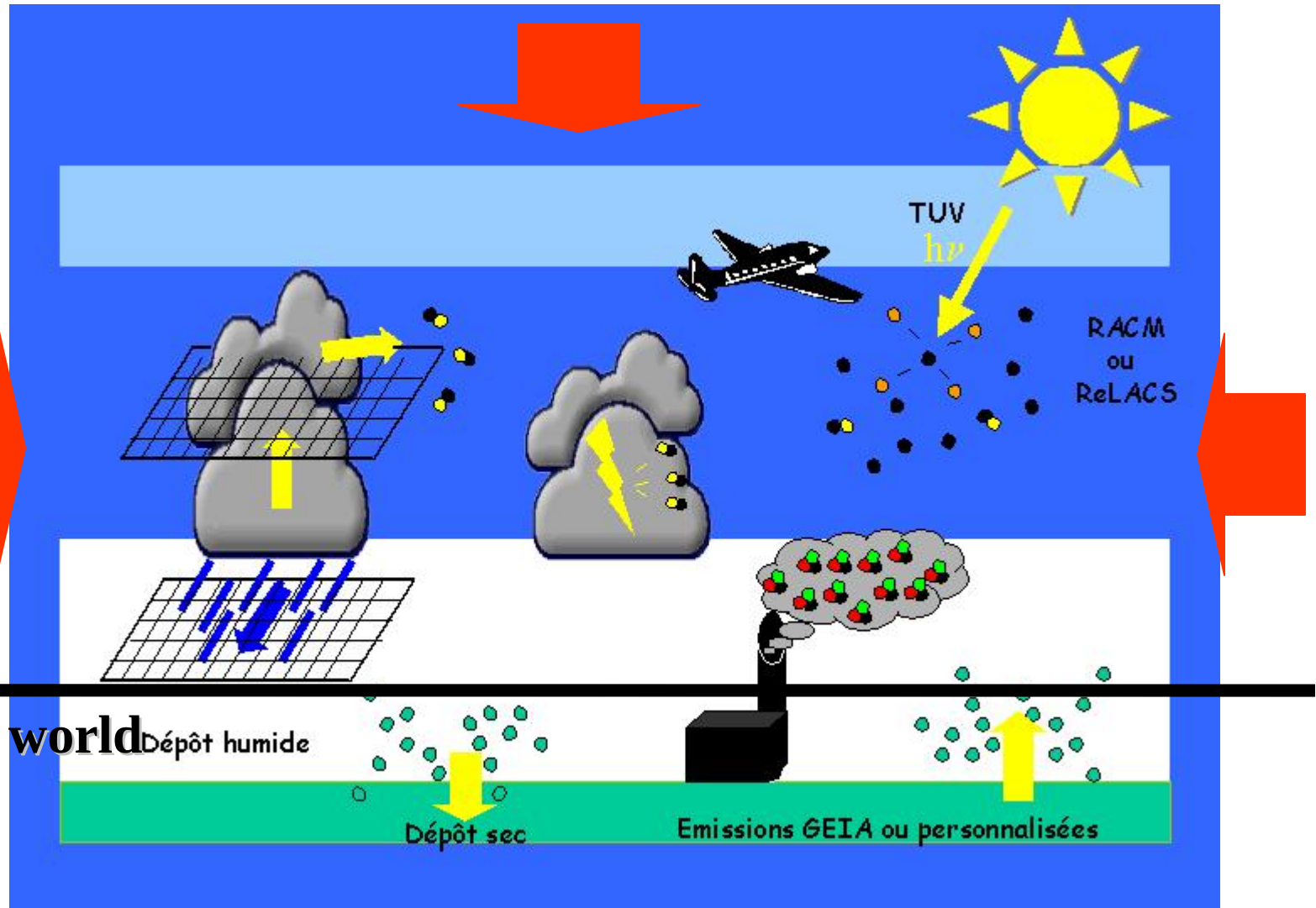


# Modelling of atmospheric chemistry from local (dx=1 km) to synoptic scale (dx=50 km)

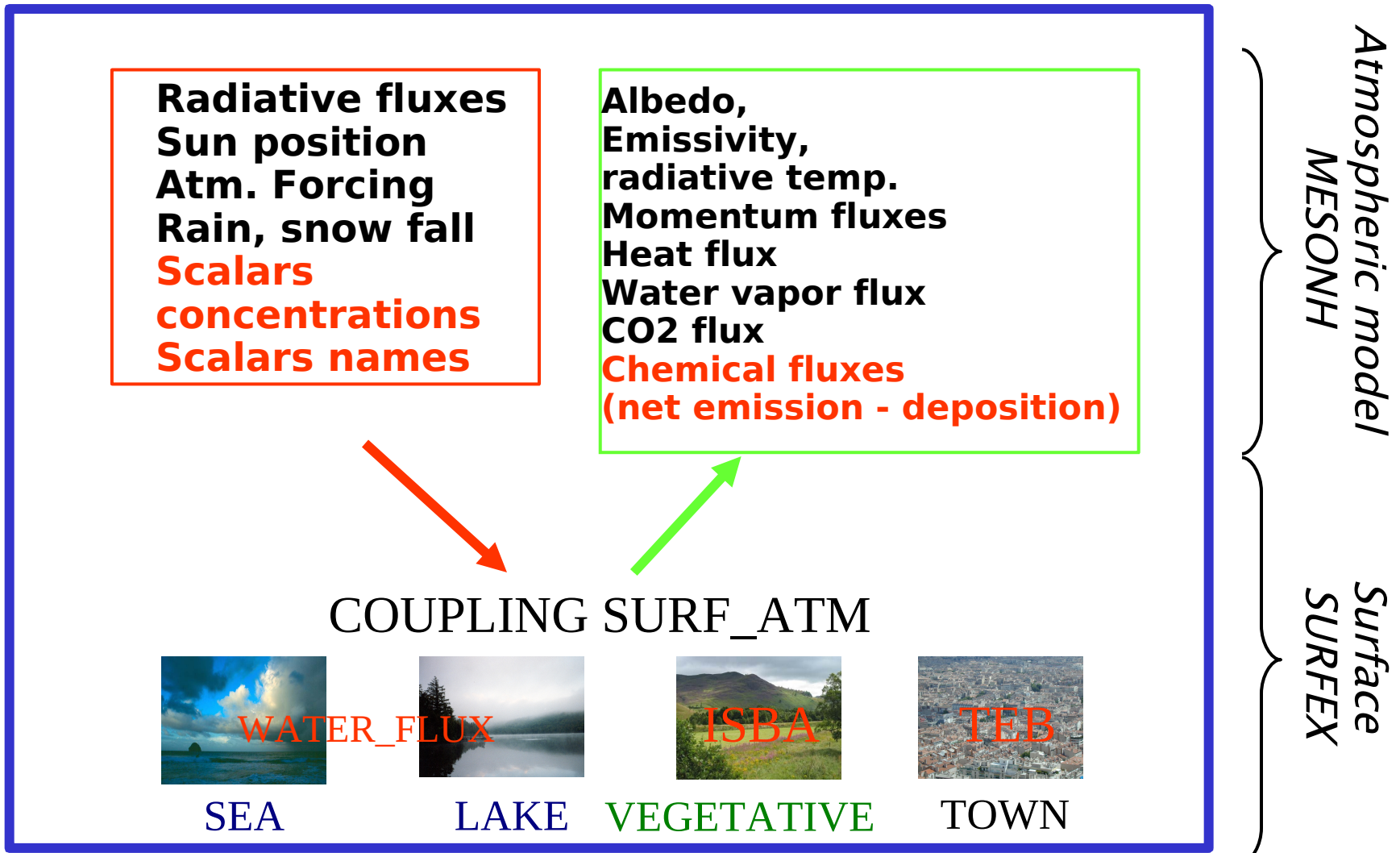
<http://www.aero.obs-mip.fr/mesonh>

Scale:  
NOAGE,  
ECMWF, ...



# 1. Surface (SURFEX) – Atmosphere (MESONH-AROME) exchanges

During run, at each timestep



## 2. Initiatilisation of Surfex scalar variable

at the first time step

Surfex recognize the parameterisation uses in the atmospheric model using a name convention for scalar variable :

Name started by « # » : Gaseous chemical variable => surface gas chemistry

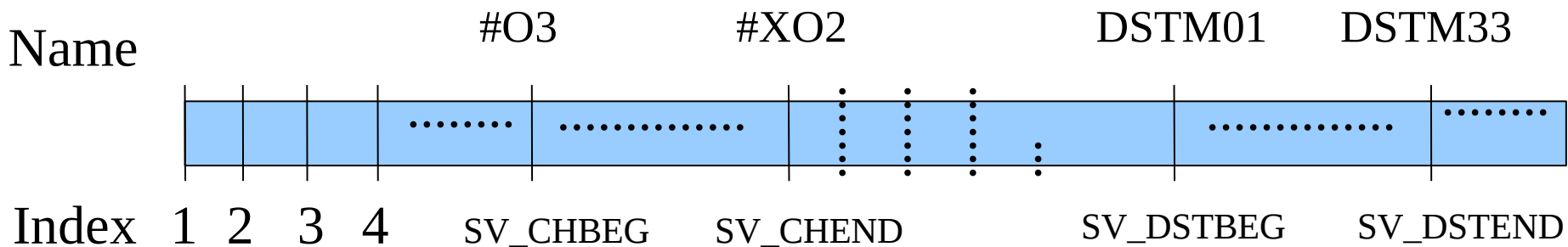
Name started by « @ » : Aerosol chemical variable => surface aerosol chemistry

Name started by « DST » : Dust aerosol variable => surface dust flux

Name started by « SLT » : Sea Salt aerosol variable => surface sea salt flux

Routines `ch_init_names`, `dst_init_names` and `slt_init_names` will recognize these conventions and initialize the begin and the ending indexes of each type of variable.

### Scalar variable vector

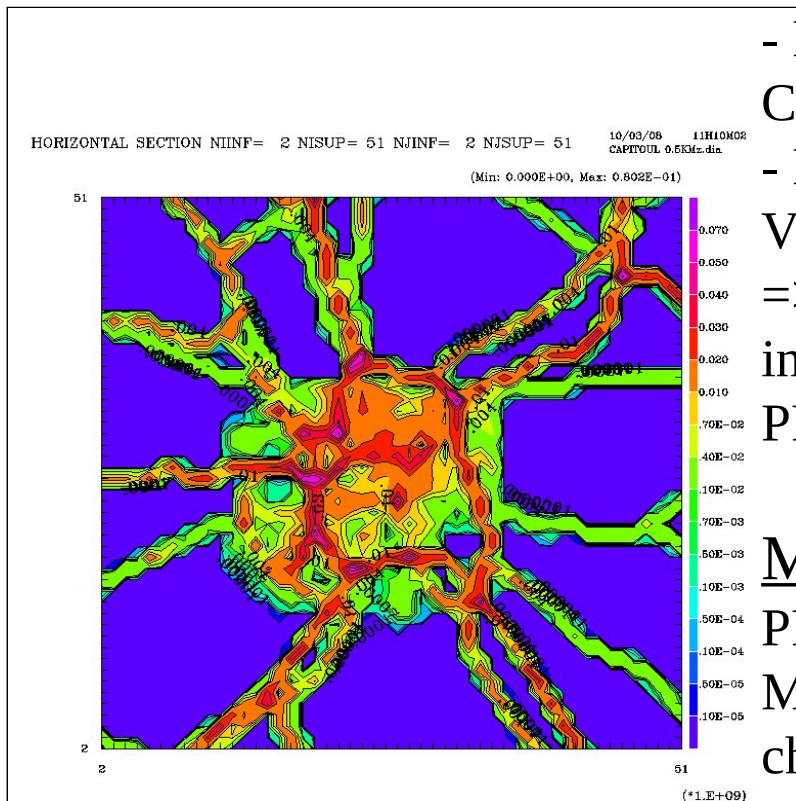


### 3. Off-line emission

Type : anthropogenic chemical emission.

Pollutants do not stictly depends on meteorological conditions: traffic, industries,...  
 So they need to be introduced using chemical emission database such as  
 GENEMIS, EDGAR, GEMS, GEIA....

Ex: Toulouse NO<sub>x</sub> flux



- Large number of primary pollutant: NO<sub>x</sub>, CO, VOCs, NH<sub>3</sub>, SO<sub>2</sub>, aérosols, ..

- High spacial and temporal

Variability

=> large number of fields have been interpolated in the simulation domain (done in the PREP\_PGD step).

Main ccde:

PREP\_PGD: pgd\_chemistry.f90

MesoNH or AROME: ch\_buidemissn.f90,

ch\_emission\_fluxn.f90

### 3. Off-line emission

**WHERE?** in namelist **PRE\_PGD1.nam** (surface field step) :

#### **HOW?**

&NAM\_CH\_EMIS\_PGD

NEMIS\_PGD\_NBR = n , ! Total number of chemical emission fields

CEMIS\_PGD\_NAME(1) = 'SO2', ! Name of emitted chemical species

CEMIS\_PGD\_AREA(1) = 'LAN', ! Interpolation on LAND only...

NEMIS\_PGD\_TIME(1) = 3600, ! Time of the day in seconds

CEMIS\_PGD\_FILE(1) = 'name of the 1st file',

CEMIS\_PGD\_FILESTYPE = 'BINLLV', ! File format binary (lat,lon,value)

CEMIS\_PGD\_NAME(2) = 'SO2', ! Name of emitted chemical species

CEMIS\_PGD\_AREA(2) = 'SEA', ! Interpolation on SEA only...

NEMIS\_PGD\_TIME(2) = 7200, ! Time of the day in seconds

CEMIS\_PGD\_FILE(2) = 'name of the 2nd file',

CEMIS\_PGD\_FILESTYPE = 'ASCLLV' ! File format ascii (lat, lon, value)

.....

CEMIS\_PGD\_NAME(n) = 'NOX', ! Name of emitted chemical species

CEMIS\_PGD\_AREA(n) = 'ALL', ! Interpolation on all pgd domain...

NEMIS\_PGD\_TIME(n) = 86400, ! Time of the day in seconds

CEMIS\_PGD\_FILE(n) = 'name of the n file',

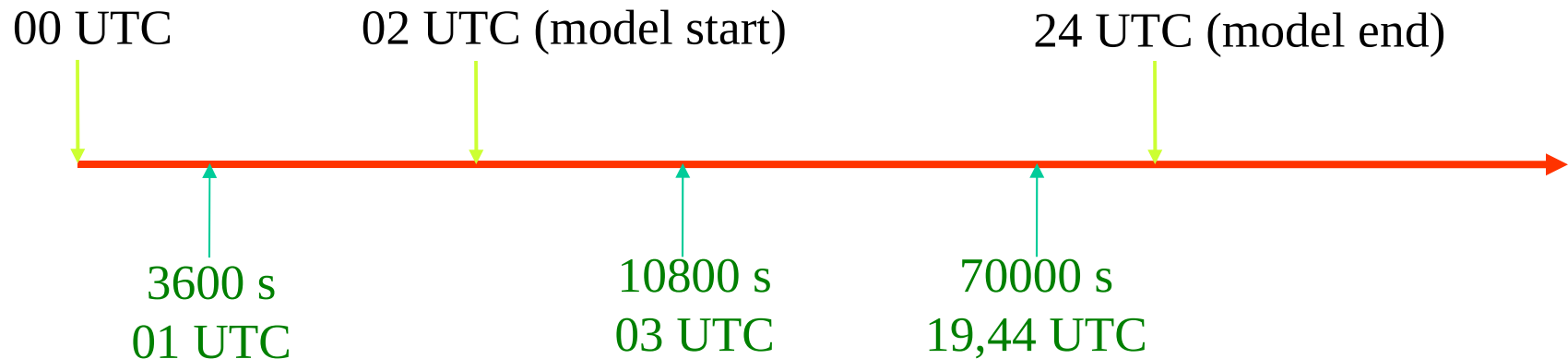
CEMIS\_PGD\_FILESTYPE = 'BINLLV' ! File format

.....

FILESGET\_LIST=« emis.tar » ! tar file containing all the emission files (CEMIS\_PGD\_FILE)

### 3. Off-line emission

*Example: You provide 3 emissions fields at 3600 s, 10800 s and 70000 s*



*The model will interpolate as follow:*

02 UTC < RUN < 03 UTC: temporal interpolation between fields at 3600 s and 7200 s

03 UTC < RUN < 19.44 UTC : temporal interpolation between fields at 7200 s and 70000 s

19.44 UTC < RUN < 24 UTC : temporal interpolation between 70000 s and 3600 s  
(return to the first values, if no time values are given after 70000s)

### 3. Off-line emission

How can I specify the link between emitted and prognostic chemicals ?

**WHERE?** By an ascii filename which the name is given by  
**CCHEM\_SURF\_FILE** of **&NAM\_CH\_CONTROL** in **EXSEG1.nam**

**HOW?**

**EMISUNIT** ! Keyword for off-line emission unit

**GENEMIS emission** ! Comment

**MIX** ! Unit (here ppp m/s)

**AGREGATION** ! Keyword for off-line emission agregation

For ReLACS prognostic species ! Comment

**NO 0.8 NOX** ! Flux of prognostic NO = 0.8 x prescribed flux of NOX

**NO2 0.2 NOX** ! Flux of the prognostic NO2 = 0.2 x prescribed flux of NOX

**CO CO** ! Flux of prognostic CO = prescribed flux of CO

**ALKA 1 HC3H03 1.3 HC8H30 ....**

! Flux of prognostic ALKA = 1 x prescribed flux of HC3H03 + 1.3 x prescribed flux of HC8H30

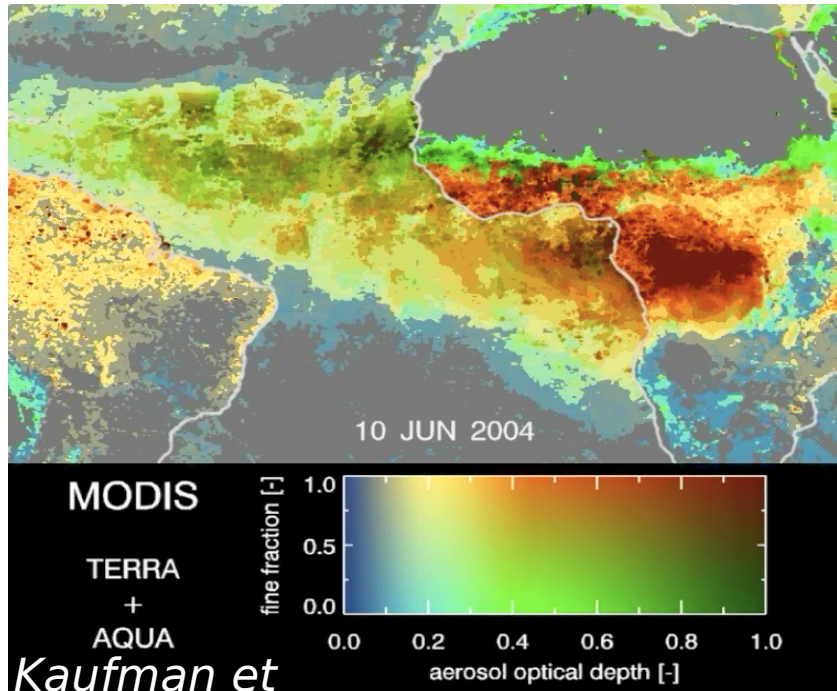
**END\_AGREGATION**



## 4. On-line emission

Type: emission that are dependant to meteorology and surface properties. Case of Dust and sea salt aerosols, Biogenics fluxes...  
 On-line Emission are parameterized are computed at each time step.

Example of dust observed by MODIS



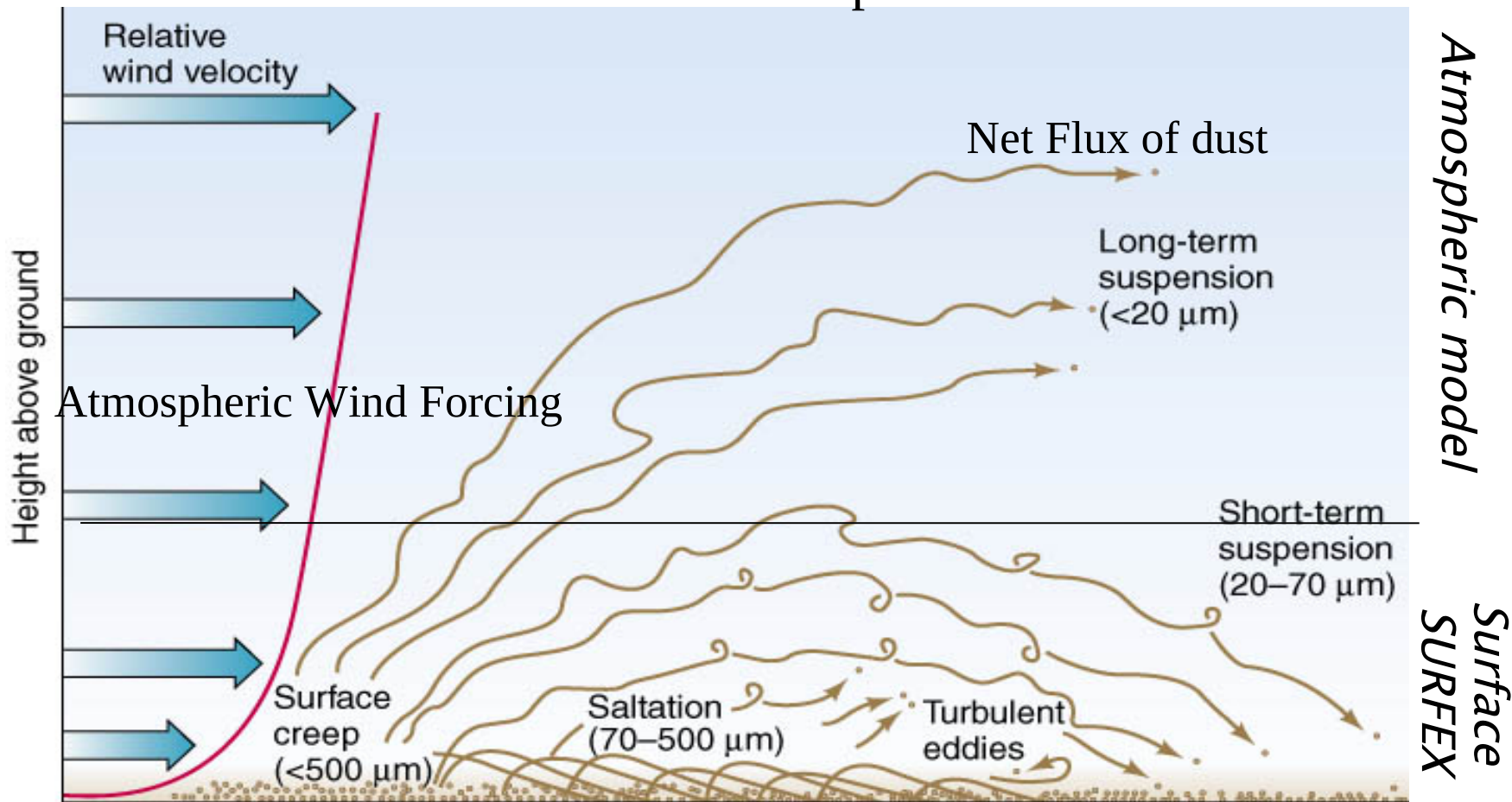
- Dust and Sea Salt parameterization are activated if Dust or Sea Salt names recognize the scalar variables names. But different parameterizations are introduced.

- For biogenics flux a flag exist (see follow)



## 4. On-line emission

### Dust emission case process



Surface properties (soil humidity, soil type, rugosity, vegetation,;..)

Main codes: coupling\_dstn.f90, mode\_dstmbl.f90, mode\_dstmblutl.f90, mode\_dst\_surf.f90, modd\_dst.f90

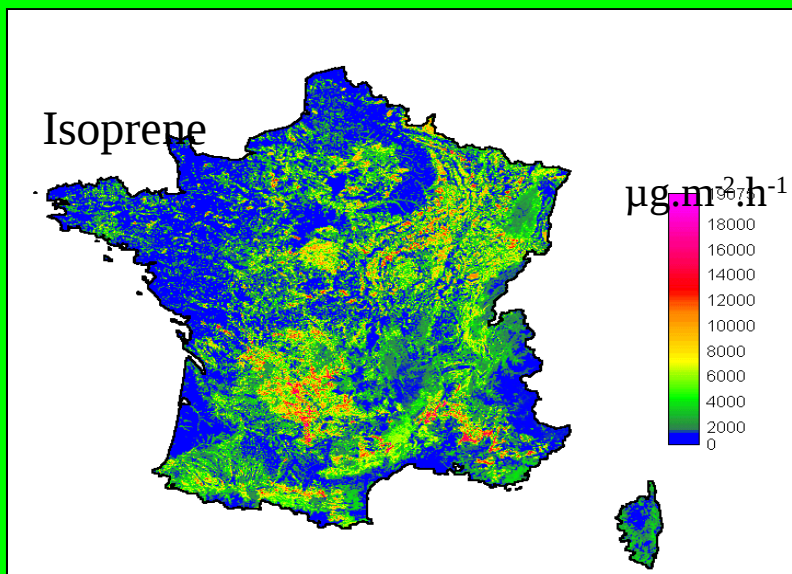
## 4. On-line emission

From Guenther et al. and  
Solmon et al, 2004

$$\text{FLUX bio} : \mathbf{F(t)} = \mathbf{EP} \times \mathbf{ECF(t)}$$

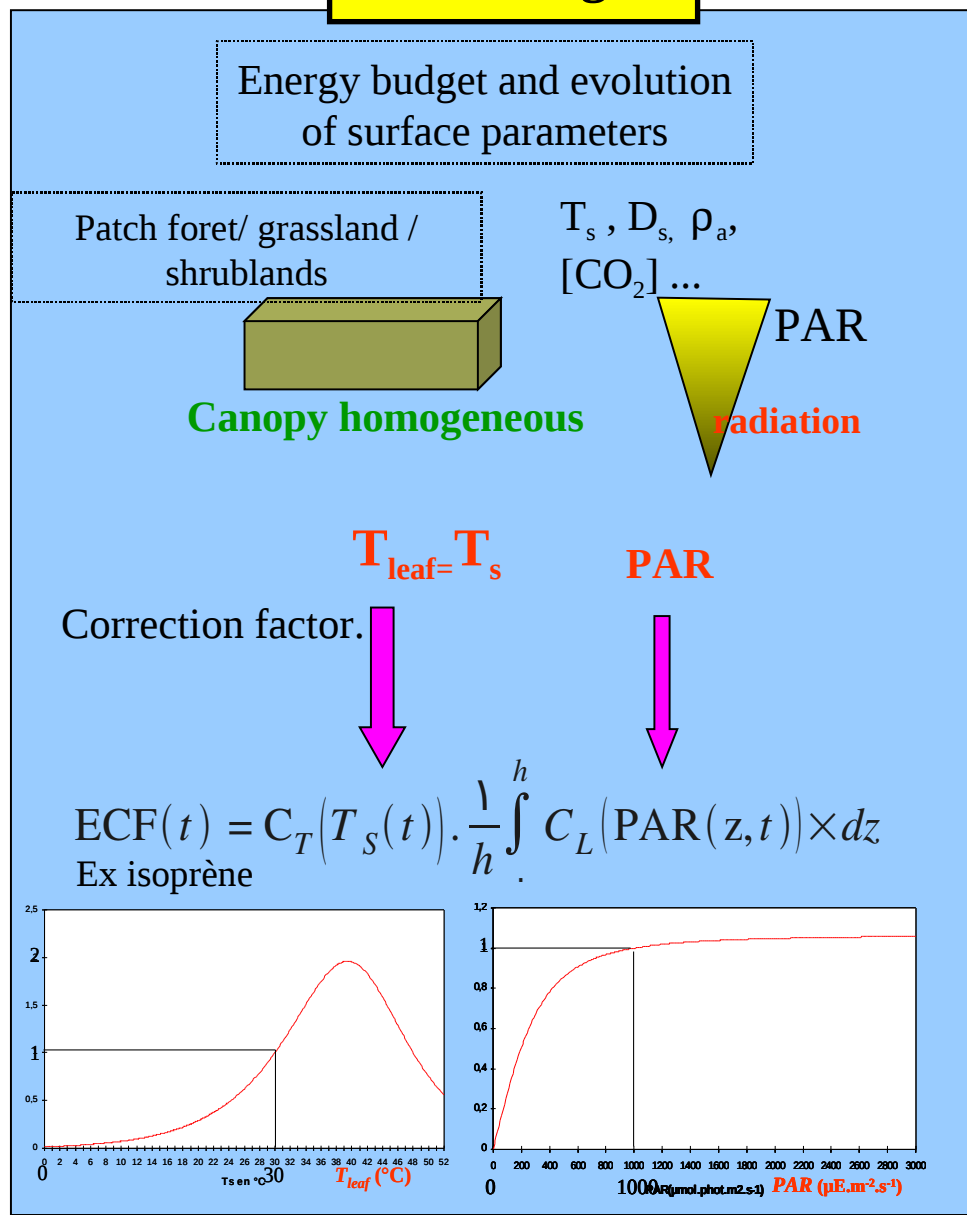
EP = Potentiel Emission (offline)

- + Corinne land cover
- + Statistics: Inventory from .French Forest. National.
- + Emission factor for ISOPRENE and MONOTERPENE



Main code: `ch_bvocemmn.f90`

### ISBA-Ags



## 4. On-line emission

### *On-line chemical emissions*

*Example: You provide potential emission factors for isoprene (ISOPOT) and monoterpene (MONOPOT) that will be used for the calculation of biogenic fluxes at each time step during the model run.*

**WHERE?** in namelist **PRE\_PGD1.nam** (surface field step) :

**HOW?**

```
&NAM_CH_EMIS_PGD
```

```
.....
```

```
CEMIS_PGD_NAME(n-1) = 'ISOPOT', ! Name of parameters for use in flux calculations
```

```
CEMIS_PGD_AREA(n-1) = 'LAN', ! Interpolation on all pgd domain...
```

```
NEMIS_PGD_TIME(n-1) = -1, ! Néegative values => no time interpolation
```

```
CEMIS_PGD_FILE(n-1) = 'isopot.txt', ! Name of the file containing 2D map of parameters
```

```
CEMIS_PGD_FILESTYPE = 'ASCLLV',
```

```
CEMIS_PGD_NAME(n) = 'MONOPOT', ! Name of parameters for use in flux calculations
```

```
CEMIS_PGD_AREA(n) = 'LAN', ! Interpolation on all pgd domain...
```

```
NEMIS_PGD_TIME(n) = -1,
```

```
CEMIS_PGD_FILE(n) = 'monopot.txt',
```

```
CEMIS_PGD_FILESTYPE = 'ASCLLV'
```

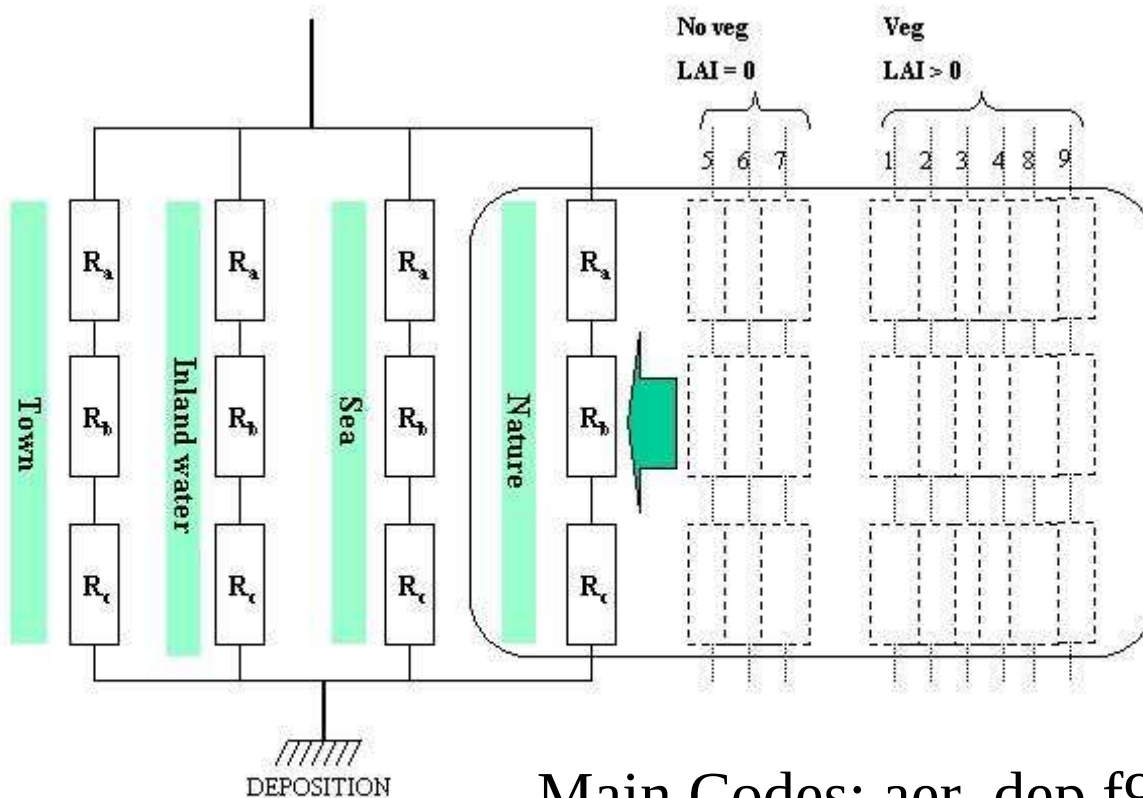
```
.....
```

```
FILESGET_LIST=« emis.tar isopot.txt monopot.txt»
```

## 4. Dry deposition

Deposition of aerosol and gas on the surface have been parameterized upon Wesely (1989) scheme.

Characteristics of gas must be introduced in namelist such as, their solubility (Henry constant), their molecular mass, and their interaction with the biology. See Seinfeld and Pandis, (2001) for more explanations.



Example of combination of dry deposition resistance for each surface type of surfex

Main Codes: aer\_dep.f90, and ch\_dep.f90

## 4. Dry deposition:

How can I specify parameters for dry deposition ?

WHERE? **MNHC\_INPUT.nam**

HOW?

**SURFVALU** ! Keyword for prescribed surface value (flux = surfvalu x exchgvel)

value at the surface are set to zero expect DMS (unit are ppp) ! Comment

1 ! number of prognostic species with prescribed surf value

(1X, A12,1X,F5.1) ! Format

'DMS ' 1E-12 ! Name of prognostic variable and associated surf value

**EXCHGVEL** ! Keyword for prescribed exchange velocity (flux = surfvalu x exchgvel)

Exchange velocity in m/s ! Comment

1 ! Number of prognostic variable with prescribed exchange velocity

(1X, A12,1X,F5.1) ! Format

'O3 ' 0.004 ! Name of prognostic variable and corresponding exchange velocity

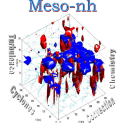
**SURF\_RES** ! Keyword for surface resistance (for use with Wesely dep scheme)

Wesely surface resistance values (if none default values are used) ! Comment

1 ! Number of surface resistance

(1X, A12,1X,F5.1) ! Format

'LANDEXT ' 150. ! Keyword for surface resistance and corresponding value



## 4. Dry deposition:

How can I specify parameters for dry deposition – (continuing) ?

WHERE? MNHC\_INPUT.nam

HOW?

### MASS\_MOL

Molecular mass (g/mol)

1

(1X, A12,1X,F5.1)

'O3            ' 48.00

! Keyword for molecular mass

! Comment

! Number of prognostic species for which molecular mass is prescribed

! Format

! Name of prognostic variable and corresponding molecular mass

### REA\_FACT

Reactivity factor with biology

1

(1X, A12,1X,F5.1)

'O3            ' 1.

! Keyword for reactivity factor

! Comment

! Number of prognostic species

! Format

! Name of prognostic variable and corresponding reactivity factor

### HENRY\_SP

Henry specific constant

1

(1X, A12,1X,E15.2,1X,F8.0)

'O3            ' 1.1E-2   ' -2300.

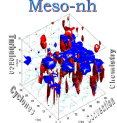
! Keyword for Henry's law constant

! Comment

! Number of prognostic species

! Format

! Name of prognostic variable and corresponding values



## 5. Surface (SURFEX) chemical parameterization: How do I specify options for dry deposition and emissions?

**WHERE?** EXSEG1.nam splits into **SURF1.nam**:

**HOW?**

&NAM\_CH\_SURF<sub>n</sub> LCH\_SURF\_EMIS = .TRUE. , Switch for off-line chemistry emissions  
CCHEM\_SURF\_FILE = « **MNHC\_INPUT.nam** »

Input file containing chemistry parameters (see after)

&NAM\_CH\_ISB<sub>n</sub> CCH\_DRY\_DEP = « WES89 » , Wesely deposition scheme on biosphere  
LCH\_BIO\_FLUX=.TRUE. Switch for on-line biogenic emission

&NAM\_CH\_WATFLUX<sub>n</sub> CCH\_DRY\_DEP = « WES89 » Wesely deposition scheme on lakes

&NAM\_CH\_SEAFLUX<sub>n</sub> CCH\_DRY\_DEP = « WES89 » Wesely deposition scheme on sea

&NAM\_CH\_TEB<sub>n</sub> CCH\_DRY\_DEP = « WES89 » Wesely deposition scheme on town

&NAM\_SURF\_DST CEMISPARAM = « AMMA »

Parameterization type for on-line emission of dust aerosols

&NAM\_CHS\_ORILAM LCH\_AERO\_FLUX=.TRUE., Switch for on-line aerosol emission - ORILAM

XEMISRADIUSI = 0.005, First mode: emission mean radius in micrometers

XEMISRADIUSJ = 0.1, 2nd mode: emission mean radius in micrometers

XEMISSIGI = 1.89, First mode: emission standard deviation

XEMISSIGJ = 2.02, 2nd mode: emission standard deviation

LCO2PM=.TRUE

Switch for conversion of CO emission to BC/POM emission

(in case no off-line emissions are provided for BC and POM precursors) .