

# Pseudo-code for CG routines and algorithms

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# 1 Time-stepping pseudo code

## 1.1 DomainSolver::advanceAdamsPredictorCorrector

Here is an overview of the DomainSolver::advanceAdamsPredictorCorrector function (cg/common/src/advancePC.bC) This is an explicit Adams predictor-corrector time stepper.

```
DomainSolver::advanceAdamsPredictorCorrector( ..., numberOfSubSteps, ..)
{
    initialize
    for( int mst=1; mst<=numberOfSubSteps; mst++ ) take time steps
        if adapt grids
            adaptGrids( ... )
        if move grids
            moveGrids( ... ); (Sec. 2.1)
            exposedPoints.interpolate(...)

        predictor step:
        getUt( ... )
        interpolateAndApplyBoundaryConditions( ... );
        solveForTimeIndependentVariables( ... )
        correctMovingGrids( ... )

    for( int correction=0; correction<numberOfCorrections; correction++ )
        corrector step:
        getUt( ... )
        solveForTimeIndependentVariables( ... )
        interpolateAndApplyBoundaryConditions( ... );
        correctMovingGrids( ... )
}
```

Figure 1: Pseudo-code outline of the advanceAdamsPredictorCorrector function.

## 2 Moving grid pseudo code

### 2.1 DomainSolver::moveGrids

Pseudo-code for DomainSolver::moveGrids (cg/common/src/move.C)

```
DomainSolver::moveGrids( t1,t2,t3,dt0, cgf1,cgf2,cgf3 )
{
    setInterfacesAtPastTimes( t1,t2,t3,dt0,cgf1,cgf2,cgf3 ); initialize interfaces
    parameters.dbase.get<MovingGrids>("movingGrids").moveGrids(t1,t2,t3,dt0,cgf1,cgf2,cgf3 );
    gridGenerator->updateOverlap( cg, mapInfo ); regenerate the grid with Ogen
}
```

### 2.2 MovingGrids::moveGrids

Pseudo-code for MovingGrids::moveGrids (cg/common/moving/src/MovingGrids.C)

```
MovingGrids::moveGrids( t1,t2,t3,dt0, cgf1,cgf2,cgf3 )
{
    First move the bodies (but not the grids):
    detectCollisions(cgf1);
    rigidBodyMotion( t1,t2,t3,dt0,cgf1,cgf2,cgf3 );
    moveDeformingBodies( t1,t2,t3,dt0,cgf1,cgf2,cgf3 );
    userDefinedMotion( t1,t2,t3,dt0,cgf1,cgf2,cgf3 );
    Apply any matrix motions: rotate, shift, scale

    getGridVelocity( cgf2,t2 );
    Now move the grids:
    for( grid=0; grid<numberOfBaseGrids; grid++ )
        MatrixTransform & transform = *cgf3.transform[grid];
        if( moveOption(grid)==matrixMotion )
            apply specified rotation and/or shift
            transform.rotate(...)
        else if( moveOption(grid)==rigidBody )
            rotate and shift the rigid body
            transform.rotate(...)
    for( int b=0; b<numberOfDeformingBodies; b++ )
        deformingBodyList[b]->regenerateComponentGrids( newT, cgf3.cg );

    getGridVelocity( cgf3,t3 );
}
```

### 2.3 MovingGrids::moveDeformingBodies

Pseudo-code for MovingGrids::moveDeformingBodies (cg/common/moving/src/MovingGrids.C)

```
MovingGrids::moveDeformingBodies( t1,t2,t3,dt0, cgf1,cgf2,cgf3 )
{
    for( int b=0; b<numberOfDeformingBodies; b++ )
        deformingBodyList[b]->integrate( t1,t2,t3,cgf1,cgf2,cgf3, stress);
}
```

### 3 DeformingBodyMotion pseudo code

The DeformingBodyMotion class handles deforming bodies.

#### 3.1 DeformingBodyMotion::integrate

Pseudo-code for DeformingBodyMotion::integrate (cg/common/moving/src/DeformingBodyMotion.C)  
This function is called by MovingGrids::movingGrids to move the deforming body (but not the grid associated with the deforming body).

```
MovingGrids::integrate( t1,t2,t3,dt0, cgf1,cgf2,cgf3, stress )
{
    if( elasticShell )
        advanceElasticShell(t1,t2,t3,cgf1,cgf2,cgf3,stress,option);
    else if( ... )

    for( int face=0; face<numberOfFaces; face++ )
        if( ... )
        else if( userDefinedDeformingBodyMotionOption==interfaceDeform )
            The deformed surface is obtained from the boundaryData array:
            RealArray & bd = parameters.getBoundaryData(side,axis,grid,cg[grid]);
            x0 = bd;
}
```

#### 3.2 DeformingBodyMotion::regenerateComponentGrids

Pseudo-code for DeformingBodyMotion::regenerateComponentGrids (cg/common/moving/src/DeformingBodyMotion.C)  
This function is called by MovingGrids::movingGrids (after calling DeformingBodyMotion::integrate) to actually generate the grid associated with the deforming body.

```
DeformingBodyMotion::regenerateComponentGrids( const real newT, CompositeGrid & cg)
{
    for( int face=0; face<numberOfFaces; face++ )
        hyp.generate(); Call the hyperbolic grid generator.
        Save the grid in the GridEvolution list:
        gridEvolution[face]->addGrid(dpm.getDataPoints(),newT);
}
```

#### 3.3 DeformingBodyMotion::correct

Pseudo-code for DeformingBodyMotion::correct (cg/common/moving/src/DeformingBodyMotion.C)  
This function is called by MovingGrids::correctGrids.

```
DeformingBodyMotion::correct( t1, t2, GridFunction & cgf1,GridFunction & cgf2 )
{
    This function currently does nothing.
}
```

## 4 Cgmp

### 4.1 Driver code 'main' for Cgmp

Pseudo-code for the main driver code for Cgmp (found in cg/mp/src/cgmpMain.C) This function reads in the overset grid, sets up the problem, solves the problem and finishes.

```
main( int argc, char *argv[] )
{
    Overture::start(argc,argv); // initialize Overture and A++/P++
    // Read command line arguments ...
    GenericGraphicsInterface & ps = *Overture::getGraphicsInterface("cgmp",false,argc,argv);

    CompositeGrid cg; // Object to hold the master overset grid for all domains.
    nameOfGridFile = readOrBuildTheGrid(ps, cg, loadBalance, ...); // Get the overset grid.

    Cgmp & mpSolver = *new Cgmp(cg,&ps,show,plotOption);

    mpSolver.setParametersInteractively(); // Setup the problem and PDE solvers.

    mpSolver.solve(); // Time-step the PDE to completion.

    mpSolver.printStatistics();
    Overture::finish();
}
```

### 4.2 Cgmp::setParametersInteractively

Pseudo-code for Cgmp::setParametersInteractively (cg/mp/src/setParametersInteractively.C) This function reads commands to setup the parameters for each DomainSolver (Cgad, Cgins, Cgsm, Cgcns,...). It initializes the list of interfaces and then request Cgmp run-time parameters.

```
Cgmp::setParametersInteractively( bool callSetup )
{
    while(true)
        gi.getAnswer("");
        if( answer.matches("setup" ) ) // Look for command: setup 'domainName'
            setupDomainSolverParameters( domain,modelNames ); // Setup a domain.
        end
    end

    initializeInterfaces(gfIndex); // Create list of interfaces.

    DomainSolver::setParametersInteractively(callSetup); // Get Cgmp run-time parameters.
}
```

### 4.3 Cgmp::setupDomainSolverParameters

Pseudo-code for Cgmp::setupDomainSolverParameters (cg/mp/src/setParametersInteractively.C) This function builds the PDE solvers for the different domains and sets all the run-time parameters (e.g. coefficient of thermal conductivity, coefficient of viscosity etc.) for each domain solver.

```
Cgmp::setupDomainSolverParameters( int domain, vector<aString> & modelNames )
// modelNames (input): list of available models (PDE solvers such as Cgins, Cgad, Cgcns, Cgsm)
{
    while(true)
```

```

gi.getAnswer("");
if( answer.matches("set solver") ) // Look for command: set solver 'solverType'

    // Construct the PDE solver for a given domain:
    domainSolver[domain] = buildModel( solverType, cg.domain[domain],...);

    else if( answer.matches("solver parameters") )
        domainSolver[domain]->setParametersInteractively(false); // Set run-time parameters for
one domain.
    end
end
}

```

#### 4.4 Cgmp::buildModel

Pseudo-code for Cgmp::buildModel (cg/mp/src/cgmpMain.C) This function builds a particular PDE solver such as Cgad, Cgins, Cgcns, etc.

```

DomainSolver* Cgmp::buildModel( const aString & modelName, CompositeGrid & cg, ... )
// modelNames (input): list of available models (PDE solvers such as Cgins, Cgad, Cgcns, Cgsm)
{
    DomainSolver *solver=NULL;
    if( modelName == "Cgins" )
        solver = new Cgins(cg,ps,show,plotOption);
    else if( modelName == "Cgcns" )
        solver = new Cgcns(cg,ps,show,plotOption);
    else if( modelName == "Cgad" )
        solver = new Cgad(cg,ps,show,plotOption);
    else if( modelName == "Cgsm" )
        solver = new Cgsm(cg,ps,show,plotOption);
    ...
    end
    solver->parameters.dbase.get<DomainSolver*>("multiDomainSolver")=this;
    return solver;
}

```

#### 4.5 Cgmp::initializeInterfaces

Pseudo-code for Cgmp::initializeInterfaces (cg/mp/src/assignInterfaceBoundaryConditions.C) This function locates interfaces (by matching grid faces on different domains) and builds a list of information about the interfaces. The values of interfaceType(side,dir,axis) can currently be noInterface, heatFluxInterface, or tractionInterface. These are set when assigning domain boundary conditions.

```

Cgmp::initializeInterfaces( std::vector<int> & gfIndex )
{
    ForDomain( d1 )
        interfaceType1 = domainSolver[d1]->parameters.dbase.get<IntegerArray>("interfaceType");
        for( grid1, dir1,side1 )
            if( interfaceType1(side1,dir1,grid1) != Parameters::noInterface ) //T his face is on an
interface
                if( This face matches an existing face of an interface )
                    GridList & gridList = interfaceDescriptor.gridListSide1
OR interfaceDescriptor.gridListSide2;
                    // Create a new (matched) face on the interface:
                    gridList.push_back(GridFaceDescriptor(d1,grid1,side1,dir1));

```

```

else
    interfaceList.push_back(InterfaceDescriptor()); // Add a new interface to the list.
    InterfaceDescriptor & interface = interfaceList.back();
    // Create a new (unmatched) face on the interface:
    interface.gridListSide1.push_back(GridFaceDescriptor(d1,grid1,side1,dir1));
end
end
end
end
}

```

## 4.6 Cgmp::solve

Pseudo-code for Cgmp::solve (cg/mp/src/solve.C)

```

Cgmp::solve()
{
    cycleZero(); // Call domain solvers before time-steps start.
    buildRunTimeDialog();

    for( int step=0; step<maximumNumberOfSteps && !finish; )

        if( t ≥ nextTimeToPrint )
            printTimeStepInfo(step,t,cpuTime);
            saveShow( gf[current] );
            finish=plot(t, optionIn, tFinal);
            if( finish ) break;
        end

        dtNew = getTimeStep( gf[current] ); // choose time step
        computeNumberOfStepsAndAdjustTheTimeStep(t,tFinal,nextTimeToPrint,numberOfSubSteps,dtNew);

        advance(tFinal); // advance to t=nextTimeToPrint

    end
}

```

## 4.7 Cgmp::multiDomainAdvance

Pseudo-code for Cgmp::multiDomainAdvance (cg/mp/src/multiDomainAdvance.C) This function may call multiDomainAdvanceNew or multiStageAdvance, depending on the options.

```
Cgmp::multiDomainAdvance( real & t, real & tFinal )
{
    if( multiDomainAlgorithm==MpParameters::stepAllThenMatchMultiDomainAlgorithm )
        // This new algorithm supports AMR:
        return multiDomainAdvanceNew(t,tFinal);
    else if( multiDomainAlgorithm==MpParameters::multiStageAlgorithm )
        // User-defined multi-stage algorithm:
        return multiStageAdvance(t,tFinal);
    end

    if( initialize )
        initializeInterfaceBoundaryConditions( t,dt,gfIndex );
        ForDomain( d ) assignInterfaceRightHandSide( d, t, dt, correct, gfIndex );
        ForDomain( d ) domainSolver[d]->initializeTimeStepping( t,dt );
    end

    // Take some time steps:
    for( int i=0; i<numberOfSubSteps; i++ )
        ForDomain( d )
            domainSolver[d]->startTimeStep( t,dt,... );
            numberOfRequiredCorrectorSteps=...; gridHasChanged=...;
        end
        if( gridHasChanged )
            initializeInterfaces(gfIndex); initializeInterfaceBoundaryConditions(...);
        end

        for( int correct=0; correct<=numberOfCorrectorSteps; correct++ )
            ForDomain( d )
                assignInterfaceRightHandSide( d, t+dt, dt, correct, gfIndex ); // (Sec. 5.2)
                domainSolver[d]->takeTimeStep( t,dt,correct,advanceOptions[d] );
            end
            if( hasConverged = checkInterfaceForConvergence( .. ) ) break;
        end

        ForDomain( d )
            domainSolver[d]->endTimeStep( td,dt,advanceOptions[d] );
        end
        t+=dt;
    end
}
```



## 4.8 Cgmp::multiDomainAdvanceNew

Pseudo-code for Cgmp::multiDomainAdvanceNew (cg/mp/src/multiDomainAdvanceNew.bC). This is the new version of the multi-domain advance routine that supports more general time stepping and the use of AMR.

```
Cgmp::multiDomainAdvanceNew( real &t, real &tFinal )
{
    if( initialize )
        initializeInterfaceBoundaryConditions( t,dt,gfIndex );
    ForDomain( d ) assignInterfaceRightHandSide( d, t, dt, correct, gfIndex );
    ForDomain( d ) domainSolver[d]->initializeTimeStepping( t,dt );
    // Take some time steps:
    for( int i=0; i<numberOfSubSteps; i++ )
        ForDomain( d )
            domainSolver[d]->startTimeStep( t,dt,... );
            numberOfRequiredCorrectorSteps=...; gridHasChanged=...;
        if( gridHasChanged )
            initializeInterfaces(gfIndex); initializeInterfaceBoundaryConditions(...);
        // Get current interface residual and save current interface values :
        getInterfaceResiduals( t, dt, gfIndex, maxResidual, saveInterfaceTimeHistoryValues );

        // Stage I: advance the solution but do not apply BC's:
        ForDomain( d )
            assignInterfaceRightHandSide( d, t+dt, dt, correct, gfIndex );
            domainSolver[d]->takeTimeStep( t,dt,correct,step but no BC's );

        // Stage II: Project interface values part 1:
        interfaceProjection( t+dt, dt, correct, gfIndex,set interface values );

        // Stage III: evaluate the interface conditions and apply the boundary conditions:
        ForDomain( d )
            assignInterfaceRightHandSide( d, t+dt, dt, correct, gfIndex ); //(Sec. 5.2)
            domainSolver[d]->takeTimeStep( t,dt,correct,apply BC's );

        // Stage IV: Project interface values part 2:
        interfaceProjection( t+dt, dt, correct, gfIndex,set interface ghost values );

        if( hasConverged = checkInterfaceForConvergence( .. ) ) break;

    ForDomain( d ) domainSolver[d]->endTimeStep( t,dt,advanceOptions[d] );
    t+=dt;
}
```

## 4.9 Cgmp::multiStageAdvance

Pseudo-code for `Cgmp::multiStageAdvance` (cg/mp/src/multiStageAdvance.bC). This is yet a newer version of the multi-domain advance algorithm. This version was developed to deal with the FSI-AMP schemes involving incompressible fluids and elastic bodies. It uses `interfaceCommunicationMode==Parameters::requestInterfaceDataWhenNeeded` so that domain solvers request interface data when they need it. Each multi-domain time-step is separated into **stages**. Associated with each stage are a subset of domains and a set of operations. For example,

**Stage 1** : Time-step Cgins domains but do not apply boundary conditions.

**Stage 2** : Time-step and apply BC's to Cgsm domains.

**Stage 3** : Apply BC's to Cgins domains.

```
Cgmp::multiStageAdvance( real & t, real & tFinal )
{
  if( initialize )
    initializeInterfaceBoundaryConditions( t,dt,gfIndex ); // Choose heatFlux conditions.
    ForDomain( d ) domainSolver[d]->initializeTimeStepping( t,dt );
  end

  // Take some time steps:
  for( int i=0; i<numberOfSubSteps; i++ )
    ForDomain( d )
      // Start the time-step and return required number of correction steps.
      domainSolver[d]->startTimeStep( t,dt,gfIndexCurrent[d],gfIndexNext[d],advanceOptions[d] );
    end

    // Each time-step consists of a predictor and zero or more correction steps.
    for( int correct=0; correct<=numberOfCorrectorSteps; correct++ )

      for( int stage=0; stage<numberOfStages; stage++ ) // Execute each stage
        StageInfo & stageInfo = stageInfoList[stage]; // Holds info on this stage.
        for( domains d involved in this stage )
          // stageInfo.action : takeStep and/or applyBoundaryConditions
          advanceOptions[d].takeTimeStepOption=stageInfo.action;

          domainSolver[d]->takeTimeStep( t,dt,correct,advanceOptions[d] );

        end
      end
      // Check if the corrections have completed.
      if( relaxCorrectionSteps && correctionIterationsHaveConverged &&
          correct>=minimumNumberOfCorrections ) break;
    end

    ForDomain( d )
      domainSolver[d]->endTimeStep( t,dt,advanceOptions[d] );
    end
    t+=dt;
  end
}
```

## 4.10 Cgmp::checkInterfaceForConvergence

Pseudo-code for `Cgmp::checkInterfaceForConvergence` (in `cg/mp/src/multiDomainAdvance.C`) This function checks the residual in the interface equations for convergence of the sub time-step iterations.

```
bool Cgmp::checkInterfaceForConvergence( int correct, int numberOfCorrectorSteps, ... )
{
    if( check residuals for convergence )

        // Evaluate the max residuals in the conditions at each interface
        // NOTE: the history of interface iterates are saved here:
        getInterfaceResiduals( tNew, dt, gfIndex, maxResidual, saveInterfaceIterateValues );

        // check if the interface iterations have converged:
        interfaceIterationsHaveConverged=true;
        for( int inter=0; inter<interfaceList.size(); inter++ )
            if( correct==0 )
                initialResidual[inter]=maxResidual[inter]; // Save initial residual
            else if( correct==1 )
                firstResidual[inter]=maxResidual[inter]; // Save first residual
            end
                interfaceIterationsHaveConverged = interfaceIterationsHaveConverged &&
                    maxResidual[inter] < interfaceList[inter].interfaceTolerance;
            if( !interfaceIterationsHaveConverged ) break;
        end

        if( interfaceIterationsHaveConverged && correct >= numberOfRequiredCorrectorSteps )
            Save statistics about interface iterations ...
            return true; Iterations have completed.
        else
            return false;
        end

    end
}
```

## 4.11 Cgmp::getInterfaceResiduals

Pseudo-code for `Cgmp::getInterfaceResiduals` (in `cg/mp/src/assignInterfaceBoundaryConditions.C`)  
This function evaluates the residual in the jump conditions at each interface.

- This function should be cleaned up.
- The interface residuals can probably be more efficiently computed within the domain solvers.

```
int Cgmp::getInterfaceResiduals( real t, real dt, vector<int> & gfIndex, vector<real> &
maxResidual,
    InterfaceValueEnum saveInterfaceValues =doNotSaveInterfaceValues )
{
    for( int inter=0; inter < interfaceList.size(); inter++ ) // Loop over interfaces.
        InterfaceDescriptor & interfaceDescriptor = interfaceList[inter];
        for( int interfaceSide=0; interfaceSide<=1; interfaceSide++ ) // two sides
            for( int face=0; face<gridList.size(); face++ ) // faces on this interface
                if( interfaceType==Parameters::heatFluxInterface )
                    // Mixed BC is a0*T + a1*T.n
                    info.a[0]=1.; info.a[1]=0.; // eval T
                    domainSolver[domain]->interfaceRightHandSide( get ,interfaceDataOptions,info,... );
                    info.a[0]=0.; info.a[1]=ktc; // eval k*T.n
                    domainSolver[domain]->interfaceRightHandSide( get,interfaceDataOptions,info, ...);
                else if( interfaceType==Parameters::tractionInterface )
                    FINISH ME..
                end

                if( saveInterfaceValues==saveInterfaceTimeHistoryValues ||
                    saveInterfaceValues==saveInterfaceIterateValues )
                    Save time-history or sub-time-step iterations.
                end

            end// end for face
        end
    end
    // ---- Transfer data to the opposite side of the interface ----
    // ---- and evaluate the jump conditions ----
    for( int interfaceSide=0; interfaceSide<=1; interfaceSide++ )

        interfaceTransfer.transferData( domainSource, domainTarget, ... )

        if( interfaceType==Parameters::heatFluxInterface )
            // Compute the maximum error in [T] and [KTn]
        end
    end
}
```

## 5 Interfaces

### 5.1 Cgmp::initializeInterfaceBoundaryConditions

Pseudo-code for `Cgmp::initializeInterfaceBoundaryConditions` (`cg/mp/src/assignInterfaceBoundaryConditions.C`). This function determines how the boundary conditions on a heat-flux interface should be assigned. For example, for a partitioned Dirichlet-Neumann (DN) approach, which sides is Dirichlet/Neumann depends on the material parameters  $\mathcal{K}$  and  $\mathcal{D}$ .

```
Cgmp::initializeInterfaceBoundaryConditions( real t, real dt, std::vector<int> & gfIndex )
{
    InterfaceList & interfaceList = parameters.dbase.get<InterfaceList>("interfaceList");
    if( interfaceList.size()==0 )
        initializeInterfaces(gfIndex);
    end

    for( int inter=0; inter < interfaceList.size(); inter++ )
        InterfaceDescriptor & interfaceDescriptor = interfaceList[inter];
        for( face ) // Loop over faces on this interface
            if( interfaceType1(side1,dir1,grid1)==Parameters::heatFluxInterface )
                 $K_r \stackrel{\text{def}}{=} (\mathcal{K}_1/\mathcal{K}_2)\sqrt{\mathcal{D}_2/\mathcal{D}_1}$ ; // See discussion in [?]
                if( solveCoupledEquatons ) // Explicit time-stepping
                    // Apply Neumann BC on both sides when solving the coupled interface equations.
                else if( useMixedInterfaceConditions )
                    // Choose mixed BC coefficients ...
                else if(  $K_r > 1$  ) // For partitioned DN approach
                    // Domain 1 is Neumann, domain 2 is Dirichlet.
                    gridDescriptor1.interfaceBC=neumannInterface;
                    gridDescriptor1.a[0]=0.; gridDescriptor1.a[1]= $\mathcal{K}_1$ ;
                    gridDescriptor2.interfaceBC=dirichletInterface;
                    gridDescriptor2.a[0]=1.; gridDescriptor2.a[1]=0.;
                else
                    // Domain 1 is Dirichlet, domain 2 is Neumann.
                    gridDescriptor1.interfaceBC=dirichletInterface;
                    gridDescriptor1.a[0]=1.; gridDescriptor1.a[1]=0.;
                    gridDescriptor2.interfaceBC=neumannInterface;
                    gridDescriptor2.a[0]=0.; gridDescriptor2.a[1]= $\mathcal{K}_2$ ;
                end
                // Save the info about the interface condition in the domain solver: (e.g. in bcData)
                domainSolver[d1]->setInterfaceBoundaryCondition( gridDescriptor1 );
                domainSolver[d2]->setInterfaceBoundaryCondition( gridDescriptor2 );
            end
        end
    end
}
```

### 5.2 Cgmp::assignInterfaceRightHandSide

The `Cgmp::assignInterfaceRightHandSide` function is used to get interface values from a source domain and set interface values on a target domain. It is used in the `Cgmp::multiDomainAdvance` routine 4.7.

Here is `Cgmp::assignInterfaceRightHandSide` (`cg/mp/src/assignInterfaceBoundaryConditions.C`)

```
int Cgmp::assignInterfaceRightHandSide( int d, real t, real dt, int correct, std::vector<int> &
gfIndex )
// d : target domain, assign the interface RHS for this domain.
{
```

```

if( interfaceList.size()==0 )
    initializeInterfaces(gfIndex); // Create the list of interfaces.

for( each interface on domain d )
    InterfaceDescriptor & interfaceDescriptor = interfaceList[inter];
    // Target and source grid functions:
    GridFunction & gfTarget = domainSolver[domainTarget]->gf[gfIndex[domainTarget]];
    GridFunction & gfSource = domainSolver[domainSource]->gf[gfIndex[domainSource]];

    // Get source data:
    for( int face=0; face<gridListSource.size(); face++ )
        domainSolver[domainSource]->interfaceRightHandSide( getInterfaceRightHandSide, ... );

    // Transfer the source arrays to the target arrays:
    interfaceTransfer.transferData(... );

    // Adjust the target data before assigning:
    for( int face=0; face<gridListTarget.size(); face++ )
        // Extrapolate the initial guess.
        // Under-relaxed iteration.

    // Assign the target data:
    for( int face=0; face<gridListTarget.size(); face++ )
        domainSolver[domainTarget]->interfaceRightHandSide( setInterfaceRightHandSide,...);
}

```

### 5.3 DomainSolver::interfaceRightHandSide

The DomainSolver::interfaceRightHandSide function is used to get or set interface values. Each DomainSolver (cgad, cgcns, cgins, cgsm,...) has a version of this routine. The generic version appears in cg/common/src/interfaceBoundaryConditions.C.

Here is Cgcns::interfaceRightHandSide (cg/cns/src/interface.bC)

```
Cgcns::interfaceRightHandSide( InterfaceOptionsEnum option, int interfaceDataOptions,
    GridFaceDescriptor & info, GridFaceDescriptor & gfd, int gfIndex, real t )
{
    RealArray & bd = parameters.getBoundaryData(side,axis,grid,mg); // Interface data on this domain.
    RealArray & f = *info.u; // Interface data from another domain.

    if( interfaceType(side,axis,grid)==Parameters::heatFluxInterface )
        // Set RHS for a heatFlux interface
        if( option==setInterfaceRightHandSide )
            bd(I1,I2,I3,tc)=f(I1,I2,I3,tc); // Set T (using values from another domain).
        else if( option==getInterfaceRightHandSide )
            // Evaluate  $a_0T + a_1T_n$  (to send to another domain).
            f(I1,I2,I3,tc) = a[0]*uLocal(I1,I2,I3,tc) + a[1]*( normal(I1,I2,I2,0)*ux + ... );
        end

    else if( interfaceType(side,axis,grid)==Parameters::tractionInterface )
        // Set RHS for a traction interface.
        if( option==setInterfaceRightHandSide )
            bd(I1,I2,I3,V)=f(I1,I2,I3,V); // Set positions of the interface.
        else if( option==getInterfaceRightHandSide )
            parameters.getNormalForce( gf[gfIndex].u,traction,ipar,rpar );
            f(I1,I2,I3,V)=traction(I1,I2,I3,D);

            // Optionally save a time history of some quantities.
            InterfaceDataHistory & idh = gfd.interfaceDataHistory; // Holds interface history.
            if( interfaceDataOptions & Parameters::tractionRateInterfaceData )
                RealArray & f0 = idh.interfaceDataList[prev].f;
                f(I1,I2,I3,Vt)= (f(I1,I2,I3,V) - f0(I1,I2,I3,V))/dt; // Time derivative of the traction.
            end
        end
    end
}
```