

The pop_in namelist

For the ITAMOC climatologic run with closed ITF and the meaning of important parameters (in red)

```

&domain_nml
  nprocs_clinic = 3840      <- number of processors to be used for most of the code
  nprocs_tropic = 3840     <- number of processors to be used for barotropic solver
  clinic_distribution_type = 'cartesian' <- method for distributing blocks across processors
  tropic_distribution_type = 'cartesian' <- method for distributing blocks across processors
  ew_boundary_type = 'cyclic' <-type of boundary in the logical east-west direction for global domain
  ns_boundary_type = 'tripole' <-type of boundary in the logical north-south direction for gl. domain
/

&context_nml
/

&io_nml
  num_iotasks = 20          <- number of I/O processes for parallel binary I/O
  lredirect_stdout = .false. <- flag to write stdout to log file
  log_filename = 'pop.out'  <- root filename (with path) of optional output log file
  luse_pointer_files = .true. <- flag to turn on use of pointer files
  pointer_filename = 'pointer'
/

&time_manager_nml
  runid      = 'clim_closeditf'
  stop_option = 'eom'          <- units of time for 'stop count', eom = end of month
  stop_count  = 1 <- how long in above units to run this segment (use yyymmdd for date)
  time_mix_opt = 'avgfit'      <- Method to suppress leapfrog computational mode
  fit_freq    = 1 <-When using 'avgfit', the intervals per day into which full and half steps must fit
  time_mix_freq = 17 <- Requested frequency (in steps) for taking mixing steps
  dt_option   = 'steps_per_day' <- units for determining timestep (combined with dt count)
  dt_count    = 170 <- number of timesteps in above units to compute timestep
  impcor      = .true. <- If .true., the Coriolis terms treated implicitly
  laccel      = .false. <- if .true., tracer timesteps increase with depth
  accel_file  = 'unknown_accel_file' <- file containing vertical profile of timestep acceleration factor
  dtuxcel     = 1.0 <- factor to multiply momentum timestep for different momentum and tracer tsteps
  allow_leapyear = .false. <- use leap years in calendar
  iyear0      = 75 <- year (yyyy) at start of full run sequence
  imonth0     = 1 <- Month at start of sequence
  iday0       = 1 <- day at start of sequence
  ihour0      = 0 <- etc.
  iminute0    = 0
  iseccond0   = 0
  date_separator = '' <- Character to separate yyyy mm dd in date ( ' ' means no separator)
/

```

```

&grid_nml
  horiz_grid_opt   = 'file'      read horizontal grid from a file OR create simple lat/lon grid
  horiz_grid_file  = 'path_to_file/grid.3600x2400.fob.da'
  sfc_layer_opt    = 'varthick' <- surface layer is variable thickness OR rigid lid OR old free
                                surface formulation
  vert_grid_opt     = 'file' <- read vertical grid structure from file OR compute vertical grid
                                internally
  vert_grid_file   = 'path_to_file/in_depths.42.dat'
  topography_opt   = 'file'
  topography_file  = 'path_to_file /kmt_noITF.big_endian'
  partial_bottom_cells = .true. <- use partial bottom cells
  bottom_cell_file = 'path_to_file/dzbc_pbc.p1_tripole.s2.0-og.20060315.no_caspian_or_black'
  region_mask_file = 'unknown_region_mask' <- file containing region number @ each gridpoint
  topo_smooth      = .false. <- if .true., smooth topography using 9-point averaging stencil
  flat_bottom       = .false. <- if .true., flat bottom is used
  lremove_points    = .false. <- if .true., remove iso- lated or disconnected ocean points
/

```

```

&init_ts_nml
  init_ts_option = 'restart' <- start from restart OR read initial ocean conditions from a file OR
                                create conditions from an input mean ocean profile OR create
                                initial conditions based on 1992 Levitus mean ocean profile
                                computed internally
  init_ts_file   = 'path_to_file/r.t0.1_42l_nccs01.00750101_fixedU' <- restart file OR file
                                                                    containing 3D potential
                                                                    temperature and salinity
                                                                    at grid points OR file
                                                                    containing depth pro-
                                                                    file of potential tempera-
                                                                    ture and salinity OR (ig-
                                                                    nored for 'internal' or
                                                                    when luse pointer files
                                                                    is enabled)

  init_ts_file_fmt = 'bin' <- data format (binary or netCDF) for input init ts file ('file' and 'restart'
                                                                    options only)
/

```

```

&diagnostics_nml
  diag_global_freq_opt = 'nday'
  diag_global_freq     = 1 <- how often (in above units) to compute and print global diagnostics
  diag_cfl_freq_opt    = 'nday'
  diag_cfl_freq        = 1 <- how often (in above units) to compute and print CFL stability
                                diagnostics
  diag_transp_freq_opt = 'nday'
  diag_transp_freq     = 1 <- how often (in above units) to compute and print transport
                                diagnostics
  diag_transport_file  = 'transport_file_141lines'
  diag_outfile         = 'diag'
  diag_transport_outfile = 'transp'
  diag_all_levels      = .false. <- if true, tracer mean diagnostics at all vertical levels are output
  cfl_all_levels       = .false.
/

```

```

&restart_nml
restart_freq_opt = 'nmonth' <- units of time for 'restart freq'
restart_freq     = 1 <- number of units between output of restart files
restart_outfile  = 'path_to_file/restart/r' <- root filename (with path prepended, if necessary)
                                     for restart files ('runid' and suf- fixes will be added)
restart_fmt      = 'bin' <- data format (binary or netCDF) for restart output files
leven_odd_on     = .false. <- create alternating even/odd restart outputs
                                     which over- write each other
even_odd_freq    = 3840 <- frequency (in steps) for even/odd output
pressure_correction = .false. <- if true, corrects surface pressure error due to (possible)
                                     different timestep. use .false. for exact restart
/

```

```

&tavg_nml
tavg_freq_opt = 'nmonth'
tavg_freq     = 1 <- interval in above units for computation & output of time average history files
tavg_start_opt = 'nstep'
tavg_start    = 0 <- time in above units after which to start accumulating time average
tavg_infile   = '' <- restart file for partial tavg sums if starting from restart (ignored if luse pointer
                                     files is enabled)
tavg_fmt_in   = 'bin' <- format for tavg restart file
tavg_outfile  = 'path_to_file /tavg/t'
tavg_fmt_out  = 'bin' <- format for tavg output files
tavg_contents = 'tavg_contents' <- file name for input file containing names of fields
                                     requested for tavg output
/

```

```

&history_nml
history_freq_opt = 'never' <- This namelist makes snapshot history files possible, we do not need
                                     this, we want monthly mean history files so it's set to never
history_freq     = 100000
history_outfile  = 'unknown_history'
history_fmt      = 'nc'
history_contents = 'sample_history_contents'
/

```

```

&movie_nml
movie_freq_opt = 'nday'
movie_freq     = 1 <- number of units (movie_freq_opt) between output of movie files
movie_outfile  = 'path_to_file'/movie/m'
movie_fmt      = 'bin'
movie_contents = 'movie_contents' <- file containing names of fields requested for movie
                                     output
/

```

```

&solvers
solverChoice    = 'ChronGear'
convergenceCriterion = 1.e-12 <- convergence criterion:
                                      $10^X/XI < \text{convergenceCriterion}$ 
maxIterations    = 1000 <- upper limit on number of iterations allowed
convergenceCheckFreq = 25 <- check for convergence every convergenceCheckFreq
                                     iterations
preconditionerChoice = 'diagonal'
preconditionerFile  = 'unknownPrecondFile' <- file containing preconditioner coefficients
                                     for solver

```

```
&vertical_mix_nml
vmix_choice = 'kpp' <- method of computing vertical diffusion
aidif      = 1.0    <- time-centering parameter for implicit vertical mixing; use of the default
                    value [1.0] is recommended
bottom_drag = 1.0e-3 <- (dimensionless) coefficient used in quadratic bottom drag formula
implicit_vertical_mix = .true.
convection_type = 'diffusion' <- Convection treated by adjustment or by large mixing
                                coefficients
nconvad = 2 <- number of passes through the convec- tive adjustment algorithm
convect_diff = 1000.0 <- tracer mixing coefficient to use with diffusion option
convect_visc = 1000.0 <- momentum mixing coefficient to use with diffusion option
bottom_heat_flux = 0.0 <- constant (geothermal) heat flux (W/m2) to apply to bottom layers
bottom_heat_flux_depth = 100000.00 <- depth (cm) below which to apply bot- tom heat flux
/

&vmix_const_nml <- Constant vertical mixing namelist
const_vvc = 0.25 <- vertical viscosity coefficient (momentum mixing) (cm2/s)
const_vdc = 0.25 <- vertical diffusivity coefficient (tracer mix- ing) (cm2/s)
/

&vmix_rich_nml <- Richardson-number vertical mixing namelist
bckgrnd_vvc = 1.0 <- background vertical viscosity (cm2/s)
bckgrnd_vdc = 0.1 <- background vertical diffusivity (cm2/s)
rich_mix    = 50.0 <- Coefficient for Richardson-number function
/

&vmix_kpp_nml
bckgrnd_vdc1 = 0.55 <- base background    vertical  diffusivity (cm2 /s)
bckgrnd_vdc2 = 0.303615 <- variation in background vertical diffusivity (cm2 /s)
bckgrnd_vdc_dpth= 2500.0e2 <- depth (cm) at which background vertical diffusivity is vdc1
bckgrnd_vdc_linv= 4.5e-5 <- inverse of the length scale (1/L in cm-1) over which diffusivity
                        transition takes place
Prandtl      = 10.0 <- (unitless) ratio of background vertical vis- cosity and diffusivity
rich_mix      = 50.0 <- Coefficient for Richardson-number function
lrich         = .true. <- use Richardson-number for interior mixing
ldbl_diff     = .true. <- add double-diffusive parameterization
lshort_wave   = .true. <- use penetrative shortwave forcing
lcheckekmo    = .false.<- check whether boundary layer exceeds Ekman or Monin-Obukhov
                    limit
num_v_smooth_Ri = 1 <- Number of passes to smooth Richardson number
/

&advect_nml
tadvect_ctype = 'centered' <- centered differences OR 3rd-order up- winding
/

&hmix_nml
hmix_momentum_choice = 'del4' <- method for horizontal mixing of momentum (Laplacian,
                                biharmonic or anisotropic)
hmix_tracer_choice   = 'del4' <- method for horizontal mixing of tracers (Laplacian, biharmonic
                                or Gent-McWilliams)
/
```

```

&hmix_del2u_nml
  lauto_hmix      = .true. <- computes mixing coefficient based on resolution
  lvariable_hmix   = .false. <- scales mixing coeff by grid cell area
  am              = 1.e8 <- momentum mixing coefficient (cm2/s)
/

&hmix_del2t_nml
  lauto_hmix      = .true. <- computes mixing coefficient based on resolution
  lvariable_hmix   = .false. <- scales mixing coeff by grid cell area
  ah              = 1.e8 <- tracer mixing coefficient (cm2/s)
/

&hmix_del4u_nml
  lauto_hmix      = .false. <- compute mixing coefficient based on resolution
  lvariable_hmix   = .true. <- scale mixing coeff by grid cell area
  am              = -27.0e17 <- momentum mixing coeff (cm2/s)
/

&hmix_del4t_nml
  lauto_hmix      = .false. <- compute mixing coefficient based on resolution
  lvariable_hmix   = .true. <- scale mixing coeff by grid cell area
  ah              = -3.0e17 <- tracer mixing coefficient (cm2/s)
/

&hmix_gm_nml <- Gent-McWilliams horizontal mixing namelist
/

&hmix_aniso_nml <- Anisotropic viscosity namelist
/

&state_nml <- Equation of state namelist
  state_choice = 'mwjf' <- McDougall et al. eos OR Jackett and McDougall eos OR polynomial fit
                           to UN- ESCO eos OR linear eos
  state_file = 'internal' <- compute polynomial coefficients internally OR read from file filename
  state_range_opt = 'enforce' <- ignore (ignore) when T,S outside valid polynomial range OR
                           check (check) and report OR compute (enforce) eos as if T,S
                           were in valid range (but don't alter T,S)
  state_range_freq = 100000 <- frequency (steps) for checking T,S range
/

&baroclinic_nml
  reset_to_freezing = .true. <- if .true. and Tsurf(i,j) < Tfreezing, Tsurf(i,j) is reset to Tfreezing
/

&ice_nml
  ice_freq_opt = 'never' <- frequency units for computing ice formation
  ice_freq     = 100000 <- frequency in above units for computing ice formation
  kmxice       = 1 <- compute ice formation above this vertical level
/

```

```

&pressure_grad_nml
  lpressure_avg = .true. <- use pressure averaging to increase time step
  lbouss_correct = .false. <- applies depth-dependent factor to correct for assumed constant
                             density
/

&topostress_nml
  ltopostress = .false. <- true if topographic stress enabled
  nsmooth_topo = 0 <- number of passes to smooth topography
/

&xdisplay_nml
  lxdisplay = .false. <- if .true., enable x-display
  nstep_xdisplay = 1 <- frequency (in steps) for updating x-display
/

&forcing_ws_nml <- Windstress forcing namelist
  ws_data_type = 'monthly' <- type or periodicity of wind stress forcing
  ws_data_inc = 1.e20 <- increment (in hours) between forcing times if ws data type='n-hour'
  ws_interp_freq = 'every-timestep' <- how often to temporally interpolate wind stress data to
                                     current time
  ws_interp_type = 'linear' <- type of temporal interpolation for wind stress data
  ws_interp_inc = 1.e20 <- increment (in hours) between interpolation times if
                          ws_interp_freq = 'n-hour'
  ws_filename =
'/work/e24/sar00059/sar00059/itamoc/scripts/prod_run3_0.5Sv/files_mat/forcing/
ws.o_n_avg.mon' <- name of file containing wind stress, or root of filenames if
                    ws_data_type='n-hour'
  ws_file_fmt = 'bin' <- format of wind stress file
  ws_data_renorm(1) = 10. <- renormalization constants for the components in the wind stress
                           forcing file
  ws_data_renorm(2) = 10.
/

&forcing_shf_nml <- Surface heat flux forcing namelist
  shf_formulation = 'normal-year' <- surface heat flux formulation
  shf_data_type = 'monthly' <- type or periodicity of surface heat flux forcing
  shf_data_inc = 1.e20 <- increment (in hours) between forcing times if shf data type='n-hour'
  shf_interp_freq = 'every-timestep' <- how often to temporally in- terpolate surface heat flux
                                     data to current time
  shf_interp_type = 'linear' <- type of temporal interpola- tion for surface heat flux data
  shf_interp_inc = 1.e20 <- increment (in hours) between interpolation times if
                          shf_interp_freq = 'n-hour'
  shf_restore_tau = 1.e20 <- restoring timescale (days) if type restoring
  shf_weak_restore = 0.0 <- restoring flux for weak restor- ing in bulk-NCEP
  shf_strong_restore = 15.8 <- restoring flux for strong restoring in bulk-NCEP
  shf_filename =
'/work/e24/sar00059/sar00059/itamoc/scripts/run_clim_closeditf/files_mat/forcing/
shf.normal_year+Hurrell.monthly' <- name of file containing surface heat flux data, or root of
                                     filenames if shf data type='n-hour'
  shf_file_fmt = 'bin' <- format (binary or netCDF) of shf file
  shf_data_renorm(3) = 1. <- renormalization constants for the components in the sur- face heat
                           flux forcing file
  shf_data_renorm(4) = 1.

```

```

&forcing_sfwf_nml <- Surface fresh water flux forcing namelist
sfwf_formulation = 'bulk-NCEP' <- surface fresh water flux formulation. Bulk-NCEP means:
                                calculate fluxes based on atmospheric state variables and
                                radiation similar to a fully-coupled model (and using bulk
                                flux formulations extracted from the NCAR flux coupler)
sfwf_data_type = 'monthly' <- type or periodicity of surface fresh water flux forcing
sfwf_data_inc = 1.e20 <- increment (hours) between forcing times if sfwf data type='n-hour'
sfwf_interp_freq = 'every-timestep' <- how often to temporally in- terpolate surface fresh
                                water flux data to current time
sfwf_interp_type = 'linear' <- type of temporal interpola- tion for surface fresh water flux data
sfwf_interp_inc = 1.e20 <- increment (hours) between interpolation times if
                                sfwf interp freq = 'n-hour'
sfwf_restore_tau = 1.e20 <- restoring timescale (days) if restoring
sfwf_weak_restore = 0.009 <- restoring flux for weak restor- ing in bulk-NCEP
sfwf_strong_restore = 0.11 <- restoring flux for strong restoring in bulk-NCEP
sfwf_filename =
'/work/e24/sar00059/sar00059/itamoc/scripts/run_clim_closeditf/files_mat/forcing/
sfwf.CORE+runoff.monthly' <- name of file containing surface fresh water flux data, or root of
                                filenames if sfwf data type='n-hour'
sfwf_file_fmt = 'bin' <- format (binary or netCDF) for sfwf file
sfwf_data_renorm(1) = 0.001 <- renormalization constants for components in
                                sfwf forcing file
sfwf_data_renorm(2) = 1.
ladjust_precip = .true. <- adjust precipitation to balance water budget
lwf_as_salt_flux = .true. <- treat fresh water flux as virtual salt flux
                                when using varthick sfc layer
runoff = .true.
/

```

```

&forcing_pt_interior_nml <- Interior potential temperature forcing namelist
pt_interior_formulation = 'restoring' <- interior pt formulation
pt_interior_data_type = 'none' <- type or periodicity of in- terior pt forcing
pt_interior_data_inc = 1.e20 <- increment (hours) between forcing times if data type 'n-hour'
pt_interior_interp_freq = 'never' <- how often to temporally interpolate interior pt data to
                                current time
pt_interior_interp_type = 'nearest' <- type of temporal interpo- lation for interior pt data
pt_interior_interp_inc = 1.e20 <- increment (hours) between interpolation times if
                                interp freq = 'n-hour'
pt_interior_restore_tau = 1.e20 <- restoring timescale (days) if restoring
pt_interior_filename = 'unknown-pt_interior' <- file containing interior pt data, or root of
                                filenames if data type='n-hour'
pt_interior_file_fmt = 'bin' <- file format (binary or netCDF)
pt_interior_data_renorm = 1. <- renormalization constants for components in
                                interior pt forcing file
pt_interior_restore_max_level = 0 <- maximum level for inte- rior pt restoring
pt_interior_variable_restore = .false. <- enable variable interior pt restoring
pt_interior_restore_filename = 'unknown-pt_interior_restore' <- name of file contain- ing
                                variable interior pt restoring data
pt_interior_restore_file_fmt = 'bin' <- file format (binary or netCDF)
/

```

```

&forcing_s_interior_nml <- Interior salinity restoring namelist
  s_interior_formulation = 'restoring' <- forcing formulation
  s_interior_data_type = 'none' <- type or periodicity of interior salinity forcing
  s_interior_data_inc = 1.e20 <- increment (hours) between forcing times if data type 'n-hour'
  s_interior_interp_freq = 'never' <- how often to temporally interpolate interior S data to
                                current time
  s_interior_interp_type = 'nearest' <- type of temporal interpolation for interior S data
  s_interior_interp_inc = 1.e20 <- increment (in hours) between interpolation times if
                                interp freq 'n-hour'
  s_interior_restore_tau = 1.e20 <- restoring timescale (days) if restoring
  s_interior_filename = 'unknown-s_interior' <- name of file containing interior S data, or
                                root of filenames if data type 'n-hour'
  s_interior_file_fmt = 'bin' <- format (binary or netCDF) of s interior file
  s_interior_data_renorm = 1. <- renormalization constants for components in interior S forcing
                                file
  s_interior_restore_max_level = 0 <- maximum level for interior S restoring
  s_interior_variable_restore = .false. <- enable variable interior S restoring
  s_interior_restore_filename = 'unknown-s_interior_restore' <- name of file containing
                                variable interior S restoring data
  s_interior_restore_file_fmt = 'bin'
/

&forcing_ap_nml <- Atmospheric pressure forcing namelist
  ap_data_type = 'none' <- type or periodicity of atmospheric forcing
  ap_data_inc = 1.e20 <- increment (in hours) between forcing times if ap data type='n-hour'
  ap_interp_freq = 'never'
  ap_interp_type = 'nearest'
  ap_interp_inc = 1.e20
  ap_filename = 'unknown-ap'
  ap_file_fmt = 'bin'
  ap_data_renorm = 1.
/

&coupled_nml
  coupled_freq_opt = 'never' <- unit of time for coupled freq
  coupled_freq = 100000
/

&tidal_nml
/

&passive_tracers_on_nml
  dye_on = .false.
  iage_on = .false.
/

```



```
&dye_nml
  init_dye_option = 'restart'
  init_dye_init_file = 'same_as_TS'
  dye_region_file = 'blablabla'
  dye_region_file_fmt = 'bin'
  tracer_init_ext(1)%mod_varname = 'DYE'
  tracer_init_ext(1)%filename = 'unknown'
  tracer_init_ext(1)%default_val = 0.0
  dye_tadvect_ctype = 'lw_lim'
/

&sw_absorption_nml
/

&float_nml
/
```

For more information about parameters that are not described check:
<http://climate.lanl.gov/Models/POP/UsersGuide.pdf>