

Numbat

High-resolution simulations of density-driven convective mixing in porous media

Christopher Green

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CSIRO Energy
71 Normanby Road, Clayton VIC, 3168, Australia
Private Bag 10, Clayton South VIC, 3169, Australia
Telephone: +61 3 9545 2777
Fax: +61 3 9545 8380

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1 Numbat

Numbat is an open-source MOOSE¹ application for high-resolution simulations of buoyancy-driven convection in porous media in both two and three dimensions.

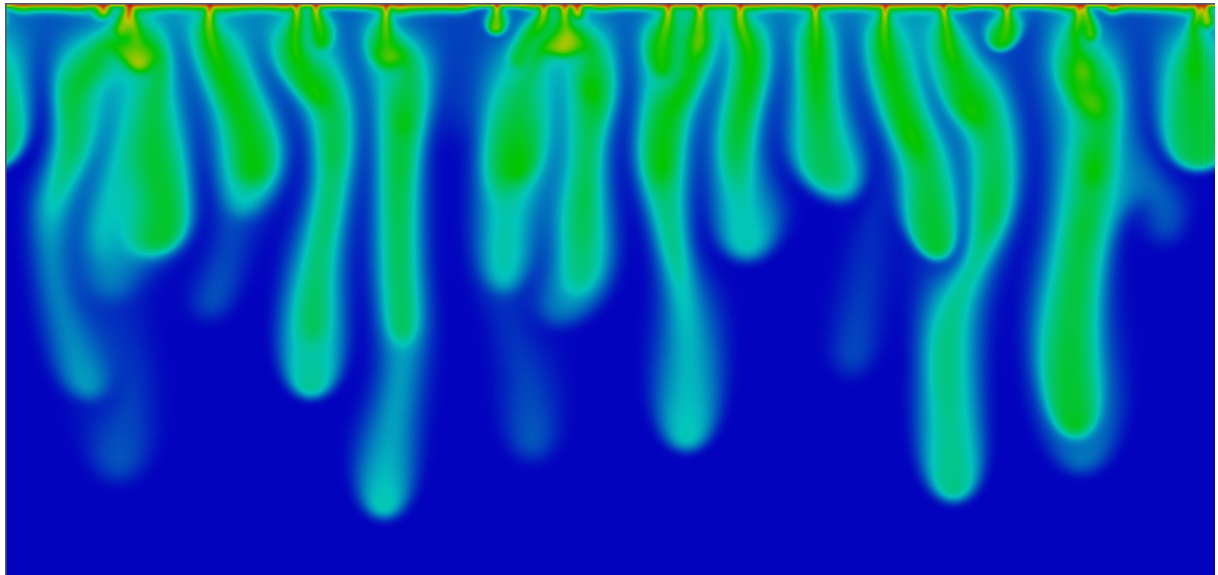


Figure 1.1: Density-driven convective mixing in porous media

As a MOOSE app, it provides access to powerful MOOSE features such as adaptive mesh refinement, hybrid parallelism, both continuous and discontinuous Galerkin methods, and much more, all wrapped in a simple interface.

Nubat solves the coupled convection-diffusion and Darcy equations with the Boussinesq approximation using the finite element method.

Several formulations are available: from the full, dimensional governing equations, to a dimensionless streamfunction formulation.

1.1 Development and testing

Nubat is developed on GitHub² by CSIRO³. It follows the MOOSE continuous integration/-continuous development philosophy, where changes are merged into the master branch of the repository only after being tested successfully against the automatic test suite provided by Nubat.

Nubat is also part of the upstream MOOSE testing procedure, where all changes to MOOSE are tested against Nubat (as well as other MOOSE applications) to ensure no conflicts.

1.2 User manual version

Due to this development philosophy, Nubat does not feature the concept of software versions. A consequence of this is that there are no version label applied to this user manual. This

¹www.mooseframework.org

²www.github.com/cpgr/numbat

³www.csiro.au

documentation is updated as new features and examples are added to the Numbat application. The most up-to-date version of this document is always available on the Numbat website.

2 Introduction

Convective mixing of fluids in porous media is a physical process that manifests in complex flow patterns. In solutal convective mixing, the presence of a fluid component in the background fluid results in a density contrast between fluid containing the solute and fluid without. This density difference gives rise to a pressure gradient which drives motion.

Recently, a significant body of scientific literature has been concerned with density-driven convective mixing in porous media due to its applicability to geological storage of carbon dioxide, see Emami-Meybodi *et al.* (2015) for an extensive literature review. In the case where CO₂ is injected into a saline aquifer with a bounding cap rock, buoyancy drives vertical migration of the mobile CO₂ (CO₂ in the supercritical gas phase), which then spreads beneath the cap rock to form a thin, laterally extensive plume. In time, the gaseous CO₂ begins to dissolve into the local formation water, leading to a small increase in density of the saturated brine at the top of the aquifer of approximately 1%. Diffusion of the dissolved CO₂ allows further dissolution, a process that leads to a gravitational instability whereby a denser fluid lies atop a less dense one. After a sufficient period of time, vertical fluid motion is induced as vertical acceleration overcomes diffusion, and CO₂-rich water descends to the lower part of the reservoir.

The process of convective mixing can significantly increase the rate of dissolution of CO₂ in the formation water and hence reduce the amount of mobile CO₂. This can significantly reduce the risk of leakage into overlying aquifers, increasing the security of storage.

3 Background theory

3.1 Governing equations

Numbat implements the Boussinesq approximation to model density-driven convective mixing in porous media. To reduce the computational burden, only a single fluid phase is considered. This is a simplification to the actual physical process, where a gas phase may be present. This simplification is often used in practice, see Emami-Meybodi *et al.* (2015) for a discussion about the use of this simplifying assumption.

Note:

The more complicated two-phase model can be implemented using the `porous_flow` module

The governing equations for density-driven flow in porous media are Darcy's law

$$\mathbf{u} = -\frac{\mathbf{K}}{\mu} (\nabla P - \rho(c)g\hat{\mathbf{k}}), \quad (3.1)$$

where $\mathbf{u} = (u, v, w)$ is the velocity vector, \mathbf{K} is permeability, μ is the fluid viscosity, P is the fluid pressure, $\rho(c)$ is the fluid density as a function of solute concentration c , g is gravity, and $\hat{\mathbf{k}}$ is the unit vector in the z direction.

The fluid velocity must also satisfy the continuity equation

$$\nabla \cdot \mathbf{u} = 0, \quad (3.2)$$

and the solute concentration is governed by the convection - diffusion equation

$$\phi \frac{\partial c}{\partial t} + \mathbf{u} \cdot \nabla c = \phi D \nabla^2 c, \quad (3.3)$$

where ϕ is the porosity, t is time and D is the diffusivity.

Darcy's law and the convection-diffusion equations are coupled through the fluid density, which is given by

$$\rho(c) = \rho_0 + \frac{c}{c_0} \Delta\rho, \quad (3.4)$$

where c_0 is the equilibrium concentration, and $\Delta\rho$ is the increase in density of the fluid at equilibrium concentration.

The boundary conditions are

$$\begin{aligned} w &= 0, & z &= 0, -H, \\ \frac{\partial c}{\partial z} &= 0, & z &= -H, \\ c &= c_0, & z &= 0, \end{aligned} \quad (3.5)$$

which correspond to impermeable boundary conditions at the top and bottom boundaries, given by $z = 0$ and $z = -H$, respectively, and a saturated condition at the top boundary.

Initially, there is no solute in the model

$$c = 0, \quad t = 0. \quad (3.6)$$

Numbat solves Eq. (3.1) and Eq. (3.3) with density coupled to concentration as in Eq. (3.4).

3.2 Dimensionless formulation

The governing equations can also be solved using a streamfunction formulation in 2D and a vector potential formulation in 3D. As a result, we shall consider the two cases separately.

3.2.1 2D solution

If we consider an anisotropic model, with vertical and horizontal permeabilities given by k_z and k_x , respectively, we can non-dimensionalise the governing equations in 2D following Ennis-King & Paterson (2005) Defining the anisotropy ratio γ as

$$\gamma = \frac{k_z}{k_x}, \quad (3.7)$$

we scale the variables using

$$\begin{aligned} x &= \frac{\phi\mu D}{k_z\Delta\rho g\gamma^{1/2}}\hat{x}, & z &= \frac{\phi\mu D}{k_z\Delta\rho g}\hat{z}, & u &= \frac{k_z\Delta\rho g}{\mu\gamma^{1/2}}\hat{u}, & w &= \frac{k_z\Delta\rho g}{\mu}\hat{w} \\ t &= \left(\frac{\phi\mu}{k_z\Delta\rho g}\right)^2\hat{t}, & c &= c_0\hat{c}, & P &= \frac{\mu\phi D}{k_z}\hat{P}, \end{aligned} \quad (3.8)$$

where \hat{x} refers to a dimensionless variable. The governing equations in dimensionless form are then

$$\mathbf{u} = -(\nabla P + c\hat{\mathbf{k}}), \quad (3.9)$$

$$\mathbf{u} = 0, \quad (3.10)$$

$$\frac{\partial c}{\partial t} + \mathbf{u} \cdot \nabla c = \gamma \frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial z^2}, \quad (3.11)$$

where we have dropped the hat on the dimensionless variables for brevity.

The dimensionless boundary conditions are

$$\begin{aligned} w &= 0, & z &= 0, -Ra, \\ \frac{\partial c}{\partial z} &= 0, & z &= -Ra, \\ c &= 1, & z &= 0, \end{aligned} \quad (3.12)$$

where Ra is the Rayleigh number, defined as

$$Ra = \frac{k_z \Delta \rho g H}{\phi \mu D}. \quad (3.13)$$

In this form, the Rayleigh number only appears in the boundary conditions as the location of the lower boundary. Therefore, Ra can be interpreted in this formalism as a dimensionless model height, and can be varied in simulations by simply changing the height of the mesh.

Finally, the dimensionless initial condition is

$$c = 0, \quad t = 0. \quad (3.14)$$

For isotropic models, where $k_x = k_z$ and hence $\gamma = 1$, we recover the dimensionless equations given by Slim (2014)

The coupled governing equations must be solved numerically. To simplify the numerical analysis, we introduce the streamfunction $\psi(x, z, t)$ such that

$$u = -\frac{\partial \psi}{\partial z}, \quad w = \frac{\partial \psi}{\partial x}. \quad (3.15)$$

This definition satisfies the continuity equation, Eq. (3.10), immediately.

The pressure P is removed from Eq. (3.9) by taking the curl of both sides and noting that $\nabla \times \nabla P = 0$ for any P , to give

$$\nabla^2 \psi = -\frac{\partial c}{\partial x}, \quad (3.16)$$

where we have introduced the streamfunction ψ using Eq. (3.15).

The convection-diffusion equation, Eq. (3.11) becomes

$$\frac{\partial c}{\partial t} - \frac{\partial \psi}{\partial z} \frac{\partial c}{\partial x} + \frac{\partial \psi}{\partial x} \frac{\partial c}{\partial z} = \gamma \frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial z^2}. \quad (3.17)$$

The boundary conditions become

$$\begin{aligned} \frac{\partial \psi}{\partial x} &= 0, \quad z = 0, -Ra, \\ \frac{\partial c}{\partial z} &= 0, \quad z = -Ra, \\ c &= 1, \quad z = 0, \end{aligned} \quad (3.18)$$

while the initial condition is still given by Eq. (3.14).

In two dimensions, Numbat solves Eq. (3.16) and Eq. (3.17).

3.2.2 3D solution

We now consider the case of a three-dimensional model. For simplicity, we consider the case where all lateral permeabilities are equal ($k_y = k_x$). The governing equations for the 3D model are identical to the 2D model. In dimensionless form, they are given by Eq. (3.9) to Eq. (3.11), with boundary conditions given by Eq. (3.12), and initial condition given by Eq. (3.14).

To solve these governing equations in 3D, a different approach must be used as the streamfunction ψ is not defined in three dimensions. Instead, we define a vector potential $\Psi = (\psi_x, \psi_y, \psi_z)$ such that

$$\mathbf{u} = \nabla \times \Psi. \quad (3.19)$$

It is important to note that the vector potential is only known up to the addition of the gradient of a scalar ζ as

$$\nabla \times (\Psi + \nabla\zeta) = \nabla \times \Psi \quad \forall \zeta, \quad (3.20)$$

as $\nabla \times \nabla\zeta = 0$ for any scalar ζ . This uncertainty is referred to as gauge freedom, and is common in electrodynamics. Taking the curl of Eq. (3.9) and substituting Eq. (3.19), we have

$$\nabla(\nabla \cdot \Psi) - \nabla^2 \Psi = \left(-\frac{\partial c}{\partial y}, \frac{\partial c}{\partial x}, 0 \right), \quad (3.21)$$

where we have again used the fact that $\nabla \times \nabla P = 0$. If we choose $\nabla \cdot \Psi = 0$ to specify the gauge condition, this simplifies to

$$\nabla^2 \Psi = \left(\frac{\partial c}{\partial y}, -\frac{\partial c}{\partial x}, 0 \right). \quad (3.22)$$

As shown in E & Liu (1997) $\nabla \cdot \Psi = 0$ is satisfied throughout the domain if

$$\begin{aligned} \psi_x = \psi_y = 0, \quad z = 0, -Ra, \\ \frac{\partial \psi_z}{\partial z} = 0, \quad z = 0, -Ra. \end{aligned} \quad (3.23)$$

The governing equations are then

$$\nabla^2 \Psi = \left(\frac{\partial c}{\partial y}, -\frac{\partial c}{\partial x}, 0 \right), \quad (3.24)$$

$$\frac{\partial c}{\partial t} + \mathbf{u} \cdot \nabla c = \gamma \left(\frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial y^2} \right) + \frac{\partial^2 c}{\partial z^2}, \quad (3.25)$$

where the continuity is satisfied automatically because $\nabla \cdot (\nabla \times \Psi) = 0$ for any Ψ .

Finally, it is straightforward to show that $\psi_z = 0$ in order to satisfy $\nabla^2 \psi_z = 0$ and $\frac{\partial \psi_z}{\partial z} = 0$, which means that the vector potential has only x and y components,

$$\Psi = (\psi_x, \psi_y, 0), \quad (3.26)$$

and therefore the fluid velocity $\mathbf{u} = (u, v, w)$ is

$$\mathbf{u} = \left(-\frac{\partial\psi_y}{\partial z}, \frac{\partial\psi_x}{\partial z}, \frac{\partial\psi_y}{\partial x} - \frac{\partial\psi_x}{\partial y} \right). \quad (3.27)$$

Note that if there is no y dependence, Eq. (3.24) and (3.25) reduce to

$$\nabla^2\Psi = \left(0, -\frac{\partial c}{\partial x}, 0 \right), \quad (3.28)$$

$$\frac{\partial c}{\partial t} + \mathbf{u} \cdot \nabla c = \gamma \frac{\partial^2 c}{\partial x^2} + \frac{\partial^2 c}{\partial z^2}.$$

It is simple to show that $\nabla^2\psi_x = 0$ and $\psi_x = 0$ at $z = 0, -Ra$ are only satisfied if $\psi_x = 0$ in the entire domain. In this case, the governing equations reduce to the two-dimensional formulation, as expected.

In three dimensions, Numbat solves Eq. (3.24) and Eq. (3.25).

4 Installation instructions

To install Numbat, follow these simple instructions.

4.1 Install MOOSE

Numbat is based on the MOOSE framework, so the first step is to install MOOSE. For detailed installation instructions depending on your hardware, see www.mooseframework.com.

4.2 Clone Numbat

The next step is to clone the Numbat repository to your local machine.

In the following, it is assumed that MOOSE was installed to the directory `~/projects`. If MOOSE was installed to a different directory, the following instructions must be modified accordingly.

To clone Numbat, use the following commands

```
cd ~/projects
git clone https://github.com/cpgr/numbat.git
cd numbat
git checkout master
```

4.3 Compile Numbat

Next, compile Numbat using

```
make -jn
```

where n is the number of processing cores on the computer. If everything has gone well, Numbat should compile without error, producing a binary named `numbat-opt`.

4.4 Test Numbat

Finally, to test that the installation worked, the test suite can be run using

```
./run_tests -jn
```

where n is the number of processing cores on the computer. At this stage, all of the Numbat tests should have run successfully, and you are ready to run more complicated simulations, see the [2D examples](#) and [3D examples](#) for more details.

4.5 Keep up to date

New features and changes to Numbat may be implemented from time to time. To ensure that Numbat continues to work without issue, you should make sure that you update your installation periodically. This can be done using

```
git fetch origin
git rebase origin/master
make -jn
./run_tests -jn
```

5 Implementation details

5.1 Rayleigh number

The Rayleigh number relates the relative importance of buoyancy forces to diffusive forces, and is an important dimensionless number in convection. Slim (2014) showed that increasing the Rayleigh number increases the number of flow regimes that the convective mixing process goes through.

In the dimensionless model implemented in Numbat, the Rayleigh number appears in the vertical position of the bottom boundary only (Slim, 2014). As a result, it is simple to vary the Rayleigh number in the dimensionless model by simply extending the mesh in the vertical direction.

In the dimensional model, the Rayleigh number may be varied by changing the various physical parameters present, such as permeability, fluid viscosity etc.

5.2 Mesh considerations

The process of density-driven convective mixing in porous media begins initially as diffusive mass transfer across the interface between a saturated fluid lying atop an unsaturated fluid. After a sufficient time, the gravitational instability grows until small downwelling fingers of saturated fluid descend into the unsaturated fluid. As time continues, these fingers grow and merge, forming larger fingers. The temporal evolution of this process is shown in Figure 5.1. The top image shows an initial diffusive profile. As time increases, fingers form, merge and grow in a complex manner.

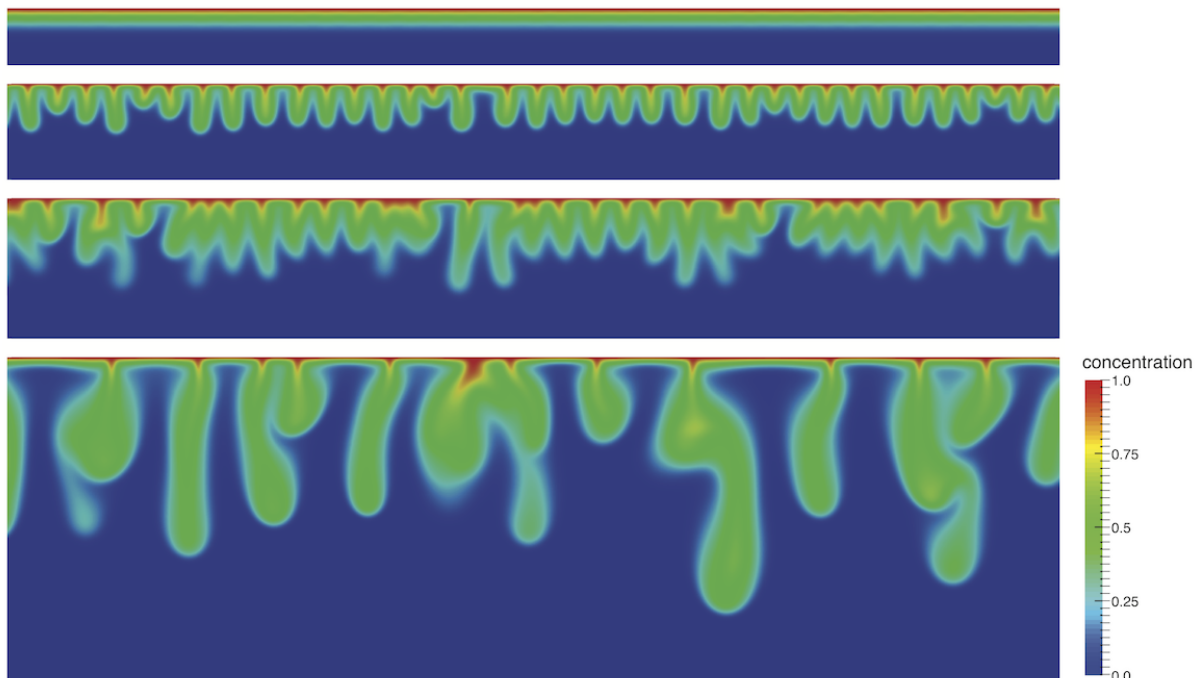


Figure 5.1: Evolution of convective dissolution (time increasing downwards). Note that the scale is identical for all images.

It is apparent from the temporal evolution above that the mesh requirements differ in different regimes. Early on and near the top of the mesh, the elements must be sufficiently refined to capture the initial formation of small fingers. Later on, and away from the top boundary, larger elements may adequately capture the large downwelling fingers, and as such, the mesh may be coarser in this region.

There are two ways that the user may deal with these disparate mesh requirements: use a mesh that is more refined in the vicinity of the top boundary; or use adaptive mesh refinement. Both of these are possible in Numbat.

Numbat provides a type of GeneratedMesh where the elements are biased towards the top boundary, NumbatBiasedMesh, so that it is more refined in the vicinity of the constant concentration boundary. This is particularly useful in two dimensions.

For three-dimensional models, an external meshing code may be used to generate an unstructured mesh. MOOSE can read a number of mesh formats natively, see the MOOSE documentation for details. One open-source option for generating unstructured meshes that MOOSE can read is Gmsh. An example of a fully unstructured 3D mesh that is refined near the top boundary and coarse near the bottom boundary is shown in Figure 5.2.

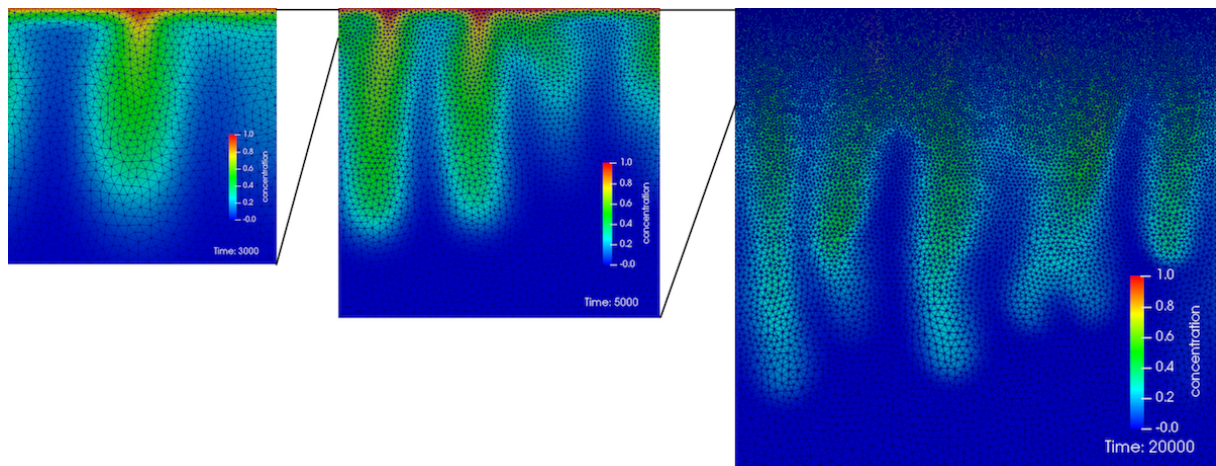


Figure 5.2: Example of unstructured mesh used in Numbat that was generated by Gmsh. Leftmost image shows small section near the top with the concentration profile at an early time; middle image shows top quarter of full mesh and concentration profile at an intermediate time; rightmost image shows full mesh and concentration profile at late time.

5.3 Seeding the perturbation

Numerical simulations of solutal convection rely on seeding the gravitational instability to initiate the convective mixing process. Numerical roundoff will eventually initiate convective mixing, but can significantly overestimate the critical time for the onset of convection.

Instead, it is usual to seed the instability using a prescribed random perturbation in the model. Several types of perturbation have been considered in the literature, see Emami-Meybodi *et al.* (2015) for a detailed review.

In Numbat, an initial perturbation to seed the instability can be applied in several ways:

- In the dimensional formulation, a random noise sampled from a uniform distribution can

be applied to the porosity;

- In the dimensionless streamfunction formulation, an initial perturbation to the diffusive concentration profile can be applied;
- A random fluctuation to the concentration boundary condition can be applied.

The strength of the initial perturbation has been shown to affect the initial onset time for convection, see Emami-Meybodi *et al.* (2015) for details. This should be taken into consideration when modelling convective mixing in porous media.

6 Input file syntax

The input file for a Numbat simulation is a simple, block-structured text file based on the input files for a plain MOOSE input file.

6.1 Essential input

Details of the minimum input file requirements are given below.

6.1.1 Mesh

All simulations must feature a mesh. For the basic model with a rectangular mesh, the built-in `NumbatBiasedMesh` can be used to create a suitable mesh. This is a type of `GeneratedMesh` that provides the option to refine the mesh near one boundary. The size of the initial element can be specified, after which the elements are progressively coarser, see `NumbatBiasedMesh` for details. This can be extremely useful in simulations of density-driven convection, where it is necessary to have a finer mesh in the vicinity of the boundary where the fingers form in order to capture the process adequately. Away from this region, the fingers grow and merge, so that a coarser mesh is sufficient to simulate them. Having a biased mesh such as that produced by `NumbatBiasedMesh` can minimise the number of elements necessary, reducing the overall computational demands.

In 2D, the input block looks like:

```
[Mesh]
  type = NumbatBiasedMesh
  dim = 2
  ymax = 1.5
  nx = 100
  ny = 50
  refined_edge = top
  refined_resolution = 0.001
[]
```

In 3D, the Mesh block would look like:

```
[Mesh]
  type = NumbatBiasedMesh
  dim = 3
  zmax = 1.5
  nx = 20
  ny = 20
  nz = 500
  refined_edge = front
  refined_resolution = 0.001
[]
```

Note:

In contrast to the normal `GeneratedMesh` provided by MOOSE, `NumbatBiasedMesh` renames the boundaries of the three dimensional mesh so that the boundaries `top` and `bottom` are at the extrema of the `z` axis.

6.1.2 Variables

The number and type of variables required depends on the chosen formulation. For the dimensional formulation, two nonlinear variables must be provided, representing the fluid pressure and solute concentration.

```
[Variables]
  [./concentration]
    initial_condition = 0
    scaling = 1e6
  [./]
  [./pressure]
    initial_condition = 1e6
  [./]
[]
```

For the dimensionless streamfunction formulation, the nonlinear variables for a 2D simulations are solute concentration and streamfunction:

```
[Variables]
  [./concentration]
    order = FIRST
    family = LAGRANGE
  [./]
  [./streamfunction]
    order = FIRST
    family = LAGRANGE
    initial_condition = 0.0
  [./]
[]
```

In 3D, an additional streamfunction variable must also be added:

```
[Variables]
  [./concentration]
    order = FIRST
    family = LAGRANGE
    [./InitialCondition]
      type = NumbatPerturbationIC
      variable = concentration
      amplitude = 0.1
      seed = 1
    [./]
  [./]
  [./streamfunctionx]
    order = FIRST
    family = LAGRANGE
    initial_condition = 0.0
  [./]
  [./streamfunctiony]
    order = FIRST
    family = LAGRANGE
    initial_condition = 0.0
  [./]
[]
```

6.1.3 Materials

For the dimensional formulation, several material and fluid properties are required: porosity, permeability, fluid density and viscosity, and diffusivity. These can be added using the following Numbat materials:

```
[Materials]
  [./porosity]
    type = NumbatPorosity
    porosity = 0.3
    noise = noise
  [./]
  [./permeability]
    type = NumbatPermeability
    permeability = '1e-11 0 0 0 1e-11 0 0 0 1e-11'
  [./]
  [./diffusivity]
    type = NumbatDiffusivity
    diffusivity = 2e-9
  [./]
  [./density]
    type = NumbatDensity
    concentration = concentration
    zero_density = 995
    delta_density = 10.5
    saturated_concentration = 0.049306
  [./]
  [./viscosity]
    type = NumbatViscosity
    viscosity = 6e-4
  [./]
[]
```

Note:

No material properties are required in the dimensionless streamfunction formulation

6.1.4 Kernels

Four kernels are required for a dimensional model: NumbatTimeDerivative, NumbatDiffusion, NumbatConvection, and NumbatDarcy.

```
[Kernels]
  [./time]
    type = NumbatTimeDerivative
    variable = concentration
  [./]
  [./diffusion]
    type = NumbatDiffusion
    variable = concentration
  [./]
  [./convection]
    type = NumbatConvection
    variable = concentration
    pressure = pressure
  [./]
  [./darcy]
```

```

type = NumbatDarcy
variable = pressure
concentration = concentration
[../]
[]

```

For the dimensionless streamfunction formulation, four kernels are required for a 2D model: a NumbatDarcySF kernel, a NumbatDiffusionSF kernel, a NumbatConvectionSF kernel, and a TimeDerivative kernel.

```

[Kernels]
[./Darcy]
type = NumbatDarcySF
variable = streamfunction
concentration = concentration
[../]
[./Convection]
type = NumbatConvectionSF
variable = concentration
streamfunction = streamfunction
[../]
[./Diffusion]
type = NumbatDiffusionSF
variable = concentration
[../]
[./TimeDerivative]
type = TimeDerivative
variable = concentration
[../]
[]

```

For 3D models, an additional NumbatDarcySF kernel is required for the additional streamfunction variable. An example of the kernels block for a 3D isotropic model is

```

[Kernels]
[./Darcy_x]
type = NumbatDarcySF
variable = streamfunctionx
concentration = concentration
component = x
[../]
[./Darcy_y]
type = NumbatDarcySF
variable = streamfunctiony
concentration = concentration
component = y
[../]
[./Convection]
type = NumbatConvectionSF
variable = concentration
streamfunction = 'streamfunctionx streamfunctiony'
[../]
[./Diffusion]
type = NumbatDiffusionSF
variable = concentration
[../]
[./TimeDerivative]

```

```

    type = TimeDerivative
    variable = concentration
  [../]
[]

```

In the 3D case, it is important to note that the NumbatDarcySF kernel must specify the component that it applies to, and that the `streamfunction` keyword in the NumbatConvectionSF kernel must contain both streamfunction variables ordered by the x component then the y component.

Note:

For the streamfunction formulation, a TimeDerivative kernel is used, rather than a NumbatTimeDerivative kernel, as porosity has been scaled out of the problem.

6.1.5 Boundary conditions

Appropriate boundary conditions must be prescribed. Typically, these will be constant concentration at the top of the model domain, periodic boundary conditions on the lateral sides (to mimic an infinite model), and no-flow boundary conditions at the top and bottom surfaces.

In the 2D dimensional formulation, this can be achieved using the following input block:

```

[BCs]
  [./conctop]
    type = PresetBC
    variable = concentration
    boundary = top
    value = 0.049306
  [../]
  [./Periodic]
    [./x]
      variable = concentration
      auto_direction = x
    [../]
  [../]
[]

```

while in 3D

```

[BCs]
  [./conctop]
    type = PresetBC
    variable = concentration
    boundary = front
    value = 0.049306
  [../]
  [./Periodic]
    [./x]
      variable = concentration
      auto_direction = 'x y'
    [../]
  [../]
[]

```

In this case, the `conctop` boundary condition fixes the value of concentration at the top boundary, while the `Periodic` boundary condition enforces periodicity of concentration along

the boundaries in the directions specified in the `auto_direction` parameter.

It is useful to note that a MOOSE `GeneratedMesh` provides descriptive names for the sides of the model (top, bottom, left, right) which can be referenced in the input file.

For the dimensionless streamfunction formulation, no-flow boundary conditions are prescribed on the top and bottom surfaces by holding the streamfunction variable constant (in this case 0).

```
[BCs]
  [./conctop]
    type = DirichletBC
    variable = concentration
    boundary = top
    value = 1.0
  [./]
  [./streamfuntop]
    type = DirichletBC
    variable = streamfunction
    boundary = top
    value = 0.0
  [./]
  [./streamfunbottom]
    type = DirichletBC
    variable = streamfunction
    boundary = bottom
    value = 0.0
  [./]
  [./Periodic]
    [./x]
      variable = 'concentration streamfunction'
      auto_direction = x
    [./]
  [./]
[]
```

6.1.6 Executioner

Each MOOSE simulation must use an Executioner, which provides parameters for the solve.

```
[Executioner]
  type = Transient
  l_max_its = 200
  end_time = 3e5
  solve_type = NEWTON
  petsc_options = -ksp_snes_ew
  petsc_options_iname = '-pc_type -sub_pc_type -ksp_atol'
  petsc_options_value = 'bjacobi ilu 1e-12'
  nl_abs_tol = 1e-10
  nl_max_its = 25
  dtmax = 2e3
  [./TimeStepper]
    type = IterationAdaptiveDT
    dt = 1
  [./]
[]
```

Executioners are a standard MOOSE feature that are well documented on the MOOSE, so no

further detail is provided here.

6.1.7 Preconditioning

A default preconditioning block is used that provides all Jacobian entries to aid convergence.

```
[Preconditioning]
  [./smp]
    type = SMP
    full = true
  [./]
[]
```

This is a standard MOOSE feature that is documented on the MOOSE website, so no further detail is provided here.

6.1.8 Outputs

To provide output from the simulation, an Outputs block must be specified. An example is

```
[Outputs]
  perf_graph = true
  [./exodus]
    type = Exodus
    file_base = 2Dddc
    execute_on = 'INITIAL TIMESTEP_END '
  [./]
  [./csvoutput]
    type = CSV
    file_base = 2Dddc
    execute_on = 'INITIAL TIMESTEP_END '
  [./]
[]
```

There are a large number of output options available in MOOSE, see the MOOSE website for further details.

6.2 Action system

To avoid having to enter several of these input file blocks each time, and ensuring that the correct parameters are provided to each object in the correct order, Numbat provides some powerful actions that automatically add most of the required objects.

The NumbatAction adds all of the nonlinear variables, kernels, aux variables, aux kernels and postprocessors typically required in a dimensional Numbat simulation.

This action is called in the input file simply as

```
[Numbat]
  [./Dimensional]
  [./]
[]
```

The use of this action is exactly equivalent to the following input file syntax

```
[Variables]
  [./concentration]
    initial_condition = 0
  [./]
```

```

[./pressure]
  initial_condition = 1e6
[./]
[]

[AuxVariables]
[./u]
  order = CONSTANT
  family = MONOMIAL
[./]
[./v]
  order = CONSTANT
  family = MONOMIAL
[./]
[]

[Kernels]
[./time]
  type = NumbatTimeDerivative
  variable = concentration
[./]
[./diffusion]
  type = NumbatDiffusion
  variable = concentration
[./]
[./convection]
  type = NumbatConvection
  variable = concentration
  pressure = pressure
[./]
[./darcy]
  type = NumbatDarcy
  variable = pressure
  concentration = concentration
[./]
[]

[AuxKernels]
[./uAux]
  type = NumbatDarcyVelocity
  pressure = pressure
  variable = u
  component = x
[./]
[./vAux]
  type = NumbatDarcyVelocity
  pressure = pressure
  variable = v
  component = y
[./]
[]

[BCs]
[./conctop]
  type = DirichletBC
  variable = concentration

```

```

    boundary = top
    value = 1.0
  [../]
  [./Periodic]
    [./x]
      variable = 'concentration pressure'
      auto_direction = x
    [../]
  [../]
[]

[Postprocessors]
  [./boundary_flux]
    type = NumbatSideFlux
    variable = concentration
    boundary = top
  [../]
  [./total_mass]
    type = NumbatTotalMass
    variable = concentration
  [../]
[]

```

A specific value for the saturated boundary concentration can optionally be provided

```

[Numbat]
  [./Dimensional]
    boundary_concentration = 0.05
  [../]
[]

```

Similarly, the NumbatSFAction adds all of the nonlinear variables, kernels, aux variables, aux kernels and postprocessors typically required in a dimensionless Numbat simulation.

This action is called in the input file simply as

```

[Numbat]
  [./Dimensionless]
  [../]
[]

```

The use of this action is exactly equivalent to the following input file syntax for a 2D simulation.

```

[Variables]
  [./concentration]
    order = FIRST
    family = LAGRANGE
    initial_condition = 0.0
  [../]
  [./streamfunction]
    order = FIRST
    family = LAGRANGE
    initial_condition = 0.0
  [../]
[]

[AuxVariables]
  [./u]

```

```

    order = CONSTANT
    family = MONOMIAL
[../]
[./v]
    order = CONSTANT
    family = MONOMIAL
[../]
[]

[Kernels]
[./Darcy]
    type = NumbatDarcySF
    variable = streamfunction
    concentration = concentration
[../]
[./Convection]
    type = NumbatConvectionSF
    variable = concentration
    streamfunction = streamfunction
[../]
[./Diffusion]
    type = NumbatDiffusionSF
    variable = concentration
[../]
[./TimeDerivative]
    type = TimeDerivative
    variable = concentration
[../]
[]

[AuxKernels]
[./uAux]
    type = NumbatDarcyVelocitySF
    variable = u
    component = x
    streamfunction = streamfunction
[../]
[./vAux]
    type = NumbatDarcyVelocitySF
    variable = v
    component = y
    streamfunction = streamfunction
[../]
[]

[BCs]
[./conctop]
    type = DirichletBC
    variable = concentration
    boundary = top
    value = 1.0
[../]
[./streamfuntop]
    type = DirichletBC
    variable = streamfunction
    boundary = top

```

```

    value = 0.0
  [./]
  [./streamfunbottom]
    type = DirichletBC
    variable = streamfunction
    boundary = bottom
    value = 0.0
  [./]
  [./Periodic]
    [./x]
      variable = 'concentration streamfunction'
      auto_direction = x
    [./]
  [./]
[]

[Postprocessors]
  [./boundary_flux]
    type = NumbatSideFluxSF
    variable = concentration
    boundary = top
  [./]
  [./total_mass]
    type = NumbatTotalMassSF
    variable = concentration
  [./]
[]

```

The use of these actions is recommended for all users, as they reduce the possibility of input file errors.

6.3 Optional input

While the above required blocks will enable a Numbat simulation to run, there are a number of optional input blocks that will improve the simulations and increase the amount of results provided.

6.3.1 Mesh adaptivity

MOOSE features built-in mesh adaptivity that is extremely useful in Numbat simulations to reduce computational expense. This can be included using:

```

[Adaptivity]
  max_h_level = 1
  initial_marker = boxmarker
  initial_steps = 1
  marker = errormarker
  [./Indicators]
    [./gradjumpindicator]
      type = GradientJumpIndicator
      variable = concentration
    [./]
  [./]
  [./Markers]
    [./errormarker]
      type = ErrorToleranceMarker
      refine = 0.05
  [./]

```

```

    indicator = gradjumpindicator
  [../]
  [./boxmarker]
    type = BoxMarker
    bottom_left = '0 0 -10'
    top_right = '500 500 0'
    inside = refine
    outside = dont_mark
  [../]
[../]
[]

```

For details about mesh adaptivity, see the MOOSE website.

6.3.2 Initial condition

To seed the instability, a random perturbation to the initial concentration can be prescribed using the NumbatPerturbationIC initial condition.

```

[ICs]
  [./concentration]
    type = NumbatPerturbationIC
    variable = concentration
    amplitude = 0.1
    seed = 1
  [../]
[]

```

The NumbatPerturbationIC initial condition applies the diffusive concentration profile to the nodes for (scaled) time $t = 1$,

$$c_d(z, t = 1) = 1 + \operatorname{erf}(z/2), \quad (6.1)$$

for $z < 0$, where $\operatorname{erf}(z)$ is the error function.

A uniform random perturbation is then added to the diffusive concentration profile, where the amplitude of the perturbation is specified by the input file value `amplitude`.

6.3.3 Postprocessors

The flux over the top boundary or the total mass of solute in the model is of particular interest in many cases (especially convective mixing of CO_2). These can be calculated at each time step using the NumbatSideFlux and NumbatTotalMass Postprocessors.

```

[Postprocessors]
  [./boundaryfluxint]
    type = NumbatSideFlux
    variable = concentration
    boundary = top
  [../]
  [./mass]
    type = NumbatTotalMass
    variable = concentration
  [../]
[]

```

Versions of these Postprocessors for the dimensionless streamfunction formulation are also provided, see NumbatSideFluxSF and NumbatTotalMassSF for details.

Numbat also provides a simple Postprocessor to calculate the Rayleigh number for dimensional simulations, see NumbatRayleighNumber for details.

6.3.4 AuxKernels

The velocity components in the x and y directions (in 2D), and x , y , and z directions in 3D can be calculated using the auxiliary system. These velocity components are calculated using the streamfunction(s), see the governing equations for details.

In the 2D case, two auxiliary variables, u and w , can be defined for the horizontal and vertical velocity components, respectively.

Note:

Importantly, these auxiliary variables **must** have monomial shape functions (these are referred to as elemental variables, as the value is constant over each mesh element). This restriction is due to fact that the gradient of variables is undefined for nodal auxiliary variables (for example, those using linear Lagrange shape functions).

An example of the input syntax for the 2D case is

```
[AuxVariables]
  [./u]
    order = CONSTANT
    family = MONOMIAL
  [./]
  [./w]
    order = CONSTANT
    family = MONOMIAL
  [./]
[]
```

For the 3D case, there is an additional horizontal velocity component (v), so the input syntax is

```
[AuxVariables]
  [./u]
    order = CONSTANT
    family = MONOMIAL
  [./]
  [./v]
    order = CONSTANT
    family = MONOMIAL
  [./]
  [./w]
    order = CONSTANT
    family = MONOMIAL
  [./]
[]
```

The velocity components are calculated by NumbatDarcyVelocity AuxKernels (or NumbatDarcyVelocitySF AuxKernels for the dimensionless streamfunction formulation). Each velocity component is computed by an AuxKernel.

For the 2D case, two AuxKernels are required:

```
[AuxKernels]
[./uAux]
  type = NumbatDarcyVelocitySF
  variable = u
  component = x
  streamfunction = streamfunction
[./]
[./wAux]
  type = NumbatDarcyVelocitySF
  variable = w
  component = y
  streamfunction = streamfunction
[./]
[]
```

while for 3D, three AuxKernels are necessary:

```
[AuxKernels]
[./uAux]
  type = NumbatDarcyVelocitySF
  variable = u
  component = x
  streamfunction = 'streamfunctionx streamfunctiony'
[./]
[./vAux]
  type = NumbatDarcyVelocitySF
  variable = v
  component = y
  streamfunction = 'streamfunctionx streamfunctiony'
[./]
[./wAux]
  type = NumbatDarcyVelocitySF
  variable = w
  component = z
  streamfunction = 'streamfunctionx streamfunctiony'
[./]
[]
```

Note:

For the 3D case, both streamfunction variables must be given, in the correct order (eg. x then y)

7 Running the Numbat application

7.1 Commandline

Most often, Numbat will be run from the commandline using

```
./numbat-opt -i input.i
```

Numbat (and all MOOSE applications) have a large number of commandline options available to the user. The complete list can be viewed using the `--help` option

```
./numbat-opt --help
```

7.1.1 Recovering

If you output checkpoint files (using `checkpoint = true` in your Outputs block) then the `--recover` option will allow you to continue a solve that died in the middle of the solve, perhaps because the job ran out of time on the cluster you were using.

We recommend that all input files for large Numbat simulations enable checkpointing. This can be enabled using

```
[Outputs]
  perf_graph = true
  [./csvoutput]
    type = CSV
    file_base = 2DSF
    execute_on = 'INITIAL TIMESTEP_END '
  [./]
  [./checkpoint]
    type = Checkpoint
    num_files = 2
  [./]
[]
```

For all of the options available for checkpointing, see the MOOSE documentation.

If a long-running simulation does fail to complete, it can be recovered by running

```
./numbat-opt --recover checkpoint_dir/XXXX -i input.i
```

where `checkpoint_dir` is the subdirectory where the checkpoint files are saved, and `XXXX` is the number of one of the available checkpoint files.

7.1.2 Overriding parameters

MOOSE provides a handy feature where any parameter in the input file can be overridden from the commandline, making it possible to script studies where only parameters are changed from simulation to simulation.

For example, assume that the anisotropy γ of the porous medium is set as 1 in the input file

```
[Kernels]
  [./Darcy]
```

```
type = NumbatDarcySF
variable = streamfunction
concentration = concentration
gamma = 1
[../]
[]
```

This value can be changed to 0.5 by running Numbat with the following commandline option

```
./numbat-opt -i input.i Kernels/Darcy/gamma=0.5
```

7.2 Graphical user interface (Peacock)

MOOSE provides a graphical user interface, Peacock, which can be used to both run simulations and create input files. Starting Peacock from within the base Numbat directory allows Peacock to extract the Numbat syntax, so that all Numbat objects are available in the menus.

Peacock can be run using

```
$MOOSE_DIR/python/peacock/peacock -i input.i
```

where \$MOOSE_DIR is the directory where the MOOSE repository is located.

Note:

It is not recommended to use Peacock to run very large models (e.g. three dimensional simulations) that require lots of memory.

8 Two-dimensional examples

Complete input files for 2D models using the dimensional and dimensionless streamfunction formulations are provided, for both isotropic and anisotropic porous media. These examples are provided in the Numbat *examples* folder.

8.1 Isotropic models

The first 2D examples are for an isotropic porous medium ($\gamma = 1$).

8.1.1 Input file

The complete input file for this problem is

```
# 2D density-driven convective mixing
# The dimensional Numbat action is used to add all variables, kernels
  etc.
# Instability is seeded by small perturbation to porosity.
# A coarse grid and time stepping used to allow this model to run in ~
  15 minutes

[Mesh]
  type = NumbatBiasedMesh
  dim = 2
  nx = 150
  ny = 50
  ymax = 0.5
  refined_edge = top
  refined_resolution = 0.001
[]

[Numbat]
  [./Dimensional]
    concentration_scaling = 1e4
    boundary_concentration = 0.049306
    periodic_bcs = true
  [../]
[]

[AuxVariables]
  [./noise]
    family = MONOMIAL
    order = CONSTANT
  [../]
[]

[ICs]
  [./noise]
    type = RandomIC
    variable = noise
    max = 0.003
    min = -0.003
  [../]
  [./pressure]
    type = ConstantIC
    variable = pressure
    value = 10e6
```

```

[../]
[]

[Materials]
[./porosity]
  type = NumbatPorosity
  porosity = 0.3
  noise = noise
[../]
[./permeability]
  type = NumbatPermeability
  permeability = '1e-11 0 0 0 1e-11 0 0 0 1e-11'
[../]
[./diffusivity]
  type = NumbatDiffusivity
  diffusivity = 2e-9
[../]
[./density]
  type = NumbatDensity
  concentration = concentration
  zero_density = 994.56
  delta_density = 10.45
  saturated_concentration = 0.049306
[../]
[./viscosity]
  type = NumbatViscosity
  viscosity = 0.5947e-3
[../]
[]

[Preconditioning]
[./smp]
  type = SMP
  full = true
[../]
[]

[Functions]
[./timesteps]
  type = PiecewiseConstant
  x = '0 100 500 1e3 1e4 2e5'
  y = '10 50 100 500 1e3 1e3'
[../]
[]

[Executioner]
  type = Transient
  l_max_its = 100
  end_time = 2e5
  solve_type = NEWTON
  petsc_options = -ksp_snes_ew
  petsc_options_iname = '-pc_type -sub_pc_type -ksp_atol'
  petsc_options_value = 'asm ilu 1e-12'
  nl_abs_tol = 1e-8
[./TimeStepper]
  type = FunctionDT

```

```

function = timesteps
[./]
[]

[Outputs]
perf_graph = true
[./exodus]
type = Exodus
file_base = 2Dddc
execute_on = 'INITIAL TIMESTEP_END '
[./]
[./csvoutput]
type = CSV
file_base = 2Dddc
execute_on = 'INITIAL TIMESTEP_END '
[./]
[]

```

8.1.2 Running the example

This example can be run on the commandline using

```
numbat-opt -i 2Dddc.i
```

Alternatively, this file can be run using the *Peacock* gui provided by MOOSE using

```
peacock -i 2Dddc.i
```

in the directory where the input file *2Dddc.i* resides.

8.1.3 Results

This 2D example should take only a few minutes to run to completion, producing a concentration profile similar to that presented in Figure 8.1, where several downwelling plumes of high concentration can be observed after 3528 s:

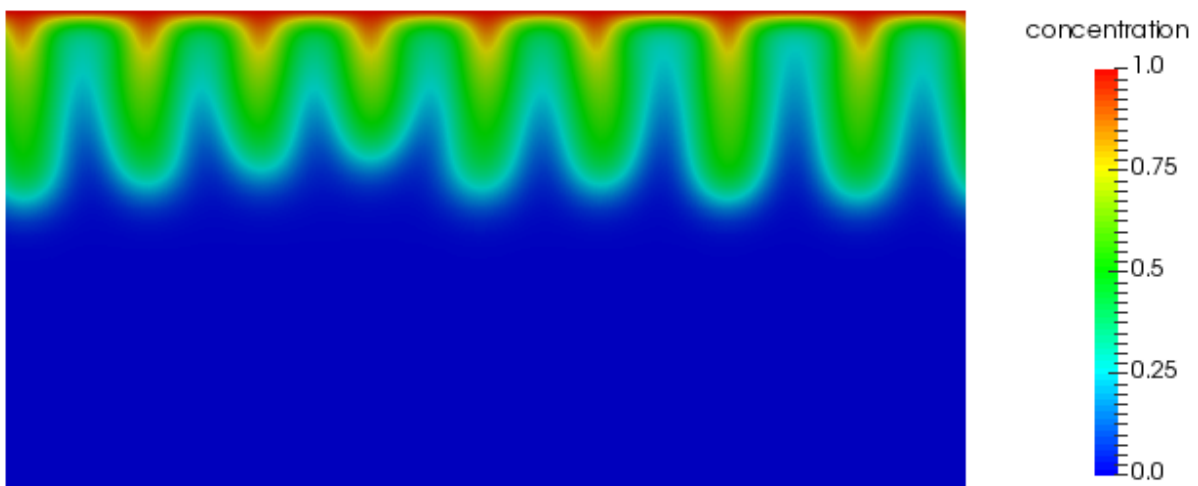


Figure 8.1: 2D concentration profile ($t = 3528$ s)

The flux per unit width over the top boundary is of particular interest in many cases (especially

convective mixing of CO₂). This is calculated using the *boundaryfluxint* postprocessor in the input file, and presented in 8.2.

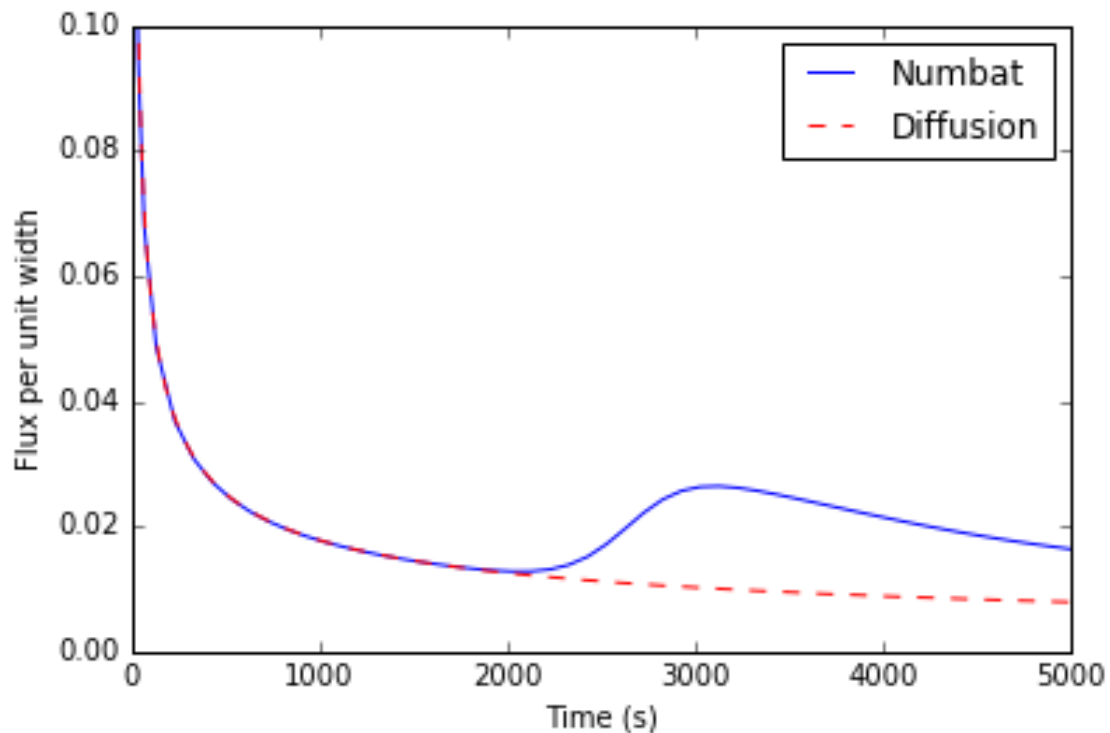


Figure 8.2: 2D flux across the top boundary

Initially, the flux is purely diffusive, and scales as $1/\sqrt{(\pi t)}$, where t is time (shown as the dashed red line). After some time, the convective instability becomes sufficiently strong, at which point the flux across the top boundary rapidly increases (at a time of approximately 2000 seconds).

8.2 Anisotropic models

The second 2D example is for an anisotropic porous medium with $\gamma = 0.75$ (ie., the vertical permeability is three quarters of the horizontal permeability).

8.2.1 Input file

```
# 2D density-driven convective mixing with permeability anisotropy gamma
= 0.5
# The dimensional Numbat action is used to add all variables, kernels
etc.
# Instability is seeded by small perturbation to porosity.
# A coarse grid and time stepping used to allow this model to run in ~
15 minutes

[Mesh]
type = NumbatBiasedMesh
dim = 2
```

```

nx = 150
ny = 50
ymax = 0.5
refined_edge = top
refined_resolution = 0.001
[]

[Numbat]
[./Dimensional]
  concentration_scaling = 1e4
  boundary_concentration = 0.049306
  periodic_bcs = true
[./]
[]

[AuxVariables]
[./noise]
  family = MONOMIAL
  order = CONSTANT
[./]
[]

[ICs]
[./noise]
  type = RandomIC
  variable = noise
  max = 0.003
  min = -0.003
[./]
[./pressure]
  type = ConstantIC
  variable = pressure
  value = 10e6
[./]
[]

[Materials]
[./porosity]
  type = NumbatPorosity
  porosity = 0.3
  noise = noise
[./]
[./permeability]
  type = NumbatPermeability
  permeability = '1e-11 0 0 0 5e-12 0 0 0 1e-11'
[./]
[./diffusivity]
  type = NumbatDiffusivity
  diffusivity = 2e-9
[./]
[./density]
  type = NumbatDensity
  concentration = concentration
  zero_density = 994.56
  delta_density = 10.45
  saturated_concentration = 0.049306

```

```

[../]
[./viscosity]
    type = NumbatViscosity
    viscosity = 0.5947e-3
[../]
[]

[Preconditioning]
[./smp]
    type = SMP
    full = true
[../]
[]

[Functions]
[./timesteps]
    type = PiecewiseConstant
    x = '0 100 500 1e3 1e4 4e5'
    y = '10 50 100 500 1e3 2e3'
[../]
[]

[Executioner]
    type = Transient
    l_max_its = 100
    end_time = 4e5
    solve_type = NEWTON
    petsc_options = -ksp_snes_ew
    petsc_options_iname = '-pc_type -sub_pc_type -ksp_atol'
    petsc_options_value = 'asm ilu 1e-12'
    nl_abs_tol = 1e-8
[./TimeStepper]
    type = FunctionDT
    function = timesteps
[../]
[]

[Outputs]
    perf_graph = true
[./exodus]
    type = Exodus
    file_base = 2Dddc
    execute_on = 'INITIAL TIMESTEP_END'
[../]
[./csvoutput]
    type = CSV
    file_base = 2Dddc
    execute_on = 'INITIAL TIMESTEP_END'
[../]
[]

```

Note that the permeability anisotropy is introduced in the kernels using the *gamma* and *anisotropic_tensor* input parameters.

8.2.2 Running the example

This example can be run on the commandline using


```
numbat-opt -i 2Dddc_anisotropic.i
```

Alternatively, this file can be run using the *Peacock* gui provided by MOOSE using

```
peacock -i 2Dddc_anisotropic.i
```

in the directory where the input file *2Dddc_anisotropic.i* resides.

8.2.3 Results

This 2D example should take only a few minutes to run to completion, producing a concentration profile similar to that presented in Figure 8.3, where several downwelling plumes of high concentration can be observed after 5000 s:

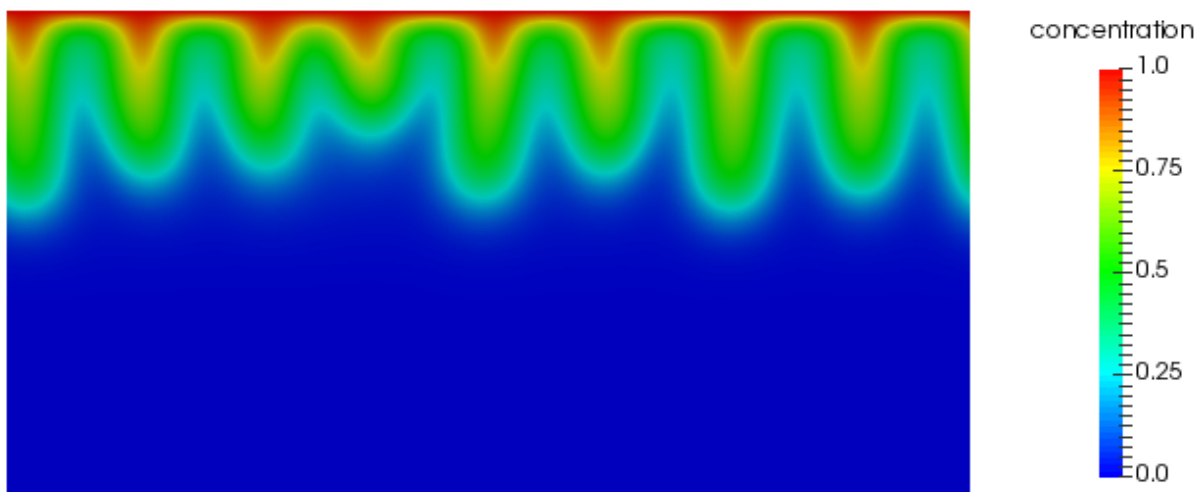


Figure 8.3: 2D concentration profile for $\gamma = 0.75$ ($t = 5000$ s)

In comparison to the isotropic example (with $\gamma = 1$) presented in Figure 8.1, we note that the concentration profile in the anisotropic example has only reached a similar depth after 5000 s (compared to 3528 s). The effect of the reduced vertical permeability in the anisotropic example slows the convective transport.

This observation can be quantified by comparing the flux per unit width over the top boundary of both examples, see Figure 8.4.

The inclusion of permeability anisotropy delays the onset of convection in comparison to the isotropic example, from a time of approximately 2000 seconds in the isotropic example to approximately 3500 seconds in the anisotropic example.

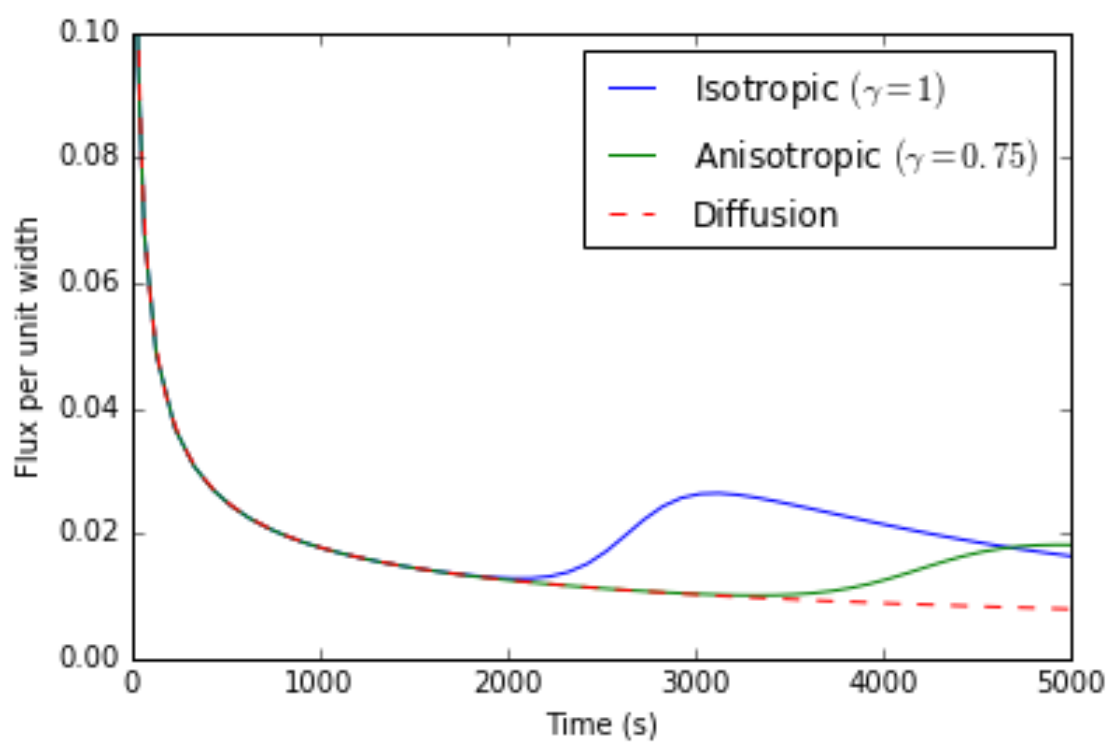


Figure 8.4: Comparison of the 2D flux across the top boundary

9 Three-dimensional examples

Complete input files for 2D models using the dimensional and dimensionless streamfunction formulations are provided, for both isotropic and anisotropic porous media. These examples are provided in the Numbat *examples* folder.

9.1 Isotropic models

The first examples are for an isotropic porous medium ($\gamma = 1$).

9.1.1 Input file

The complete input file for this problem is

```
# 3D density-driven convective mixing. Instability is seeded by small
  perturbation
# to porosity. Don't try this on a laptop!

[Mesh]
  type = NumbatBiasedMesh
  dim = 3
  zmax = 1.5
  nx = 20
  ny = 20
  nz = 500
  refined_edge = front
  refined_resolution = 0.001
[]

[Numbat]
  [./Dimensional]
    concentration_scaling = 1e6
    boundary_concentration = 0.049306
    periodic_bcs = true
  [../]
[]

[AuxVariables]
  [./noise]
    family = MONOMIAL
    order = CONSTANT
  [../]
[]

[ICs]
  [./noise]
    type = RandomIC
    variable = noise
    max = 0.003
    min = -0.003
  [../]
  [./pressure]
    type = ConstantIC
    variable = pressure
    value = 1e6
  [../]
[]
```

```

[Materials]
  ./porosity
    type = NumbatPorosity
    porosity = 0.3
    noise = noise
  ./]
  ./permeability
    type = NumbatPermeability
    permeability = '1e-11 0 0 0 1e-11 0 0 0 1e-11'
  ./]
  ./diffusivity
    type = NumbatDiffusivity
    diffusivity = 2e-9
  ./]
  ./density
    type = NumbatDensity
    concentration = concentration
    zero_density = 995
    delta_density = 10.5
    saturated_concentration = 0.049306
  ./]
  ./viscosity
    type = NumbatViscosity
    viscosity = 6e-4
  ./]
[]

[Preconditioning]
  ./smp]
    type = SMP
    full = true
  ./]
[]

[Executioner]
  type = Transient
  l_max_its = 200
  end_time = 3e5
  solve_type = NEWTON
  petsc_options = -ksp_snes_ew
  petsc_options_iname = '-pc_type -sub_pc_type -ksp_atol -pc_asm_overlap
  ,
  petsc_options_value = 'asm ilu 1e-12 4'
  nl_abs_tol = 1e-10
  nl_max_its = 25
  dtmax = 2e3
  ./TimeStepper]
    type = IterationAdaptiveDT
    dt = 1
  ./]
[]

[Outputs]
  perf_graph = true
  ./exodus]

```

```

type = Exodus
file_base = 3Dddc
execute_on = 'INITIAL TIMESTEP_END '
[./]
[./csvoutput]
type = CSV
file_base = 3Dddc
execute_on = 'INITIAL TIMESTEP_END '
[./]
[]

```

9.1.2 Running the example

Note:

This example should **not** be run on a laptop or workstation due to the large computational requirements. Do **not** run this using the *Peacock* gui provided by MOOSE.

Examples of the total run times for this problem on a cluster are over 27 hours for a single processor down to only 30 minutes using 100 processors in parallel.

9.1.3 Results

This 3D example should produce a concentration profile similar to that presented in Figure 9.1, where several downwelling plumes of high concentration can be observed:

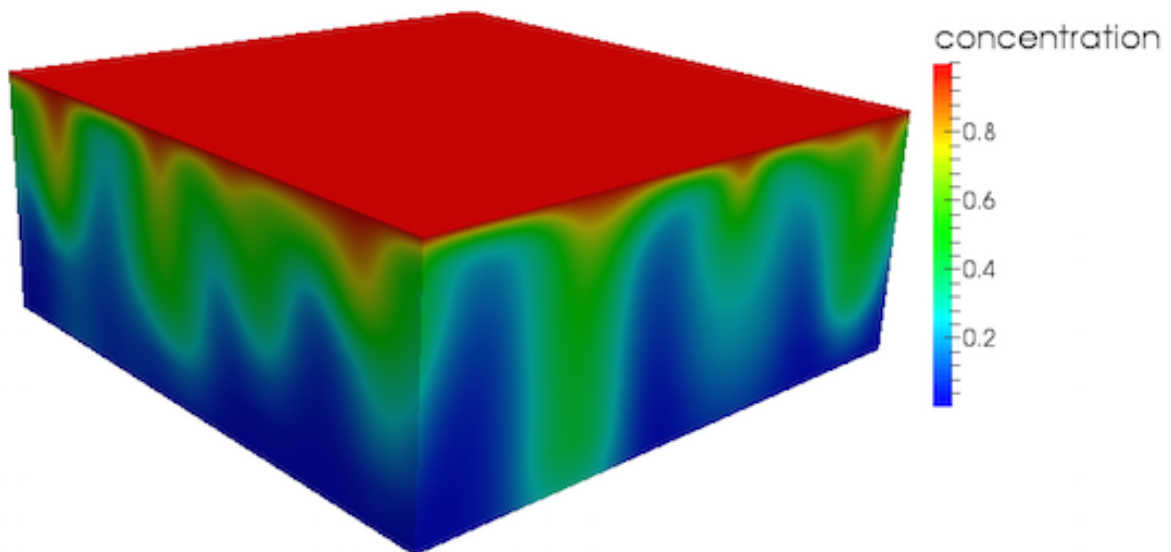


Figure 9.1: 3D concentration profile

Note that due to the random perturbation applied to the initial concentration profile, the geometry of the concentration profile obtained will differ from run to run.

The flux over the top surface is of particular interest in many cases (especially convective mixing of CO₂). This is calculated in this example file using the *NumbatSideFlux* in the input file, and presented in Figure 9.2.

Initially, the flux is purely diffusive, and scales as $1/\sqrt{(\pi t)}$, where t is time (shown as the

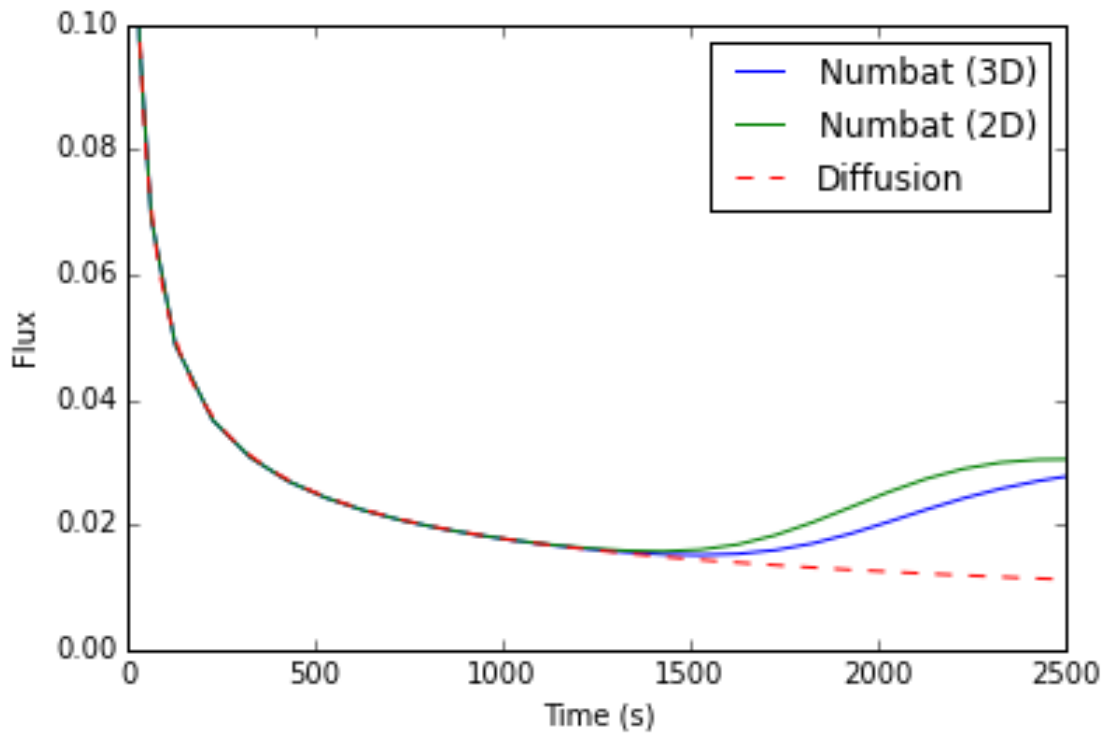


Figure 9.2: 3D flux across the top boundary

dashed green line). After some time, the convective instability becomes sufficiently strong, at which point the flux across the top boundary rapidly increases (at a time of approximately 1,700 seconds). Also shown for comparison is the flux for the 2D example. It is apparent that the 3D model leads in a slower onset of convection (the time where the flux first increases from the diffusive rate).

9.2 Large model

Increasing the size of the mesh enables more of the flow regimes to be modelled (at the cost of increased computational expense). Consider a dimensionless model with Rayleigh number $Ra = 5000$. Lateral model dimensions are chosen so that approximately twenty fingers in the lateral directions are present at the onset of convection.

A suitable fully unstructured mesh for this model that is sufficiently refined near the top boundary with increasing element size with depth to minimise the number of elements can be constructed in an external meshing code. In this example, the open-source mesh generator Gmsh is used. The following geometry file describes the dimensions of the model and the target resolutions.

```
// Gmsh geometry file for a 3D mesh corresponding to
// Ra = 5000 (dimensionless model).
// The mesh is refined at the top boundary with element
// size increasing with depth.
//
// This geometry file can be converted to a mesh using
```

```

// gmesh -3 Ra5000.geo

// Mesh width in each dimension
xmax = 2000;
ymax = 2000;
zmax = 5000;

// Resolution at the top and bottom
lctop = 10;
lcbot = 200;

// Vertices of mesh
Point(1) = {0, 0, 0, lcbot};
Point(2) = {xmax, 0, 0, lcbot};
Point(3) = {0, ymax, 0, lcbot};
Point(4) = {xmax, ymax, 0, lcbot};
Point(5) = {xmax, ymax, zmax, lctop};
Point(6) = {xmax, 0, zmax, lctop};
Point(7) = {0, ymax, zmax, lctop};
Point(8) = {0, 0, zmax, lctop};

// Lines connecting vertices
Line(1) = {3, 7};
Line(2) = {7, 5};
Line(3) = {5, 4};
Line(4) = {4, 3};
Line(5) = {3, 1};
Line(6) = {2, 4};
Line(7) = {2, 6};
Line(8) = {6, 8};
Line(9) = {8, 1};
Line(10) = {1, 2};
Line(11) = {8, 7};
Line(12) = {6, 5};

// Surfaces defined by lines
Line Loop(13) = {7, 8, 9, 10};
Plane Surface(14) = {13};
Line Loop(15) = {6, 4, 5, 10};
Plane Surface(16) = {15};
Line Loop(17) = {3, 4, 1, 2};
Plane Surface(18) = {17};
Line Loop(19) = {12, -2, -11, -8};
Plane Surface(20) = {19};
Line Loop(21) = {7, 12, 3, -6};
Plane Surface(22) = {21};
Line Loop(23) = {9, -5, 1, -11};
Plane Surface(24) = {23};
Surface Loop(25) = {14, 22, 20, 18, 16, 24};

// Volume defined by surfaces
Volume(26) = {25};

// Make the sides suitable for periodic boundary conditions
Periodic Surface {18} = {14} Translate {0, ymax, 0};
Periodic Surface {22} = {24} Translate {xmax, 0, 0};

```

```
// Name the faces for easy application of boundary conditions
Physical Surface("top") = {20};
Physical Surface("bottom") = {16};
Physical Surface("front") = {14};
Physical Surface("back") = {18};
Physical Surface("left") = {24};
Physical Surface("right") = {22};
Physical Volume("0") = {26};
```

This mesh geometry file can be used to generate a mesh using Gmsh, either through its graphical user interface, or alternatively, on the commandline

```
gmsht -3 Ra5000.geo
```

which results in a mesh file with approximately 8.4 million tetrahedral elements.

9.2.1 Input file

The complete input file for this problem is

```
# Density-driven convective mixing in a 3D model using the
  streamfunction
# formulation for Ra = 5000
#
# Uses an unstructured mesh generated by Gmsh
# To generate the mesh, run 'gmsht -3 Ra5000.geo'
#
# Note: this mesh has approximately 8.4 million elements and takes about
# 8 hours to run using 200 processors

[Mesh]
  type = FileMesh
  file = Ra5000.e
[]

[Numbat]
  [./Dimensionless]
  [../]
[]

[ICs]
  [./concentration]
    type = NumbatPerturbationIC
    variable = concentration
    amplitude = 0.1
    seed = 1
  [../]
  [./streamfunctionx]
    type = ConstantIC
    variable = streamfunction_x
    value = 0.0
  [../]
  [./streamfunctiony]
    type = ConstantIC
    variable = streamfunction_y
    value = 0.0
```



```

[../]
[]

[Functions]
  [./timesteps]
    type = PiecewiseConstant
    x = '0 10 500 1e3 1e4 6e4'
    y = '9 10 50 100 200 200'
  [../]
[]

[Executioner]
  type = Transient
  end_time = 6e4
  start_time = 1
  solve_type = NEWTON
  petsc_options = -snes_ksp_ew
  petsc_options_iname = '-ksp_type -pc_type -pc_asm_overlap -sub_pc_type
    -pc_factor_levels -ksp_atol'
  petsc_options_value = 'gmres asm 10 ilu 4 1e-12'
  nl_abs_tol = 1e-9
  l_max_its = 200
  [./TimeStepper]
    type = FunctionDT
    function = timesteps
  [../]
[]

[Preconditioning]
  [./smp]
    type = SMP
    full = true
  [../]
[]

[Outputs]
  perf_graph = true
  [./exodus]
    type = Exodus
    execute_on = 'INITIAL TIMESTEP_END'
    interval = 4
  [../]
  [./csvoutput]
    type = CSV
    execute_on = 'INITIAL TIMESTEP_END'
  [../]
  [./checkpoint]
    type = Checkpoint
    num_files = 2
    interval = 10
  [../]
[]

```

An example of the evolution of the convective fingers in this model are presented in Figure 9.3 and Figure 9.4 for an isotropic model. The concentration profile just after the onset of convection is shown in Figure 9.3 for a dimensionless time $t = 1400$. As the isosurface shows,

there are a large number of small fingers at this stage. As time increases, these small structures merge, forming larger fingers in a process that continues as time proceeds, until only a few large fingers are present, see Figure 9.4. This merging behaviour is very complicated and difficult to characterise in any quantitative manner

Many interesting observations can be made from large-scale 3D models. For example, the temporal evolution of the fingers shown in Figure 9.3 and Figure 9.4 can also be examined through a horizontal slice through the model, see Figure 9.5. In this example, a horizontal slice is taken at a dimensionless distance of 100 from the top surface (where the CO_2 concentration is 1). As the fingers approach this depth, they are initially observed as circular regions of higher concentration, cf Figure 9.5 (a) and (b), where we can see that the fingers have just reached this depth at dimensionless time 1000. As time increases, the complexity of the fingering process can be observed, with merging of adjacent fingers and growth observed. Like Pau *et al.* (2010) and Fu *et al.* (2013) we observe that the fingers arrange themselves into polygonal structures with thin profiles surrounded by large regions of unsaturated fluid, see Figure 9.5 (d), (e) and (f).

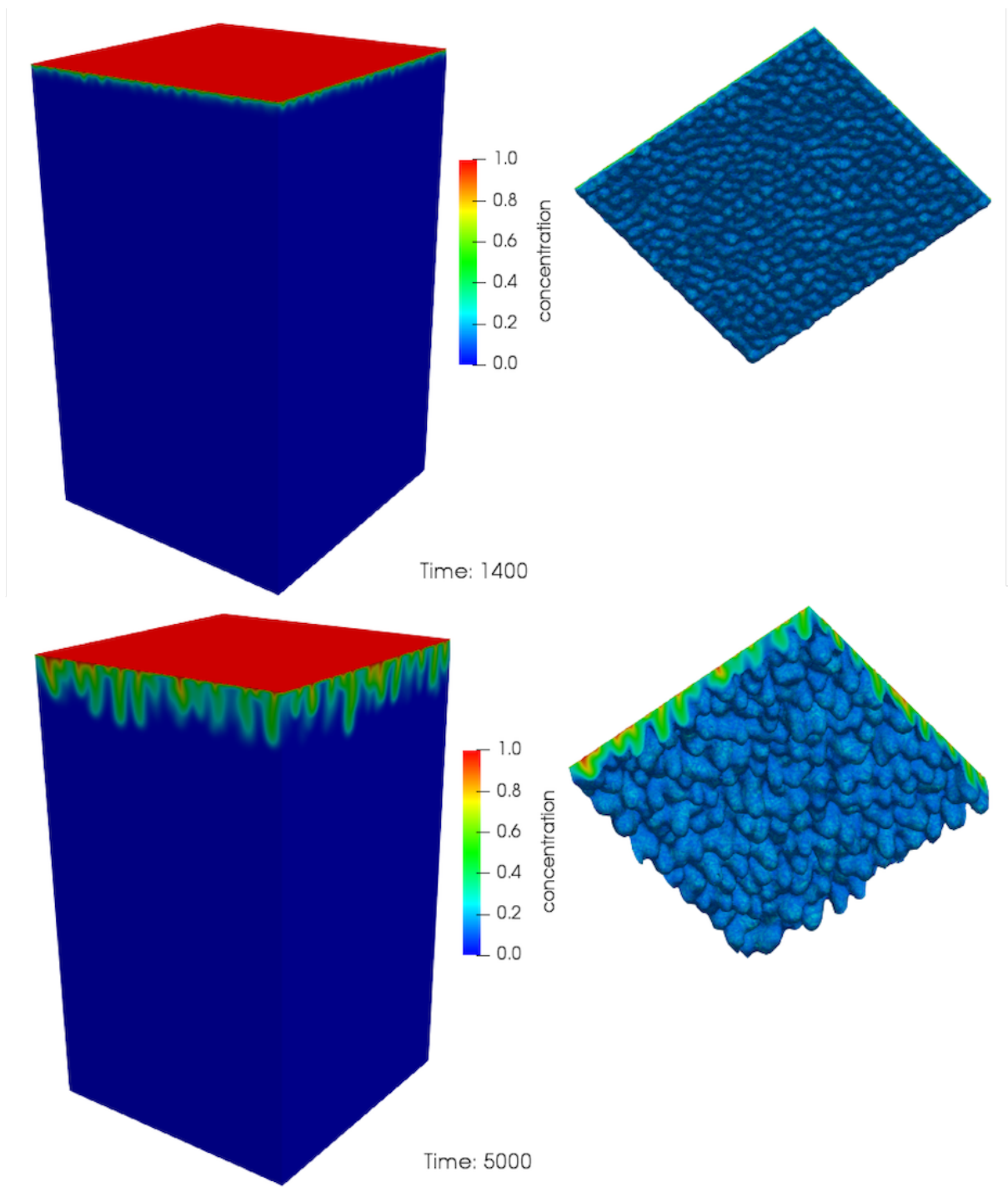


Figure 9.3: Evolution of convective mixing in 3D. Time is dimensionless.

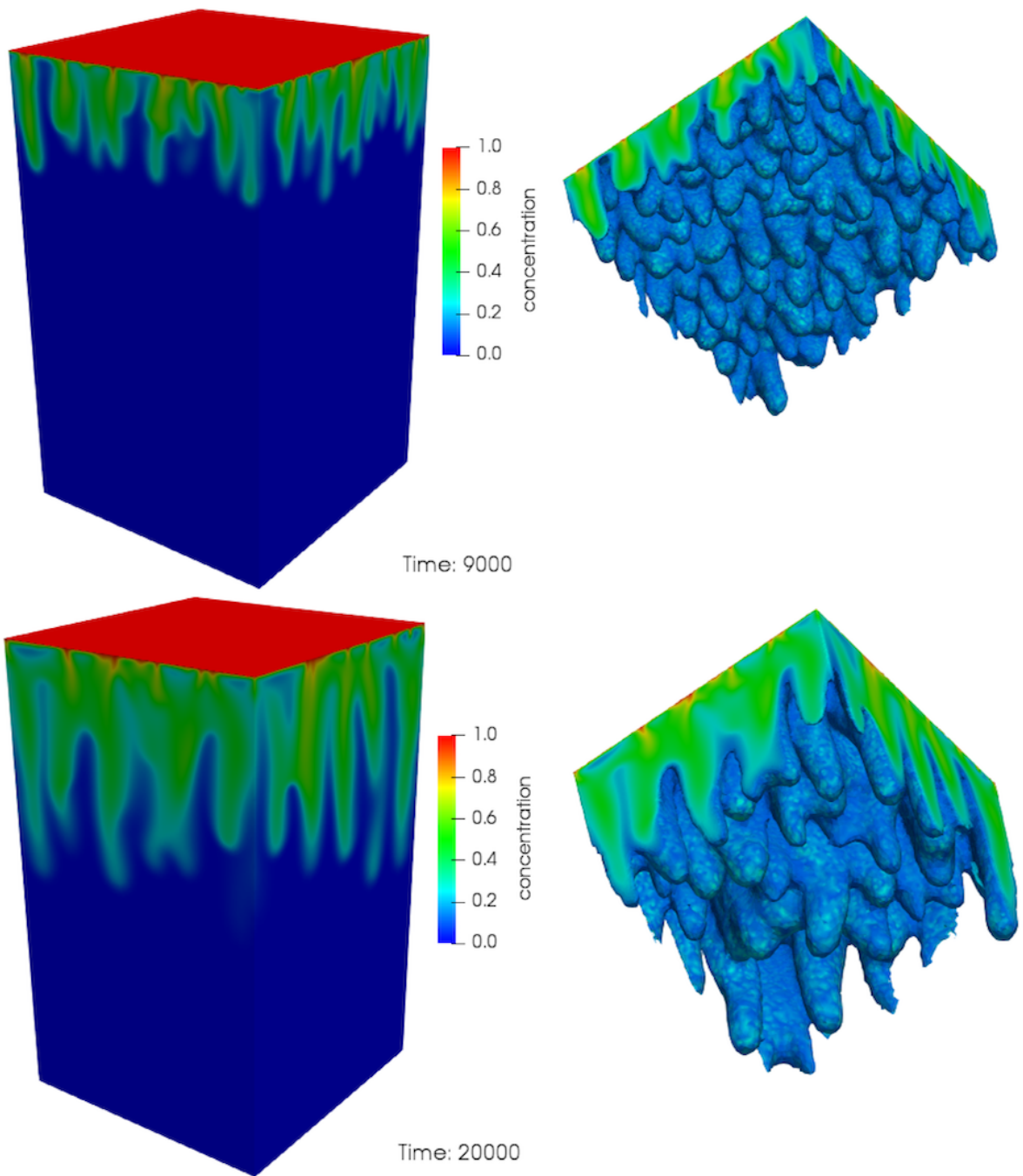


Figure 9.4: Evolution of convective mixing in 3D (continued). Time is dimensionless.

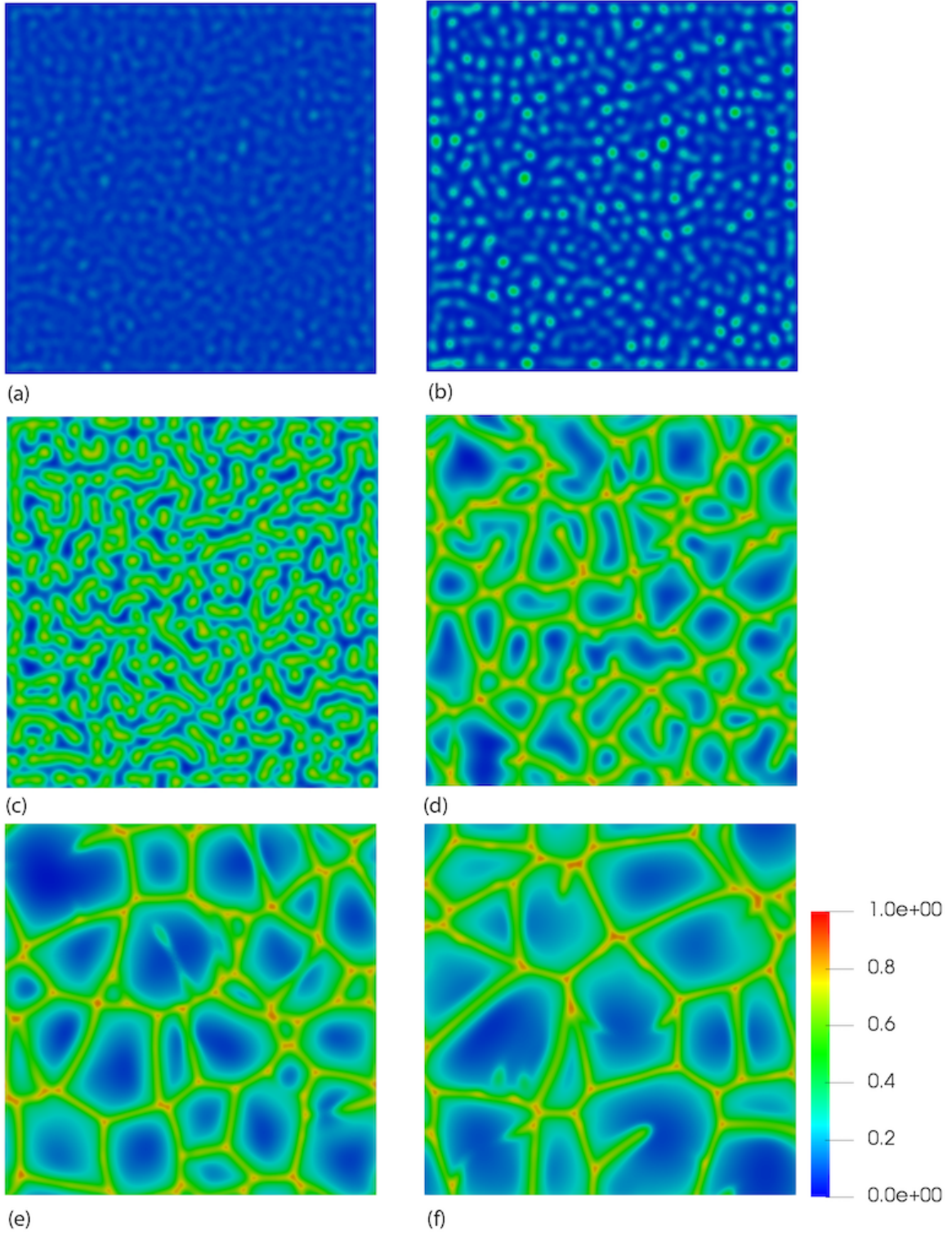


Figure 9.5: Horizontal slice at dimensionless depth 10 showing the evolution of the convective fingers in 3D. Time increasing from (a) to (f).

10 Contributing

Numbat is an open-source application built using the MOOSE framework. It is developed on GitHub, and is written using C++. We welcome new contributions to the code base, which can be submitted using the workflow outlined below.

This guide is based on the MOOSE contributing guide, where you can find lots of information about the code standards and workflow.

10.1 Fork numbat

The first step is to create your own fork of Numbat where you can commit your set of changes.

- Navigate to <https://github.com/cpgr/numbat>
- Click the “Fork” button in the upper right corner
- Clone your fork to your local machine (replace “username” with your GitHub username).

+Note:+ We recommend that you use SSH URLs instead of HTTPS. Generally you will have fewer problems with firewalls and authentication this way. It does however require an additional step of setting up keys. Please follow the instructions provided by Github to setup your SSH keys.

```
git clone git@github.com:username/numbat.git
```

10.2 Add the upstream Remote:

Add the main Numbat repository as a remote named “upstream”:

```
cd moose
git remote add upstream git@github.com:cpgr/numbat.git
```

10.3 Make Modifications

Create a branch for your work:

```
git checkout -b branch_name upstream/master
```

Make your modifications and commit them to a branch (be sure to reference an issue number in your commit messages).

```
git add your_file.h your_file.C
git commit -m "A message about the commit

closes #12345"
```

See `git add` and `git commit` for more assistance on these commands.

Before contributing your changes you should rebase them on top of the current set of patches in the master branch in the main Numbat repository:

```
git fetch upstream
git rebase upstream/master
```

10.4 Push Modifications Back to GitHub

Push your branch back into your fork on GitHub:

```
git push origin branch_name
```

10.5 Create a Pull Request

GitHub uses Pull Requests (PRs) to allow you to submit changes stored in your Fork back to the main Numbat repository. If you are generally interested in how PRs work, see the official GitHub documentation.

References

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CONTACT US

t 1300 363 400
+61 3 9545 2176
e csiroenquiries@csiro.au
w www.csiro.au

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Chris Green
t +61 3 9545 8371
e chris.green@csiro.au
w www.csiro.au