 48MPI, 12OMP)
```

### 2.3. Restart files

For LMDZ and ORCHIDEE you can use restart files from any simulation of v6.1.10, or v6.2 family (even coupled model). For Inca you can start without any restart, or use one from a simulation using a version with a revision number bigger than 838

You can also use default restart files (initial states files) :

With the LMDZ model you can create initial files (`start.nc`, `startphy.nc` and `limit.nc`) for each simulation (for this, use the experiments :

`EXPERIMENTS/LMDZORINCA/CREATE_era/`). However, `start.nc` file only contain zero-values for tracers when the model is coupled to INCA. You must

therefore work on those values or use output files from existing simulations. By default, we provide one initial state per configuration.

They are defined in the [InitialStateFiles] section of lmdz.card. For example:

```
[InitialStateFiles]
List=  ( ${R_INIT} /CHM/LMDZORINCA/INCA${RESOL_CHM}/start_o2a_noTer.nc, start.nc ) \
       ( ${R_INIT} /CHM/LMDZORINCA/INCA${RESOL_CHM}/startphy_o2a_noTer.nc, startphy.nc )
```

## 2.4. Output frequencies

### 2.4.1. Output frequencies for LMDZORINCA\_v6.2 configurations

The output frequencies and the files to write is set in inca.card (idem for LMDZ in lmdz.card, ORCHIDEE in orchidee.card and stomate.card). See comments in the corresponding files. The [WriteFrequency?](#) option in config.card is not used any more.

### 2.4.2. Output frequencies for LMDZORINCA\_v6.1 configurations

for Inca output frequencies, you can choose by default between 1 day and / or 1 month. For this you need to fill in your choice in section [WriteFrequency?](#) for chemistry part

```
#D-- CHM -
[CHM]
#D-- choose inca ...
#D-- you can also ...
WriteFrequency="1D"
```

or

```
[CHM]
#D-- choose inca ...
#D-- you can also ...
WriteFrequency="1D 1M"
```

or

```
[CHM]
#D-- choose inca ...
#D-- you can also ...
WriteFrequency="1M"
```

If you want another output frequency you need to modify the file\_def\_inca\_\*.xml file in modeles/INCA/src/INCA\_XML/ directory. You will find two files for each chemistry configurations (ex: file\_def\_inca\_NMHC\_daily.xml and file\_def\_inca\_NMHC\_monthly.xml).

## 2.5. Flags to manage LMDZORINCA configuration

In inca.card, lmdz.card and orchidee.card you can modify several parameters in sections [UsersChoices]

### INCA

LMDZ_10m_winds	y (use LMDZ 10 meters winds)	n (use offline winds)		
wind_parameter	0.85 (default parameter value for threshold scheme to adjust 10 meters winds calculated by LMDZ)			

<b>feedb</b>	<b>0</b> (no feedback from aerosol on climate with Old Physics scheme)	<b>1</b> (aerosol effects selected by ok_ade and ok_aie in lmdz.card)		
<b>CoupOrchInca</b>	<b>y</b> (coupled model with ORCHIDEE for VOC)	<b>n</b>		
<b>calcul_flux</b>	<b>y</b> (calcul and write in output the chemistry flux of reactions)	<b>n</b>		
<b>flag_plane</b>	<b>0</b> (no aircraft)	<b>1</b> (old inca aircraft scheme)	<b>2</b> (new subsonic inca aircraft scheme)	<b>3</b> (subsonic + hypersonic inca aircraft scheme)
<b>emi_interp_time</b>	<b>0</b> (no time interpolation when using emission files)	<b>1</b> (point to point time interpolation when using emissions files - default)		

### 2.5.10. meters winds

If you choose to use offline winds, you need to define their path in the BoundaryFiles part of inca.card  
For Example :

```
[BoundaryFiles]
List=      ($CCCWORKDIR/../../subipsl/subipsl/ECMWF320x160/AN${year}/165_${year}${month}.nc      , u10mec.nc)\
          ($CCCWORKDIR/../../subipsl/subipsl/ECMWF320x160/AN${year}/166_${year}${month}.nc      , v10mec.nc)\
          (${R_BC}/ATM/LMDZORINCA/${RESOL_ATM}/NUDGE_FILES/ERA/AN${year}/u10m_ecmwf_${year}${month}.nc , u10m.nc )\
          (${R_BC}/ATM/LMDZORINCA/${RESOL_ATM}/NUDGE_FILES/ERA/AN${year}/v10m_ecmwf_${year}${month}.nc , v10m.nc )
```

### 2.5.11. The BVOC and Surface coupling

If you choose to activate the bvoc calcul in Orchidee (CHEMISTRY\_BVOC=y in Orchidee.def) you can copy the new flux value in Inca. For this you need to modify the parameter **CoupOrchInca** in inca.card. By default it will copy all bvoc flux : iso - mono - ORVOC - MBO - methanol - acetone - acetal - formal - acetic - formic - no\_soil - nox - fertil\_no - apinen - bpinen - limonen - myrcen - sabinen - camphen - 3caren - tbocimen - othermono - sesquiter.

And iso and mono are use in inca instead of the value read in sflx.nc.

**WARNING** it can only work with NMHC\_AER configuration.

snow, lai, veget, vegetfrac are also copy from Orchidee to Inca. You can choose to copy only these variables, for this you need to add in PARAM/inca.def the line :

```
nbFlux_FromOrch=0
```

If you want copy some flux, but not all the list, you need to modify your inca.def

```
nbFlux_FromOrch=3
emi_FromOrch=ORVOC acetone bpinen
```

You need to modify orchidee.card

```
CHEMISTRY_BVOC=y

# to decomment if we want to calcul bvoc - for this we need to activate CHEMISTRY_BVOC in orchidee.def_Choi
ListNonDel= (${R_IN}/SRF/chemistry/orchidee_fertilizer_1995.nc, .), \
            (${R_IN}/SRF/chemistry/orchidee_bbg_clim.nc, .)
```

and inca.card

```
CoupOrchInca=y
#
```

## 2.6. Lengths, frequencies

We advice you to use a PeriodLength of 1 month or 1 year, and a Pack Frequency of 1 year.

## 2.7. Computing centers

LMDZORINCA\_v6.1.10is available on TGCC and IDRIS computers

### 2.7.1. IDRIS-JeanZay

Because of JeanZay architecture (computing nodes based on sockets of 20 cores), we advice you to use 2, 4, 5, 10 or 20 OpenMP threads for LMDZ-ORCHIDEE component. By default, the number of OpenMP threads defined in config.card is 8. To change that, you have to edit and modify config.card as follows :

```
[Executable]
#D- For each component, Real name of executable, Name of executable for oasis
ATM= (gcm.e, lmdz.x, 48MPI, 10OMP)
```

Do not forget to install a new Job (command ins\_job) in order to take into account these modifications.