Additional document

Conjugate gradient algorithm to solve the primal interface problem

In practice, the interface problem is never assembled to be solved by a direct solver. The use of an iterative solver is much less expensive and more efficient since it only involves the product of local matrix-vector products which can be done in parallel in a distributed way.

In this document, the application of the conjugate gradient algorithm is proposed in order to solve the primal interface problem :

$$\mathbf{S}_{p}\mathbf{u}_{b} = \mathbf{b}_{p}$$

with:

$$\mathbf{S}_{p} = \mathbb{A}^{\diamondsuit} S_{p}^{\diamondsuit} \mathbb{A}^{\diamondsuit}^{T}$$

$$\mathbf{b}_{p} = \mathbb{A}^{\diamondsuit} b_{p}^{\diamondsuit}$$

1 Conjugate gradient algorithm

The conjugate gradient algorithm to solve a linear system Ax = b is given in **Algorithm 1**.

Algorithm 1 Conjugate gradient

Initialization : Choose x_0 (generally $x_0 = 0$)

- $\Box r_{(0)} = b A x_{(0)}$
- $\Box d_{(0)} = r_{(0)}$

for For i = 0 to m **do**

- 1. Compute optimal step: $\alpha_{(i)} = \frac{r_{(i)}^T r_{(i)}}{d_{(i)}^T A d_{(i)}} \left(= \frac{r_{(i)}^T d_{(i)}}{d_{(i)}^T A d_{(i)}} \right)$
- 2. Compute iterate : $x_{(i+1)} = x_{(i)} + \alpha_{(i)} d_{(i)}$
- 3. Compute residual : $r_{(i+1)} = r_{(i)} \alpha_{(i)} A d_{(i)}$
- 4. Compute orthogonalization parameter : $\beta_{(i+1)} = \frac{r_{(i+1)}^T r_{(i+1)}}{r_{(i)}^T r_{(i)}} \left(= -\frac{r_{(i+1)}^T A d_{(i)}}{d_{(i)}^T A d_{(i)}} \right)$
- 5. Update search direction : $d_{(i+1)} = r_{(i+1)} + \beta_{(i+1)} d_{(i)}$

end for

2 Conjugate gradient algorithm to solve the primal interface problem

By considering **Algorithm 1** to solve a linear system Ax = b, one can see that matrix-vector product Ad is required. When applied to solve the primal interface problem, computing matrix-vector product $\mathbf{S}_p \mathbf{r}_b$ can be done thanks to local (subdomain) matrix-vector products. One recall that bold characters refer to assembled quantities.

Let consider the initialization step of the algorithm. Knowing the initialization $\mathbf{u}_{b(0)}$, one deduces that for each subdomain:

$$\left\{ \begin{array}{ll} \forall \, s \in \mathbf{S}, & \begin{pmatrix} K_{ii}^{(s)} & K_{ib}^{(s)} \\ K_{bi}^{(s)} & K_{bb}^{(s)} \end{pmatrix} \begin{pmatrix} u_{i(0)}^{(s)} \\ u_{b(0)}^{(s)} \end{pmatrix} & = & \begin{pmatrix} f_{i}^{(s)} \\ f_{b}^{(s)} \end{pmatrix} \end{array} \right.$$

where $u_{b(0)}^{\diamondsuit} = \mathbb{A}^{\diamondsuit}^T \mathbf{u}_{b(0)}$ is known. In this case, from the first line of the previous system, one can obtain the internal

$$u_{i(0)}^{(s)} = K_{ii}^{(s)-1} \left(f_i^{(s)} - K_{ib}^{(s)} u_{b(0)}^{(s)} \right)$$

which corresponds to the solution of a Dirichlet problem with prescribed displacement $u_{b(0)}^{(s)}$. One deduces that :

$$\left\{ \begin{array}{ll} \forall \, s \in \mathbf{S}, & \begin{pmatrix} K_{ii}^{(s)} & K_{ib}^{(s)} \\ K_{bi}^{(s)} & K_{bb}^{(s)} \end{pmatrix} \begin{pmatrix} u_{i(0)}^{(s)} \\ u_{b(0)}^{(s)} \end{pmatrix} - \begin{pmatrix} f_{i}^{(s)} \\ f_{b}^{(s)} \end{pmatrix} = \begin{pmatrix} 0 \\ -r_{b(0)}^{(s)} \end{pmatrix} \\ &= \begin{pmatrix} 0 \\ S_{p}^{(s)} u_{b(0)}^{(s)} - b_{p}^{(s)} \end{pmatrix} \right.$$

The second line of the system reflects the equilibrium residual of the subdomain and represents exactly the contribution $r_{b(0)}^{(s)} = b_p^{(s)} - S_p^{(s)} u_{b(0)}^{(s)}$ to the assembled residual $\mathbf{r}_{b(0)} = \mathbf{b}_p - \mathbf{S}_p \mathbf{u}_{b(0)}$. Thus, the assembled residual can be subsequently computed as follows:

$$\mathbf{r}_{b(0)} = \mathbb{A}^{\diamondsuit} r_{b(0)}^{\diamondsuit}$$

It is worth noting at the fact that it is consequently not necessary to compute explicitly the Schur complement $S_p^{(s)}$. Only the computation of the substructure equilibrium residual $r_{b(0)}^{(s)}$ is required. One can also easily check that one finally computes:

$$\mathbf{r}_{b(0)} = \mathbb{A}^{\diamondsuit} \ r_{b(0)}^{\diamondsuit} = \mathbb{A}^{\diamondsuit} \left(b_p^{\diamondsuit} - S_p^{\diamondsuit} \ u_b^{\diamondsuit} \right) = \mathbf{b}_p - (\mathbb{A}^{\diamondsuit} \ S_p^{\diamondsuit} \ \mathbb{A}^{\diamondsuit}^T) \mathbf{u}_{b(0)} = \mathbf{b}_p - \mathbf{S}_p \mathbf{u}_{b(0)}$$

which represents the residual of the primal interface problem. Initial search direction is defined as $\mathbf{d}_{b(0)} = \mathbf{r}_{b(0)}$.

Let now consider a given iteration k + 1. Quantities $\mathbf{u}_{b(k)}$, $\mathbf{d}_{b(k)}$ and $\mathbf{r}_{b(k)}$ at previous iteration k are known. Following the same procedure as before, one can compute the product by the Schur complement of any vector \mathbf{d}_b thanks to local (subdomain) computations as follows:

- Compute local (concatenated) vector : $d_{b(k)} = \mathbb{A}^{\diamondsuit} \mathbf{d}_{b(k)}$ Solve local Dirichlet problem : $\forall s \in \mathbf{S}$, $d_{i(k)}^{(s)} = -K_{ii}^{(s)-1}K_{ib}^{(s)}d_{b(k)}^{(s)}$
- Compute local matrix-vector product :

$$\left\{ \begin{array}{ll} \forall \, s \in \mathbf{S}, & \begin{pmatrix} K_{ii}^{(s)} & K_{ib}^{(s)} \\ K_{bi}^{(s)} & K_{bb}^{(s)} \end{pmatrix} \begin{pmatrix} d_{i(k)}^{(s)} \\ d_{b(k)}^{(s)} \end{pmatrix} & = & \begin{pmatrix} 0 \\ S_p^{(s)} d_{b(k)}^{(s)} \end{pmatrix} \end{array} \right.$$

— Assemble and compute global matrix-vector product : $\mathbf{S}_{p}\mathbf{d}_{b(k)} = \mathbb{A}^{\diamondsuit} \left(S_{p}^{\diamondsuit} d_{b(k)}^{\diamondsuit} \right)$

Once again, one can note that only local vectors $S_p^{(s)}d_{b(k)}^{(s)}$ have to be computed and exchanged between subdomains/proessors which represents a small amount of data. The local primal Schur complement $S_p^{(s)}$ does not have to be computed or exchanged between subdomains/processors. Once matrix-vector product $\mathbf{S}_p \mathbf{d}_{b(k)}$ is known, one can proceed to next steps of the the conjugate gradient algorithm. This leads to Algorithm 2.

Remark 1 New iterate is computed as follows:

$$\mathbf{u}_{b(k+1)} = \mathbf{u}_{b(k)} + \alpha_{(k)} \mathbf{d}_{b(k)} \quad \Rightarrow \quad u_{b(k+1)} ^{\diamondsuit} = \mathbb{A}^{\diamondsuit} \mathbf{u}_{b(k+1)}$$

Internal degrees of freedom within subdomains can be easily updated if needed:

$$\begin{array}{lcl} K_{ii}^{(s)} u_{i(k+1)}^{(s)} & = & f_i^{(s)} - K_{ib}^{(s)} u_{b(k+1)}^{(s)} \\ & = & f_i^{(s)} - K_{ib}^{(s)} \left(u_{b(k)}^{(s)} + \alpha_{(k)} d_{b(k)}^{(s)} \right) \end{array}$$

Thus,

$$\begin{array}{ll} u_{i(k+1)}^{(s)} & = & K_{ii}^{(s)^{-1}} \left(f_i^{(s)} - K_{ib}^{(s)} u_{b(k)}^{(s)} \right) + \alpha_{(k)} \left(-K_{ii}^{(s)^{-1}} d_{b(k)}^{(s)} \right) \\ & = & u_{i(k)}^{(s)} + \alpha_{(k)} d_{i(k)}^{(s)} \end{array}$$

where quantities $u_{i(k)}^{(s)}$ and $d_{i(k)}^{(s)}$ have already been computed.

Remark 2 Residual of the initial (unsubstructured) problem is directly linked to the residual of the interface problem:

$$\begin{aligned} \mathbf{r}_{(k)} &= & \mathbf{f} - \mathbf{K} \mathbf{u}_{(k)} = \sum_{s} \left[\begin{pmatrix} f_{i(s)}^{(s)} \\ f_{b}^{(s)} \end{pmatrix} - \begin{pmatrix} K_{ii}^{(s)} & K_{ib}^{(s)} \\ K_{bi}^{(s)} & K_{bb}^{(s)} \end{pmatrix} \begin{pmatrix} u_{i(0)}^{(s)} \\ u_{b(0)}^{(s)} \end{pmatrix} \right] \\ &= & \sum_{s} \begin{pmatrix} 0 \\ b_{p}^{(s)} - S_{p}^{(s)} u_{b}^{(s)} \end{pmatrix} = \sum_{s} \begin{pmatrix} 0 \\ r_{b(0)}^{(s)} \end{pmatrix} \end{aligned}$$

Thus: $\|\mathbf{r}_{(k)}\| = \|\mathbf{f} - \mathbf{K}\mathbf{u}_{(k)}\| = \|\mathbf{r}_{b(k)}\|_{\Gamma} = \|\mathbf{b}_p - \mathbf{S}_p\mathbf{u}_{b(k)}\|_{\Gamma}$

Algorithm 2 Conjugate gradient algorithm to solve the primal interface problem

Initialization : Choose $\mathbf{u}_{b(0)}$ (generally $\mathbf{u}_{b(0)} = 0$)

 \Box Compute local (concatenated) vector : $u_{b(0)}^{\diamondsuit} = \mathbb{A}^{\diamondsuit}^T \mathbf{u}_{b(0)}$

for $\overline{\text{For } s = 1}$ to N_s do

- Solve local Dirichlet problem : $u_{i(0)}^{(s)} = K_{ii}^{(s)-1} \left(f_i^{(s)} K_{ib}^{(s)} u_{b(0)}^{(s)} \right)$ 1.
- 2. Compute local matrix-vector product:

$$\begin{pmatrix} K_{ii}^{(s)} & K_{ib}^{(s)} \\ K_{bi}^{(s)} & K_{bb}^{(s)} \end{pmatrix} \begin{pmatrix} u_{i(0)}^{(s)} \\ u_{b(0)}^{(s)} \end{pmatrix} - \begin{pmatrix} f_{i}^{(s)} \\ f_{b}^{(s)} \end{pmatrix} = \begin{pmatrix} 0 \\ -r_{b(0)}^{(s)} \end{pmatrix} = \begin{pmatrix} 0 \\ S_{p}^{(s)} u_{b(0)}^{(s)} - b_{p}^{(s)} \end{pmatrix}$$

end for

- \square Compute global residual : $\mathbf{r}_{b(0)} = \mathbb{A}^{\diamondsuit} r_{b(0)}^{\diamondsuit} (= \mathbf{b}_p \mathbf{S}_p \mathbf{u}_{b(0)})$
- $\Box \ \overline{\mathbf{d}_{b(0)} = \mathbf{r}_{b(0)}}$

for For k = 0 to m **do**

Compute local (concatenated) vector : $d_{b(k)} = \mathbb{A}^{\diamondsuit}^T \mathbf{d}_{b(k)}$

for For s = 1 to N_s do

- Solve local Dirichlet problem : $\forall s \in \mathbf{S}, d_{i(k)}^{(s)} = -K_{ii}^{(s)^{-1}} K_{ib}^{(s)} d_{b(k)}^{(s)}$
- Compute local matrix-vector product :

$$\left\{ \begin{array}{ll} \forall \, s \in \mathbf{S}, & \begin{pmatrix} K_{ii}^{(s)} & K_{ib}^{(s)} \\ K_{bi}^{(s)} & K_{bb}^{(s)} \end{pmatrix} \begin{pmatrix} d_{i(k)}^{(s)} \\ d_{b(k)}^{(s)} \end{pmatrix} & = & \begin{pmatrix} 0 \\ S_p^{(s)} d_{b(k)}^{(s)} \end{pmatrix} \end{array} \right.$$

end for

- Compute global matrix-vector product : $\mathbf{S}_{p}\mathbf{d}_{b(k)} = \mathbb{A}^{\diamondsuit}\left(S_{p}^{\diamondsuit} d_{b(k)}^{\diamondsuit}\right)$ Compute optimal step : $\alpha_{(k)} = \frac{\mathbf{r}_{b(k)}^{T}\mathbf{r}_{b(k)}}{\mathbf{d}_{b(k)}^{T}\mathbf{S}_{p}\mathbf{d}_{b(k)}} \left(= \frac{\mathbf{r}_{b(k)}^{T}\mathbf{d}_{b(k)}^{T}\mathbf{d}_{b(k)}}{\mathbf{d}_{b(k)}^{T}\mathbf{S}_{p}\mathbf{d}_{b(k)}}\right)$
- Compute iterate : $\mathbf{u}_{b(k+1)} = \mathbf{u}_{b(k)} + \alpha_{(k)} \mathbf{d}_{b(k)}$
- Compute residual : $\mathbf{r}_{b(k+1)} = \mathbf{r}_{b(k)} - \alpha_{(k)} \mathbf{S}_p \mathbf{d}_{b(k)}$
- Compute orthogonalization parameter:

$$\beta_{(k+1)} = \frac{\mathbf{r}_{b(k+1)}^T \mathbf{r}_{b(k)} \mathbf{r}_{b(k)}}{\mathbf{r}_{h(k)}^T \mathbf{r}_{b(k)}} \left(= -\frac{\mathbf{r}_{b(k+1)}^T \mathbf{S}_p \mathbf{d}_{b(k)}}{\mathbf{d}_{h(k)}^T \mathbf{S}_p \mathbf{d}_{b(k)}} \right)$$

Update search direction : $\mathbf{d}_{b(k+1)} = \mathbf{r}_{b(k+1)} + \beta_{(k+1)} \mathbf{d}_{b(k)}$ end for