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# Preconditioners with projectors for mixed hybrid finite element methods

**Abstract:** We propose and investigate numerically two new preconditioners for the matrices, which arise in the mixed-hybrid finite element methods for diffusion equation in strongly heterogeneous media. Both preconditioners include special projectors on the vector spaces orthogonal to the vectors with constant components. We give general description of the preconditioners and discuss numerical results which demonstrate their efficiency compared to the classical diagonal preconditioner.

**Keywords:** Preconditioned conjugate gradient method, preconditioners with projectors, mixed-hybrid finite element method, diffusion equation.

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In the present paper, we propose and investigate numerically two new preconditioners for the condensed matrices, arising in mixed-hybrid finite element approximations of the diffusion equation in strongly heterogeneous media. The construction of preconditioners is based on the ideas originally proposed and investigated in [4]. The preconditioners contain special projectors on the vector spaces orthogonal to vectors with constant components.

The paper is organized as follows. In Section 1, we describe the differential mixed macro-hybrid formulation of the diffusion equation. To simplify the exposition, we consider only the case of polygonal/polyhedral domains and subdomains, and the case of homogeneous Neumann conditions. For the condensed matrix arising in mixed hybrid or mixed macro-hybrid finite element methods we describe the general idea for construction of preconditioners with special projectors [4].

In Section 2, we consider the model diffusion problem in the square domain partitioned in square subdomains with constant diffusion and reaction coefficients in subdomains. Based on the results of Section 1 we propose a variant of a preconditioner with projector which we refer to as a 'coarse mesh preconditioner'. We briefly describe the algorithm of the preconditioner implementation and the respective numerical results.

In Section 3 we consider the domain from Section 2 and assume the checkerboard ordering of square subdomains with large values of the diffusion coefficient. For this particular case we propose a simple block-diagonal preconditioner with projectors. The computational arithmetical cost of this preconditioner is proportional to its dimension. We compare the efficiency of this preconditioner with the classical diagonal (point Jacobi) preconditioner in context of the Preconditioned Conjugate Gradient (PCG) method.

#### 1 Problem formulation

We consider the diffusion problem

$$-\nabla (K\nabla p) + c p = f \quad \text{in} \quad \Omega$$
  
-  $(K\nabla p) \cdot \mathbf{n} = 0 \quad \text{on} \quad \partial \Omega$  (1.1)

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where  $\Omega$  is a simply connected polygon/polyhedron in  $\mathbb{R}^2/\mathbb{R}^3$  with boundary  $\partial \Omega$ ,  $\boldsymbol{n}$  is the outward unit normal to  $\partial \Omega$ , K = K(x) is the diffusion tensor, c is a nonnegative coefficient, and f = f(x) is a given source function. We partition  $\Omega$  into m>1 nonoverlapping polygons/polyhedrons  $E_s$ ,  $s=1,\ldots,m$ , and denote by  $\Gamma_{s,t}$  the interfaces between  $E_s$  and  $E_t$ , i.e.,  $\Gamma_{s,t} = \partial E_s \cap \partial E_t$ ,  $s < t, s, t = 1, \ldots, m$ . For the sake of simplicity we assume that subdomains  $E_s$  are convex,  $K \equiv K_s = k_s I_s$  are constant scalar tensors in  $E_s$ , where  $I_s$  is the identity matrix, and  $c \equiv c_s$  are nonnegative constants,  $s = 1, \dots, m$ .

The differential mixed macro-hybrid formulation of (1.1) based on the partitioning of  $\Omega$  into subdomains  $E_s$ , s = 1, ..., m, reads as follows:

$$K_{s}^{-1}\boldsymbol{u}_{s} + \nabla p_{s} = 0 \quad \text{in } E_{s}$$

$$\nabla \boldsymbol{u}_{s} + c_{s} p_{s} = f \quad \text{in } E_{s}$$

$$p_{s} = p_{t} \quad \text{on } \Gamma_{s,t}, \ s < t$$

$$\boldsymbol{u}_{s} \cdot \boldsymbol{n}_{s} + \boldsymbol{u}_{t} \cdot \boldsymbol{n}_{t} = 0 \quad \text{on } \Gamma_{s,t}, \ s < t$$

$$\boldsymbol{u}_{s} \cdot \boldsymbol{n} = 0 \quad \text{on } \partial \Omega \cap \partial E_{s}$$

$$(1.2)$$

where  $n_s$  are the outward normals to  $\partial E_s$  and  $\partial E_t$ , s,  $t = 1, \dots, m$ . The variational mixed hybrid formulation of (1.2) is well known [1].

Let  $\Omega_h$  be a triangular/tetrahedral mesh in  $\Omega$ , conforming with  $\partial E_s$ , i.e.,  $\partial E_s$  is the union of the sides/faces of mesh cells in  $\Omega_h$ ,  $s=1,\ldots,m$ . We approximate (1.2) by the lowest order mixed hybrid Raviart–Thomas finite element method ( $RT_0$  method) on the mesh  $\Omega_h$ . The method results in the linear algebraic system [1, 2]:

$$\mathcal{A}\begin{pmatrix} \overline{u} \\ \overline{p} \\ \overline{\lambda} \end{pmatrix} = \begin{pmatrix} 0 \\ -\overline{F} \\ 0 \end{pmatrix} \tag{1.3}$$

with the saddle point matrix

$$\mathcal{A} = \begin{pmatrix} M & B^T & C^T \\ B & -\Sigma & 0 \\ C & 0 & 0 \end{pmatrix}. \tag{1.4}$$

Under the appropriate ordering of the flux variables (DOFs) the matrix M is block diagonal with  $3 \times 3/4 \times 4$ blocks. Eliminating the vector  $\overline{u}$  in (1.3) we get the system:

$$S\begin{bmatrix} \overline{p} \\ \overline{\lambda} \end{bmatrix} \equiv \begin{bmatrix} S_{pp} & S_{p\lambda} \\ S_{\lambda p} & S_{\lambda \lambda} \end{bmatrix} \begin{bmatrix} \overline{p} \\ \overline{\lambda} \end{bmatrix} = \begin{bmatrix} \overline{F} \\ 0 \end{bmatrix}$$
 (1.5)

with the symmetric positive definite matrix *S* (semidefinite, if  $c_S = 0$ ,  $s = 1, \ldots, m$ ):

$$S = \begin{bmatrix} B \\ C \end{bmatrix} M^{-1} \begin{bmatrix} B^T & C^T \end{bmatrix} + \begin{bmatrix} \Sigma & 0 \\ 0 & 0 \end{bmatrix}. \tag{1.6}$$

The matrix *S* in (1.5) can be defined by subassembling matrices

$$S_t = A_t + \begin{bmatrix} \Sigma & 0 \\ 0 & 0 \end{bmatrix} \tag{1.7}$$

associated with subdomains  $E_t$ , t = 1, ..., m, i.e.,

$$S = \sum_{t=1}^{m} N_t S_t N_t^T$$

where  $N_t$  are the subassembling matrices,  $A_t$  are condensed matrices for subdomains  $E_t$ , and  $\Sigma_t$  are the diagonal matrices, t = 1, ..., m. It is obvious that the matrices  $A_t$  are singular, and ker(A) consists of vectors with constant components, i.e., consists of the vectors, which are proportional to the vector  $\overline{e}$  with all components equal to one.

Let  $M_t$  be diagonal  $m_t \times m_t$  matrices with positive diagonal entries, where  $m_t$  is the dimension of  $A_t$ . Let us consider the eigenvalue problem

$$A_t \overline{W} = \lambda M_t \overline{W}. \tag{1.8}$$

We denote by  $0 = \lambda_{t,1} < \lambda_{t,2} \le \ldots \le \lambda_{t,m_t}$  the eigenvalues in (1.8) and by  $\overline{w}_{t,1},\ldots,\overline{w}_{t,m_t}$  the set of  $M_t$ orthogonal eigenvectors in (1.8), t = 1, ..., m. Then we get the spectral decompositions

$$A_t = M_t W_t \Lambda_t W_t^T M_t \tag{1.9}$$

where  $\Lambda_t = \text{diag}\{\lambda_{t,1}, \lambda_{t,2}, \dots, \lambda_{t,m_t}\}$  and

$$W_t = \left[ \overline{W}_{t,1}, \dots, \overline{W}_{t,m_t} \right]. \tag{1.10}$$

It is obvious that

$$W_{t}^{T} M_{t} W_{t} = M_{t} W_{t} W_{t}^{T} = W_{t} W_{t}^{T} M_{t} = I_{t}$$
(1.11)

with the  $m_t \times m_t$  identity matrix  $I_t$ , t = 1, ..., m. We also remind that

$$\overline{w}_{t,1} = \frac{1}{\|\overline{e}_t\|_{M_t}} \overline{e}_t \tag{1.12}$$

where  $\overline{e}_t$  is  $m_t \times m_t$  vector with the components equal to one and  $\|\overline{e}_t\|_{M_t}$  is the  $M_t$ -norm of  $\overline{e}_t$ ,  $t = 1, \ldots, m$ . Following [4], we replace the matrices  $A_t$  in (1.9) with matrices

$$B_t = \alpha_t M_t W_t \left( I_t - \hat{I}_t \right) W_t M_t = \alpha \left( M_t - M_t \overline{W}_{t,1} \overline{W}_{t,1}^T M_t \right)$$
(1.13)

where  $\alpha_t$  are some positive numbers and  $\hat{I}_t$  are  $m_t \times m_t$  matrix with the only nonzero entry in the position (1,1) equals to one, t = 1, ..., m. It is obvious that  $B_t = B_t^T \ge 0$ ,  $\ker(B_t) = \ker(A_t)$  and

$$\frac{\lambda_{t,2}}{\alpha_t} B_t \leqslant A_t \leqslant \frac{\lambda_{t,m_t}}{\alpha_t} B_t, \quad t = 1, \dots, m.$$
 (1.14)

We define the matrices

$$B = \sum_{t=1}^{m} N_t \begin{bmatrix} B_t + \begin{bmatrix} \Sigma_t & 0 \\ 0 & 0 \end{bmatrix} \end{bmatrix} N_t^T.$$
 (1.15)

Then from [4] we get the inequalities

$$\beta_1 B \leqslant S \leqslant \beta_2 B \tag{1.16}$$

with

$$\beta_1 = \min \left\{ \min_{t=1,...,m} \frac{\lambda_{t,2}}{\alpha_t}; 1 \right\}$$
 (1.17)

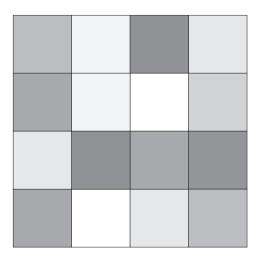
and

$$\beta_2 = \max \left\{ \max_{t=1,\dots,m} \frac{\lambda_{t,m_t}}{\alpha_t}; 1 \right\}$$
 (1.18)

**Remark 1.1.** In this section triangular/tetrahedral meshes and  $RT_0$  approximations were chosen to illustrate the algebraic operations (1.9)-(1.15) and estimates (1.16)-(1.18). In fact, we can repeat the same transformations and estimates for many other meshes and approximations. In particular, for numerical results in Sections 2 and 3 we shall use square meshes and macro-hybrid mixed discretizations with piece-wise constant fluxes [5] which practically coincide with the macro-hybridized finite-volume method.

### Coarse mesh preconditioner

In this section, we consider the diffusion equation in the unit square partitioned into  $m = \sqrt{m} \times \sqrt{m}$ subsquares as shown in Fig. 1. We assume that the diffusion coefficients c are constant in each subdomain/subsquare  $E_t$ , i.e.,  $k_t = \text{const} > 0$  and  $c \equiv c_t = \text{const} \ge 0$  in  $E_t$ ,  $t = 1, \ldots, m$ . In  $\Omega = [0; 1] \times [0; 1]$  we



**Figure 1.** Domain  $\Omega$  partitioned in subsquares  $E_s$ ,  $s=1,\ldots,m, m=16$ .

define the square mesh  $\Omega_h$  with the step-size h = 1/n,  $n \ge 2$ . We assume that n is a multiple of  $\sqrt{m}$ . Thus, we defined the macro-mesh (coarse mesh) with the step-size  $H = 1/\sqrt{m}$ , and the mini-mesh  $\Omega_h$ .

We approximate the diffusion problem (1.2) by the mixed macro-hybrid finite element method with piecewise constant fluxes [5]. It can be easily shown that the proposed discretization is equivalent to the finite volume discretization of (1.1) on  $\Omega_h$  hybridized on the interfaces  $\Gamma_{s,t}$  between macrocells  $E_s$  and  $E_t$ , and on the boundary of  $\Omega$ . This approximation results in system (1.3)–(1.4) with a diagonal matrix M. We observe, in this method DOFs (degrees of freedom) for  $\lambda$  represent the mean values of the solution function p on the boundaries of mesh cells in  $\Omega_h$  belonging to the boundaries of  $E_t$ , t = 1, ..., m.

To define the matrices  $B_t$ ,  $t = 1, \dots, m$ , in (1.13) we choose  $M_t$  to be a diagonal matrix with the diagonal entries equal to  $h^2$  for the components of subvector  $\overline{p}$ , and equal to h for the components of subvector  $\overline{\lambda}$ ,  $t=1,\ldots,m$ . Thus each  $M_t$  has  $(H/h)^2$  diagonal entries equal to  $h^2$ , and 4H(H/h) diagonal entries equal to h, t = 1, ..., m.

For numerical experiments we choose

$$\alpha_t = ||M_t^{-1}A_t|| \equiv \alpha = \frac{8k_t}{h^2}, \quad t = 1, \dots, m.$$
 (2.1)

Then, in inequalities (1.16) we get

$$\beta_1 = \min_{t=1,...,m} \frac{\lambda_{t,2}}{\alpha_t}, \qquad \beta_2 = 1.$$
(2.2)

We observe, that the value of  $\beta_1$  does not depend on the values  $k_t$  and  $c_t$ ,  $t = 1, \ldots, m$ .

The numerical results are given in Tables 1–4. Abbreviation PCG stays for the preconditioned conjugate gradient method with preconditioner equal to the diagonal of S, and abbreviation CMP (Coarse Mesh Preconditioner) stays for the Preconditioned Conjugate Gradient method with the above preconditioner B in (1.15). The algorithm for solving a system with the matrix *B* is described in [4].

In Tables 1 and 2 we give the numbers of iterations of PCG and CMP methods sufficient for minimization of S-norm of the initial error in  $\frac{1}{c} = 10^6$  times in the case when the reaction coefficient  $c \equiv 1$  in  $\Omega$ . In Tables 3 and 4 we give the number of iterations of the same methods with the same tolerance when the values of  $c_s$ are chosen randomly in the interval [0.01; 100].

We can clearly observe that the number of iterations in the PCG method essentially depends on the values of the diffusion coefficients. The number of iterations in CMP method does not depend on the values of coefficients  $k_t$  and  $c_t$ , t = 1, ..., m. We can also observe that the number of iterations is proportional to H/h. This statement correlates with theoretical estimates in [4].

**Table 1.** H = 1/8,  $m = 8 \times 8 = 64$  subsquares.

n×n	<i>k</i> <sub>max</sub> = 1		$k_{\text{max}} = 10^2$		$k_{\text{max}} = 10^4$		$k_{\text{max}} = 10^6$	
	PCG	CMP	PCG	CMP	PCG	CMP	PCG	CMP
128 × 128	502	310	927	310	1220	309	1227	307
$256 \times 256$	910	652	1617	676	2317	657	3741	651
512 × 512	1705	1373	3419	1398	4767	1377	5615	1381

**Table 2.** H = 1/16,  $m = 16 \times 16 = 256$  subsquares.

n × n	$k_{\text{max}} = 1$		$k_{\text{max}} = 10^2$		k <sub>max</sub> :	= 10 <sup>4</sup>	$k_{\text{max}} = 10^6$	
	PCG	CMP	PCG CMP		PCG	CMP	PCG	CMP
128 × 128	600	144	1267	145	2499	144	5309	142
$256 \times 256$	944	330	1958	331	4536	330	9139	326
512 × 512	1678	744	3219	744	8591	741	18044	733

**Table 3.** H = 1/8,  $m = 8 \times 8 = 64$  subsquares.

n × n	$k_{max} = 1$		$k_{\text{max}} = 10^2$		<i>k</i> <sub>max</sub>	= <b>10</b> <sup>4</sup>	$k_{\text{max}} = 10^6$		
	PCG	CMP	PCG	CMP	PCG	CMP	PCG	CMP	
128 × 128	276	377	701	317	1278	307	1520	306	
$256 \times 256$	543	736	1324	691	2145	655	2655	650	
512 × 512	970	1497	2217	1421	4508	1392	6680	1303	

**Table 4.** H = 1/16,  $m = 16 \times 16 = 256$  subsquares.

n×n	$k_{max} = 1$		$k_{max} = \mathbf{10^2}$		k <sub>max</sub> :	= <b>10</b> <sup>4</sup>	$k_{max} = \mathbf{10^6}$	
	PCG	CMP	PCG	CMP	PCG	CMP	PCG	CMP
128 × 128	278	199	798	158	2009	147	4726	143
$256 \times 256$	527	388	1323	339	3524	331	8938	323
512 × 512	907	794	2277	755	7502	736	15326	733

## 3 Block diagonal preconditioner

In this section, we partition the macrocells into two groups and assume that any two macrocells  $E_s$  and  $E_t$  in the first group do not have interfaces, i.e.,  $|\Gamma_{s,t}| = 0$ ,  $1 \le s < t \le m_0$ ,  $m_0 < m$ .

We assume that the diffusion coefficients  $k_s$  for the first group of macrocells are much larger than one  $(k_s \gg 1)$ ,  $s = 1, \ldots, m_0$ , and  $k_s \approx 1$  for the second group of macrocells,  $s = m_0 + 1, \ldots, m$ . We represent the matrix S as the sum of two matrices  $S_1$  and  $S_2$ , where the matrix  $S_1$  is defined by

$$S_1 = \sum_{t=1}^{m_0} N_t \left[ (k_t - 1) A_t + \begin{bmatrix} \Sigma_t & 0 \\ 0 & 0 \end{bmatrix} \right] N_t^T.$$
(3.1)

Then, the block diagonal preconditioner *B* for the matrix *S* is defined by

$$B = D + \sum_{t=1}^{m_0} N_t \left[ (k_t - 1) B_t + \begin{bmatrix} \Sigma_t & 0 \\ 0 & 0 \end{bmatrix} \right] N_t^T$$
(3.2)

where the matrix D is a diagonal matrix with positive diagonal entries. The matrix B is a block diagonal matrix

$$B = \sum_{t=1}^{m} N_t B_{tt} N_t^T (3.3)$$

where the blocks

$$B_{tt} = D_{tt} + (k_t - 1) M_t \overline{W}_{t,1} \overline{W}_{t,1}^T M_t$$
(3.4)

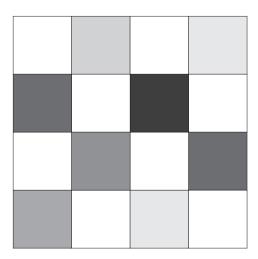


Figure 2. Example of 'checkerboard distribution'.

**Table 5.** H = 1/8,  $m = 8 \times 8 = 64$  subsquares.

n × n	$k_{\text{max}} = 10^2$			k	$k_{max} = \mathbf{10^4}$			$k_{max} = \mathbf{10^6}$		
	PCG	CMP	BDP	PCG	CMP	BDP	PCG	CMP	BDP	
128 × 128	1141	306	809	2101	265	747	4804	245	625	
$256 \times 256$	2222	642	1517	4206	558	1347	8337	542	1155	
$512 \times 512$	4455	1375	2826	9111	1219	2487	13796	1151	2188	

**Table 6.** H = 1/16,  $m = 16 \times 16 = 256$  subsquares.

n × n	$k_{\text{max}} = 10^2$			$k_{\text{max}} = 10^4$			$k_{max} = 10^6$		
	PCG	CMP	BDP	PCG	CMP	BDP	PCG	CMP	BDP
128 × 128	1508	143	692	6588	128	625	23093	112	541
$256 \times 256$	2479	325	1300	11095	276	1153	32797	267	987
512 × 512	4662	733	2357	16405	639	2080	58545	598	1711

are rank one perturbations of diagonal matrices  $D_{tt}$  and the blocks  $B_{tt}$ ,  $t = m_0 + 1, ..., m$ , are diagonal matrices. Let us assume that

$$v_1 D \leqslant S_2 \leqslant v_2 D \tag{3.5}$$

with some positive  $v_1 < v_2$ . Then

$$\beta_1 B \leqslant S \leqslant \beta_2 B \tag{3.6}$$

where

$$\beta_1 = \min \left\{ \min_{t=1,...,m} \frac{\lambda_{t,2}}{\alpha_t}; \, \nu_1; \, 1 \right\}$$
 (3.7)

and

$$\beta_2 = \max \left\{ \max_{t=1,...,m} \frac{\lambda_{t,m_t}}{\alpha_t}; \nu_2; 1 \right\}.$$
 (3.8)

It is obvious that the computational cost of solving a system with matrix B in (3.3)–(3.4) is lower then the computational cost for a residual with the matrix S.

For numerical experiments we choose the domain and partitioning from Section 2 shown in Fig. 1. We partition macrocells into two subgroups based on checkerboard ordering shown in Fig. 2.

The diffusion coefficient for macrocells in the first group is chosen randomly on the segment [10;  $k_{\text{max}}$ ] and the diffusion coefficient equals to one for the macrocells in the second group. We choose  $c \equiv 1$  in  $\Omega$  and

define the preconditioner B by

$$B = 8I + \sum_{t=m_0+1}^{m} N_t \begin{bmatrix} \Sigma_t & 0 \\ 0 & 0 \end{bmatrix} N_t^T$$
(3.9)

where I is the identity matrix of the same size as B in (3.9).

The numerical results are given in Tables 5–6. Abbreviation BDP is used for the Preconditioned Conjugate Gradient method with preconditioner (3.9). An algorithm for inverting a diagonal matrix with rank one perturbation can be found in [3].

The numerical results clearly show that the number of iterations of BDP is  $2 \div 2.5$  times larger than the number of iterations for the PCG method in Tables 1–2 for  $k_{\text{max}} = 1$ , and does not depend on the values of  $k_s$ ,  $s = 1, ..., m_0.$ 

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