Compute Fluxes of Gradient Fields

We regularly need to compute flux es of the gradients of scalar potential fields.

h = scalar potential

$$\Rightarrow \nabla \times q = 0$$

Discrete approximation: q = - K & h

This works in the interior of the domain, but on boundary Gh is zero by construction.

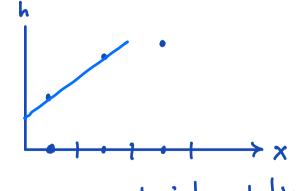
> need to reconstruct boundary flux

Option 1: Extrapolate to bud

Equivalent to using a

one-sided derivative

Problem: lose discrete



conservation because error in interpolation

Option 2: Reconstruct from discrete balance ldea: Use the discrete balance in the bud cell to compute the exact bud flux required for conservation.

Consider a discrete linear system

Lu = fs

u = unknown

Discrete residual of equation

 $\mathbb{L}(\vec{n}) = \vec{F}\vec{n} - \vec{t}^2$

If the discrete equations are satisfied $\Gamma = 0$. In the bud cells $\Gamma \neq 0$ because G arbitrarily sets the gradient/flux to zero.

⇒ non-zero residual in the bond eells contains information about the boundary flux ?

Consider a system if flux boundary

Lu = fs + fu

with residual: $\Gamma = \underline{L}u - \underline{f}s = \underline{f}u$ $\Rightarrow \Gamma = \underline{f}u$

The residual on bnd is equal to the i.h.s. vector due to the boundary fluxes?

Entries of f_{b} on bud are: $f_{n} = q_{b} \frac{A}{V}$ If we are given $\Gamma = f_{n}$ we can reverse this argument and solve for flux: $q_{b} = f_{n} \frac{V}{A} = r \frac{V}{A}$

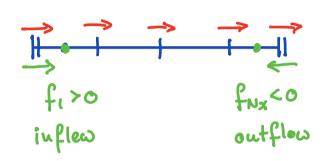
This also works on Dirichlet bud so that the boundary flux is generally given by:

||a|| = |r| \frac{1}{A} note: up to a sign ?

Note: We assume only 1 face of bud cell has flux!

Sign change:

We want q_b to have sign that fits with the rest of fluxes computed as $q = -K\nabla h$. There q's are positive if they point in x-dir.



Need to change sign ou xmax bud.

Implementation

In function compeflux-res.m we will compute boundary fluxes as follows:

Define two vectors:

dof-cell: column vector containing all bnol cells

dof-face: column vector containing all associated bud faces

These vectors are same length because we assume only one face is associated with each bud cell.

Compute all bud fluxes togeather in one line

q(dof-face) = sign. * [(dof-cell, u). * V(dof-cells)./A(dof-face)

You can use ismember. in to detect bind.