TOP

Tracer in Ocean Paradigm –The NEMO Passive Tracer engine

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Introduction

TOP (Tracers in the Ocean Paradigm) handles oceanic passive tracers in NEMO. At present, this component provides the physical constraints and boundaries conditions for oceanic tracers transport and represents a generalized, hardwired interface toward biogeochemical models to enable a seamless coupling.

It includes three independent components:

- a transport code TRP sharing the same advection/diffusion routines with the dynamics, with specific treatment of some features like the surface boundary conditions, or the positivity of passive tracers concentrations
- sources and sinks SMS models that can be typically biogeochemical, biological or radioactive
- an offline option which is a simplified OPA 9 model using fields of physics variables that are previously stored to disk

There is two ways of coupling TOP to the dynamics:

- *online coupling*: the evolution of passive tracers is computed along with the dynamics
- *offline coupling*: the fields of physics variables are read from files and interpolated at each model time step, with no constraints on the time sampling in the input files

TOP is designed to handle multiple oceanic tracers through a modular approach and it includes different sub-modules:

- the ocean water age module (AGE) tracks down the time-dependent spread of surface waters into the ocean interior
- inorganic carbon (e.g. CFCs, SF6) and radiocarbon (C14) passive tracers can be modeled to assess ocean absorption timescales of anthropogenic emissions and further address water masses ventilation
- a built-in biogeochemical model (PISCES) to simulate lower trophic levels ecosystem dynamics in the global ocean
- a prototype tracer module (MY_TRC) to enable user-defined cases or the coupling with alternative biogeochemical models (e.g. BFM, MEDUSA, ERSEM, BFM, ECO3M)

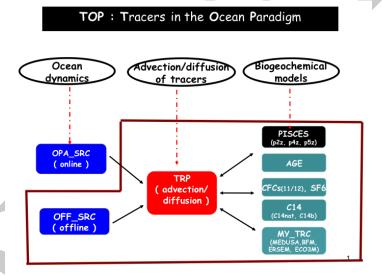


Figure 1: A schematic view of NEMO-TOP component



Model Description

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1.1 Basics

The time evolution of any passive tracer ${\cal C}$ follows the transport equation, which is similar to that of active tracer - temperature or salinity :

$$\frac{\partial C}{\partial t} = S(C) - \frac{1}{b_t} \left[\frac{\partial e_{2u} e_{3u} u C}{\partial i} + \frac{\partial e_{1v} e_{3v} u v, C}{\partial i} \right] + \frac{1}{e_{3t}} \frac{\partial w C}{\partial k} + D^{lC} + D^{vC}$$
(1.1)

where expressions of D^{lC} and D^{vC} depend on the choice for the lateral and vertical subgrid scale parameterizations, see equations 5.10 and 5.11 in (Madec, 2008)

S(C), the first term on the right hand side of 1.1; is the SMS - Source Minus Sink inherent to the tracer. In the case of biological tracer such as phytoplankton, S(C) is the balance between phytoplankton growth and its decay through mortality and grazing. In the case of a tracer comprising carbon, S(C) accounts for gas exchange, river discharge, flux to the sediments, gravitational sinking and other biological processes. In the case of a radioactive tracer, S(C) is simply loss due to radioactive decay.

The second term (within brackets) represents the advection of the tracer in the three directions. It can be interpreted as the budget between the incoming and outgoing tracer fluxes in a volume T-cells $b_t=e_{1t}\,e_{2t}\,e_{3t}$

The third term represents the change due to lateral diffusion.

The fourth term is change due to vertical diffusion, parameterized as eddy diffusion to represent vertical turbulent fluxes :

$$D^{vC} = \frac{1}{e_{3t}} \frac{\partial}{\partial k} \left[A^{vT} \frac{\partial C}{\partial k} \right]$$
 (1.2)

where A^{vT} is the vertical eddy diffusivity coefficient of active tracers

1.2 The NEMO-TOP interface

TOP is the NEMO hardwired interface toward biogeochemical models and provide the physical constraints/boundaries for oceanic tracers. It consists of a modular framework to handle multiple ocean tracers, including also a variety of built-in modules.

This component of the NEMO framework allows one to exploit available modules and further develop a range of applications, spanning from the implementation of a dye passive tracer to evaluate dispersion processes (by means of MY_TRC), track water masses age (AGE module), assess the ocean interior penetration of persistent chemical compounds (e.g., gases like CFC or even PCBs), up to the full set of equations involving marine biogeochemical cycles.

TOP interface has the following location in the code repository: <repository>
/src/TOP/

and the following modules are available:

- TRP: Interface to NEMO physical core for computing tracers transport
- CFC: Inert carbon tracers (CFC11,CFC12, SF6)
- C14 : Radiocarbon passive tracer
- AGE: Water age tracking
- MY_TRC : Template for creation of new modules and external BGC models coupling
- **PISCES**: Built in BGC model. See (Aumont et al., 2015) for a throughout description.

1.3 The transport component: TRP

The passive tracer transport component shares the same advection/diffusion routines with the dynamics, with specific treatment of some features like the surface boundary conditions, or the positivity of passive tracers concentrations.

1.3.1 Advection

```
! advection scheme for passive tracer (default: NO selection)
&namtrc_adv
  ln_trcadv_OFF = .false. ! No passive tracer advection
ln_trcadv_cen = .false. ! 2nd order centered scheme
                              ! =2/4, horizontal 2nd order CEN / 4th order CEN
! =2/4, vertical 2nd order CEN / 4th order COMP
     nn\_cen\_h = 4
                = 4
     nn_cen_v
  ln_trcadv_fct = .false. ! FCT scheme
               nn fct h
     nn_fct_v
  ln_trcadv_mus = .false. ! MUSCL scheme
     ln_mus_ups = .false.
                               ! use upstream scheme near river mouths
  ln_trcadv_ubs = .false. ! UBS scheme
     nn_ubs_v
                               ! =2 , vertical 2nd order FCT
  ln_trcadv_qck = .false. ! QUICKEST scheme
```

The advection schemes used for the passive tracers are the same than the ones for T and S and described in section 5.1 of (Madec, 2008). The choice of an advection scheme can be selected independently and can differ from the ones used for active tracers. This choice is made in the $namtrc_adv$ namelist, by setting to true one and only one of the logicals ln_trcadv_xxx , the same way of what is done for dynamics. cen2, MUSCL2, and UBS are not positive schemes meaning that negative values can appear in an initially strictly positive tracer field which is advected, implying that false extrema are permitted. Their use is not recommended on passive tracers

1.3.2 Lateral diffusion

```
!

&namtrc_ldf ! lateral diffusion scheme for passive tracer (default: NO

→ selection)
!

! Type of the operator:

ln_trcldf_OFF = .false. ! No explicit diffusion

ln_trcldf_tra = .false. ! use active tracer setting

! Coefficient (defined with namtra_ldf coefficient)

rn_ldf_multi = 1. ! multiplier of aht for TRC mixing coefficient

rn_fact_lap = 1. ! Equatorial enhanced zonal eddy diffusivity(lap only)
```

In NEMO v4.0, the passive tracer diffusion has necessarily the same form as the active tracer diffusion, meaning that the numerical scheme must be the same. However the passive tracer mixing coefficient can be chosen as a multiple of the active ones by changing the value of *rn_ldf_multi* in namelist *namtrc_ldf*. The choice of numerical scheme is then set in the namtra_ldf namelist for the dynamic described in section 5.2 of (Madec, 2008).

1.3.3 Tracer damping

The use of newtonian damping to climatological fields or observations is also coded, sharing the same routine dans active tracers. Boolean variables are defined in the namelist_top_ref to select the tracers on which restoring is applied Options are defined through the namtrc_dmp namelist variables. The restoring term is added when the namelist parameter ln_trcdmp is set to true. The restoring coefficient is a three-dimensional array read in a file, which name is specified by the namelist variable cn_resto_tr. This netcdf file can be generated using the DMP_TOOLS tool.

1.3.4 Tracer positivity

Sometimes, numerical scheme can generates negative values of passive tracers concentration that must be positive. For exemple, isopycnal diffusion can created extrema. The trerad routine artificially corrects negative concentrations with a very crude solution that either sets negative concentration to zero without adjusting the tracer budget, or by removing negative concentration and keeping mass conservation. The treatment of negative concentrations is an option and can be selected in the namelist namtre_rad by setting the parameter ln_trerad to true.

1.4 The SMS modules

1.4.1 Ideal Age

An 'ideal age' tracer is integrated online in TOP when $ln_age = .true.$ in namelist namtrc. This tracer marks the length of time in units of years that fluid has spent in the interior of the ocean, insulated from exposure to the atmosphere. Thus, away from the surface for $z < -H_{Age}$ where H_{Age} is specified by the namage namelist

variable rn_age_depth , whose default value is 10 m, there is a source SMS_{Age} of the age tracer A:

$$SMS_{Age} = 1yr^{-1} = 1/T_{year},$$
 (1.3)

where the length of the current year $T_{\rm year} = 86400 * N_{\rm days\ in\ current\ year}\ {
m s}$, where $N_{\rm days\ in\ current\ year}$ may be 366 or 365 depending on whether the current year is a leap year or not. Near the surface, for $z > -H_{\rm Age}$, ideal age is relaxed back to zero:

$$SMS_{Age} = -\lambda_{Age}A, \tag{1.4}$$

where the relaxation rate $\lambda_{\rm Age}$ (units s $^{-1}$) is specified by the *namage* namelist variable $rn_age_kill_rate$ and has a default value of 1/7200 s. Since this relaxation is applied explicitly, this relaxation rate in principle should not exceed $1/\Delta t$, where Δt is the time step used to step forward passive tracers (2 * nn_dttrc * rn_rdt when the default leapfrog time-stepping scheme is employed).

Currently the 1-dimensional reference depth of the grid boxes is used rather than the dynamically evolving depth to determine whether the age tracer is incremented or relaxed to zero. This means that the tracer only works correctly in z-coordinates. To ensure that the forcing is independent of the level thicknesses, where the tracer cell at level k has its upper face z=-depw(k) above the depth $-H_{\rm Age}$, but its lower face z=-depw(k+1) below that depth, then the age source

$$SMS_{Age} = -f_{kill}\lambda_{Age}A + f_{add}/T_{year}, \qquad (1.5)$$

where

$$f_{\text{kill}} = e3t_k^{-1}(H_{\text{Age}} - depw(k)),$$
 (1.6)

$$f_{\text{add}} = 1 - f_{\text{kill}}.\tag{1.7}$$

This implementation was first used in the CORE-II intercomparison runs described e.g. in Danabasoglu and Co-authors (2014).

1.4.2 Inert carbons tracer

Chlorofluorocarbons 11 and 12 (CFC-11 and CFC-12), and sulfur hexafluoride (SF6), are synthetic chemicals manufactured for industrial and domestic applications from the early 20th century onwards. CFC-11 (CCl₃F) is a volatile liquid at room temperature, and was widely used in refrigeration. CFC-12 (CCl₂F₂) is a gas at room temperature, and, like CFC-11, was widely used as a refrigerant, and additionally as an aerosol propellant. SF6 (SF₆) is also a gas at room temperature, with a range of

applications based around its property as an excellent electrical insulator (often replacing more toxic alternatives). All three are relatively inert chemicals that are both non-toxic and non-flammable, and their wide use has led to their accumulation within the Earth's atmosphere. Large-scale production of CFC-11 and CFC-12 began in the 1930s, while production of SF6 began in the 1950s, and their atmospheric concentration time-histories are shown in Figure 1.1. As can be seen in the figure, while the concentration of SF6 continues to rise to the present day, the concentrations of both CFC-11 and CFC-12 have levelled off and declined since around the 1990s. These declines have been driven by the Montreal Protocol (effective since August 1989), which has banned the production of CFC-11 and CFC-12 (as well as other CFCs) because of their role in the depletion of stratospheric ozone (O₃), critical in decreasing the flux of ultraviolet radiation to the Earth's surface. Separate to this role in ozone-depletion, all three chemicals are significantly more potent greenhouse gases than CO₂ (especially SF6), although their relatively low atmospheric concentrations limit their role in climate change.

The ocean is a notable sink for all three gases, and their relatively recent occurrence in the atmosphere, coupled to the ease of making high precision measurements of their dissolved concentrations, has made them valuable in oceanography. Because they only enter the ocean via surface air-sea exchange, and are almost completely chemically and biologically inert, their distribution within the ocean interior reveals its ventilation via transport and mixing. Measuring the dissolved concentrations of the gases – as well as the mixing ratios between them – shows circulation pathways within the ocean as well as water mass ages (i.e. the time since last contact with the atmosphere). This feature of the gases has made them valuable across a wide range of oceanographic problems. One use lies in ocean modelling, where they can be used to evaluate the realism of the circulation and ventilation of models, key for understanding the behaviour of wider modelled marine biogeochemistry (e.g. (Dutay et al., 2002; Palmiéri et al., 2015)).

Modelling these gases (henceforth CFCs) in NEMO is done within the passive tracer transport module, TOP, using the conservation state equation 1.1

Advection and diffusion of the CFCs in NEMO are calculated by the physical module, OPA, whereas sources and sinks are done by the CFC module within TOP. The only source for CFCs in the ocean is via air-sea gas exchange at its surface, and since CFCs are generally stable within the ocean, we assume that there are no sinks (i.e. no loss processes) within the ocean interior. Consequently, the sinks-minus-sources term for CFCs consists only of their air-sea fluxes, F_{cfc} , as described in the Ocean Model Inter-comparison Project (OMIP) protocol (Orr et al., 2017):

$$F_{cfc} = K_w \cdot (C_{sat} - C_{surf}) \cdot (1 - f_i) \tag{1.8}$$

Where K_w is the piston velocity (in m s⁻¹), as defined in Equation 1.10; C_{sat} is the saturation concentration of the CFC tracer, as defined in Equation 1.9; C_{surf} is the local surface concentration of the CFC tracer within the model (in mol m⁻³); and f_i is the fractional sea-ice cover of the local ocean (ranging between 0.0 for ice-free ocean, through to 1.0 for completely ice-covered ocean with no air-sea exchange).

The saturation concentration of the CFC, C_{sat} , is calculated as follows:

$$C_{sat} = Sol \cdot P_{cfc} \tag{1.9}$$

Where Sol is the gas solubility in mol m⁻³ pptv⁻¹, as defined in Equation 1.12; and P_{cfc} is the atmosphere concentration of the CFC (in parts per trillion by volume, pptv). This latter concentration is provided to the model by the historical time-series of Bullister (2015). This includes bulk atmospheric concentrations of the CFCs for both hemispheres – this is necessary because of the geographical asymmetry in the production and release of CFCs to the atmosphere. Within the model, hemispheric concentrations are uniform, with the exception of the region between 10°N and 10° in which they are linearly interpolated.

The piston velocity K_w is a function of 10 m wind speed (in m s⁻¹) and sea surface temperature, T (in °C), and is calculated here following Wanninkhof (1992):

$$K_w = X_{conv} \cdot a \cdot u^2 \cdot \sqrt{\frac{Sc(T)}{660}}$$
(1.10)

Where $X_{conv} = \frac{0.01}{3600}$, a conversion factor that changes the piston velocity from cm h⁻¹ to m s⁻¹; a is a constant re-estimated by Wanninkhof (2014) to 0.251 (in $\frac{cm\ h^{-1}}{(m\ s^{-1})^2}$); and u is the 10 m wind speed in m s⁻¹ from either an atmosphere model or reanalysis atmospheric forcing. Sc is the Schmidt number, and is calculated as follow, using coefficients from Wanninkhof (2014) (see Table 1.2).

$$Sc = a0 + (a1 \cdot T) + (a2 \cdot T^2) + (a3 \cdot T^3) + (a4 \cdot T^4)$$
 (1.11)

The solubility, Sol, used in Equation 1.9 is calculated in mol l^{-1} atm⁻¹, and is specific for each gas. It has been experimentally estimated by Warner and Weiss (1985) as a function of temperature and salinity:

$$\ln(Sol) = a_1 + \frac{a_2}{T_X} + a_3 \cdot \ln T_X + a_4 \cdot T_X^2 + S \cdot (b_1 + b_2 \cdot T_X + b_3 \cdot T_X^2)$$
 (1.12)

Where T_X is $\frac{T+273.16}{100}$, a function of temperature; and the a_x and b_x coefficients are specific for each gas (see Table 1.1). This is then converted to mol m⁻³ pptv⁻¹ assuming a constant atmospheric surface pressure of 1 atm. The solubility of CFCs thus decreases with rising T while being relatively insensitive to salinity changes. Consequently, this translates to a pattern of solubility where it is greatest in cold, polar regions (see Figure 1.2).

The standard outputs of the CFC module are seawater CFC concentrations (in mol m⁻³), the net air-sea flux (in mol m⁻² d⁻¹) and the cumulative net air-sea flux (in mol m⁻²). Using XIOS, it is possible to obtain outputs such as the vertical integral of CFC concentrations (in mol m⁻²; see Figure 1.3). This property, when divided by the surface CFC concentration, estimates the local penetration depth (in m) of the CFC.

Table 1.1: Coefficients for fit of the CFCs solubility (Eq. 1.12).

Gas	a1	a2	a3	a4	b1	b2	b3
CFC-11	-218.0971	298.9702	113.8049	-1.39165	-0.143566	0.091015	-0.0153924
CFC-12	-229.9261	319.6552	119.4471	-1.39165	-0.142382	0.091459	-0.0157274
SF6	-80.0343	117.232	29.5817	0.0	0.0335183	-0.0373942	0.00774862

Table 1.2: Coefficients for fit of the CFCs Schmidt number (Eq. 1.11).

Gas	a0	a1	a2	a3	a4
CFC-11	3579.2	-222.63	7.5749	-0.14595	0.0011874
CFC-12	3828.1	-249.86	8.7603	-0.1716	0.001408
SF6	3177.5	-200.57	6.8865	-0.13335	0.0010877

Notes

In comparison to the OMIP protocol, the CFC module in NEMO has several differences:

For instance, C_{sat} is calculated for a fixed surface pressure of 1atm, what could be corrected in a further version of the module.

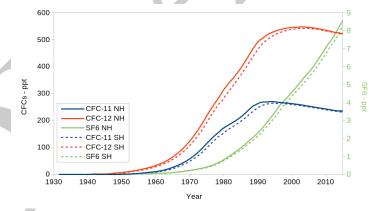


Figure 1.1: Atmospheric CFC11, CFC12 and SF6 partial pressure evolution in both hemispheres.

1.4.3 Radiocarbon

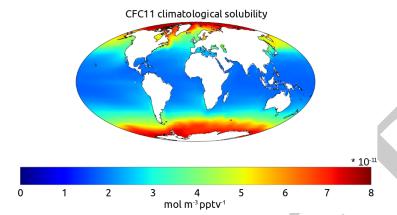


Figure 1.2: CFC11 solubility in mol m⁻³ pptv⁻¹, calculated from the World Ocean Atlas 2013 temperature and salinity annual climatology.

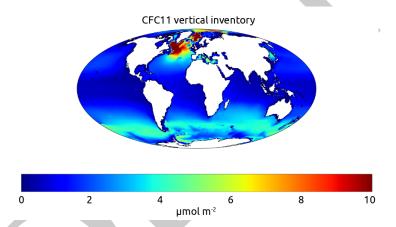


Figure 1.3: CFC11 vertical inventory in μ mol m⁻², from one of the UK Earth System Model 1 model (UKESM1 - which uses NEMO as ocean component, with TOP for the passive tracers) historical run at year 2000.

The C14 package implemented in NEMO by Anne Mouchet models ocean $\Delta^{14}\mathrm{C}$. It offers several possibilities: $\Delta^{14}\mathrm{C}$ as a physical tracer of the ocean ventilation (natural $^{14}\mathrm{C}$), assessment of bomb radiocarbon uptake, as well as transient studies of paleohistorical ocean radiocarbon distributions.

Method

Let $^{14}\mathrm{R}$ represent the ratio of $^{14}\mathrm{C}$ atoms to the total number of carbon atoms in the sample, i.e. $^{14}\mathrm{C/C}$. Then, radiocarbon anomalies are reported as

$$\Delta^{14}C = \left(\frac{^{14}R}{^{14}R_{ref}} - 1\right)10^3, \tag{1.13}$$

where $^{14}R_{ref}$ is a reference ratio. For the purpose of ocean ventilation studies $^{14}R_{ref}$ is set to one.

Here we adopt the approach of Fiadeiro (1982) and Toggweiler et al. (1989a,b) in which the ratio $^{14}\mathrm{R}$ is transported rather than the individual concentrations C and $^{14}\mathrm{C}$. This approach calls for a strong assumption, i.e., that of a homogeneous and constant dissolved inorganic carbon (DIC) field (Toggweiler et al., 1989a; Mouchet, 2013). While in terms of oceanic $\Delta^{14}\mathrm{C}$, it yields similar results to approaches involving carbonate chemistry, it underestimates the bomb radiocarbon inventory because it assumes a constant air-sea CO_2 disequilibrium (Mouchet, 2013). Yet, field reconstructions of the ocean bomb $^{14}\mathrm{C}$ inventory are also biased low (Naegler, 2009) since they assume that the anthropogenic perturbation did not affect ocean DIC since the pre-bomb epoch. For these reasons, bomb $^{14}\mathrm{C}$ inventories obtained with the present method are directly comparable to reconstructions based on field measurements.

This simplified approach also neglects the effects of fractionation (e.g., air-sea exchange) and of biological processes. Previous studies by Bacastow and Maier-Reimer (1990) and Joos et al. (1997) resulted in nearly identical Δ^{14} C distributions among experiments considering biology or not. Since observed ¹⁴R ratios are corrected for the isotopic fractionation when converted to the standard Δ^{14} C notation (Stuiver and Polach, 1977) the model results are directly comparable to observations.

Therefore the simplified approach is justified for the purpose of assessing the circulation and ventilation of OGCMs.

The equation governing the transport of ¹⁴R in the ocean is

$$\frac{\partial}{\partial t}^{14} \mathbf{R} = - \nabla \cdot (\mathbf{u}^{14} \mathbf{R} - \mathbf{K} \cdot \nabla^{14} \mathbf{R}) - \lambda^{14} \mathbf{R}, \tag{1.14}$$

where λ is the radiocarbon decay rate, \mathbf{u} the 3-D velocity field, and \mathbf{K} the diffusivity tensor.

At the air-sea interface a Robin boundary condition (Haine, 2006) is applied to (1.14), i.e., the flux through the interface is proportional to the difference in the ratios between the ocean and the atmosphere

$$\mathcal{F} = \kappa_R(^{14}R - {}^{14}R_a), \tag{1.15}$$

where \mathcal{F} is the flux out of the ocean, and $^{14}\mathrm{R}_a$ is the atmospheric $^{14}\mathrm{C/C}$ ratio. The transfer velocity κ_R for the radiocarbon ratio in (1.15) is computed as

$$\kappa_R = \frac{\kappa_{\text{CO}_2} K_0}{\overline{\text{C}_{\text{T}}}} \, \text{p}_{\text{CO}_2}^{\text{a}} \tag{1.16}$$

with $\kappa_{\rm CO_2}$ the carbon dioxide transfer or piston velocity, K_0 the ${\rm CO_2}$ solubility in seawater, ${\rm p_{\rm CO_2}^a}$ the atmospheric ${\rm CO_2}$ pressure at sea level, and $\overline{\rm C_T}$ the average seasurface dissolved inorganic carbon concentration.

The CO₂ transfer velocity is based on the empirical formulation of Wanninkhof (1992) with chemical enhancement (Wanninkhof and Knox, 1996; Wanninkhof, 2014). The original formulation is modified to account for the reduction of the air-sea exchange rate in the presence of sea ice. Hence

$$\kappa_{\text{CO}_2} = (K_W \,\mathrm{w}^2 + b) \,(1 - f_{\text{ice}}) \,\sqrt{660/Sc},$$
(1.17)

with w the wind magnitude, f_{ice} the fractional ice cover, and Sc the Schmidt number. K_W in (1.17) is an empirical coefficient with dimension of an inverse velocity. The chemical enhancement term b is represented as a function of temperature T (Wanninkhof, 1992)

$$b = 2.5(0.5246 + 0.016256T + 0.00049946 * T^{2}). \tag{1.18}$$

Model setup

To activate the C14 package, set the parameter $ln_{-}c14 = .true$. in namelist namtrc.

Parameters and formulations The radiocarbon decay rate (RLAM14; in trcnam_c14 module) is set to $\lambda = (1/8267) \text{ yr}^{-1}$ (Stuiver and Polach, 1977), which corresponds to a half-life of 5730 yr.

The Schmidt number Sc, Eq. (1.17), is calculated with the help of the formulation of Wanninkhof (2014). The CO_2 solubility K_0 in (1.16) is taken from Weiss (1974). K_0 and Sc are computed with the OGCM temperature and salinity fields (tresms_c14 module).

The following parameters intervening in the air-sea exchange rate are set in namelist_c14:

• The reference DIC concentration $\overline{C_T}$ (XDICSUR) intervening in (1.16) is classically set to 2 mol m⁻³ (Toggweiler et al., 1989a; Orr et al., 2001; Butzin et al., 2005).

- The value of the empirical coefficient K_W (XKWIND) in (1.17) depends on the wind field and on the model upper ocean mixing rate (Toggweiler et al., 1989a; Wanninkhof, 1992; Naegler, 2009; Wanninkhof, 2014). It should be adjusted so that the globally averaged CO_2 piston velocity is $\kappa_{CO_2} = 16.5 \pm 3.2$ cm/h (Naegler, 2009).
- Chemical enhancement (term *b* in Eq. 1.18) may be set on/off by means of the logical variable LN_CHEMH.

Experiment type The type of experiment is determined by the value given to KC14TYP in namelist_c14. There are three possibilities:

- 1. natural Δ^{14} C: KC14TYP=0
- 2. bomb Δ^{14} C: KC14TYP=1
- 3. transient paleo-historical Δ^{14} C: KC14TYP=2

Natural or Equilibrium radiocarbon KC14TYP=0

Unless otherwise specified in namelist_c14, the atmospheric $^{14}R_a$ (RC14AT) is set to one, the atmospheric CO_2 (PCO2AT) to 280 ppm, and the ocean ^{14}R is initialized with RC14INIT=0.85, i.e., $\Delta^{14}C$ =-150‰(typical for deep-ocean, Fig 6 in Key et al., 2004).

Equilibrium experiment should last until 98% of the ocean volume exhibit a drift of less than 0.001%/year (Orr et al., 2000); this is usually achieved after few kyr (Fig. 1.4).

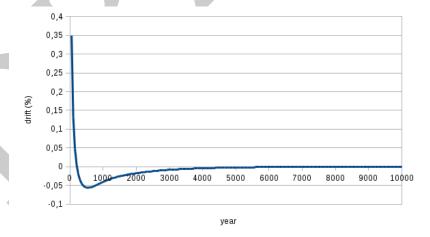


Figure 1.4: Time evolution of 14 R inventory anomaly for equilibrium run with homogeneous ocean initial state. The anomaly (or drift) is given in % change in total ocean inventory per 50 years. Time on x-axis is in simulation year.

Transient: Bomb KC14TYP=1

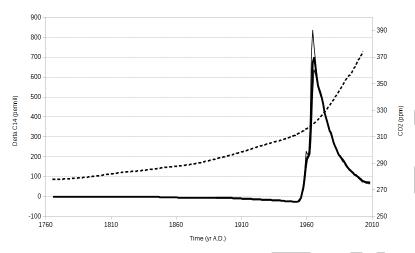


Figure 1.5: Atmospheric $\Delta^{14}C$ (solid; left axis) and CO_2 (dashed; right axis) forcing for the ^{14}C -bomb experiments. The $\Delta^{14}C$ is illustrated for the three zonal bands (upper, middle, and lower curves correspond to latitudes > 20N, $\in [20S, 20N]$, and < 20S, respectively.

Performing this type of experiment requires that a pre-industrial equilibrium run be performed beforehand (LN_RSTTR should be set to .TRUE.).

An exception to this rule is when wishing to perform a perturbation bomb experiment as was possible with the package C14b. It is still possible to easily set-up that type of transient experiment for which no previous run is needed. In addition to the instructions as given in this section it is however necessary to adapt the atmc14.dat file so that it does no longer contain any negative $\Delta^{14}\mathrm{C}$ values (Suess effect in the pre-bomb period).

The model is integrated from a given initial date following the observed records provided from 1765 AD on (Fig. 1.5). The file atmc14.dat (Enting et al., 1994, & I. Levin, personal comm.) provides atmospheric $\Delta^{14}\mathrm{C}$ for three latitudinal bands: 90S-20S, 20S-20N & 20N-90N. Atmospheric CO_2 in the file splc02.dat is obtained from a spline fit through ice core data and direct atmospheric measurements (Orr et al., 2000, & J. Orr, personal comm.). Dates in these forcing files are expressed as yr AD.

To ensure that the atmospheric forcing is applied properly as well as that output files contain consistent dates and inventories the experiment should be set up carefully:

- Specify the starting date of the experiment: NN_DATE0 in namelist. NN_DATE0 is written as Year0101 where Year may take any positive value (AD).
- Then the parameters NN_RSTCTL in namelist (on-line) and NN_RSTTR in namelist_top (off-line) must be set to 0 at the start of the experiment (force the date to NN_DATEO for the first experiment year).
- These two parameters (NN_RSTCTL and NN_RSTTR) have then to be **set to 2** for the following years (the date must be read in the restart file).

If the experiment date is outside the data time span then the first or last atmospheric concentrations are prescribed depending on whether the date is earlier or later. Note that TYRC14_BEG (namelist_c14) is not used in this context.

Transient: Past KC14TYP=2

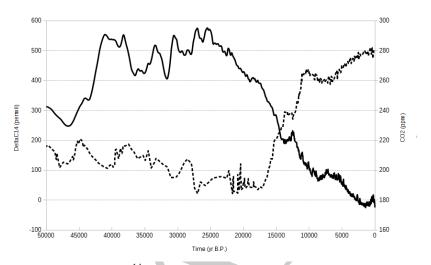


Figure 1.6: Atmospheric $\Delta^{14}C$ (solid) and CO_2 (dashed) forcing for the Paleo experiments. The CO_2 scale is given on the right axis.

This experiment type does not need a previous equilibrium run. It should start 5–6 kyr earlier than the period to be analyzed. Atmospheric $^{14}\mathrm{R}_a$ and CO_2 are prescribed from forcing files. The ocean $^{14}\mathrm{R}$ is initialized with the value attributed to RC14INIT in namelist_c14.

The file intcall3.14c (Reimer et al., 2013) contains atmospheric $\Delta^{14}C$ from 0 to 50 kyr cal BP¹. The CO_2 forcing is provided in file ByrdEdcCO2.txt. The content of this file is based on the high resolution record from EPICA Dome C (Monnin and et al., 2004) for the Holocene and the Transition, and on Byrd Ice Core CO2 Data for 20–90 kyr BP (Ahn and Brook, 2008). These atmospheric values are reproduced in Fig. 1.6. Dates in these files are expressed as yr BP.

To ensure that the atmospheric forcing is applied properly as well as that output files contain consistent dates and inventories the experiment should be set up carefully. The true experiment starting date is given by TYRC14_BEG (in yr BP) in namelist_c14. In consequence, NN_DATEO in namelist MUST be set to 00010101.

Then the parameters NN_RSTCTL in namelist (on-line) and NN_RSTTR in namelist_top (off-line) must be set to 0 at the start of the experiment (force the date to NN_DATEO for the first experiment year). These two parameters have then to be set to 2 for the following years (read the date in the restart file).

If the experiment date is outside the data time span then the first or last atmospheric concentrations are prescribed depending on whether the date is earlier or later.

¹cal BP: number of years before 1950 AD

Model output All output fields in Table 1.3 are routinely computed. It depends on the actual settings in iodef.xml whether they are stored or not.

Field	Type	Dim	Units	Description
RC14	ptrc	3-D	-	Radiocarbon ratio
DeltaC14	diad	3-D	%0	$\Delta^{14}\mathrm{C}$
C14Age	diad	3-D	yr	Radiocarbon age
RAge	diad	2-D	yr	Reservoir age
qtr_c14	diad	2-D	$\mathrm{m}^{-2}~\mathrm{yr}^{-1}$	Air-to-sea net ¹⁴ R flux
qint_c14	diad	2-D	m^{-2}	Cumulative air-to-sea ¹⁴ R flux
AtmCO2	scalar	0-D	ppm	Global atmospheric CO_2
AtmC14	scalar	0-D	%0	Global atmospheric $\Delta^{14}\mathrm{C}$
K_CO2	scalar	0-D	${ m cm}~{ m h}^{-1}$	Global CO_2 piston velocity $(\overline{\kappa_{CO_2}})$
K_C14	scalar	0-D	${ m m~yr^{-1}}$	Global $^{14}\mathrm{R}$ transfer velocity $(\overline{\kappa_R})$
C14Inv	scalar	0-D	10^{26} atoms	Ocean radiocarbon inventory

Table 1.3: Standard output fields for the C14 package.

The radiocarbon age is computed as $(-1/\lambda) \ln (^{14}R)$, with zero age corresponding to $^{14}R=1$.

The reservoir age is the age difference between the ocean uppermost layer and the atmosphere. It is usually reported as conventional radiocarbon age; i.e., computed by means of the Libby radiocarbon mean life (8033 yr;?)

$$^{14}\tau_{\rm c} = -8033 \ln\left(1 + \frac{\Delta^{14}C}{10^3}\right),$$
 (1.19)

where $^{14}\tau_{c}$ is expressed in years B.P. Here we do not use that convention and compute reservoir ages with the mean lifetime $1/\lambda$. Conversion from one scale to the other is readily performed. The conventional radiocarbon age is lower than the radiocarbon age by $\simeq 3\%$.

The ocean radiocarbon inventory is computed as

$$N_A^{14} R_{\text{oxa}} \overline{C_T} \left(\int_{\Omega}^{14} R d\Omega \right) / 10^{26},$$
 (1.20)

where N_A is the Avogadro's number ($N_A=6.022\times 10^{23}$ at/mol), $^{14}{\rm R}_{\rm oxa}$ is the oxalic acid radiocarbon standard ($^{14}{\rm R}_{\rm oxa}=1.176\times 10^{-12}$; Stuiver and Polach, 1977), and Ω is the ocean volume. Bomb $^{14}{\rm C}$ inventories are traditionally reported in units of 10^{26} atoms, hence the denominator in (1.20).

All transformations from second to year, and inversely, are performed with the help

of the physical constant RSIYEA the sideral year length expressed in seconds².

The global transfer velocities represent time-averaged³ global integrals of the transfer rates:

$$\overline{\kappa_{\rm CO_2}} = \int_{\mathcal{S}} \kappa_{\rm CO_2} d\mathcal{S} \text{ and } \overline{\kappa_R} = \int_{\mathcal{S}} \kappa_R d\mathcal{S}$$
(1.21)

1.4.4 PISCES biogeochemical model

PISCES is a biogeochemical model which simulates the lower trophic levels of marine ecosystem (phytoplankton, microzooplankton and mesozooplankton) and the biogeochemical cycles of carbonand of the main nutrients (P, N, Fe, and Si). The model is intended to be used for both regional and global configurations at high or low spatial resolutions as well as for short-term (seasonal, interannual) and long-term (climate change, paleoceanography) analyses. Two versions of PISCES are available in NEMO v4.0:

PISCES-v2, by setting in namelist_pisces_ref_ln_p4z to true, can be seen as one of the many Monod models (Monod, 1942). It assumes a constant Redfield ratio and phytoplankton growth depends on the external concentration in nutrients. There are twenty-four prognostic variables (tracers) including two phytoplankton compartments (diatoms and nanophytoplankton), two zooplankton size-classes (microzooplankton and mesozooplankton) and a description of the carbonate chemistry. Formulations in PISCES-v2 are based on a mixed Monod/Quota formalism: On one hand, stoichiometry of C/N/P is fixed and growth rate of phytoplankton is limited by the external availability in N, P and Si. On the other hand, the iron and silicium quotas are variable and growth rate of phytoplankton is limited by the internal availability in Fe. Various parameterizations can be activated in PISCES-v2, setting for instance the complexity of iron chemistry or the description of particulate organic materials.

PISCES-QUOTA has been built on the PISCES-v2 model described in Aumont et al. (2015). PISCES-QUOTA has thirty-nine prognostic compartments. Phytoplankton growth can be controlled by five modeled limiting nutrients: Nitrate and Ammonium, Phosphate, Silicate and Iron. Five living compartments are represented: Three phytoplankton size classes/groups corresponding to picophytoplankton, nanophytoplankton and diatoms, and two zooplankton size classes which are microzooplankton and mesozooplankton. For phytoplankton, the prognostic variables are the carbon, nitrogen, phosphorus, iron, chlorophyll and silicon biomasses (the latter only for diatoms). This means that the N/C, P/C, Fe/C and Chl/C ratios of both phytoplankton groups as well as the Si/C ratio of diatoms are prognostically predicted by the model. Zooplankton are assumed to be strictly homeostatic (e.g., Sterner and Elser, 2002; Woods and Wilson, 2013; Meunier and Boersma, 2014). As a consequence, the C/N/P/Fe ratios of these groups are maintained constant and are not allowed to vary. In PISCES, the Redfield ratios C/N/P are set to 122/16/1 (Takahashi et al., 1985) and the -O/C ratio is set to 1.34 (Kortzinger et al., 2001). No silicified zooplankton is assumed. The bacterial pool is not yet explicitly modeled.

²The variable (NYEAR_LEN) which reports the length in days of the previous/current/future year (see oce_trc.F90) is not a constant.

³the actual duration is set in iodef.xml

There are three non-living compartments: Semi-labile dissolved organic matter, small sinking particles, and large sinking particles. As a consequence of the variable stoichiometric ratios of phytoplankton and of the stoichiometric regulation of zooplankton, elemental ratios in organic matter cannot be supposed constant anymore as that was the case in PISCES-v2. Indeed, the nitrogen, phosphorus, iron, silicon and calcite pools of the particles are now all explicitly modeled. The sinking speed of the particles is not altered by their content in calcite and biogenic silicate ("The ballast effect", (Honjo, 1996; Armstrong et al., 2002)). The latter particles are assumed to sink at the same speed as the large organic matter particles. All the non-living compartments experience aggregation due to turbulence and differential settling as well as Brownian coagulation for DOM.

1.4.5 MY_TRC interface for coupling external BGC models

The NEMO-TOP has only one built-in biogeochemical model - PISCES - but there are several BGC models - MEDUSA, ERSEM, BFM or ECO3M - which are meant to be coupled with the NEMO dynamics. Therefore it was necessary to provide to the users a framework for easily add their own BGCM model, that can be a single passive tracer. The generalized interface is pivoted on MY_TRC module that contains template files to build the coupling between NEMO and any external BGC model. The call to MY_TRC is activated by setting $ln_my_trc = .true$. in namelist namtrc

The following 6 fortran files are available in MY_TRC with the specific purposes here described.

- par_my_trc.F90: This module allows to define additional arrays and public variables to be used within the MY_TRC interface
- trcini_my_trc.F90: Here are initialized user defined namelists and the call to the
 external BGC model initialization procedures to populate general tracer array
 (trn and trb). Here are also likely to be defined suport arrays related to system
 metrics that could be needed by the BGC model.
- trcnam_my_trc.F90: This routine is called at the beginning of trcini_my_trc and should contain the initialization of additional namelists for the BGC model or user-defined code.
- *trcsms_my_trc.F90*: The routine performs the call to Boundary Conditions and its main purpose is to contain the Source-Minus-Sinks terms due to the biogeochemical processes of the external model. Be aware that lateral boundary conditions are applied in trcnxt routine. IMPORTANT: the routines to compute the light penetration along the water column and the tracer vertical sinking should be defined/called in here, as generalized modules are still missing in the code.
- trcice_my_trc.F90: Here it is possible to prescribe the tracers concentrations
 in the seaice that will be used as boundary conditions when ice melting occurs
 (nn_ice_tr =1 in namtrc_ice). See e.g. the correspondent PISCES subroutine.

• *trcwri_my_trc.F90*: This routine performs the output of the model tracers using IOM module (see Manual Chapter Output and Diagnostics). It is possible to place here the output of additional variables produced by the model, if not done elsewhere in the code, using the call to *iom_put*.

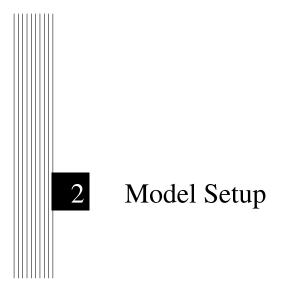
1.5 The Offline Option

```
&namdta_dyn
                   offline ocean input files
                                                        (OFF_SRC only)
   cn_dir = './'
                       root directory for the ocean data location
          !file name!freq(hours)! variable
                                                  (T/F)
                    !(<0 months)!
                                    name
                                                              (T/F)
   sn_tem = 'grid_T',
                                  'votemper',
                                                .true.
                                                            .true.,
  sn_sal = 'grid_T',
                                  'vosaline'
   sn_mld = 'grid_T',
                                  'somixhgt'
                                                 .true.
                                                            .true.
  sn_emp = 'grid_T',
                         120
                                  'sowaflup',
                                                 .true.
  sn_fmf = 'grid_T',
                                , 'iowaflup',
                                                 .true.
                                                             .true.
  sn_ice = 'grid_T'
                                  'soicecov'
                                                 .true.
                                                          , .true.
  sn_qsr = 'grid_T'
                                  'soshfldo',
                                                 .true.
                                                            .true.
  sn_wnd = 'grid_T'
                                  'sowindsp',
                                                 .true.
                                                          , .true. ,
  sn_uwd = 'grid_U'
                                  'uocetr'
                                                 .true.
                                                           , .true. ,
                                                                       vearly
  sn_vwd = 'grid_V'
                                  'vocetr'
                                                 .true.
                                                            .true.,
  sn_wwd = 'grid_W',
                                  'wocetr'
                                                 .true.
                                                           , .true. ,
                                                                       yearly
  sn_avt = 'grid_W',
                                  'voddmavs'
                                                 .true.
                                                            .true.,
                                                                       vearly
  sn_ubl = 'grid_U',
                                   sobblcox',
                                                                      'yearly
                                                 .true.
                                                          , .true. ,
                                                                      'yearly
  sn_vbl = 'grid_V',
                                   sobblcoy',
                                                 .true.
                                                            .true.,
```

Coupling passive tracers offline with NEMO requires precomputed physical fields from OGCM. Those fields are read from files and interpolated on-the-fly at each model time step At least the following dynamical parameters should be absolutely passed to the transport: ocean velocities, temperature, salinity, mixed layer depth and for ecosystem models like PISCES, sea ice concentration, short wave radiation at the ocean surface, wind speed (or at least, wind stress). The so-called offline mode is useful since it has lower computational costs for example to perform very longer simulations - about 3000 years - to reach equilibrium of CO2 sinks for climate-carbon studies.

The offline interface is located in the code repository: CPF/. It is activated by adding the CPP key key_offline to the CPP keys list. There are two specifics routines for the Offline code:

- dtadyn.F90: this module allows to read and compute the dynamical fields at each model time-step
- nemogcm.F90: a degraded version of the main nemogcm.F90 code of NEMO to manage the time-stepping



2.1 Setting up a passive tracer configuration

```
&namtrc
                      tracers definition
                                  ! Number of passive tracers of the BGC model
   jp_bgc
  ln_pisces
                    .false.
                                    Run PISCES BGC model
  ln_my_trc
                    .false.
                                    Run MY_TRC BGC model
                     .false.
                                    Run the sea water age tracer
  ln_cfc11
                   .false.
                                   Run the CFC11 passive tracer
                                 ! Run the CFC12 passive tracer
! Run the SF6 passive tracer
   ln_cfc12
   ln_sf6
                    .false.
   ln_c14
                     .false.
                                    Run the Radiocarbon passive tracer
                     .false.
                                 Initialisation from data input file (T) or not
   ln_trcdta
                                 add a damping termn (T) or not (F)
   ln_trcdmp
                     .false.
   ln_trcdmp_clo =
                                 damping term (T) or not (F) on closed seas
                               title of the field
                                                       ! units ! init from file !
                      name
                              'Tracer Concentration', ' - ',
                    'tracer',
```

The usage of TOP is activated

- by including in the configuration definition the component TOP_SRC
- by adding the macro key_top in the configuration cpp file

As an example, the user can refer to already available configurations in the code, GYRE_PISCES being the NEMO biogeochemical demonstrator and GYRE_BFM to see the required configuration elements to couple with an external biogeochemical model (see also section §1.4).

22 Model Setup

Note that, since version 4.0, TOP interface core functionalities are activated by means of logical keys and all submodules preprocessing macros from previous versions were removed.

There are only three specific keys remaining in TOP

- *key_top* : to enables passive tracer module
- key_trdtrc and key_trdmxl_trc: trend computation for tracers

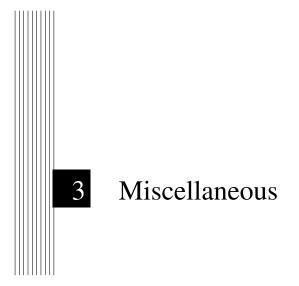
For a remind, the revisited structure of TOP interface now counts for five different modules handled in namelist_top:

- PISCES, default BGC model
- MY_TRC, template for creation of new modules couplings (maybe run a single passive tracer)
- CFC, inert carbon tracers dynamics (CFC11,CFC12,SF6) Updated with OMIP-BGC guidelines (Orr et al, 2016)
- C14, radiocarbon passive tracer
- AGE, water age tracking revised implementation

The modular approach was implemented also in the definition of the total number of passive tracers (jptra). This results from to user setting from the namelist *namtrc*

2.2 TOP Tracer Initialisation

2.3 TOP Boundaries Conditions



3.1 TOP synthetic Workflow

- 3.1.1 Model initialization
- 3.1.2 Time marching procedure

3.2 Coupling an external BGC model using NEMO framework

The coupling with an external BGC model through the NEMO compilation framework can be achieved in different ways according to the degree of coding complexity of the Biogeochemical model, like e.g., the whole code is made only by one file or it has multiple modules and interfaces spread across several subfolders.

Beside the 6 core files of MY_TRC module, see (see , let's assume an external BGC model named "MYBGC" and constituted by a rather essential coding structure, likely few Fortran files. The new coupled configuration name is NEMO_MYBGC.

The best solution is to have all files (the modified MY_TRC routines and the BGC model ones) placed in a unique folder with root <MYBGCPATH> and to use the *makenemo* external readdressing of MY_SRC folder.

The coupled configuration listed in **cfg.txt** will look like

NEMO_MYBGC OPA_SRC TOP_SRC

and the related cpp_MYBGC.fcm content will be

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```
bld::tool::fppkeys key_iomput key_mpp_mpi key_top
```

the compilation with *makenemo* will be executed through the following syntax

```
makenemo -n NEMO_MYBGC -m <arch_my_machine> -j 8 -e <MYBGCPATH>

bld::tool::fppkeys key_iomput key_mpp_mpi key_top

src::MYBGC::initialization <MYBGCPATH>/initialization
src::MYBGC::pelagic <MYBGCPATH>/pelagic
src::MYBGC::benthic <MYBGCPATH>/benthic

bld::pp::MYBGC 1
bld::tool::fppflags::MYBGC %FPPFLAGS
bld::tool::fppkeys %bld::tool::fppkeys MYBGC_MACROS
```





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