

# Unstructured FVM Mesh Format Definition

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Passed to nearly all FVM functions

Defined as follows:

```
mesh =
[
  cells,          #(list of lists, each containing the indices of the faces that
make up the cell)
  cellVolumes,    #(list of scalar cell volumes)
  cellCenters,    #(list of position vectors of cell centers)
  faces,          #(list of lists, each sublist containing two cell indices: the
owner cell and the neighbour cell)
  fAVectors,      #(list of face area vectors)
  fCenters,       #(list of position vectors of face centers)
  boundaryFaces   #(list of lists, each containing the indices of faces on the
ith boundary)
]
```

## Notes

- Cells and faces are numbered according to their storage location in the mesh arrays
- Faces are numbered such that boundary faces come last
- Face area vector point outward from the owner cells, into the neighbour cell
- Cells must be convex, composed of planar faces
- "List" above, in the context of Julia, means a 1-D Array

OpenFOAM meshes can be parsed into the format above using the "OpenFOAMMesh" function in "mesh.jl". The current parse is slightly less flexible than OpenFOAM in terms of formatting, so if problems occur, try running the OpenFOAM utility "renumberMesh -overwrite" to ensure the mesh format is exactly as expected for mesh.jl.

## Solution State Definition

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Passed around FVM functions, intended to contain all universally-applicable info for FVM computations. Examples offered are all for 1D. In a multidimensional computation, the states/residuals include an additional momentum term for each additional dimension, and the cell/face flux matrices will include one flux per coordinate direction per conserved variable. For multidimensional computations, fluxes are ordered first by flux type, then by flux direction (ex: x-dirMassFlux, y-dirMassFlux, z-dirMassFlux, x-dirXMomentumFlux, y-dirXMomentumFlux, etc...)

```
solutionState =
[
  cellState,      # rho, rhoU, eV2 (conserved variables) at cell centers
  cellFluxes,     # massFluxes, momentemFluxes, energyFluxes at cell centers
```

```

    cellPrimitives,    # P, T, U at cell centers
    fluxResiduals,     # d/dt of each "cellState" variable (flux balances for each
cell)
    faceFluxes         # massFluxes, momentemFluxes, and energyFluxes at face
centers
]

```

Where

### Cell State definition

```

# CellState =
# Cell      rho  x-Momentum  Total Energy
# Cell 1    rho_1  xM_1      eV2_1
# Cell 2    rho_2  xM_2      eV2_2
# ...

```

### Cell Fluxes definition

```

# CellFluxes =
# Cell      x_dirMassFlux  x-dir_x-MomentumFlux  x-dir_TotalEnergyFlux
# Cell 1    rhoU_1        (rho*U^2 + P)_1      (eV2*U + P*U)_1
# Cell 2    rhoU_2        (rho*U^2 + P)_2      (eV2*U + P*U)_2
# ...

```

### Cell Primitives definition

```

# CellPrimitives =
# Cell      P      T      Ux
# Cell 1    P_1    T_1    Ux_1
# Cell 2    P_2    T_2    Ux_2
# ...

```

### Flux Residuals definition

Note that "residual" is taken to mean flux balance, which is only strictly true in a steady-state computations. Term used loosely here.

```

# fluxResiduals =
# Cell      rho      x-Momentum  Total Energy
# Cell 1    d(rho_1)/dt  d(xM_1)dt  d(eV2_1)/dt
# Cell 2    d(rho_2)/dt  d(xM_2)dt  d(eV2_2)/dt
# ...

```

## Face Fluxes definition

```
# faceFluxes =  
# Face      rho      x-Momentum      Total Energy  
# Face 1    rhoU_f1   (rho*U^2 + P)_f1   (eV2*U + P*U)_f1  
# Face 2    rhoU_f2   (rho*U^2 + P)_f2   (eV2*U + P*U)_f2  
# ...
```