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Matrix-free multigrid block-preconditioners for higher order Discontinuous Galerkin discretisations

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Abstract

Efficient and suitably preconditioned iterative solvers for elliptic partial differential equations (PDEs) of the convection-diffusion type are used in all fields of science and engineering, including for example computational fluid dynamics, nuclear reactor simulations and combustion models. To achieve optimal performance, solvers have to exhibit high arithmetic intensity and need to exploit every form of parallelism available in modern manycore CPUs. This includes both distributed- or shared memory parallelisation between processors and vectorisation on individual cores.

The computationally most expensive components of the solver are the repeated applications of the linear operator and the preconditioner. For discretisations based on higher-order Discontinuous Galerkin methods, sum-factorisation results in a dramatic reduction of the computational complexity of the operator application while, at the same time, the matrix-free implementation can run at a significant fraction of the theoretical peak floating point performance. Multigrid methods for high order methods often rely on block-smoothers to reduce high-frequency error components within one grid cell. Traditionally, this requires the assembly and expensive dense matrix solve in each grid cell, which counteracts any improvements achieved in the fast matrix-free operator application. To overcome this issue, we present a new matrix-free implementation of block-smoothers. Inverting the block matrices iteratively avoids storage and factorisation of the matrix and makes it possible to harness the full power of the CPU. We implemented a hybrid multigrid algorithm with matrix-free block-smoothers in the high order Discontinuous Galerkin (DG) space combined with a low order coarse grid correction using algebraic multigrid where only low order components are explicitly assembled. The effectiveness of this approach is demonstrated by solving a set of representative elliptic PDEs of increasing complexity, including a convection dominated problem and the stationary SPE10 benchmark.

Keywords: multigrid, elliptic PDE, Discontinuous Galerkin, matrix-free methods, preconditioners, DUNE

1. Introduction

Second order elliptic PDEs of the convection-diffusion-reaction form

$$-\nabla \cdot (K \nabla u) + \nabla \cdot (bu) + cu = f \quad (1)$$

with spatially varying coefficients play an important role in many areas of science and engineering. The Discontinuous Galerkin (DG) method [1, 2, 3, 4, 5, 6, 7, 8] is a popular discretisation scheme for various reasons: it allows local mass conservation and (when used with an appropriate basis) leads to diagonal mass matrices which avoid global solves in explicit time integrators. It also has several computational advantages as will be discussed below. Problems of the form in (1) arise for example in the simulation of subsurface flow phenomena [9, 10, 11, 12]. Another important application area is fluid dynamics. When the incompressible Navier Stokes equations are solved with Chorin's projection method [13, 14], a Poisson equation has to be solved for the pressure correction and the momentum equation needs to

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be solved for the tentative velocity, see [15] for a DG-based version. Implicit DG solvers for the fully compressible atmospheric equations of motion are described in [16, 17].

The fast numerical solution of equation (1) requires algorithmically optimal algorithms, such as multigrid methods (see [18] for an overview), which have a numerical complexity that typically grows linearly with the number of unknowns. Equally important is the efficient implementation of those methods on modern computer hardware. While low order discretisation methods such as finite volume approximations and linear conforming finite element methods are very popular, from a computational point of view one of their main drawbacks is the fact that their performance is limited by main memory bandwidth. Modern computer architectures such as the Intel Haswell/Broadwell CPUs, Xeon Phi co-processors and graphics processing units (GPUs) may only achieve their peak performance floating point performance if data loaded into the CPU is reused, i.e. the arithmetic intensity of the algorithm (ratio of floating point operations executed per byte loaded, in short: flop-to-byte-ratio) is sufficiently high. For example the 16 core Haswell chip used for the computations in this work has a theoretical peak performance of $243.2 \cdot 10^9$ fused multiply-add (FMA) operations per second and a main memory bandwidth (as measured with the STREAM benchmark [19]) of 40 – 50 GB/s. Hence about 40 FMA operations can be carried out in the time it takes to read one double precision number from memory. Low order discretisation methods require frequent sparse matrix-vector products. If the matrix is stored in the standard compressed sparse row storage (CSR) format, this multiplication has a very low arithmetic intensity of about 1 FMA operation per matrix entry loaded from memory (assuming perfect caching of the vectors). Hence traditional solvers which first assemble the system matrix and then invert it with sparse matrix-vector operations will only use a fraction of the system's peak performance.

Matrix-free methods, on the other hand, recompute each matrix-element on-the-fly instead of storing an assembled matrix. In addition to eliminating setup costs for the matrix assembly, this raises the computational intensity and allows utilisation of the full performance of modern manycore chip architectures. An additional advantage are the reduced storage requirements: since the matrix does not have to be kept in memory, this allows the solution of much larger problems and has been quantified by estimating the memory requirements for different implementations in this paper. While high arithmetic intensity is a desirable property of a good implementation, the quantity which really needs to be minimised is the total computation time. A naive matrix-free implementation can result in an increase in the run-time since a large number of floating point operations (FLOPs) is executed for each unknown. Careful matrix-free implementations of higher-order finite element discretisations avoid this problem. If p is the polynomial degree of the basis function in each d -dimensional grid cell, a naive implementation requires $\mathcal{O}(p^{2d})$ FLOPs per cell. Sum factorisation, which exploits the tensor-product structure of the local basis functions, reduces this to $\mathcal{O}(d \cdot p^{d+1})$ [20, 21, 22, 23], making the matrix-free implementation particularly competitive if the order p is sufficiently high. Numerical results indicate that sum factorisation reduces the runtime even for low polynomial degrees $p > 1$ [23, 24]. Another advantage of the DG method is that all data associated with one grid cell is stored contiguously in memory. This avoids additional gather/scatter operations which are necessary if data is also held on other topological entities such as vertices and edges as in conforming high order finite elements.

The efficient matrix-free implementation of higher-order DG methods was described by some of us in a previous paper [24] which concentrates on the efficient application of the DG operator. Here we extend this approach to iterative solvers for elliptic PDEs. While matrix-vector products are one key ingredient in Krylov-subspace solvers, efficient preconditioners are equally important and often form the computational bottleneck. Many preconditioners such as algebraic multigrid (AMG) [25, 26] or incomplete factorisations (ILU) [27] require the assembly of the system matrix, which can offset the advantage of the matrix-free sparse matrix-vector product.

In this work we describe the matrix-free implementation of block-smoothers, which are the key ingredient of multigrid methods such as those described in [28, 12]. The idea behind the multigrid algorithm in [28] is that the block-smoother eliminates high-frequency errors within one grid cell whereas an AMG solver in a lower order subspace of the full DG space reduces long range error components. Due to the much smaller size of the coarse level system, the cost for one AMG V-cycle on the low order subspace (which uses a traditional matrix-based implementation) is negligible in comparison to the block smoother on the high order system. In the simplest case the smoother is a block-Jacobi iteration applied to the system $Au = f$ which arises from high-order DG discretisation of (1). In each cell T of the grid this requires the following operations:

- 1: Calculate local defect $\mathbf{d}_T = \mathbf{f}_T - \sum_{T', \partial T' \cap \partial T \neq \emptyset} A_{T,T'} \mathbf{u}_{T'}^{(k)}$
- 2: Calculate correction $\delta \mathbf{u}_T = A_{T,T}^{-1} \mathbf{d}_T$
- 3: Update solution $\mathbf{u}_T^{(k+1)} = \mathbf{u}_T^{(k)} + \delta \mathbf{u}_T$

Here \mathbf{u}_T is the solution on cell T and the matrix $A_{T,T'}$ describes the coupling between cells T and T' which share at least one common face. In addition to dense matrix-vector products to calculate the local defect \mathbf{d}_T , this requires the solution of a small dense system

$$A_{T,T} \delta \mathbf{u}_T = \mathbf{d}_T \quad (2)$$

with $n_T = (p+1)^d$ unknowns in the second step. Matrix based methods use an LU- or Cholesky-factorisation of the matrix $A_{T,T}$ to solve (2) in each grid cell. The cost of the back-substitution step, which has to be carried out for every block-Jacobi iteration, is $\mathcal{O}(n_T^2) = \mathcal{O}(p^{2d})$. There is an additional setup cost of $\mathcal{O}(n_T^3) = \mathcal{O}(p^{3d})$ for the initial factorisation of $A_{T,T}$, which may be relevant if the number of preconditioner applications is small. In contrast, the matrix-free method which we present here uses an iterative method to solve this equation and if the implementation uses sum-factorisation, this reduces the complexity to $\mathcal{O}(d \cdot n_{\text{iter}}(\epsilon) \cdot p^{d+1})$. An important factor in this expression is $n_{\text{iter}}(\epsilon)$, the number of iterations which are required to solve the system in (2) to accuracy ϵ . Solving (2) exactly, i.e. iterating until the error is reduced to machine precision, will result in a relatively large n_{iter} and might not lead to a reduction of the total solution time for the full system. However, since the inverse is only required in the preconditioner, an approximate inversion with a small number of iterations might be sufficient and can increase the computational efficiency of the overall method. This is one of the key optimisations which will be explored below. The overall efficiency of the implementation also depends on how computationally expensive the evaluation of the functions K , \mathbf{b} and c in (1) is. At first sight this might put a matrix-free preconditioner at a disadvantage, since those functions have to be re-computed at every quadrature point in each iteration. However, it is often sufficient to use a simpler functional dependency in the preconditioner, for example by approximating K , \mathbf{b} and c by their average value on each cell. To demonstrate the efficiency of this approach we solve a set of linear problems of increasing complexity. This includes advection dominated problems which require more advanced SOR-type smoothers and the stationary SPE10 benchmark. The latter is a hard problem since the permeability field $K(\mathbf{x})$ varies over several magnitudes and has high-contrast jumps between grid cells.

A similar fully matrix-free approach for the solution of elliptic problems is described in [29]. The results there are obtained with the `deal.II` [30] implementation discussed in [23]. As for our code, the matrix-free operator application in [23] is highly optimised and utilises a significant fraction the peak floating point performance. In contrast to our implementation, the authors of [29] use a (matrix-free) Chebyshev-iteration as a smoother on the DG space (a Chebyshev smoother is also used for the AMG solver on the lower order subspace). One drawback of this approach is that it requires estimates of the largest eigenvalues, which will result in additional setup costs. While the authors of [29] solve a Laplace equation on complex geometries, it is not clear how the same Chebyshev smoother can be applied to non-trivial convection-dominated problems or problems with large, high contrast jumps of the coefficients.

Another approach for preconditioning higher order finite element discretisations was originally developed for spectral methods in [20] and used for example in [31, 32]. In [32], a preconditioner is constructed by approximating the higher-order finite element operator with a low-order discretisation on the finer grid which is obtained by subdividing each grid cell into elements defined by the nodal points. The resulting sparse system is then solved directly with an AMG preconditioner. A similar approach is used in [31], where a two-level Schwarz preconditioner is constructed for a spectral element discretisation of the pressure correction equation in the Navier Stokes system. Instead of using an AMG method for the sparse low-order system, the author employs a two-level Schwarz preconditioner. As in our method, the coarse level is obtained from the linear finite element discretisation on the original grid and this is combined with an iterative solution of the low order system on the finer nodal grid. Finally, instead of re-discretising the equation on a finer grid, the authors of [33] construct a sparse approximation with algebraic methods, which requires additional setup costs. Compared to our approach, the drawback of all those methods is that the fine grid problem is obtained from a lowest-order discretisation (or an algebraic representation leading to a sparse matrix), and will therefore result in a memory-bound implementation which can not use the full capacity of modern manycore processors.

Structure. This paper is organised as follows: in Section 2 we describe the higher order DG discretisation and our hybrid multigrid solver algorithm which is used throughout this work. We introduce the principles of the matrix-free smoother implementation and analyse its computational complexity as a function of the order of the DG discretisation. The implementation of those algorithms based on the DUNE software framework [34, 35] is described in Section 3, where we also briefly review the principles of sum-factorised operator application. Numerical results and comparisons between matrix-based and matrix-free solvers are presented in Section 4, which also contains an in-depth discussion of the memory requirements of different implementations. We conclude in Section 5. Some more technical details are relegated to the appendices. In Appendix A we discuss the re-discretisation of the operator in the low-order subspace which is used in the coarse level correction of the multigrid algorithm. A detailed breakdown of setup costs for the multigrid algorithm can be found in Appendix B.

2. Methods

We consider linear second order elliptic problems of the form

$$\nabla \cdot (\mathbf{b}u - K\nabla u) + cu = f \quad \text{in } \Omega \quad (3)$$

with boundary conditions

$$u = g \quad \text{on } \Gamma_D \subset \partial\Omega, \quad (\mathbf{b}u - K\nabla u) \cdot \boldsymbol{\nu} = j \quad \text{on } \Gamma_N = \partial\Omega \setminus \Gamma_D \quad (4)$$

in the d -dimensional domain Ω . Here the unit outer normal vector on the boundary $\partial\Omega$ is denoted by $\boldsymbol{\nu} = \boldsymbol{\nu}(\mathbf{x})$. $K = K(\mathbf{x}) \in \mathbb{R}^{d \times d}$ is the diffusion tensor, $\mathbf{b} = \mathbf{b}(\mathbf{x}) \in \mathbb{R}^d$ is the advection vector and $c = c(\mathbf{x})$ is a scalar reaction coefficient. All those terms can depend on the position $\mathbf{x} \in \Omega$; the fields K , \mathbf{b} and c are not necessarily smooth and can have large, high contrast jumps. A typical example for a problem with discontinuous coefficients is the SPE10 benchmark [36]. The problem in (3) could also appear in the linearisation of a non-linear problem such as the Navier Stokes equations [15]. In this case it needs to be solved at every iteration of a Newton- or Picard- iteration. The reaction term often arises from the implicit time discretisation of an instationary problem.

2.1. Higher-order DG discretisation

To discretise (3) and (4) we use the weighted symmetric interior penalty (WSIPG) method derived in [28] based on [1, 2, 6, 37, 38, 39] together with an upwinding flux for the advective part. The full method is described in [24] and the most important properties are recalled here. We work on a grid \mathcal{T}_h consisting of a set of cuboid cells T with smallest size $h = \min_{T \in \mathcal{T}_h} \{\text{diam}(T)\}$. Each cell T is the image of a diffeomorphic map $\mu_T : \hat{T} \rightarrow T$ where \hat{T} is the reference cube in d dimensions. Given \mathcal{T}_h , the Discontinuous Galerkin space $V_h^p \subset L^2(\Omega)$ is the subspace of piecewise polynomials of degree p

$$V_h^p = \{v \in L^2(\Omega) : \forall T \in \mathcal{T}_h, v|_T = q \circ \mu_T^{-1} \text{ with } q \in \mathbb{Q}_p^d\}. \quad (5)$$

Here $\mathbb{Q}_p^d = \{P : P(x_1, \dots, x_d) = \sum_{0 \leq \alpha_1, \dots, \alpha_d \leq p} c_{\alpha_1, \dots, \alpha_d} x_1^{\alpha_1} \dots x_d^{\alpha_d}\}$ is the set of polynomials of at most degree p . It is worth stressing that, in contrast to conforming discretisations, functions in V_h^p are not necessarily continuous between cells. We also define the L^2 inner product of two (possibly vector-valued) functions on any domain Q as

$$(v, w)_Q := \int_Q v \cdot w \, dx. \quad (6)$$

For a function $u \in V_h^p$ the discrete weak form of (3) and (4) is given by

$$a_h(u, v) = \ell_h(v) \quad \forall v \in V_h^p \quad (7)$$

with the bilinear form

$$a_h(u, v) = a_h^v(u, v) + a_h^{\text{if}}(u, v) + a_h^{\text{bf}}(u, v) \quad (8)$$

which has been split into a volume (v), interface (if) and boundary (b) term defined by

$$\begin{aligned}
a_h^v(u, v) &= \sum_{T \in \mathcal{T}_h} [(K \nabla u - \mathbf{b}u, \nabla v)_T + (cu, v)_T] \\
a_h^{\text{if}}(u, v) &= \sum_{F \in \mathcal{F}_h^i} [(\Phi(u^-, u^+, \boldsymbol{\nu}_F \cdot \mathbf{b}), \llbracket v \rrbracket)_F - (\boldsymbol{\nu}_F \cdot \{K \nabla u\}_\omega, \llbracket v \rrbracket)_F - (\boldsymbol{\nu}_F \cdot \{K \nabla v\}_\omega, \llbracket u \rrbracket)_F + \gamma_F (\llbracket u \rrbracket, \llbracket v \rrbracket)_F] \\
a_h^{\text{bf}}(u, v) &= \sum_{F \in \mathcal{F}_h^D} [(\Phi(u, 0, \boldsymbol{\nu}_F \cdot \mathbf{b}), v)_F - (\boldsymbol{\nu}_F \cdot (K \nabla u), v)_F - (\boldsymbol{\nu}_F \cdot K \nabla v, u)_F + \gamma_F (u, v)_F].
\end{aligned} \tag{9}$$

The right hand side functional is

$$\ell_h(v) = \sum_{T \in \mathcal{T}_h} (f, v)_T - \sum_{F \in \mathcal{F}_h^N} (j, v)_F - \sum_{F \in \mathcal{F}_h^D} [(\Phi(0, g, \boldsymbol{\nu}_F \cdot \mathbf{b}), v)_F + (\boldsymbol{\nu}_F \cdot (K \nabla v), g)_F - \gamma_F (g, v)_F]. \tag{10}$$

In those expressions the set of interior faces is denoted as \mathcal{F}_h^i . The sets of faces on the Neumann- and Dirichlet boundary are \mathcal{F}_h^N and \mathcal{F}_h^D respectively. With each face $F \in \mathcal{F}_h = \mathcal{F}_h^i \cup \mathcal{F}_h^N \cup \mathcal{F}_h^D$ we associate an oriented unit normal vector $\boldsymbol{\nu}_F$. For any point $\mathbf{x} \in F \in \mathcal{F}_h^i$ on an internal face we define the jump

$$\llbracket v \rrbracket(\mathbf{x}) = v^-(\mathbf{x}) - v^+(\mathbf{x}) \tag{11}$$

and the weighted average

$$\{v\}_\omega(\mathbf{x}) = \omega^-(\mathbf{x})v^-(\mathbf{x}) + \omega^+(\mathbf{x})v^+(\mathbf{x}) \quad \text{for weights} \quad \omega^-(\mathbf{x}) + \omega^+(\mathbf{x}) = 1, \omega^\pm(\mathbf{x}) \geq 0 \tag{12}$$

where $v^\pm = \lim_{\epsilon \downarrow 0} v(\mathbf{x} \pm \epsilon \boldsymbol{\nu}_F)$ is the value of the field on one side of the facet. In the expressions $\ell_h(v)$ and $a_h^{\text{bf}}(u, v)$ we implicitly assume that u and v are evaluated on the interior side of the boundary facets. To account for strongly varying diffusion fields, we choose the weights introduced in [39]:

$$\omega^\pm(\mathbf{x}) = \frac{\delta_{K\nu}^\mp(\mathbf{x})}{\delta_{K\nu}^-(\mathbf{x}) + \delta_{K\nu}^+(\mathbf{x})}, \quad \text{with} \quad \delta_{K\nu}^\pm(\mathbf{x}) = \nu^T(\mathbf{x})K^\pm(\mathbf{x})\nu(\mathbf{x}). \tag{13}$$

Defining the harmonic average of two numbers as $\langle a, b \rangle = 2ab/(a + b)$, the penalty factor γ_F is chosen based on combination of choices in [37, 38, 39] as in [28]

$$\gamma_F = \alpha \cdot p(p + d - 1) \cdot \begin{cases} \frac{\langle \delta_{K\nu_F}^-, \delta_{K\nu_F}^+ \rangle |F|}{\min(|T^-(F)|, |T^+(F)|)} & \text{for } F \in \mathcal{F}_h^i \\ \frac{\delta_{K\nu_F}^- |F|}{|T^-(F)|} & \text{for } F \in \mathcal{F}_h^D. \end{cases} \tag{14}$$

In this expression α is a free parameter chosen to be $\alpha = 1.25$ in all the computations reported below. To discretise the advection terms, we use the upwind flux on the face F which is given by

$$\Phi(u^-, u^+, b_\nu) = \begin{cases} b_\nu u^- & \text{if } b_\nu \geq 0 \\ b_\nu u^+ & \text{otherwise} \end{cases}.$$

While the formulation in (8), (10) is valid on any grid, here we only consider grids based on hexahedral elements. To use sum-factorisation techniques, we also assume that the basis functions on the reference element as well as the quadrature rules have tensor-product structure. Similar techniques can be used on simplicial elements [40, 41]. A basis $\Psi_h = \{\psi_1^p, \dots, \psi_{N_h}^p\}$ is chosen for V_h^p , i.e. every function $u \in V_h^p$ can be written as

$$u(\mathbf{x}) = \sum_{i \in I} u_i \psi_i^p(\mathbf{x}), \quad I = \{1, \dots, N_h\} \subset \mathbb{N}. \tag{15}$$

Here N_h is the total number of degrees of freedom and $\mathbf{u} = (u_1, u_2, \dots, u_{N_h})^T \in \mathbb{R}^n$ is the vector of unknowns. With this basis the weak formulation in (7) is equivalent to a matrix equation

$$\mathbf{A}\mathbf{u} = \mathbf{f} \quad \text{with} \quad A_{ij} = a_h(\psi_j^p, \psi_i^p) \quad \text{and} \quad \mathbf{f} \in \mathbb{R}^n \quad \text{with} \quad f_i := \ell_h(\psi_i^p). \tag{16}$$

It is this equation which we will solve in the following. Since we consider a discontinuous function space, the basis can be chosen such that the support of any basis function $\psi_i^p \in \Psi_h$ is restricted to a single element $T \in \mathcal{T}_h$ and this implies that the matrix A is block-structured. This property is important for the algorithms developed in the following. Throughout this work we use a basis which is constructed from the tensor-product of one-dimensional Gauss-Lobatto basis functions (i.e. Lagrange polynomials with nodes at the Gauss-Lobatto points) on each grid cell. We also assume that the basis is nodal, i.e. there is a set of points ζ_j , $j \in I$ such that $\psi_i^p(\zeta_j) = \delta_{ij}$.

Block notation of linear systems. To efficiently deal with block-structured matrices we introduce the following notation. For any finite index set $I \subset \mathbb{N}$ we define the vector space \mathbb{R}^I to be isomorphic to $\mathbb{R}^{|I|}$ with components indexed by $i \in I$. Thus any vector of unknowns $\mathbf{u} \in \mathbb{R}^I$ can be interpreted as a mapping $\mathbf{u} : I \rightarrow \mathbb{R}$ and $\mathbf{u}(i) = u_i$. In the same way, for any two finite index sets $I, J \subset \mathbb{N}$ we write $A \in \mathbb{R}^{I \times J}$ with the interpretation $A : I \times J \rightarrow \mathbb{R}$ and $A(i, j) = A_{i,j}$. Finally, for any subset $I' \subseteq I$ we define the restriction matrix $R_{I,I'} : \mathbb{R}^I \rightarrow \mathbb{R}^{I'}$ as $(R_{I,I'} \mathbf{u})_i = u_i \forall i \in I'$. Now, let $I_T = \{i \in \mathbb{N} : \text{supp } \psi_i^p \subseteq T\}$ denote the subset of indices which are associated with basis functions that are nonzero on element T . Then

$$\bigcup_{T \in \mathcal{T}_h} I_T = I = \{1, \dots, n\} \quad \text{and} \quad I_T \cap I_{T'} = \emptyset, \quad \forall T \neq T', \quad (17)$$

where the last property follows from the fact that we consider a discontinuous function space and therefore $\{I_T\}$ forms a disjoint partitioning of the index set I . Using this partitioning and imposing an ordering $\mathcal{T}_h = \{T_1, \dots, T_m\}$ on the mesh elements, the linear system in (16) can be written in block form

$$\begin{pmatrix} A_{T_1, T_1} & \dots & A_{T_1, T_m} \\ \vdots & \ddots & \vdots \\ A_{T_m, T_1} & \dots & A_{T_m, T_m} \end{pmatrix} \begin{pmatrix} \mathbf{u}_{T_1} \\ \vdots \\ \mathbf{u}_{T_m} \end{pmatrix} = \begin{pmatrix} \mathbf{f}_{T_1} \\ \vdots \\ \mathbf{f}_{T_m} \end{pmatrix} \quad (18)$$

where $A_{T_\rho, T_\sigma} = R_{I, I_{T_\rho}} A R_{I_{T_\sigma}}^T$, $\mathbf{u}_{T_\rho} = R_{I, I_{T_\rho}} \mathbf{u}$ and $\mathbf{f}_{T_\rho} = R_{I, I_{T_\rho}} \mathbf{f}$. For each cell T and function $u \in V_h^p$ we also define the function $u_{\cap T} \in V_h^p$ as

$$u_{\cap T}(x) = \begin{cases} u(x) & \text{for } x \in T \\ 0 & \text{otherwise} \end{cases} \quad \Leftrightarrow \quad u_{\cap T} = \sum_{i \in I_T} (u_T)_i \psi_i^p = \sum_{i \in I_T} u_i \psi_i^p. \quad (19)$$

Our choice of Gauss-Lobatto basis functions with non-overlapping support leads to the so called *minimal stencil*; the matrices A_{T_ρ, T_σ} are only nonzero if $\rho = \sigma$ or if the elements T_ρ and T_σ share a face, and hence the matrix in (18) is block-sparse. In general and for the basis considered in this paper, the $(p+1)^d \times (p+1)^d$ matrix A_{T_ρ, T_σ} is dense.

This partitioning of A and the minimal stencil property form are key for the matrix-free preconditioner implementation discussed below.

2.2. Hybrid multigrid preconditioner

To solve the linear system in (16) we use the hybrid multigrid preconditioner described in [28]. However, in contrast to [28], the computationally most expensive components, namely the operator- and preconditioner application in the DG space V_h^p , are implemented in a matrix free way. As the numerical results in Section 4 confirm, this leads to significant speedups at higher polynomial orders p and has benefits even for relatively modest orders of $p = 2, 3$.

The key idea in [28] is to combine a block-smoother in the high-order DG space V_h^p with an AMG correction in a low-order subspace $\hat{V}_h \subset V_h$. In this work we will consider low order subspaces \hat{V}_h spanned by the piecewise constant P_0 and conforming piecewise linear elements Q_1 . One V-cycle of the multigrid algorithm is shown in Algorithm 1. At high polynomial orders p the computationally most expensive steps are the operator application in the defect (or residual) calculation and the pre-/post-smoothing on the DG level. Usually the multigrid algorithm is used as a preconditioner of a Krylov subspace method, which requires additional operator applications. To solve the problem in the lower-order subspace we use the aggregation based AMG solver described in [42, 43]. In the following we discuss the efficient implementation of the individual components in Algorithm 1:

Algorithm 1 Hybrid multigrid V-cycle. Input: Right hand side \mathbf{f} , initial guess \mathbf{u}_0 (often set to zero); Output: \mathbf{u} , the approximate solution to $\mathbf{A}\mathbf{u} = \mathbf{f}$

- 1: $\mathbf{u} \leftarrow S(\mathbf{f}, \mathbf{u}_0; n_{\text{pre}}, \omega)$ $\{\text{Presmooth with } n_{\text{pre}} \text{ iterations in DG space}\}$
 - 2: $\mathbf{r} \leftarrow \mathbf{f} - \mathbf{A}\mathbf{u}$ $\{\text{Calculate residual}\}$
 - 3: $\hat{\mathbf{r}} \leftarrow R(\mathbf{r})$ $\{\text{Restrict to subspace}\}$
 - 4: $\hat{\mathbf{u}} \leftarrow \text{AMG}(\hat{\mathbf{r}}) \approx \hat{\mathbf{A}}^{-1}\hat{\mathbf{r}}$ $\{\text{AMG V-cycle on subspace fields}\}$
 - 5: $\mathbf{u} \leftarrow \mathbf{u} + P(\hat{\mathbf{u}})$ $\{\text{Prolongate and add to DG solution}\}$
 - 6: $\mathbf{u} \leftarrow S(\mathbf{f}, \mathbf{u}; n_{\text{post}}, \omega)$ $\{\text{Postsmooth with } n_{\text{post}} \text{ iterations in DG space}\}$
-

1. Matrix-free operator application in the DG space to calculate the residual (line 2)
2. Matrix-free pre-/post-smoothing in the DG space (lines 1 and 6)
3. Transfer of grid functions between between DG space and low order subspace (lines 3 and 5)
4. Assembly of the low order subspace matrix which is used in the AMG V-cycle (line 4)

The matrix-free operator application based on sum factorisation techniques has been previously described in [24] (and is briefly reviewed in Section 3.1 below). Here we will concentrate on the other components.

2.2.1. Matrix-free smoothers

In the context of multigrid methods, iterative improvement of the solution with a stationary method is referred to as *smoothing* since a small number of iterations with a simple method reduces high-frequency error components. For a linear equation $\mathbf{A}\mathbf{u} = \mathbf{f}$, this process can be summarised as follows: given a matrix $W \approx A$, calculate the defect $\mathbf{d} = \mathbf{f} - \mathbf{A}\mathbf{u}^{(k-1)}$ for some guess $\mathbf{u}^{(k-1)}$ and solve for $\delta\mathbf{u}$ such that

$$W\delta\mathbf{u} = \mathbf{d}. \quad (20)$$

This correction $\delta\mathbf{u}$ can then be used to obtain an improved solution

$$\mathbf{u}^{(k)} = \mathbf{u}^{(k-1)} + \delta\mathbf{u} = \mathbf{u}^{(k-1)} + W^{-1}(\mathbf{f} - \mathbf{A}\mathbf{u}^{(k-1)}). \quad (21)$$

Starting from some initial guess $\mathbf{u}^{(0)}$, the iterates $\mathbf{u}^{(0)}, \mathbf{u}^{(1)}, \dots, \mathbf{u}^{(k)}$ form an increasingly better approximation to the exact solution \mathbf{u} of $\mathbf{A}\mathbf{u} = \mathbf{f}$ for a convergent method. Choosing a block-smoother guarantees that any high-frequency oscillations inside one cell are treated exactly, which is particularly important for higher polynomial orders. Lower-frequency errors with variations between the grid cells are reduced by the AMG preconditioner in the low-order subspace (line 4 in Alg. 1). Following the block partitioning in (18), let $A = D + L + U$ be decomposed into the strictly lower block triangular part L , the block diagonal D and the strictly upper block triangular part U with

$$D_{T_\rho, T_\sigma} = \begin{cases} D_{T_\rho} = A_{T_\rho, T_\rho} & \text{for } \rho = \sigma \\ 0 & \text{otherwise,} \end{cases} \quad L_{T_\rho, T_\sigma} = \begin{cases} A_{T_\rho, T_\sigma} & \text{for } \rho > \sigma \\ 0 & \text{otherwise,} \end{cases} \quad U_{T_\rho, T_\sigma} = \begin{cases} A_{T_\rho, T_\sigma} & \text{for } \rho < \sigma \\ 0 & \text{otherwise.} \end{cases} \quad (22)$$

In this work we consider the following choices for the block preconditioner W : (1) block-Jacobi $W = \omega^{-1}D$, (2) successive block over-relaxation (block-SOR) $W = \omega^{-1}D + L$ and (3) symmetric block-SOR (block-SSOR) $W = (\omega(2-\omega))^{-1}(D+\omega L)D^{-1}(D+\omega U)$. The over-relaxation parameter ω can be tuned to accelerate convergence. The explicit form of the block-Jacobi and block-SOR algorithms are written down in Algorithms 2 and 3. Note that for block-SOR, the field $\mathbf{u}^{(k)}$ can be obtained by transforming $\mathbf{u}^{(k-1)}$ in-place, which reduces storage requirements. The corresponding backward-SOR sweep is obtained by reversing the direction of the grid traversal. It can then be shown that one application of the block-SSOR smoother is equivalent to a forward-block-SOR sweep following by a backward-block-SOR iteration. Note that in all cases the computationally most expensive operations are the multiplication with the dense $(p+1)^d \times (p+1)^d$ matrices A_{T_ρ, T_σ} and the solution of the dense system $D_{T_\rho} \delta\mathbf{u}_{T_\rho} = \mathbf{d}_{T_\rho}$ in each cell. As will be described in Section 3, the defect can be calculated in a matrix-free way by using the techniques from [24]. The standard approach to calculating the correction $\delta\mathbf{u}_{T_\rho}$ is to assemble the local matrices D_{T_ρ} , factorise them with an LU-/Cholesky-approach and then use this to solve the linear system. However, the costs for the factorisation- and back-substitution steps grow like $(p+1)^{3d}$ and $(p+1)^{2d}$

Algorithm 2 Block-Jacobi smoother. Input: Right hand side \mathbf{f} , initial guess $\mathbf{u}^{(0)}$, number of smoothing steps n_{smooth} . Output: Approximate solution $\mathbf{u}^{(n_{\text{smooth}})}$

- 1: **for** $k = 1, \dots, n_{\text{smooth}}$ **do**
 - 2: Calculate defect $\mathbf{d} = \mathbf{f} - A\mathbf{u}^{(k-1)}$
 - 3: Solve $D_T \delta \mathbf{u}_T = \mathbf{d}_T$ for $\delta \mathbf{u}_T$ on all grid elements T
 - 4: Update solution $\mathbf{u}^{(k)} = \mathbf{u}^{(k-1)} + \omega \delta \mathbf{u}$
 - 5: **end for**
-

Algorithm 3 Successive block-over-relaxation, forward Block-SOR smoother. Input: Right hand side \mathbf{f} , initial guess $\mathbf{u}^{(0)}$, number of smoothing steps n_{smooth} . Output: Approximate solution $\mathbf{u}^{(n_{\text{smooth}})}$

- 1: **for** $k = 1, \dots, n_{\text{smooth}}$ **do**
- 2: **for** all grid elements T_ρ with $\rho = 1, \dots, m$ **do**
- 3: Calculate local defect

$$\mathbf{d}_{T_\rho} = \mathbf{f}_{T_\rho} - \sum_{\substack{\sigma < \rho \\ \partial T_\sigma \cap \partial T_\rho \neq \emptyset}} A_{T_\rho, T_\sigma} \mathbf{u}_{T_\sigma}^{(k)} - \sum_{\substack{\sigma > \rho \\ \partial T_\sigma \cap \partial T_\rho \neq \emptyset}} A_{T_\rho, T_\sigma} \mathbf{u}_{T_\sigma}^{(k-1)} \quad (23)$$

- 4: Solve $D_{T_\rho} \delta \mathbf{u}_{T_\rho} = \mathbf{d}_{T_\rho}$ for $\delta \mathbf{u}_{T_\rho}$
 - 5: Update solution $\mathbf{u}_{T_\rho}^{(k)} = (1 - \omega) \mathbf{u}_{T_\rho}^{(k-1)} + \omega \delta \mathbf{u}_{T_\rho}$
 - 6: **end for**
 - 7: **end for**
-

respectively, and hence this method becomes prohibitively expensive for high polynomial degrees p . To avoid this, we solve the block-system

$$D_{T_\rho} \delta \mathbf{u}_{T_\rho} = \mathbf{d}_{T_\rho} \quad (24)$$

with a Krylov-subspace method such as the Conjugate Gradient (CG) [44] or Generalised Minimal Residual (GMRES) [45] iteration. This only requires operator applications which can be implemented in a matrix-free way. If sum-factorisation techniques are used, the overall cost of this inversion is $\mathcal{O}(n_{\text{iter}}(\epsilon) \cdot dp^{d+1})$ where $n_{\text{iter}}(\epsilon)$ is the number of iterations required to solve the block-system. An important observation, which drastically effects the practical usage of the matrix-free smoothers, is that it suffices to compute an inexact solution of (24) with the Krylov subspace method. For this we iterate until the two-norm of the residual $\|\mathbf{d}_{T_\rho} - D_{T_\rho} \delta \mathbf{u}_{T_\rho}\|_2$ has been reduced by a fixed factor $\epsilon \ll 1$. The resulting reduction in n_{iter} and preconditioner cost has to be balanced against the potential degradation in the efficiency of the preconditioner. This dependence on ϵ on the overall solution time will be studied below.

Using block smoothers, the method proposed here is not strictly robust with respect to the polynomial degree p . This can be remedied by using overlapping block smoothers [42, 28] where the blocks originate from an overlapping partitioning of the mesh. However, this also increases the computational cost substantially. Since in practical applications we consider moderate polynomial degree anyway, we concentrate on simple non-overlapping block smoothers here where the blocks originate from a single element.

2.2.2. Intergrid operators

Since we use nested function spaces with $\hat{V}_h \subset V_h^p$, injection is the natural choice for the prolongation $P : \hat{V}_h \rightarrow V_h^p$. Given a function $\hat{u} \in \hat{V}_h$, we define

$$P : \hat{u} \mapsto u \in V_h^p \quad \text{such that } u(\mathbf{x}) = \hat{u}(\mathbf{x}) \text{ for all } \mathbf{x} \in \Omega. \quad (25)$$

Expanding in basis functions on element T and using (19) this leads to

$$\hat{u}_{\cap T} = \sum_{i \in \hat{I}_T} \hat{u}_i \hat{\psi}_i = \sum_{j \in I_T} u_j \psi_j^p = u_{\cap T}. \quad (26)$$

Here \hat{I}_T is the set of indices i for which the coarse basis function $\hat{\psi}_i(x)$ is non-zero on element T . For a general basis, calculation of u_j requires solution of a small system involving the local $(p+1)^d \times (p+1)^d$ mass matrix $\int_T \psi_i^p(\mathbf{x}) \psi_j^p(\mathbf{x}) dx$ in each cell T ; this is expensive for higher polynomial degrees p . However, here we only consider the special case that \hat{V}_h is a subspace of V_h^p and the basis is nodal. In this case the unknowns in the higher order space can be found much more efficiently by evaluating the low order basis functions at the nodal points ζ_i of the higher order space,

$$u_i = \sum_{j \in \hat{I}_T} \hat{u}_j \hat{\psi}_j(\zeta_i) \quad \text{for all } i \in I_T. \quad (27)$$

In other words, the j -th column of the local prolongation matrix $P|_T$ can be calculated by evaluating $\hat{\psi}_j$ at all nodal points,

$$(P|_T)_{ij} = \hat{\psi}_j(\zeta_i) \quad \text{for } i \in I_T \text{ and } j \in \hat{I}_T. \quad (28)$$

Since the DG basis functions form a partition of unity, for the P_0 subspace this simplifies even further to

$$P|_T = (1, \dots, 1)^T. \quad (29)$$

For the restriction operation $R : V_h^p \rightarrow \hat{V}_h$ we choose the transpose of the prolongation, i.e. $R = P^T$. In the absence of advection ($\mathbf{b} = 0$) this choice preserves the symmetry of the system and the WSIPG discretisation.

2.2.3. Low order subspace matrix

The low order subspace matrix \hat{A} is needed to solve the “coarse” grid correction with the AMG preconditioner. In general this matrix is given by the Galerkin product

$$\hat{A} = RAP = P^T AP. \quad (30)$$

However, calculating \hat{A} according to this formula is expensive for several reasons: first of all it requires two matrix-matrix multiplications of large matrices with different sparsity patterns. In addition, the matrix A in the DG space has to be assembled, even though in a matrix-free implementation it is never used in the DG smoother. Furthermore, for our choice of coarse function spaces, the matrix \hat{A} has lots of entries which are formally zero. However, if it is calculated according to (30) in an actual implementation those entries will not vanish exactly due to rounding errors. This leads to drastically increased storage requirements and unnecessary calculations whenever \hat{A} is used in the AMG preconditioner. All those issues can be avoided by directly assembling the coarse grid matrix \hat{A} as follows:

For the P_0 subspace it is shown in [Appendix A.1](#) that on an equidistant grid the matrix entries \hat{A}_{ij} can be calculated directly by discretising (3) with a modified finite volume scheme in which (i) all flux terms are multiplied by a factor $\alpha p(p+d-1)$ and (ii) the boundary flux is scaled by an additional factor 2. For the conforming Q_1 subspace, all contributions to \hat{A} which arise from jump terms in $A_{ij} = a_h(\psi_j^p, \psi_i^p)$ vanish. In fact, up to an additional Dirichlet boundary term $\sum_{F \in \mathcal{F}_h^D} \gamma_F \left(\hat{\psi}_i, \hat{\psi}_j \right)_F$, the matrix \hat{A}_{ij} is identical to the matrix obtained from a conforming Q_1 discretisation of (3), and it is this matrix that we use in our implementation. In both cases the sparsity pattern of \hat{A} is the same as that obtained by a discretisation in the low-order subspace. In summary this re-discretisation in the coarse level subspace instead of using the Galerkin product in (30) results in a significantly reduced sparsity pattern of the matrix \hat{A} and avoids explicit assembly of the large DG matrix A .

3. Implementation in EXADUNE

The crucial ingredient for the development of the matrix-free block smoothers described in [Section 2.2.1](#) is the efficient on-the-fly operator application of the full operator A and multiplication with the block-diagonal D_T for each element $T \in \mathcal{T}_h$. While the operator application is required in the calculation of the local defect, the frequent multiplication with D_T is necessary in the iterative inversion of the diagonal blocks. Our code is implemented in the EXADUNE [\[35\]](#) fork of the DUNE framework [\[46, 34\]](#). EXADUNE contains several optimisations for adapting DUNE to modern computer architectures.

3.1. Matrix-free operator application

The matrix-free application of the operator A itself has been implemented and heavily optimised in [24], and here we only summarise the most important ideas. Recall that the weak form $a_h(u, v)$ in (8) is split into volume-, interface- and boundary terms defined in (9). The local contributions to the system matrix A are evaluated in a `LocalOperator` class in DUNE. More specifically, let T^+, T^- be a pair of adjacent cells which share an internal face $F^i = \partial T^+(F^i) \cap \partial T^-(F^i)$ and let $F^b \in \partial \hat{T}(F^b) \cap \mathcal{F}_h^b$ be the face of a cell T adjacent to the boundary of the domain. For each cell T , internal interface F^i and boundary face F^b the `LocalOperator` class provides methods for evaluating the *local* operator application

$$\begin{aligned} (A_{T,T}^v \mathbf{u}_T)_{i_T} &:= a_h^v(\mathbf{u}_{\cap T}, \psi_{i_T}^p) & \text{with } i_T \in I_T & \quad (\text{volume}) \\ (A_{F^i; T', T''}^{\text{if}} \mathbf{u}_{T''})_{i_{T'}} &:= a_h^{\text{if}}(\mathbf{u}_{\cap T''}, \psi_{i_{T'}}^p) & \text{with } i_{T'} \in I_{T'} & \quad (\text{interface}) \\ (A_{F^b; \hat{T}}^b \mathbf{u}_T)_{i_{\hat{T}}} &:= a_h^b(\mathbf{u}_{\cap T}, \psi_{i_{\hat{T}}}^p) & \text{with } i_{\hat{T}} \in I_{\hat{T}} & \quad (\text{boundary}). \end{aligned} \quad (31)$$

For the interface term we calculate four contributions, one for each of the combinations $(T', T'') = (T^-, T^-), (T^-, T^+), (T^+, T^-)$ and (T^+, T^+) . By iterating over all elements and faces of the grid and calling the corresponding local operator evaluations, the full operator can be applied to any field $u \in V_h^p$. This iteration over the grid is carried out by an instance of a `GridOperator` class, which takes the `LocalOperator` as template parameter.

In the following we briefly recapitulate the key ideas used to optimise the matrix-free operator application with sum-factorisation techniques and concentrate on application of the operator $A_{T,T}^v$. In each cell T this proceeds in three stages (see Algorithm 1 in [24] for details)

1. Given the coefficient vector \mathbf{u} , calculate the values of the field u and its gradient at the quadrature points on the reference element \hat{T} .
2. At each quadrature point, evaluate the coefficients K , \mathbf{b} and c as well as geometric factors from the mapping between T and \hat{T} . Using this and the function values from Stage 1, evaluate the weak form $a_h(\cdot, \cdot)$ at each quadrature point.
3. Multiply by test functions (and their derivatives) to recover the entries of the coefficient vector $\mathbf{v} = A\mathbf{u}$.

We now discuss the computational complexity of the individual stages.

Stage 1. We assume that the basis functions in the reference element \hat{T} have a tensor-product structure

$$\hat{\phi}_{\mathbf{j}}(\hat{\mathbf{x}}) = \hat{\phi}_{(j_1, \dots, j_d)}(\hat{x}_1, \dots, \hat{x}_d) = \prod_{k=1}^d \hat{\theta}_{j_k}^{(k)}(\hat{x}_k). \quad (32)$$

Those basis functions are enumerated by d -tuples $\mathbf{j} = (j_1, \dots, j_d) \in \mathbf{J} = J^{(1)} \times \dots \times J^{(d)}$, $J^{(k)} = \{1, \dots, n_k\}$. For each cell T we have an index map $g(T, \cdot) \rightarrow I$ which assigns a global index to each d -tuple \mathbf{j} in the cell; this implies that in a particular cell T any function $u \in V_h^p$ can be written as

$$u(\mathbf{x}) = \hat{u}(\hat{\mathbf{x}}) = \sum_{\mathbf{j} \in \mathbf{J}} u_{g(T, \mathbf{j})} \hat{\phi}_{\mathbf{j}}(\hat{\mathbf{x}}) \quad \text{for } \mathbf{x} = \mu_T(\hat{\mathbf{x}}) \in T. \quad (33)$$

We use a tensor-product quadrature rule with points

$$\boldsymbol{\xi}_{\mathbf{i}} = \left(\xi_{i_1}^{(1)}, \dots, \xi_{i_d}^{(d)} \right) \quad (34)$$

enumerated by d -tuples $\mathbf{i} = (i_1, \dots, i_d) \in I^{(1)} \times \dots \times I^{(d)}$, $I^{(k)} = \{1, \dots, m_k\}$. In the following we only consider $n_1 = \dots = n_d =: n$ and $m_1 = \dots = m_d =: m$, in this case that there are m^d quadrature points and n^d basis functions per cell. Following Eq. (13) in [24], the evaluation of the function u at the quadrature point $\boldsymbol{\xi}_{\mathbf{i}}$ can be written as

$$\hat{u}(\boldsymbol{\xi}_{\mathbf{i}}) = \sum_{\mathbf{j} \in \mathbf{J}} u_{g(T, \mathbf{j})} \hat{\phi}_{\mathbf{j}}(\boldsymbol{\xi}_{\mathbf{i}}) = \sum_{j_1=1}^n \dots \sum_{j_d=1}^n A_{i_1, j_1}^{(1)} \dots A_{i_d, j_d}^{(d)} u_{g(T, (j_1, \dots, j_d))}. \quad (35)$$

The entries of the $m \times n$ matrices $A^{(q)}$ are given by the one-dimensional shape functions evaluated at the quadrature points, $A_{\alpha,\beta}^{(k)} = \theta_{\beta}^{(k)}(\xi_{\alpha}^{(k)})$, $\alpha \in I^{(k)}$, $\beta \in J^{(k)}$. (35) describes the multiplication of a vector of length n^d by a $m^d \times n^d$ matrix $\hat{\Phi}$ with $\hat{\Phi}_{i,j} := \hat{\phi}_j(\xi_i) = A_{i_1,j_1}^{(1)} \cdots A_{i_d,j_d}^{(d)}$. Assuming that the ratio $\rho = n/m \leq 1$ is constant, the naive cost of this operation is $\mathcal{O}((d+1)m^{2d})$ FLOPs. A significant reduction in computational complexity is achieved by applying the small $m \times n$ matrices $A^{(k)}$ recursively in d stages as

$$\begin{aligned} u_{(j_1,\dots,j_d)}^{(0)} &= u_{g(T,(j_1,\dots,j_d))} \\ u_{(i_1,\dots,i_s,j_{s+1},\dots,j_d)}^{(s)} &= \sum_{j_s=1}^n A_{i_s,j_s}^{(s)} u_{(i_1,\dots,i_{s-1},j_s,\dots,j_d)}^{(s-1)} \quad \text{for } s = 1, \dots, d \\ \hat{u}(\xi_i) &= u_{(i_1,\dots,i_d)}^{(d)}. \end{aligned} \quad (36)$$

Each step in (36) requires the multiplication with an $m \times n$ matrix $A_{i_s,j_s}^{(s)}$ for $m^{s-1}n^{d-s} \leq m^{d-1}$ vectors. Multiplication by an $m \times n$ matrix requires $2n \times m$ operations and hence the total cost is reduced to $\mathcal{O}(d \cdot m^{d+1})$. Since m grows with the polynomial degree (typically $m = n = p + 1$), for high polynomial degrees p this sum-factorised approach is much faster than the naive algorithm with complexity $\mathcal{O}((d+1)m^{2d})$. Gradients of u can be evaluated in the same way with slightly different matrices $A^{(s)}$.

The face integrals in the bilinear form require the evaluation on points of a quadrature rule of lower dimension $d - 1$. Although this is complicated by the different embeddings of a face into the neighbouring cells, the same sum factorisation techniques can be applied. While for those terms the cost is only $\mathcal{O}(d \cdot m^d)$, for low polynomial degrees the absolute cost can still be larger than the cost of the volume terms (see Fig. 1 in [24]). We conclude that - using sum factorisation - the cost of the first stage in the matrix-free operator application is $\mathcal{O}(d \cdot m^{d+1})$.

Stage 2. Since the operations can be carried out independently at the m^d quadrature points, the cost of this stage is $\mathcal{O}(m^d)$.

Stage 3. Given the values $z_{(i_1,\dots,i_d)} := \hat{u}(\xi_i)$ at the quadrature points from Stage 2, the entries of the coefficient vector $\mathbf{v} = \mathbf{A}\mathbf{u}$ can be calculated as

$$v_{g(T,(j_1,\dots,j_d))} = \sum_{i_1=1}^m \cdots \sum_{i_d=1}^m A_{j_1,i_1}^{(1)} \cdots A_{j_d,i_d}^{(d)} z_{(i_1,\dots,i_d)} + \dots \quad (37)$$

Here “...” stands for terms of a similar structure which arise due to multiplication with the derivative of a test function in the bilinear form. For simplicity those terms are not written down here; an example can be found in Eq. (10) of [24]. Note that (37) is the same as (35), but we now multiply by the transpose of the $m \times n$ matrices $A^{(k)}$. Sum factorisation can be applied in exactly the same way, resulting in a computational complexity of $\mathcal{O}(d \cdot m^{d+1})$.

Combining the total cost of all stages we find that the overall complexity of the sum-factorised operator application is $\mathcal{O}(d \cdot m^{d+1}) = \mathcal{O}(d \cdot p^{d+1})$. Finally, observe that the operator application requires reading the local dof-vector which has n^d entries. A vector of the same size has to be written back at the end of the calculation. Overall, the total amount of memory moved is $\mathcal{O}(n^d)$. The resulting arithmetic intensity is $\mathcal{O}(d \cdot m) = \mathcal{O}(d \cdot p)$ and the operation is clearly FLOP-bound for large p .

In EXADUNE [35] several optimisations are applied to speed up the operator application and exploit modern FLOP-bound architectures. The C++ vector class library [47] is used for the multiplication with the small dense matrices $A^{(k)}$. The code is vectorised by grouping the value of the function and its three spatial derivatives and evaluating the four values $[\partial_{x_1}\hat{u}(\xi), \partial_{x_2}\hat{u}(\xi), \partial_{x_3}\hat{u}(\xi), \hat{u}(\xi)]$ simultaneously in (35). Efficient transfer of data from memory is important. In the standard DUNE implementation, a copy of the local dof-vector u_T is created on each element. While this approach is necessary for conforming finite element methods, for the DG method used here the local dofs of neighbouring cells do not overlap and it is therefore possible to directly operate on the global dof-vector, thus avoiding expensive and unnecessary memory copies. As reported in [24], the operator application runs at close to 60% of the peak floating point performance of an Intel Haswell CPU in some cases.

degree	#dofs per cell	grid size $n_x \times n_y \times n_z$	#dofs	matrix size	matrix vs. vector size
1	8	$128 \times 128 \times 256$	$33.6 \cdot 10^6$	14.0 GB	$56 \times$
2	27	$56 \times 56 \times 112$	$9.5 \cdot 10^6$	13.4 GB	$189 \times$
3	64	$32 \times 32 \times 64$	$4.2 \cdot 10^6$	14.0 GB	$448 \times$
4	125	$20 \times 20 \times 40$	$2.0 \cdot 10^6$	13.0 GB	$875 \times$
5	216	$14 \times 14 \times 28$	$1.2 \cdot 10^6$	13.4 GB	$1512 \times$
6	343	$10 \times 10 \times 20$	$0.7 \cdot 10^6$	12.3 GB	$2401 \times$
7	512	$8 \times 8 \times 16$	$0.5 \cdot 10^6$	14.0 GB	$3584 \times$
8	729	$6 \times 6 \times 12$	$0.3 \cdot 10^6$	12.0 GB	$5103 \times$
9	1000	$5 \times 5 \times 10$	$0.2 \cdot 10^6$	13.0 GB	$7000 \times$
10	1331	$4 \times 4 \times 8$	$0.2 \cdot 10^6$	11.8 GB	$9317 \times$

Table 1: Problem sizes for different polynomial degrees. The last column lists the relative size of the matrix and a dof-vector. See Tabs. 4, 5 and 6 below for estimates of the memory requirements in the full implementation.

3.2. Matrix-free application and inversion of block-diagonal

Since in our smoothers the block system (24) is solved iteratively, we need to be able to apply the block-diagonal part $D_T = A_{T,T}$ of the operator A in each cell T in an efficient way. For this, note that

$$D_T \mathbf{u}_T = A_{T,T}^v \mathbf{u}_T + \sum_{F^i \in \partial T \cap \mathcal{F}_h^i} A_{F^i,T}^{\text{if}} \mathbf{u}_T + \sum_{F^b \in \partial T \cap \mathcal{F}_h^b} A_{F^b,T}^b \mathbf{u}_T. \quad (38)$$

Based on the decomposition in (38), in each cell T we first evaluate the contribution $A_{T,T}^v \mathbf{u}_T$. We then loop over all faces $F \in \partial T$ of T and – depending on whether the face is interior or on the boundary – add $A_{F^i,T}^{\text{if}} \mathbf{u}_T$ or $A_{F^b,T}^b \mathbf{u}_T$.

To (approximately) solve the system $D_T \delta \mathbf{u}_T = \mathbf{d}_T$ in line 3 of Algorithm 2 and line 4 of Algorithm 3 we use a Krylov subspace method. This in turn frequently applies the block-diagonal D_T as described above. To obtain the fastest possible element-local matrix-free solver, additionally a preconditioner for D_T in the Krylov method is required. For diffusion dominated problems we find that dividing by the diagonal of D_T is efficient. For strongly convection-dominated problems which have large off-diagonal entries, however, we use a tridiagonal preconditioner based on the Thomas-algorithm [48]. While this requires explicit assembly of the matrix D_T at the beginning of the solve, we find that the overhead of this is small. Since only the (tri-) diagonal of D_T is retained, it also does not increase memory requirements. In the future the overhead could be reduced further by only assembling the (tri-) diagonal entries of D_T .

Since ultimately the applications of the operators D_T and D_T^{-1} call the methods $A_{T,T}^v$, $A_{F^i,T}^{\text{if}}$ and $A_{F^b,T}^b$ of the heavily optimised `LocalOperator` class for A (see Section 3.1) those operations will run at a similar high performance as the matrix-free application of A itself.

4. Results

Unless stated otherwise, we work in a domain of size $\Omega = [0, L_x] \times [0, L_y] \times [0, L_z]$ with $L_x = L_y = 1$ and $L_z = 2$. The problem sizes for different polynomial degrees are shown in Tab. 1. For each degree, the number of grid cells is chosen such that all cells have the same size (i.e. the grid is isotropic) and the resulting matrix has a size of 11 – 14 GB (see Section 4.4 for a detailed discussion of memory requirements). All runs are carried out on 16 cores of an Intel Xeon E5-2698v3 (Haswell, 2.30GHz) node of the “donkey” cluster at Heidelberg University. For the problems in subsections 4.2 and 4.3 the processor layout is chosen as $2 \times 2 \times 4$, i.e. each core works on a domain of the same size and there are no load-imbalances. For the anisotropic SPE10 benchmark in subsection 4.5 the processor layout is changed to $4 \times 4 \times 1$ to avoid parallel partitioning in the strongly coupled vertical direction. The code was compiled with version 6.1 of the Gnu C++ compiler.

4.1. Solver Implementations

To quantify gains in performance from the techniques described above we use the following three implementations of the hybrid multigrid algorithm in Section 2.2:

1. **Matrix-free (MF)**. Both the DG operator and the DG smoother are applied in an entirely matrix-free way exploiting the sum-factorisation techniques discussed in Section 3. In the smoother the diagonal matrices D_T in each cell are inverted iteratively to tolerance ϵ with a matrix-free iterative solver as described in Section 2.2.1. In contrast to the following two implementations, storage requirements are drastically reduced in this matrix-free version of the code since it is not necessary to store the assembled DG matrix at all. The matrix \hat{A} in the lower-order subspace is assembled directly as described in Section 2.2.3, leading to further performance improvements in the setup phase (see results in Appendix B).
2. **Matrix-explicit (MX)**. As a reference implementation we follow the standard approach of assembling the DG matrix A explicitly and storing the Cholesky-/LU- factors of the diagonal blocks. This matrix and the Cholesky-/LU-factors are then used for the application of operator and smoother in the DG space. To allow for a fair comparison we speed up the assembly process with sum-factorisation. In the setup phase we calculate the coarse grid matrix \hat{A} in the lower order subspace via a Galerkin product $\hat{A} = P^T A P$.
3. **Partially matrix-free (PMF)**. To quantify any gains from the matrix-free inversion of the diagonal blocks, we also store the Cholesky-/LU- factorisation of those blocks in the assembled matrix but apply the DG operator itself in a matrix-free way as described in Section 3.1. In the smoother the blocks are inverted directly by using the pre-factorised block-matrices in each cell. As in the matrix-free implementation the matrix \hat{A} is assembled directly. While the storage requirements are reduced by a factor $2d + 1$ compared to the matrix-explicit code, it is still necessary to store the (factorised) diagonal blocks in each grid cell.

4.2. Diffusion equation

We first solve two purely diffusive problems on a regular grid in $d = 3$ dimensions. For the Poisson equation $-\Delta u = f$ the diffusion tensor is the identity matrix and both the advection vector and the zero order reaction term vanish. Homogeneous Dirichlet boundary conditions are chosen on $\partial\Omega$.

While the Poisson equation is an important model problem, the fact that the diffusion tensor is constant and diagonal simplifies the on-the-fly operator evaluation and might give an unrealistic advantage to the matrix-free implementation; in particular it is not necessary to calculate an 3×3 diffusion tensor at every quadrature point. To avoid this we also include a computationally more challenging test case of the form $-\nabla \cdot (K(\mathbf{x})\nabla u) + c(\mathbf{x})u = f$ with a spatially varying positive definite diffusion tensor and (small) zero order term given by

$$K(\mathbf{x}) = \sum_{k=1}^d P^{(k)}(x_k) \mathbf{n}_k \mathbf{n}_k^T, \quad c(\mathbf{x}) = \gamma \sum_{k=1}^d x_k^2 \quad \text{with } \gamma = 10^{-8}. \quad (39)$$

In this expression the vectors \mathbf{n}_k with $\mathbf{n}_k \cdot \mathbf{n}_\ell = \delta_{k\ell}$ form an orthonormal basis of \mathbb{R}^d and the polynomials $P^{(k)}(\zeta)$ are positive in the entire domain. More specifically, we choose quadratic expressions

$$P^{(k)}(\zeta) = p_0^{(k)} + p_2^{(k)} \zeta^2 \quad \text{with } p_0^{(k)}, p_2^{(k)} > 0. \quad (40)$$

In contrast to the Poisson problem, the boundary conditions are also more complex. On the face $x_0 = 1$ Neumann conditions with $j(\mathbf{x}) = \exp(-(\mathbf{x} - \mathbf{x}_0^{(N)})^2 / (2\sigma_N^2))$ are applied, whereas on all other faces we use Dirichlet boundary conditions with $g(\mathbf{x}) = \exp(-(\mathbf{x} - \mathbf{x}_0^{(D)})^2 / (2\sigma_D^2))$. The Gaussians in those expressions are centred around $\mathbf{x}_0^{(N)} = (1, 0.5, 0.5)$ and $\mathbf{x}_0^{(D)} = (0, 0.5, 0.5)$ and have a width of $\sigma_N = \sigma_D = 0.1$.

In both cases the right hand side is set to a Gaussian $f(\mathbf{x}) = \exp(-(\mathbf{x} - \mathbf{x}_0)^2 / (2\sigma^2))$ with $\mathbf{x}_0 = (0.75, 0.5, 0.3)$ and $\sigma = 0.1$. The resulting operator is symmetric positive definite and we use a Conjugate Gradient (CG) iteration to solve the problem to a relative tolerance $\|\mathbf{r}\|/\|\mathbf{r}_0\| = 10^{-8}$ in the $L_2(\Omega)$ norm. Since most of the solution time is spent in the preconditioner, we achieved a significant speedup by only evaluating $K(\mathbf{x})$ and $c(\mathbf{x})$ in the centre of each cell in the preconditioner (but the full spatially varying expressions are used in the operator application of the Krylov-solver).

The block-Jacobi smoother is used in the DG space and the low order subspace is spanned by conforming lowest order Q_1 elements. An important parameter is the tolerance ϵ to which the diagonal blocks are inverted: a relatively loose tolerance will require less iterations of the block-solver, but might increase the total iteration count of the outer Krylov-iteration. In Fig. 1 we show the number of CG

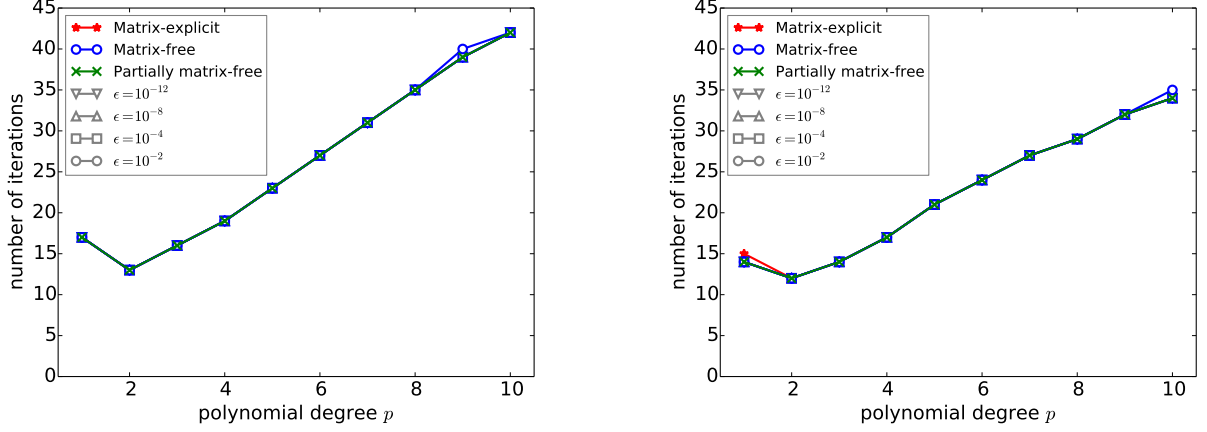


Figure 1: Number of outer CG iterations for different implementations. Results are shown both for a constant-coefficient Poisson problem (left) and for the diffusion problem with varying coefficients (right). For the matrix-free implementation the block-solver tolerance ϵ is varied between 10^{-12} and 10^{-2} .

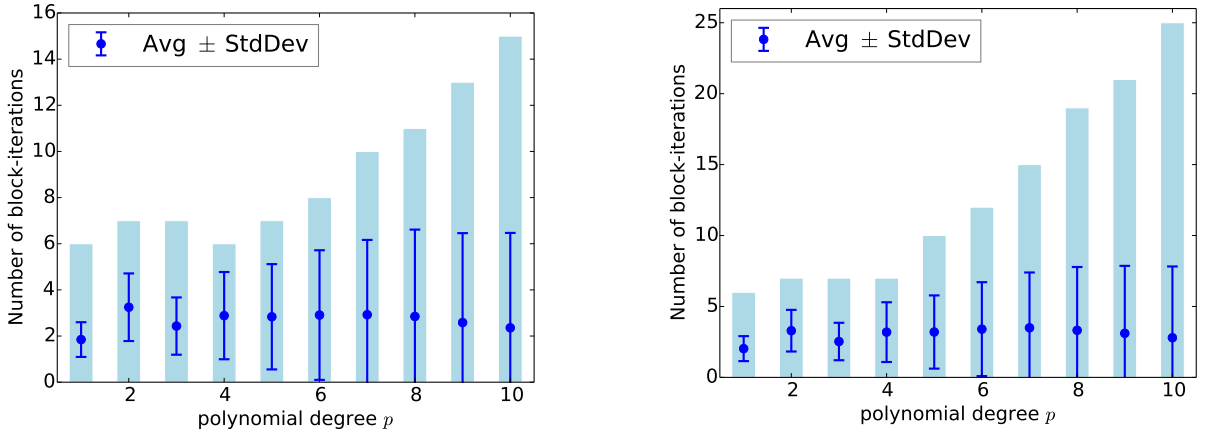


Figure 2: Average number of block-solver iterations to solve to a tolerance of $\epsilon = 10^{-2}$ for the Poisson problem (left) and the diffusion problem with spatially varying coefficients (right). The standard deviation and maximal number of solves is also shown (dark blue), the total range of iterations is indicated by the light blue bars in the background.

iterations for different tolerances ϵ . As can be seen from both figures, increasing the block-solver tolerance to $\epsilon = 10^{-2}$ has virtually no impact on the number of outer CG iterations. The average number of inner block-solver iterations which is required to reduce the residual by two orders of magnitude is shown in Fig. 2. In both cases the average number of block-solver iterations is less than four for all polynomial degrees and the maximal number of iterations never exceeds 15 (for the Poisson problem) and 25 (for the problem with spatially varying coefficients). The total solution time for a range of polynomial degrees is shown in Fig. 3 and (for the lowest polynomial degrees p) in Tab. 2. For both problems the partially matrix-free solver is superior to the matrix-based version for all polynomial degrees. As expected from the ϵ -independence of the number of iterations shown in Fig. 1, the matrix-free solver achieves the best performance if the block-solver tolerance is chosen to be very loose at $\epsilon = 10^{-2}$. For higher polynomial degrees ($p \geq 5$) the matrix-free solver gives the best overall performance and it always beats the matrix-explicit solver except for the lowest degree ($p = 1$). Although the partially matrix-free solver is fastest overall for low polynomial degrees ($p \leq 3$), Tab. 2 demonstrates that even in this case the fully matrix-free solver is only around $2 \times - 3 \times$ slower. It should also be kept in mind that using the matrix-free solver also results in dramatically reduced memory requirements, as will be discussed in Section 4.4 below.

Finally, a breakdown of the time per iteration is shown in Fig. 4. Except for very low orders where the AMG solve takes up a sizeable proportion of the runtime, this confirms that the total solution time is completely dominated by the application of the smoother and the calculation of the residual in the DG space. A breakdown of setup costs can be found in Appendix B.

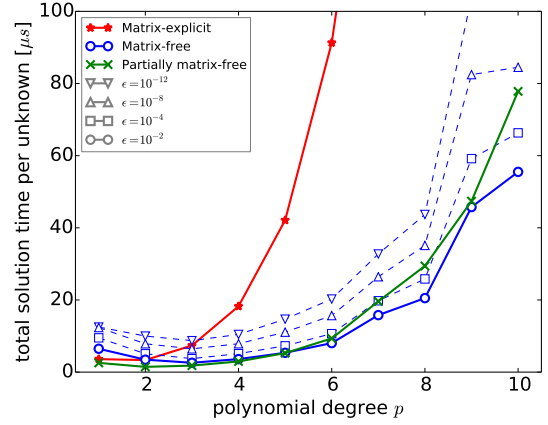
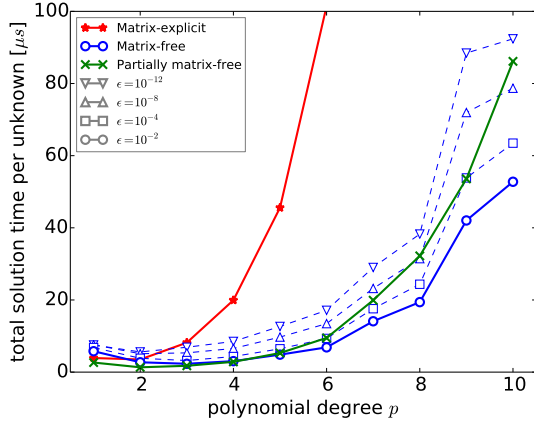


Figure 3: Total solution time for different implementations and a range of block-solver tolerances ϵ for the Poisson problem (left) and the diffusion problem with spatially varying coefficients (right).

	degree p	Poisson			Inhomogeneous		
		1	2	3	1	2	3
Matrix-explicit		3.91	3.50	8.16	3.60	3.34	7.37
Matrix-free	$\epsilon = 10^{-12}$	7.52	5.59	6.89	12.49	9.99	8.72
	$\epsilon = 10^{-8}$	7.59	5.10	5.33	12.35	7.86	6.45
	$\epsilon = 10^{-4}$	6.80	3.84	3.21	9.50	5.09	3.74
	$\epsilon = 10^{-2}$	5.79	2.73	2.31	6.47	3.45	2.60
Partially matrix-free		2.66	1.33	1.75	2.56	1.46	1.80

Table 2: Total solution time per unknown for the two diffusion problems and low polynomial degrees p . All times are given in μs .

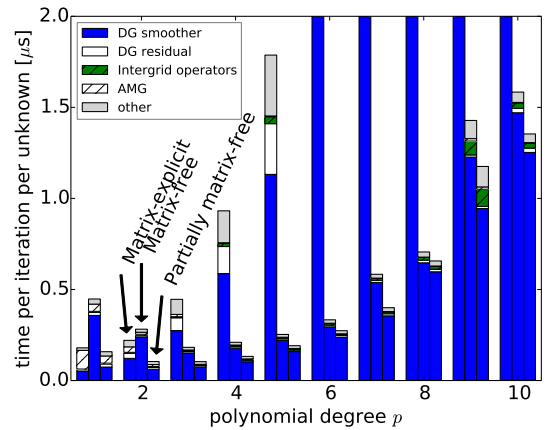
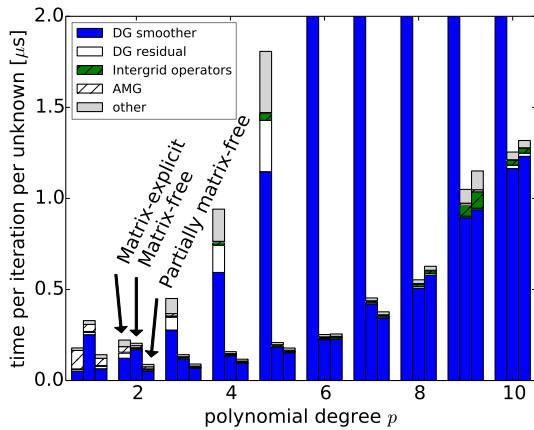


Figure 4: Breakdown of the time per iteration for different implementations and a range of block-solver tolerances ϵ for the Poisson problem (left) and the diffusion problem with spatially varying coefficients (right).

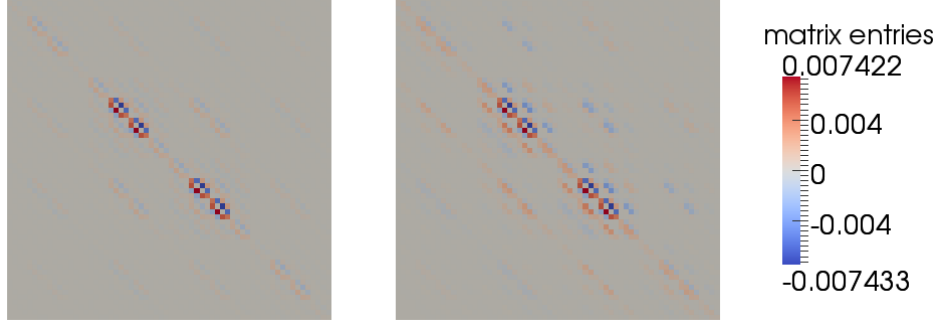


Figure 5: Block-diagonal 64×64 matrix at degree $p = 3$ for the convection dominated problem with $\mathbf{b} = (1, 0, 0)$ (left) and $\mathbf{b} = (1.0, 0.5, 0.3)$ (right). The Peclet number 2000 in both cases.

4.3. Convection dominated flow problem

While the block-Jacobi smoother is efficient for solving the diffusion equations described in the previous section, it results in poor convergence for larger values of the convection term. Here block-SOR-type smoothers are particularly efficient and a suitable preconditioner has to be chosen in the iterative inversion of the block-matrices D_T . As a further non-trivial problem we therefore consider a convection-dominated problem with constant coefficients which can be written as $-\kappa \Delta u + \nabla \cdot (\mathbf{b}u) = f$. The source term f is adjusted such that the exact solution is $u(\mathbf{x}) = \frac{x}{L_x}(1 - \frac{x}{L_x})\frac{y}{L_y}(1 - \frac{y}{L_y})\frac{z}{L_z}(1 - \frac{z}{L_z})$. We choose both an axis-parallel advection vector $\mathbf{b} = (1, 0, 0)$ and a more general vector $\mathbf{b} = (1.0, 0.5, 0.3)$. The strength of the diffusion term κ is adjusted such that the grid Peclet-number $\text{Pe} = \max_i \{b_i\} h / \kappa$ is fixed at $\text{Pe} = 2000$. This large grid-Peclet number makes the problem heavily convection dominated and we precondition the outer Krylov-solver with two iterations of a block-SSOR smoother instead of the hybrid multigrid algorithm described in Section 2.2. In the limit $\text{Pe} \rightarrow \infty$ and for a suitable ordering of the grid cells, one block-SOR sweep over the grid would solve the equation exactly. For the grid-Peclet number used here around 10-25 iterations of the flexible GMRES [49] solver are required to reduce the residual by eight orders of magnitude, see Fig. 7. A standard GMRES iteration (with restart 100) is used for inverting the block-diagonal matrices D_T in the matrix-free method.

The fast iterative inversion of the block-diagonal matrices D_T requires a suitable preconditioner. A typical 64×64 block-matrix for degree $n = 3$ is shown in Fig. 5. This matrix is clearly not diagonally dominant, in fact it has very large off-diagonal entries. The simple diagonal preconditioner, which was successfully used for all previous experiments on the diffusion problem, will therefore not be very efficient; in fact we find that the block-iteration will not converge at all. For the particular choice of an x-axis aligned advection vector $\mathbf{b} = (1, 0, 0)$ and a suitable ordering of the degrees of freedom, the first sub- and super-diagonal are particularly large. To account for the structure of the matrix we replaced the diagonal preconditioner by a tridiagonal solve with the Thomas-algorithm [48]. While this is more expensive, the cost of one preconditioner application is still proportional to number of unknowns per cell.

The structure of the matrix can be explained heuristically as follows: consider the weak form of the one-dimensional advection term

$$A_{ij} = a(\tilde{\psi}_i^p, \tilde{\psi}_j^p) \propto \int \tilde{\psi}_i^p(x) \partial_x \tilde{\psi}_j^p(x) dx. \quad (41)$$

For high polynomial degree and the nodal Gauss-Lobatto basis chosen here, each of the basis functions $\tilde{\psi}_i^p(x)$ can be approximated by a Gaussian centred at the i -th Gauss-Lobatto point and width which half the distance to the neighbouring nodal points. Simple symmetry arguments then imply that the diagonal entry A_{ii} is zero and the off-diagonal elements $A_{i,i+1}$ and $A_{i,i-1}$ are large and of approximately opposite size. Qualitatively this explains the structure seen in Fig. 5. For the non-axis-aligned cases additional off-diagonal bands appear at a distance of $p+1$ and $(p+1)^2$ from the main diagonal.

The number of block-solver iterations for the convection dominated problem is shown in Fig. 6 which should be compared to Fig. 2. On average around four iterations are required to reduce the residual by two orders of magnitude for the axis-aligned advection vector $\mathbf{b} = (1, 0, 0)$ and there is no

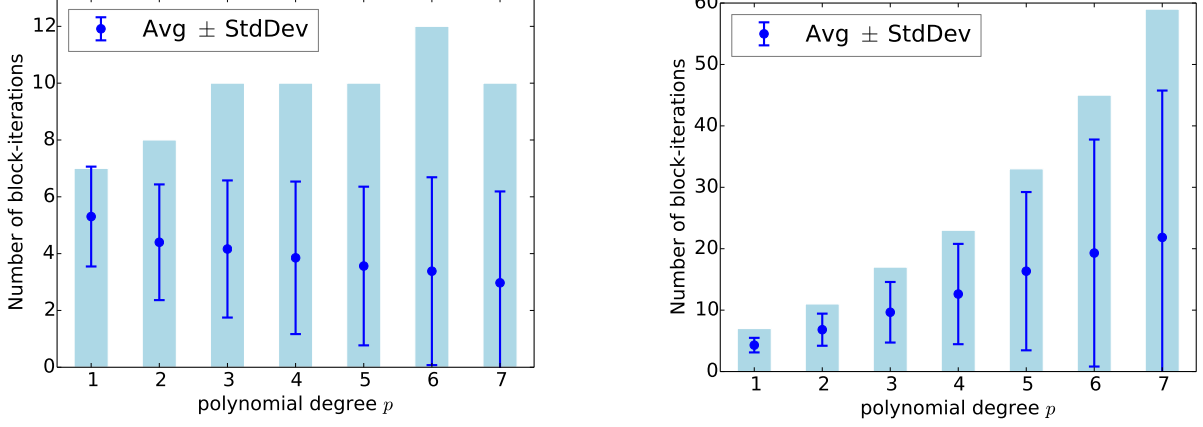


Figure 6: Number of GMRES iterations for the block solver with tri-diagonal preconditioner and tolerance $\epsilon = 10^{-2}$. The convection-dominated problem is solved with $\mathbf{b} = (1, 0, 0)$ (left) and $\mathbf{b} = (1.0, 0.5, 0.3)$ (right).

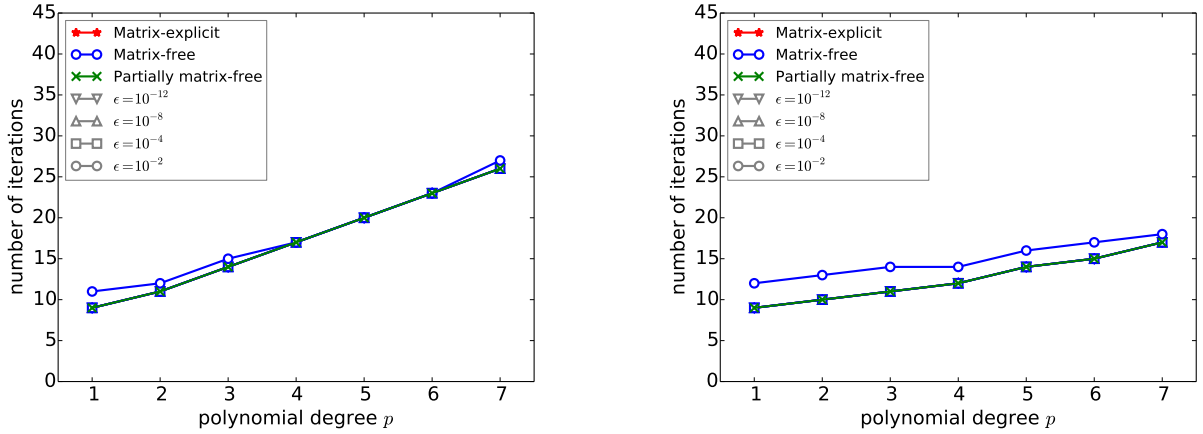


Figure 7: Number of outer flexible GMRES iterations. The convection-dominated problem is solved with $\mathbf{b} = (1, 0, 0)$ (left) and $\mathbf{b} = (1.0, 0.5, 0.3)$ (right).

strong dependence on the polynomial degree. For the more general advection vector $\mathbf{b} = (1.0, 0.5, 0.3)$ the number of iterations grows since the tridiagonal preconditioner becomes less efficient as expected due to the additional sub-diagonals in the matrix. The number of outer flexible GMRES iterations to solve the full system to a tolerance of 10^{-8} is shown in Fig. 7. For a given polynomial degree the number of iterations is virtually identical for all solvers and tolerances ϵ for the block-diagonal solver. The only exception is the matrix-free solver with tolerance $\epsilon = 10^{-2}$ which leads to a $\approx 20\%$ increase in the iteration count. Finally, the total solution time is shown in Fig. 8 and Tab. 3. In all cases the partially matrix-free solver clearly beats the matrix-based implementation. As expected from Fig. 7 the performance of the matrix-free code mainly depends on the block-solver tolerance ϵ . For the loosest tolerance $\epsilon = 10^{-2}$ it is faster than the matrix-explicit solver for moderately high polynomial degrees ($p \geq 3$ for $\mathbf{b} = (1, 0, 0)$ and $p \geq 4$ for $\mathbf{b} = (1.0, 0.5, 0.3)$). For high polynomial degree, the matrix-free solve time is in the same ballpark as the partially matrix-free time for the axis aligned advection vector $\mathbf{b} = (1, 0, 0)$. For $\mathbf{b} = (1.0, 0.5, 0.3)$ however, the matrix-free solver is substantially less efficient than the partially matrix-free version. This can be traced back to the growth in the number of iterations in the matrix-free inversion of D_T shown in Fig. 6 (right). Again it should be kept in mind that the reduced memory requirements of the matrix-free solver allow the solution of larger problems.

4.4. Memory savings in matrix-free implementation

As argued above, a key advantage of the completely matrix-free solver is its significantly lower demand on storage space. To estimate the memory requirements for all implementations carefully, first observe

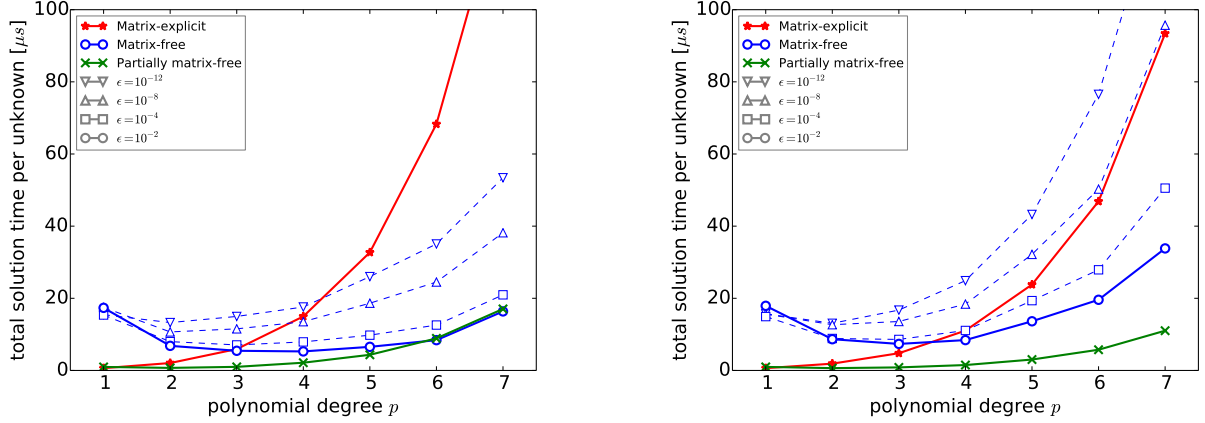


Figure 8: Total solution time per unknown. The convection-dominated problem is solved with $\mathbf{b} = (1, 0, 0)$ (left) and $\mathbf{b} = (1.0, 0.5, 0.3)$ (right).

degree p		$\mathbf{b} = (1, 0, 0)$			$\mathbf{b} = (1.0, 0.5, 0.3)$		
		2	3	4	2	3	4
Matrix-explicit		2.06	5.92	14.97	1.88	4.77	11.04
Matrix-free	$\epsilon = 10^{-12}$	13.27	14.97	17.59	13.07	16.73	24.89
	$\epsilon = 10^{-8}$	10.68	11.53	13.51	12.68	13.57	18.39
	$\epsilon = 10^{-4}$	7.99	7.07	7.91	8.86	8.60	11.08
	$\epsilon = 10^{-2}$	6.82	5.45	5.27	8.66	7.36	8.42
Partially matrix-free		0.71	1.00	2.14	0.64	0.84	1.50

Table 3: Total solution time per unknown for convection dominated problem. All times are given in μs .

that on a d -dimensional grid with N_{cell} cells an entire vector in DG space with polynomial degree p consists of $M_{\text{vector}} = N_{\text{cell}}(p+1)^d$ double precision numbers. The corresponding sizes for the full DG matrix and its blockdiagonal (used in the partially matrix-free implementation) are $M_{\text{matrix}} = 7N_{\text{cell}}(p+1)^{2d}$ and $M_{\text{blockdiag}} = N_{\text{cell}}(p+1)^{2d}$. In the following estimates we will take into account vectors and matrices on the first coarse multigrid level (i.e. in the Q_1 and P_0 subspace); we can safely neglect all data stored on the next-coarser levels. Since all grids used in the numerical experiments above contain at least 100 cells, we also ignore temporary storage for cell-local vectors.

First consider the setup for the diffusion problem in Section 4.2. The solution and right hand side, both of which are of size M_{vector} , have to be stored in every implementation. The outer CG iteration requires two additional temporaries of size M_{vector} each. Likewise, the hybrid multigrid stores three temporaries of size M_{vector} (two in the completely matrix-based implementation) and the corresponding vector storage in the Q_1 subspace is $3N_{\text{cell}}$ (three vectors with one unknown per cell). While the DG system matrix of size M_{matrix} is only stored in the matrix-based implementation, the coarse level matrix of size $27N_{\text{cell}}$, which describes the Q_1 discretisation of the problem, is required by the AMG solver in all cases. In the partially matrix-free case we store the entries of the local block-diagonal of total size $M_{\text{blockdiag}}$; in the matrix-free case we store the diagonal only (for preconditioning the cell-local iterative solver), which is equivalent to storing another vector of size M_{vector} . Overall, this results in the following storage requirements for the matrix-explicit (MF), partially matrix-free (PMF) and fully matrix-free (MF) cases:

$$M_{\text{total}}^{(\text{diffusion})} = \begin{cases} M_{\text{matrix}} + 6M_{\text{vector}} + 30N_{\text{cell}} \\ M_{\text{blockdiag}} + 7M_{\text{vector}} + 30N_{\text{cell}} \\ 8M_{\text{vector}} + 30N_{\text{cell}} \end{cases} = \begin{cases} (7(p+1)^{2d} + 6(p+1)^d + 30) N_{\text{cell}} & [\text{MX}] \\ ((p+1)^{2d} + 7(p+1)^d + 30) N_{\text{cell}} & [\text{PMF}] \\ (8(p+1)^d + 30) N_{\text{cell}} & [\text{MF}] \end{cases} \quad (42)$$

Numerical values for all considered problem sizes (see Tab. 1) in $d = 3$ dimensions are shown in Tab. 4. The table shows that even for moderate polynomial degrees the relative saving of the partially matrix-free

degree p	# dofs	memory requirements $M_{\text{total}}^{(\text{diffusion})}$			
		MX	PMF	MF	
1	$33.6 \cdot 10^6$	17.6 GB	5.0 GB	[3.5×	3.154 GB [5.6×
2	$9.5 \cdot 10^6$	14.9 GB	2.7 GB	[5.6×	0.691 GB [21.5×
3	$4.2 \cdot 10^6$	15.2 GB	2.4 GB	[6.4×	0.284 GB [53.7×
4	$2.0 \cdot 10^6$	14.1 GB	2.1 GB	[6.7×	0.132 GB [106.9×
5	$1.2 \cdot 10^6$	14.4 GB	2.1 GB	[6.8×	0.077 GB [186.5×
6	$0.7 \cdot 10^6$	13.2 GB	1.9 GB	[6.9×	0.044 GB [297.6×
7	$0.5 \cdot 10^6$	15.1 GB	2.2 GB	[6.9×	0.034 GB [445.5×
8	$0.3 \cdot 10^6$	12.9 GB	1.9 GB	[6.9×	0.020 GB [635.4×
9	$0.2 \cdot 10^6$	14.0 GB	2.0 GB	[7.0×	0.016 GB [872.5×
10	$0.2 \cdot 10^6$	12.7 GB	1.8 GB	[7.0×	0.011 GB [1162.1×

Table 4: Estimated memory requirements as defined in Eq. (42) for different implementations for the solution of the diffusion problem in Section 4.2. The savings relative to the matrix-explicit implementation are shown in square brackets.

implementation approaches a constant factor of $\approx 7\times$, since only the block-diagonal has to be stored. The relative saving for the fully matrix-free method is significantly larger, and of two- to three orders of magnitude, even for moderately large degrees. Note that asymptotically the savings grow with the third power of $p+1$ as $\approx \frac{7}{8}(p+1)^d$.

The convection-dominated problem in Section 4.3 uses a slightly different setup. The number of temporary vectors in the flexible GMRES solver with a restart value of $n_{\text{restart}}^{(\text{outer})} = 100$ is $2n_{\text{restart}}^{(\text{outer})} + 2$ for the outer solve. In contrast to the diffusion problem in Section 4.2, a block-SSOR smoother is used instead of the hybrid multigrid algorithm. This reduces the storage requirements since the SSOR smoother operates directly on the solution vector, and does not need the temporary storage for additional vectors and coarse level matrices/vectors which are required in the multigrid algorithm. However, since the local blocks are inverted iteratively with a GMRES method which uses $n_{\text{restart}}^{(\text{block})} + 1$ temporary local vectors of size $M_{\text{local vector}} = (p+1)^d$ and the restart value $n_{\text{restart}}^{(\text{block})} = 100$ in our numerical experiments is (unnecessarily) large, we choose to count those vectors in the fully matrix-free method. Furthermore, since the iterative block-diagonal solvers are preconditioned with the tridiagonal Thomas algorithm, we need two additional global vectors to store the bands of off-diagonal entries in the matrix-free algorithm. Overall, this results in the following storage requirements:

$$\begin{aligned}
M_{\text{total}}^{(\text{convection})} &= \begin{cases} M_{\text{matrix}} + (2n_{\text{restart}}^{(\text{outer})} + 4)M_{\text{vector}} & [\text{MX}] \\ M_{\text{blockdiag}} + (2n_{\text{restart}}^{(\text{outer})} + 4)M_{\text{vector}} & [\text{PMF}] \\ (2n_{\text{restart}}^{(\text{outer})} + 7)M_{\text{vector}} + (n_{\text{restart}}^{(\text{block})} + 1)M_{\text{local vector}} & [\text{MF}] \end{cases} \\
&= \begin{cases} \left(7(p+1)^{2d} + (2n_{\text{restart}}^{(\text{outer})} + 4)(p+1)^d \right) N_{\text{cell}} & [\text{MX}] \\ \left((p+1)^{2d} + (2n_{\text{restart}}^{(\text{outer})} + 4)(p+1)^d \right) N_{\text{cell}} & [\text{PMF}] \\ (2n_{\text{restart}}^{(\text{outer})} + 7)(p+1)^d N_{\text{cell}} + (n_{\text{restart}}^{(\text{block})} + 1)(p+1)^d & [\text{MF}] \end{cases} \quad (43)
\end{aligned}$$

The memory requirements and savings in the (partially-) matrix-free method for our setup (see Tab. 1) in $d = 3$ dimensions are shown in Tab. 5. As in Tab. 4, the relative saving in memory consumption of the partially matrix-free method is asymptotically $\approx 7\times$. Due to the large number of temporary vectors in the GMRES method, the saving of the fully matrix-free method is less pronounced, but still nearly two orders of magnitude for the highest polynomial degrees. In hindsight and looking at Figs. 7 and 6, the chosen restart values were probably too conservative and memory consumption could have been reduced without impact on results (except for the $\mathbf{b} = (1.0, 0.5, 0.3)$ case, where the matrix-free method is less efficient) by choosing smaller values of $n_{\text{restart}}^{(\text{outer})} = 15$ and $n_{\text{restart}}^{(\text{block})} = 12$. The corresponding numbers in this case are shown in Tab. 6 and show potential savings for the matrix-free method which are only a factor of $4\times$ smaller than those observed for the diffusion problem in Tab. 4.

degree p	# dofs	memory requirements $M_{\text{total}}^{(\text{convection})}$				
		MX	PMF			MF
1	$33.6 \cdot 10^6$	65.0 GB	53.0 GB	[1.2×	51.750 GB	[1.3×
2	$9.5 \cdot 10^6$	27.8 GB	16.3 GB	[1.7×	14.626 GB	[1.9×
3	$4.2 \cdot 10^6$	20.4 GB	8.4 GB	[2.4×	6.469 GB	[3.1×
4	$2.0 \cdot 10^6$	16.1 GB	4.9 GB	[3.3×	3.085 GB	[5.2×
5	$1.2 \cdot 10^6$	15.2 GB	3.7 GB	[4.1×	1.828 GB	[8.3×
6	$0.7 \cdot 10^6$	13.3 GB	2.8 GB	[4.8×	1.058 GB	[12.6×
7	$0.5 \cdot 10^6$	14.8 GB	2.8 GB	[5.3×	0.809 GB	[18.3×
8	$0.3 \cdot 10^6$	12.5 GB	2.2 GB	[5.7×	0.486 GB	[25.6×
9	$0.2 \cdot 10^6$	13.4 GB	2.2 GB	[6.0×	0.386 GB	[34.7×
10	$0.2 \cdot 10^6$	12.1 GB	1.9 GB	[6.2×	0.264 GB	[45.8×

Table 5: Estimated memory requirements as defined in Eq. (43) for different implementations for the solution of the convection dominated problem in Section 4.3 with $n_{\text{restart}}^{(\text{outer})} = n_{\text{restart}}^{(\text{block})} = 100$. The savings relative to the matrix-explicit implementation are shown in square brackets.

degree p	# dofs	memory requirements $M_{\text{total}}^{(\text{convection})}$				
		MX	PMF			MF
1	$33.6 \cdot 10^6$	22.5 GB	10.5 GB	[2.1×	9.250 GB	[2.4×
2	$9.5 \cdot 10^6$	15.8 GB	4.3 GB	[3.7×	2.614 GB	[6.0×
3	$4.2 \cdot 10^6$	15.1 GB	3.1 GB	[4.9×	1.156 GB	[13.0×
4	$2.0 \cdot 10^6$	13.5 GB	2.4 GB	[5.7×	0.551 GB	[24.6×
5	$1.2 \cdot 10^6$	13.7 GB	2.2 GB	[6.2×	0.327 GB	[41.8×
6	$0.7 \cdot 10^6$	12.4 GB	1.9 GB	[6.5×	0.189 GB	[65.8×
7	$0.5 \cdot 10^6$	14.1 GB	2.1 GB	[6.6×	0.145 GB	[97.8×
8	$0.3 \cdot 10^6$	12.1 GB	1.8 GB	[6.7×	0.087 GB	[138.7×
9	$0.2 \cdot 10^6$	13.1 GB	1.9 GB	[6.8×	0.069 GB	[189.8×
10	$0.2 \cdot 10^6$	11.9 GB	1.7 GB	[6.9×	0.047 GB	[252.0×

Table 6: Estimated potential memory requirements as defined in Eq. (43) for different implementations for the solution of the convection dominated problem in Section 4.3 with $n_{\text{restart}}^{(\text{outer})} = 15$, $n_{\text{restart}}^{(\text{block})} = 12$. The savings relative to the matrix-explicit implementation are shown in square brackets.

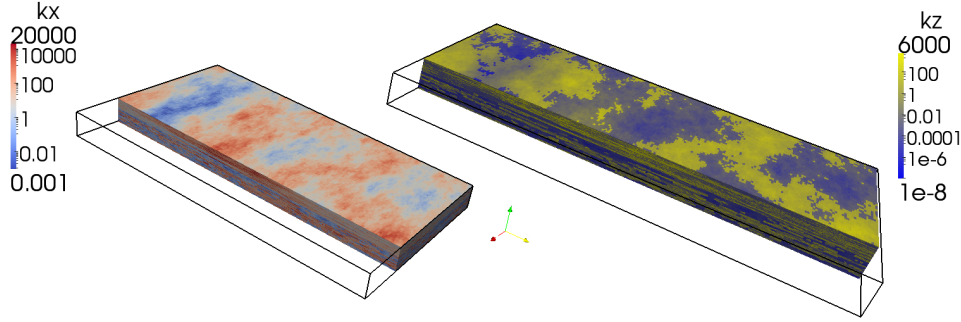


Figure 9: Permeability fields K_x and K_z from the SPE10 test case

Field	minimum	mean	maximum
K_x	$6.65 \cdot 10^{-4}$	$3.55 \cdot 10^2$	$2.0 \cdot 10^4$
K_y	$6.65 \cdot 10^{-4}$	$3.55 \cdot 10^2$	$2.0 \cdot 10^4$
K_z	$6.65 \cdot 10^{-8}$	$5.84 \cdot 10^1$	$6.0 \cdot 10^3$

Table 7: minimum, mean and maximum value of the three diagonal entries of permeability field for the SPE10 test case

4.5. SPE10 test case

We finally consider a challenging problem from a real-world application. The SPE10 test case was used to compare different reservoir modelling packages [36]. Due to the large jumps in the permeability field the solution of the SPE10 problem is not smooth. At first sight, using a high-order method seems to be questionable in this case since it will not lead to the best performance for a given error in the solution. However, in most applications the solution is smooth almost everywhere in the domain and any irregularities are confined to small, lower-dimensional subdomains. A typical example are sharp drops in pressure across fronts in atmospheric weather prediction. The main purpose of the following numerical experiments on the SPE10 benchmark is therefore to demonstrate the efficiency of our methods even for very challenging setups (in particular large, high contrast jumps in the coefficients).

The second SPE10 dataset describes the permeability tensor $K = \text{diag}(K_x, K_y, K_z)$ of a three dimensional reservoir of size $L_x \times L_y \times L_z = 1,200 \text{ ft} \times 2,200 \text{ ft} \times 170 \text{ ft}$ which is divided into $60 \times 220 \times 85 = 1.122 \cdot 10^6$ cells of size $20 \text{ ft} \times 10 \text{ ft} \times 2 \text{ ft}$. Permeability is assumed to be constant in each cell, but can vary strongly over the domain (see Fig. 9 and Tab. 7). Transport of a fluid with viscosity μ through the reservoir is described by Darcy's law which relates pressure u and flux \mathbf{q} according to

$$K \nabla u = -\mu \mathbf{q}. \quad (44)$$

For an incompressible fluid the stationary continuity equation in the absence of source terms is given by $\nabla \cdot \mathbf{q} = 0$. Inserting this into (44) and assuming constant viscosity results in the elliptic problem

$$-\nabla \cdot (K \nabla u) = 0. \quad (45)$$

This diffusion equation is solved with homogeneous Neumann boundary conditions $\boldsymbol{\nu} \cdot \nabla u = 0$ at $z = 0$ and $z = L_z$ and Dirichlet boundary conditions $u = -y$ on all other surfaces. The problem is challenging both due to the rapid variations in the diffusion tensor K and the strong anisotropy introduced by the large aspect ratio of the cells.

As for the diffusion equations in Section 4.2, the hybrid multigrid algorithm described in Section 2.2 is used as a preconditioner. However, in contrast to the conforming Q_1 space used there, for the SPE10 problem the low-order subspace is the piecewise constant P_0 space. The relative norm of the residual is not guaranteed to decrease monotonically in the CG iteration, and (particularly for the ill-conditioned matrix in this problem) this norm can be a poor indicator of convergence. We therefore use the estimator for the energy norm $\|e\|_A = \sqrt{e^T A e}$ of the error $e = \mathbf{u} - \mathbf{u}_{\text{exact}}$ given in [50, Section 3.1]. This does not add any significant overhead since only a small number of additional scalar products are required

smoother	degree	# iterations			time per unknown [μ s]		
		MX	MF	PMF	MX	MF	PMF
jacobi	1	52	53	52	5.67	38.00	8.58
	2	75	78	74	23.80	24.15	6.99
	3	—	102	—	—	22.25	—
ssor	1	37	37	37	7.44	57.67	15.84
	2	55	57	55	34.00	36.09	11.56
	3	—	77	—	—	34.14	—

Table 8: SPE10 benchmark on one node of “donkey”. The number of iterations and total solution time per unknown is shown for different polynomial degrees. For the highest polynomial degree only matrix-free solver results are reported due to memory limitations.

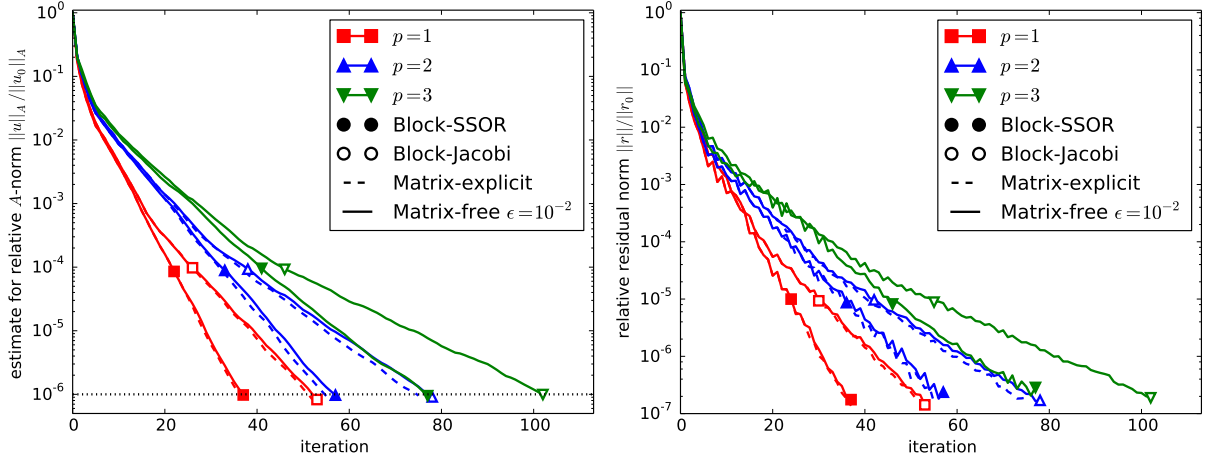


Figure 10: Convergence history for SPE10 benchmark. Both the relative energy norm (left) and the relative residual norm (right) are shown for polynomial degrees 1 (red squares), 2 (blue upward triangles) and 3 (green downward triangles). Results for the block-SSOR smoother are marked by filled symbols and results for the block-Jacobi smoother by empty symbols.

in every iteration. The quality of this estimate depends on the “error-bandwidth”, i.e. the number of terms that are used in the sum (3.5) in [50]. For all results obtained here we used an error-bandwidth of 4; increasing this further to 8 does not have any discernible impact on the results.

The SPE10 problem is solved to a relative tolerance of 10^{-6} in the energy norm. As Fig. 10 demonstrates, the error is reduced monotonically (left), whereas there are significant jumps in the relative residual (right). Solving to tighter tolerances is not of practical interest since other sources of error dominate the physical problem. The number of iterations and solution times for different polynomial degrees on one node of the “donkey” cluster are shown in Tab. 8 (run-ahead iterations for the initial estimate of the error in the energy norm are not counted but included in the reported runtimes). In the matrix-free solver the diagonal blocks were inverted to a relative tolerance of $\epsilon = 10^{-2}$. We find that in all cases a block-Jacobi smoother is more efficient than block-SSOR. For the lowest polynomial degree ($p = 1$) the matrix-explicit solver results in the shortest solution time, although it is only around $1.5\times$ faster than the partially matrix-free solver. However, for $p = 2$ the partially matrix-free solver is $3\times$ faster than the matrix-explicit version which has about the same performance as the matrix-free implementation. For the highest polynomial degree ($p = 3$) neither the full matrix nor the block-diagonal fit into memory and only results for the matrix-free solver are reported.

5. Conclusion

We described the efficient implementation of matrix-free multigrid block smoothers for higher order DG discretisations of the convection-diffusion equation. Since the implementation is FLOP-bound, this leads to significantly better utilisation of modern manycore CPUs. Sum-factorisation techniques reduce

the computational complexity from $\mathcal{O}(p^{2d})$ to $\mathcal{O}(d \cdot p^{d+1})$ and make the method particularly competitive for high polynomial degrees p . Our numerical experiments confirm that this leads to a significant reduction of the solution time even for relatively low polynomial degrees. Since the matrix does not have to be assembled and stored explicitly, our approach also leads to significantly reduced storage requirements. For moderately high polynomial degrees the totally matrix-free implementation typically gives the best overall performance. For low polynomial degrees we find that a partially matrix-free implementation, in which only the operator application is matrix-free, can reduce the runtime even further. The construction of a suitable preconditioner for the matrix-free block-inversion in convection-dominated problems is challenging if the advection field is not axis aligned, and therefore in this setup the partially matrix-free implementation (which avoids this problem) is particularly efficient.

With the SPE10 benchmark we have included a challenging, real world application. To quantify the performance for other realistic setups we will also apply the matrix-free solvers to the pressure correction equation in atmospheric models in the future.

Since the local blocks are only inverted in the preconditioner, additional approximations could improve performance further. For example, the use of lower-order Gauss-Lobatto quadrature rules for calculation of the volume integral can reduce the number of function evaluations since the quadrature points coincide with those of the surface integral.

The implementation of the highly-optimised matrix-free operators described in this work required a significant amount of low-level optimisation, leading to sophisticated code which is hard to maintain. In the future, we will explore the use of automatic code generation techniques which are currently integrated into the DUNE framework [51] and have been shown to give promising results.

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Appendix A. Discretisation in the lower- order subspaces

In this appendix we show how the coarse grid matrix $\hat{A} = P^T A P$ can be calculated efficiently by re-discretisation in the subspace \hat{V}_h . As discussed in Section 2.2.3 this avoids the expensive calculation via the explicit Galerkin product.

Appendix A.1. Piecewise constant subspace P_0

The P_0 basis functions have a vanishing gradient on each element. This simplifies the evaluation of $a_h(\hat{\psi}_j, \hat{\psi}_i)$ since the diffusion and convection term do not contribute to the volume integral $a_h^v(\hat{\psi}_j, \hat{\psi}_i)$ and the consistency and symmetrisation term vanish in the face integral $a_h^{\text{if}}(\hat{\psi}_j, \hat{\psi}_i)$. The entries of the Galerkin product with P_0 as a subspace are

$$\begin{aligned} (P^T A P)_{ij} = & \sum_{T \in \mathcal{T}_h} \int_T c \hat{\psi}_j \hat{\psi}_i dx + \sum_{F \in \mathcal{F}_h^i} \int_F \Phi(\hat{\psi}_j^-, \hat{\psi}_i^+, \nu_F \cdot \mathbf{b}) [\![\hat{\psi}_i]\!] ds + \sum_{F \in \mathcal{F}_h^D} \int_F \Phi(\hat{\psi}_j, 0, \nu_F \cdot \mathbf{b}) [\![\hat{\psi}_i]\!] ds \\ & + \alpha p(p+d-1) \left(\sum_{F \in \mathcal{F}_h^i} \int_F \frac{\langle \delta_{K\nu_F}^-, \delta_{K\nu_F}^+ \rangle |F|}{\min(|T^-(F)|, |T^+(F)|)} [\![\hat{\psi}_j]\!] [\![\hat{\psi}_i]\!] ds + \sum_{F \in \mathcal{F}_h^D} \int_F \frac{\delta_{K\nu_F}^- |F|}{|T^-(F)|} \hat{\psi}_j \hat{\psi}_i ds \right). \end{aligned} \quad (\text{A.1})$$

This should be compared to the finite volume discretisation of (3) and (4) which can be obtained as follows: let $\mathbf{x}(T)$ be the center of cell T , $\mathbf{x}(F)$ the center of face F and $\mathbf{x}^-(F)$ the center of inside cell, $\mathbf{x}^+(F)$ the center of outside cell with respect to F . The evaluation of the convection and reaction term

at $\mathbf{x}(F)$ and $\mathbf{x}(T)$, respectively, is denoted in shorthand by $\hat{\mathbf{b}} = \mathbf{b}(\mathbf{x}(F))$, $\hat{c} = c(\mathbf{x}(T))$. We approximate the diffusive normal flux with a central difference approximation and use the harmonic average of the diffusion tensor on $F \in \mathcal{F}_h^i$. On boundary faces $F \in \mathcal{F}_h^b$ the diffusive normal flux is approximated by a one-sided difference quotient. This results in the finite volume operator

$$\begin{aligned} \hat{A}_{ij}^{\text{FV}} = & \sum_{T \in \mathcal{T}_h} \hat{c} \hat{\psi}_j \hat{\psi}_i |T| + \sum_{F \in \mathcal{F}_h^i} \Phi(\hat{\psi}_j^-, \hat{\psi}_i^+, \boldsymbol{\nu}_F \cdot \hat{\mathbf{b}}) \llbracket \hat{\psi}_i \rrbracket |F| + \sum_{F \in \mathcal{F}_h^D} \Phi(\hat{\psi}_j, 0, \boldsymbol{\nu}_F \cdot \hat{\mathbf{b}}) \llbracket \hat{\psi}_i \rrbracket |F| \\ & + \sum_{F \in \mathcal{F}_h^i} \frac{\langle \delta_{K\nu_F}^-, \delta_{K\nu_F}^+ \rangle}{\|\mathbf{x}^-(F) - \mathbf{x}^+(F)\|_2} \llbracket \hat{\psi}_j \rrbracket \llbracket \hat{\psi}_i \rrbracket |F| + \sum_{F \in \mathcal{F}_h^D} \frac{\delta_{K\nu_F}}{\|\mathbf{x}^-(F) - \mathbf{x}(F)\|_2} \hat{\psi}_j \hat{\psi}_i |F|. \end{aligned} \quad (\text{A.2})$$

If in $P^T AP$ the upwind flux and $\int_T c \, ds$ are approximated by the box-rule (or c and \mathbf{b} are piecewise constant on the grid cells), the first three terms in (A.1) and (A.2) are identical. This is an admissible approximation since it is used in a subspace correction. To relate the remaining two integrals over interior- and boundary-faces, observe that on an equidistant mesh $|T^-(F)| = |T^+(F)| \, \forall F \in \mathcal{F}_h^i$. If the grid is also axi-parallel, the inverse distances in (A.2) can be written as

$$\frac{1}{\|\mathbf{x}^-(F) - \mathbf{x}^+(F)\|_2} = \frac{|F|}{\min(|T^-(F)|, |T^+(F)|)} \quad \forall F \in \mathcal{F}_h^i \quad (\text{A.3})$$

on the interior faces and as

$$\frac{1}{\|\mathbf{x}^-(F) - \mathbf{x}(F)\|_2} = \frac{|F|}{2|T^-(F)|} \quad \forall F \in \mathcal{F}_h^b \quad (\text{A.4})$$

on the boundary faces. Provided the diffusivity K is also piecewise constant on the grid cells (or, on each face is approximated by its mean value), the final two terms in (A.1) and (A.2) differ by multiplicative factors of $\alpha p(p+d-1)$ and $2\alpha p(p+d-1)$ respectively. Hence, instead of constructing \hat{A} via the Galerkin product $\hat{A} = P^T AP$, it can be obtained from a modified finite-volume discretisation, if the last two terms in (A.2) are scaled appropriately.

Appendix A.2. Conforming piecewise linear subspace Q_1

The Q_1 conforming piecewise linear basis functions are continuous over the interior edges. Exploiting this fact for $a_h(\hat{\psi}_j, \hat{\psi}_i)$ implies that all terms containing the jump operator $\llbracket \cdot \rrbracket$ that are summed over \mathcal{F}_h^i vanish. The entries of the Galerkin product with Q_1 as a subspace read

$$\begin{aligned} (P^T AP)_{ij} = & \sum_{T \in \mathcal{T}_h} \int_T \left((K \nabla \hat{\psi}_j - \mathbf{b} \hat{\psi}_j) \cdot \nabla \hat{\psi}_i + \hat{\psi}_j \hat{\psi}_i \right) dx \\ & + \sum_{F \in \mathcal{F}_h^D} \int_F \left(\Phi(\hat{\psi}_j, 0, \boldsymbol{\nu}_F \cdot \mathbf{b}) \hat{\psi}_i - \boldsymbol{\nu}_F \cdot (\omega^- K \nabla \hat{\psi}_j) \hat{\psi}_i - \hat{\psi}_j \boldsymbol{\nu}_F \cdot (\omega^- K \nabla \hat{\psi}_i) + \gamma_F \hat{\psi}_j \hat{\psi}_i \right) ds. \end{aligned} \quad (\text{A.5})$$

This is identical to the conforming piecewise linear Finite Element operator for the model problem (1). The only differences are the integrals over the Dirichlet boundary which is a remainder from the DG bilinear form. In contrast to the P_0 subspace $P^T AP$ is rediscretised by calculating $a_h(\hat{\psi}_j, \hat{\psi}_i) \, \forall i, j$ with the optimisation that the integral for all $F \in \mathcal{F}_h^i$ is skipped. Note that although \hat{A}_{ij} is calculated by assembling the DG operator $a_h(\hat{\psi}_i, \hat{\psi}_j)$ on a low-dimensional subspace, the polynomial degree which is used in the calculation of γ_F is the degree p of the DG space V_h^p , and not the degree of Q_1 .

Appendix B. Setup costs

The setup costs for the diffusion problems described in Section 4.2 are shown for a selection of polynomial degrees in in Tables B.9 and B.10. Note that the (partially) matrix-free implementation has a significantly lower setup cost since it does not require assembly of the DG matrix and construction of the coarse grid matrix \hat{A} via the Galerkin product.

	$p = 2$			$p = 3$			$p = 6$		
	MX	MF	PMF	MX	MF	PMF	MX	MF	PMF
Assemble prolongation	0.09	0.07	0.07	0.05	0.04	0.04	0.02	0.01	0.01
DG Matrix setup+assembly	2.36	—	—	2.50	—	—	3.85	—	—
DG smoother setup (block-factor)	0.25	—	1.10	0.26	—	1.09	0.43	—	1.64
Diagonal matrix assembly	—	0.91	—	—	0.91	—	—	1.43	—
Galerkin product $\hat{A} = P^T A P$	2.01	—	—	0.99	—	—	2.89	—	—
Coarse matrix assembly	—	0.18	0.18	—	0.04	0.04	—	0.00	0.00
AMG setup	1.03	0.37	0.37	0.22	0.10	0.10	0.02	0.01	0.01
total setup	5.73	1.54	1.73	4.03	1.09	1.27	7.21	1.46	1.67
total solve time	33.13	25.85	12.64	34.25	9.71	7.35	69.61	4.71	6.41

Table B.9: Breakdown of setup time for the Poisson problem for different polynomial degrees p . To put the setup costs into perspective, the total solve time (which includes the setup time) is also given

	$p = 2$			$p = 3$			$p = 6$		
	MX	MF	PMF	MX	MF	PMF	MX	MF	PMF
Assemble prolongation	0.09	0.07	0.07	0.05	0.04	0.04	0.02	0.01	0.01
DG Matrix setup+assembly	3.24	—	—	3.25	—	—	4.52	—	—
DG smoother setup (block-factor)	0.25	—	1.28	0.26	—	1.23	0.43	—	1.84
Diagonal matrix assembly	—	1.07	—	—	1.03	—	—	1.67	—
Galerkin product $\hat{A} = P^T A P$	1.97	—	—	0.98	—	—	2.86	—	—
Coarse matrix assembly	—	0.20	0.19	—	0.04	0.04	—	0.00	0.00
AMG setup	1.05	0.39	0.39	0.22	0.10	0.10	0.02	0.01	0.01
total setup	6.60	1.73	1.93	4.77	1.21	1.42	7.85	1.70	1.87
total solve time	31.70	32.74	13.82	30.95	10.89	7.51	62.58	5.52	6.37

Table B.10: Breakdown of setup time for the inhomogeneous diffusion problem for different polynomial degrees p . To put the setup costs into perspective, the total solve time (which includes the setup time) is also given

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