LOW RANK APPROXIMATION OF A SPARSE MATRIX BASED ON LU FACTORIZATION WITH COLUMN AND ROW TOURNAMENT PIVOTING*

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Abstract. In this paper we present an algorithm for computing a low rank approximation of a sparse matrix based on a truncated LU factorization with column and row permutations. We present various approaches for determining the column and row permutations that show a trade-off between speed versus deterministic/probabilistic accuracy. We show that if the permutations are chosen by using tournament pivoting based on QR factorization, then the obtained truncated LU factorization with column/row tournament pivoting, LU_CRTP, satisfies bounds on the singular values which have similarities with the ones obtained by a communication avoiding rank revealing QR factorization. Experiments on challenging matrices show that LU_CRTP provides a good low rank approximation of the input matrix and it is less expensive than the rank revealing QR factorization in terms of computational and memory usage costs, while also minimizing the communication cost. We also compare the computational complexity of our algorithm with randomized algorithms and show that for sparse matrices and high enough but still modest accuracies, our approach is faster.

Key words. rank revealing, LU and QR factorizations, column pivoting, minimize communication

AMS subject classifications. 65F25, 65F20

DOI. 10.1137/16M1074527

1. Introduction. In this paper we address the problem of computing a low rank approximation of a large sparse matrix by using a rank revealing LU factorization. This problem has numerous and diverse applications ranging from scientific computing problems such as fast solvers for integral equations to data analytics problems such as principal component analysis (PCA) or image processing. The singular value decomposition produces the best rank-k approximation, but it is expensive to compute. There are a number of less expensive approaches in the literature that approximate the singular value decomposition of a matrix, such as rank revealing QR and LU factorizations, or the Lanczos algorithm (see, e.g., [7, 35]). In recent years, several randomized algorithms have been introduced for this problem that aim at further decreasing the computational cost while obtaining accurate results with high probability. For recent surveys, see, e.g., [26, 32].

^{*}Submitted to the journal's Software and High-Performance Computing section May 9, 2016; accepted for publication (in revised form) August 11, 2016; published electronically March 15, 2018. http://www.siam.org/journals/sisc/40-2/M107452.html

Funding: The first author's work was supported by the NLAFET project as part of the European Union's Horizon 2020 research and innovation program under grant 671633. The third author's work was supported by the financial support of DOE Office of Science grants DE-SC0008700, DOE DE-SC0010200, and DOE AC02-05CH11231, DARPA grant HR0011-12-2-0016, and ASPIRE Lab industrial sponsors and affiliates Intel, Google, Hewlett-Packard, Huawei, LGE, NVIDIA, Oracle, and Samsung. Any opinions, findings, conclusions, or recommendations in this paper are solely those of the authors and do not necessarily reflect the position or the policy of the sponsors.

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In this paper we focus on sparse LU factorizations that are effective in revealing the singular values of a matrix, in terms of both accuracy and speed. For sparse matrices, direct methods of factorization lead to factors that are denser than the original matrix A. Since the R factor obtained from the QR factorization is the Cholesky factor of A^TA , it is expected that the factors obtained from a QR factorization are denser than the factors obtained from an LU factorization. Indeed, our focus is on obtaining a factorization that is less expensive than the rank revealing QR factorization in terms of computational and memory usage costs, while also minimizing the communication cost. The communication cost is one of the dominant factors for the performance of an algorithm on current and future computers [13], and classic algorithms based on row and/or column permutations are suboptimal in terms of communication.

We begin by surveying prior work in order to compare our work to it. Consider first rank revealing QR factorizations, such as QR with column pivoting (QRCP) [20, 4] or its strong version [25]. When these algorithms are executed in a distributed memory environment, the matrix A is typically distributed over P processors by using a one-dimensional or two-dimensional (block) cyclic partitioning. Finding the column of maximum norm at each step of the factorization as in QRCP requires a reduction operation among processors, which costs $O(\log P)$ messages. QRCP exchanges $O(k \log P)$ messages for computing a rank-k approximation, and if the factorization proceeds until the end, it exchanges $O(n \log P)$ messages. A lower bound on the number of messages required for computing the QR factorization of a dense $n \times n$ matrix A (under certain hypotheses and when the memory size per processor is $O(n^2/P)$) [2] is $\Omega(\sqrt{P})$. Hence QRCP is not optimal in terms of the number of messages exchanged. A communication avoiding RRQR factorization, referred to as CARRQR, was introduced in [11]. This factorization selects k linearly independent columns by using tournament pivoting which requires only $O(\log P)$ messages. CARRQR is optimal in terms of communication, modulo polylogarithmic factors, on both sequential machines with two levels of slow and fast memory and parallel machines with one level of parallelism, while performing three times more floating point operations than QRCP. Extensive numerical experiments presented in [11] show that tournament pivoting reveals the singular values of a matrix with an accuracy close to the one obtained by QRCP. We will discuss it in more detail in section 2.

In this paper we focus on computing a low rank approximation of a matrix A and we consider different possible cases in which either the desired rank k is known, all singular values larger than a tolerance τ need to be approximated, or a gap in the singular values needs to be identified. We introduce a truncated LU factorization with column and row permutations which, for a given rank k, has the form

$$(1) \qquad \qquad P_r A P_c = \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ \bar{A}_{21} & \bar{A}_{22} \end{pmatrix} = \begin{pmatrix} I \\ \bar{A}_{21} \bar{A}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ & S(\bar{A}_{11}) \end{pmatrix},$$

(2)
$$S(\bar{A}_{11}) = \bar{A}_{22} - \bar{A}_{21}\bar{A}_{11}^{-1}\bar{A}_{12},$$

where $A \in \mathbb{R}^{m \times n}$, $\bar{A}_{11} \in \mathbb{R}^{k \times k}$, $\bar{A}_{22} \in \mathbb{R}^{(m-k) \times (n-k)}$, and the rank-k approximation matrix \tilde{A}_k is

(3)
$$\tilde{A}_{k} = \begin{pmatrix} I \\ \bar{A}_{21}\bar{A}_{11}^{-1} \end{pmatrix} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \end{pmatrix} = \begin{pmatrix} \bar{A}_{11} \\ \bar{A}_{21} \end{pmatrix} \bar{A}_{11}^{-1} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \end{pmatrix}.$$

The column and row permutations are chosen such that the singular values of \bar{A}_{11}

approximate the first k singular values of A, while the singular values of $S(\bar{A}_{11})$ approximate the last min(m,n)-k singular values of A. The factorization from (3) is obtained by first selecting k "most linearly independent" columns from the matrix A, permuting them to the leading positions, and then selecting k most linearly independent rows from these columns. Depending on the methods used to select columns and rows, different algorithms can be obtained, with different bounds on the revealed singular values and on the numerical stability of the truncated LU factorization. The design space for selecting the k columns and rows can be summarized by the following list (other possibilities have been proposed, e.g., choosing the k-by-k submatrix of nearly maximum determinant [34, 33]):

- 1. Select k linearly independent columns of A (call the result B), by using
 - (a) (strong) QRCP/tournament pivoting using QR,
 - (b) LU/tournament pivoting based on LU, with some form of pivoting (column, complete, rook),
 - (c) randomization, e.g., randomly sample columns of A, or premultiply X = ZA, where random matrix Z is short and wide, and then pick k rows from X^T , by some method from 2) below,
 - (d) tournament pivoting based on randomized algorithms to select columns at each step.
- 2. Select k linearly independent rows of B, by using
 - (a) (strong) QRCP/tournament pivoting based on QR on B^T , or on Q^T , the rows of the thin Q factor of B,
 - (b) LU/tournament pivoting based on LU with some form of pivoting (row, TSLU [24], complete, rook) on B,
 - (c) randomization, e.g., randomly sample rows of B,
 - (d) tournament pivoting based on randomized algorithms to select rows.

These various approaches show a trade-off between speed versus deterministic/probabilistic accuracy and are presented in order of expected decreasing accuracy. Concerning speed, the selections based on QR are more expensive than those based on LU, in terms of floating point operations and also memory usage, but they are expected to provide more accurate approximations of the singular values. The classic pivoting strategies used in LU and QR factorizations are suboptimal in terms of communication, while tournament pivoting provides a communication optimal alternative, with worse theoretical bounds on the approximations of the singular values.

The second formulation of \tilde{A}_k from (3) is called a CUR decomposition (see [22, 36, 32] and references therein), a popular low rank approximation factorization in data analytics in which C and R are columns and rows selected from the matrix A, and hence very sparse. In many cases of interest, \tilde{A}_k is applied to a vector, and \bar{A}_{11}^{-1} should not be computed, but instead its LU or QR factorization should be used. The column/row selection design space described above can be used to obtain such a CUR decomposition. The randomized algorithms for selecting columns and rows reviewed in [32] generally fit in category (1c) and (2c) of our design space.

In this design space, we show that LU_CRTP, a factorization that uses tournament pivoting based on QR from part 1(a) above and tournament pivoting based on QR on Q^T from part 2(a) above, is a good choice in terms of both speed and accuracy. We discuss it in more detail in section 3 and show that LU_CRTP allows us not only to obtain bounds on the approximated singular values, the stability of the LU factorization, but also to minimize communication. The obtained factorization

satisfies the following properties:

(4)
$$1 \leq \frac{\sigma_i(A)}{\sigma_i(\bar{A}_{11})}, \frac{\sigma_j(S(\bar{A}_{11}))}{\sigma_{k+j}(A)} \leq q(m, n, k),$$

(5)
$$\rho_l(\bar{A}_{21}\bar{A}_{11}^{-1}) \le F_{TP}$$

for any $1 \leq l \leq m-k$, $1 \leq i \leq k$, and $1 \leq j \leq \min(m,n)-k$, where $\rho_l(B)$ denotes the 2-norm of the lth row of B, and F_{TP} is a quantity arising in the bound on singular values obtained after column tournament pivoting based on QR [11]. If a binary tree is used during tournament pivoting to select k columns from n, then $F_{TP} \leq \frac{1}{\sqrt{2k}} \left(n/k\right)^{\log_2\left(\sqrt{2}fk\right)}$, and $q(m,n,k) = \sqrt{(1+F_{TP}^2(n-k))(1+F_{TP}^2(m-k))}$, where f is a small constant related to the strong RRQR factorization used during the tournament (typically f=2). The bound on F_{TP} can be regarded as a polynomial in n for a fixed k and f. For more details see Theorem 3.1. The second inequality (5) is important for bounding element growth in the LU factorization which governs its numerical stability. All these results are obtained assuming infinite precision, and they are expected not to hold when the singular values approach machine precision, and so may be significantly changed by roundoff error. The existence of a rank revealing LU factorization has been proved by Pan in [34], and we discuss it in more detail in section 3.

Once the first k singular values of A are approximated, it might be the case that subsequent singular values need to be computed. One example is when the algorithm is used to compute a low rank approximation and no gap has been identified in the singular values. Then the block LU factorization algorithm continues recursively on $S(\bar{A}_{11})$ until an overestimation K (which is a multiple of k) of the rank has been determined. Some of the previous bounds become exponential, where the exponent is the number of steps of tournament pivoting, K/k; for more details see Theorem 3.2. But in practice the algorithm is effective in revealing the spectrum of A and provides a good low rank approximation.

In section 4 we discuss the cost of our truncated LU factorization for the case when the rank k is known, first for matrices with arbitrary sparsity structure and then for matrices with small separators. For matrices with arbitrary sparsity structure, an upper bound on the number of flops required by one use of tournament based on QR factorization to choose k columns, executed sequentially, which dominates the cost of our algorithm, is $O(k^2 nnz(A))$. When the algorithm is executed in parallel on P processors, an upper bound on the number of flops is $O(k^2 \frac{nnz(A)}{P} \log \frac{n}{k})$ and the number of messages exchanged is $O(\log P)$. Given the assumptions we make to derive the bounds on the number of flops, we expect that these bounds are pessimistic.

For comparison, one of the most efficient randomized algorithms for computing a low rank approximation of a sparse matrix is due to Clarkson and Woodruff [6]. For an $n \times n$ matrix A, it computes a rank-k approximation which satisfies $||A - LDW^T||_F \leq (1+\epsilon)||A - A_k||_F$, where L and W are of dimension $n \times k$, D is of dimension $k \times k$, and A_k is the best rank-k approximation. This algorithm requires $O(nnz(A)) + \bar{O}(nk^2\epsilon^{-4} + k^3\epsilon^{-5})$ flops, where $\bar{O}(f) = f \cdot \log^{O(1)}(f)$. By ignoring the constants in the asymptotic O() notation, if ϵ is chosen such that $nnz(A) \leq \bar{O}(n\epsilon^{-4})$, or after rearranging, $\epsilon \leq \frac{1}{(\frac{nnz(A)}{n})^{1/4}}$, our algorithm is faster than the randomized approach, while also being deterministic. For example, if $\epsilon = 0.5$ or $\epsilon = 0.1$, then our algorithm is faster if the average number of nonzeros per column of A is smaller than 16 or 10^4 , respectively. We note that even for a fixed ϵ , comparing the theoretical

bounds on accuracy of the two algorithms is difficult, since we bound the error of the approximated singular values, while Clarkson and Woodruff bound the Frobenius norm of the error of the low rank approximation. A meaningful comparison of the accuracy of the two algorithms should include varying ϵ and comparing the results on matrices from different applications. This remains future work.

We say in this paper that the singular values of A are well approximated when the ratio of the approximated singular values to the singular values of A is bounded as in (4). In practice, the singular values are well approximated if this ratio is on average close to one and in the worst case a small constant. In section 5 we present experimental results which compare LU_CRTP with results obtained by using the singular value decomposition and QRCP. On a set of 16 selected matrices that were used in previous papers on rank revealing factorizations, we show that our factorization algorithm approximates well the singular values of the matrix A. The ratio of the absolute values of the diagonal elements of the R factor obtained from tournament pivoting to the corresponding singular values of A is on average close to one and typically within a factor of 10. We also compare the number of nonzeros in the low rank approximation obtained by our algorithm with respect to a low rank approximation obtained by using QRCP or LU with partial pivoting. For the same rank, we obtain a factor of up to 208x fewer nonzeros with respect to QRCP, and in some cases fewer nonzeros than LU with partial pivoting.

- 2. Background. In this section we review some of the factorizations and their properties that will be used in our rank revealing LU factorization. We focus in particular on rank revealing and strong rank revealing QR and LU factorizations.
- **2.1. Notation.** We use MATLAB notation. Given two matrices $A \in \mathbb{R}^{m \times n}$ and $B \in \mathbb{R}^{m \times k}$, the matrix $C \in \mathbb{R}^{m \times (n+k)}$ obtained by putting the two matrices next to each other is referred to as C = [A, B]. Given two matrices $A \in \mathbb{R}^{m \times n}$ and $B \in \mathbb{R}^{k \times n}$, the matrix $C \in \mathbb{R}^{(m+k) \times n}$ obtained by putting A on top of B is referred to as C = [A; B]. The submatrix of A formed by rows i to j and columns l to k is referred to as A(i:j,l:k). The element in position (i,j) of A is referred to as A(i,j). Very often, when a matrix is partitioned into a block matrix, we give the dimensions of a few blocks and we assume the reader can deduce the dimensions of the remaining blocks. Given a matrix A partitioned as $A = [A_{00}, \ldots, A_{T,0}]$, we refer to the submatrices A_{00} to $A_{T,0}$ as $A_{0:T,0}$. Similar notation is used for a block matrix with multiple rows.

The absolute values of the matrix A are referred to as |A|. The max norm is defined as $||A||_{max} = \max_{i,j} |A(i,j)|$. We refer to the 2-norm of the jth row of A as $\rho_j(A)$, the 2-norm of the jth column of A as $\chi_j(A)$, and the 2-norm of the jth row of A^{-1} as $\omega_j(A)$.

The different methods considered in this paper are the following:

- QRCP: QR with column pivoting [4];
- RRQR: rank revealing QR factorization, a QR factorization with column permutations which satisfies inequality (7);
- strong RRQR: strong rank revealing QR factorization [25], a QR factorization with column permutations which satisifes inequality (9);
- CARRQR: communication avoiding rank revealing factorization [11], a QR factorization with column tournament pivoting which satisfies inequality (10);
- LU_CRTP: block LU factorization with column/row tournament pivoting which satisfies the inequalities described in Theorem 3.1 (after one tournament) and Theorem 3.2 (after multiple steps of tournament);

- LU_CTP: block LU factorization with column tournament pivoting and row permutations based on partial pivoting, a version of LU_CRTP which requires fewer floating point operations, but more communication;
- LU_CRQCRP: block LU factorization similar to LU_CRTP, in which column/row pivoting is based on QRCP, a factorization used for comparison purposes.
- **2.2. Rank revealing QR factorizations.** Consider first the QR factorization with column permutations of a matrix $A \in \mathbb{R}^{m \times n}$ of the form

(6)
$$AP_c = QR = Q \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix},$$

where $Q \in \mathbb{R}^{m \times m}$ is orthogonal, $R_{11} \in \mathbb{R}^{k \times k}$ is upper triangular, $R_{12} \in \mathbb{R}^{k \times (n-k)}$, and $R_{22} \in \mathbb{R}^{(m-k) \times (n-k)}$. We say that this factorization is rank revealing if

(7)
$$1 \le \frac{\sigma_i(A)}{\sigma_i(R_{11})}, \frac{\sigma_j(R_{22})}{\sigma_{k+j}(A)} \le q(k, n)$$

for any $1 \leq i \leq k$ and $1 \leq j \leq \min(m, n) - k$, where q(k, n) is a low degree polynomial in k and n. Note that definitions of a rank revealing factorization as given in [28, 5] bound only $\sigma_{max}(R_{11})$ and $\sigma_{min}(R_{22})$ with respect to the singular values of A, so that our definition is stricter. The two following theorems recall the properties of a strong rank revealing QR factorization [25].

THEOREM 2.1 (see Gu and Eisenstat [25]). Let A be an $m \times n$ matrix and let $1 \leq k \leq \min(m,n)$. For any given parameter f > 1, there exists a permutation P_c such that

(8)
$$AP_c = Q \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix},$$

where R_{11} is $k \times k$ and

(9)
$$\left(R_{11}^{-1}R_{12}\right)_{i,j}^{2} + \omega_{i}^{2}\left(R_{11}\right)\chi_{j}^{2}\left(R_{22}\right) \leq f^{2}$$

for any $1 \le i \le k$ and $1 \le j \le n - k$.

The inequality (9) defining a strong rank revealing factorization allows one to bound the singular values of R_{11} and R_{22} as in a rank revealing factorization, while also upper bounding the absolute values of $R_{11}^{-1}R_{12}$.

Theorem 2.2 (Gu and Eisenstat [25]). Let the factorization in Theorem 2.1 satisfy inequality (9). Then

$$1 \le \frac{\sigma_i(A)}{\sigma_i(R_{11})}, \frac{\sigma_j(R_{22})}{\sigma_{k+j}(A)} \le \sqrt{1 + f^2 k(n-k)}$$

for any $1 \le i \le k$ and $1 \le j \le \min(m, n) - k$.

The communication avoiding RRQR factorization, referred to as CARRQR [11], computes a rank revealing factorization by using a block algorithm that selects k columns at each iteration, permutes them to be the leading columns, computes k steps of a QR factorization with no pivoting, and then proceeds recursively on the trailing matrix. The k columns are selected by using tournament pivoting as described

later in this section. It is shown in [11] that CARRQR computes a factorization as in (8) and a permutation P_c that satisfies

(10)
$$\chi_j^2 \left(R_{11}^{-1} R_{12} \right) + \left(\chi_j \left(R_{22} \right) / \sigma_{\min}(R_{11}) \right)^2 \le F_{TP}^2, \text{ for } j = 1, \dots, n - k.$$

Here F_{TP} depends on k, f, n, the shape of reduction tree used during tournament pivoting, and the number of iterations of CARRQR. The following theorem, which is a relaxed version of Theorem 2.1, shows that CARRQR reveals the rank in a similar way to strong RRQR factorization; for more details see [11].

Theorem 2.3. Assume that there exists a permutation P_c for which the QR factorization

$$AP_c = Q \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix},$$

where R_{11} is $k \times k$ and satisfies (10). Then

(12)
$$1 \le \frac{\sigma_i(A)}{\sigma_i(R_{11})}, \frac{\sigma_j(R_{22})}{\sigma_{k+j}(A)} \le \sqrt{1 + F_{TP}^2(n-k)}$$

for any $1 \le i \le k$ and $1 \le j \le \min(m, n) - k$.

In this work we use only one step of tournament pivoting, which corresponds to one iteration of CARRQR. We refer to this factorization as QR_TP(A,k) and a detailed description of this algorithm and how it can be implemented in a parallel environment can be found in [11]. Tournament pivoting selects k columns from the matrix A by using a reduction tree. It first divides the matrix A into n/(2k) subsets of columns and at the leaves of the reduction tree, it selects from each subset k columns by using (strong) rank revealing QR factorization. At each node of the subsequent levels of the reduction tree, a new matrix is formed by adjoining next to each other the two subsets of candidate columns selected by its children. A new set of k candidate columns is selected by using (strong) rank revealing QR factorization. The columns selected at the root of the reduction tree are the final columns selected by tournament pivoting. The factorization from Theorem 2.3, which satisfies inequality (10), is obtained by permuting those columns to the first k positions. We note that in our experimental results as well as in those reported in [11], we use QRCP for the selection of k columns at each node of the reduction tree. However the bounds on F_{TP} are obtained by using strong rank revealing QR factorization.

Corollaries 2.6 and 2.7 from [11] show that the selection of k columns of the $m \times n$ matrix A using QR with binary tree based tournament pivoting reveals the rank of A in the sense of Theorem 2.3, with bound

(13)
$$F_{TP-BT} \le \frac{1}{\sqrt{2k}} \left(\sqrt{2} f k \right)^{\log_2(n/k)} = \frac{1}{\sqrt{2k}} \left(n/k \right)^{\log_2(\sqrt{2} f k)}.$$

If tournament pivoting uses a flat tree, then the bound becomes

(14)
$$F_{TP-FT} \le \frac{1}{\sqrt{2k}} \left(\sqrt{2}fk\right)^{n/k}.$$

In the case of a binary tree, the bound in (13) can be regarded as a polynomial in n in general for a fixed k and f. The bound (14) obtained for a flat tree is exponential in n, but is a rapidly decreasing function of k. Even if this suggests that the singular values approximated by the factorization might be far from the singular values of A, the extensive numerical experiments performed in [11] show that both binary tree and flat tree are effective in approximating the singular values of A.

2.3. Orthogonal matrices. Consider an orthogonal matrix $Q \in \mathbb{R}^{m \times m}$ and its partitioning as

$$(15) Q = \begin{pmatrix} Q_{11} & Q_{12} \\ Q_{21} & Q_{22} \end{pmatrix},$$

where $Q_{11} \in \mathbb{R}^{k \times k}$. The QR_TP factorization of $(Q_{11}; Q_{21})^T$ leads to the following factorization:

$$(16) P_r Q = \begin{pmatrix} \bar{Q}_{11} & \bar{Q}_{12} \\ \bar{Q}_{21} & \bar{Q}_{22} \end{pmatrix} = \begin{pmatrix} I \\ \bar{Q}_{21}\bar{Q}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{Q}_{11} & \bar{Q}_{12} \\ & S(\bar{Q}_{11}) \end{pmatrix},$$

where $S(\bar{Q}_{11}) = \bar{Q}_{22} - \bar{Q}_{21}\bar{Q}_{11}^{-1}\bar{Q}_{12} = \bar{Q}_{22}^{-T}$ (see [34, proof of Theorem 3.7]). This factorization satisfies the following bounds:

(17)
$$\rho_j(\bar{Q}_{21}\bar{Q}_{11}^{-1}) \le F_{TP},$$

(18)
$$\frac{1}{q_2(k,m)} \le \sigma_i(\bar{Q}_{11}) \le 1$$

for all $1 \le i \le k$, $1 \le j \le m - k$, where $q_2(k, m) = \sqrt{1 + F_{TP}^2(m - k)}$. F_{TP} is the bound obtained from QR with tournament pivoting, given in (13) for a binary tree and in (14) for a flat tree.

The singular values of \bar{Q}_{11} are also bounded in Lemma 3.4 from [34]. In that paper, $vol(\bar{Q}_{11})$ is the absolute value of the determinant of \bar{Q}_{11} . It is a local μ -maximum in Q if for all neighboring submatrices Q' which differ from \bar{Q}_{11} in exactly one column or one row, $\mu \cdot vol(\bar{Q}_{11}) \geq vol(Q')$ for some $\mu \geq 1$. The permutation P_r is chosen in [34] such that $vol(\bar{Q}_{11})$ is a local μ -maximum in Q and $q_2(k,m) = \sqrt{1 + k(m-k)\mu^2}$.

By using the properties of the CS decomposition of an orthogonal matrix, we also have that $\sigma_{min}(\bar{Q}_{11}) = \sigma_{min}(\bar{Q}_{22})$.

- 3. LU_CRTP: Block LU factorization with column/row tournament pivoting. In this section we describe the algebra of our low rank approximation factorization based on LU decomposition with column/row tournament pivoting. There are two different cases that we consider here: in the first the rank k of the approximation is known, and in the second the rank K of the approximation is to be determined while computing the factorization. We also discuss the numerical properties of our LU factorization. We present bounds on the singular values of the obtained factors with respect to the original matrix A. In addition we also discuss the backward stability of the LU factorization.
- **3.1. Rank-k approximation, when** k **is known.** We consider first the case in which the rank k is known. Given a matrix $A \in \mathbb{R}^{m \times n}$, we are seeking a factorization of the form

(19)
$$\bar{A} = P_r A P_c = \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ \bar{A}_{21} & \bar{A}_{22} \end{pmatrix} = \begin{pmatrix} I \\ \bar{A}_{21} \bar{A}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ & S(\bar{A}_{11}) \end{pmatrix},$$

where

(20)
$$S(\bar{A}_{11}) = \bar{A}_{22} - \bar{A}_{21}\bar{A}_{11}^{-1}\bar{A}_{12}.$$

The permutation matrices P_r, P_c are chosen such that the singular values of \bar{A}_{11} approximate well the largest k singular values of A, while the singular values of $S(\bar{A}_{11})$

approximate well the smallest n-k singular values. If this factorization is run to completion, that is, if $S(\bar{A}_{11})$ is factored by using the same procedure and the factorization obtained is $P'_rAP'_c = LU$, then it is known that the stability of the factorization depends on the ratio $||\hat{L}||_{max}||\hat{U}||_{max}$, where \hat{L} and \hat{U} are the computed factors. For more details see [14], and also [30]. We make the assumption that $||\hat{L}||_{max}||\hat{U}||_{max} \approx ||L||_{max}||U||_{max}$, i.e., that roundoff does not change the norms of the L and U factors very much, and in the following we are interested in bounding $||L||_{max}$ and $||U||_{max}$. For this the elements of $|\bar{A}_{21}\bar{A}_{11}^{-1}|$ need to be bounded.

The algorithm selects the first k columns by using QR factorization with tournament pivoting on the matrix A,

$$AP_c = Q \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix} = \begin{pmatrix} Q_{11} & Q_{12} \\ Q_{21} & Q_{22} \end{pmatrix} \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix},$$

where $Q \in \mathbb{R}^{m \times m}$, $Q_{11}, R_{11} \in \mathbb{R}^{k \times k}$. As discussed in section 2.2, R_{11} reveals the largest k singular values of A, while R_{22} reveals the smallest n-k singular values of A. However, with respect to the factorization in (19), we want \bar{A}_{11} to reveal the largest k singular values of A and $S(\bar{A}_{11})$ the smallest n-k singular values of A. For this, we select k rows from the first k columns of k0 by using k2 with tournament pivoting on k3.

(21)
$$P_r Q = \begin{pmatrix} \bar{Q}_{11} & \bar{Q}_{12} \\ \bar{Q}_{21} & \bar{Q}_{22} \end{pmatrix} = \begin{pmatrix} I \\ \bar{Q}_{21} \bar{Q}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{Q}_{11} & \bar{Q}_{12} \\ & S(\bar{Q}_{11}) \end{pmatrix},$$

where

(22)
$$S(\bar{Q}_{11}) = \bar{Q}_{22} - \bar{Q}_{21}\bar{Q}_{11}^{-1}\bar{Q}_{12} = \bar{Q}_{22}^{-T},$$

such that $\rho_j(\bar{Q}_{21}\bar{Q}_{11}^{-1}) = \rho_j(\bar{A}_{21}\bar{A}_{11}^{-1}) \leq F_{TP}$ for all $1 \leq j \leq m-k$ is upper bounded as in (17), and the singular values of \bar{Q}_{11} and \bar{Q}_{22} are bounded as in (18) (see section 2.3). This will allow us to show that \bar{A}_{11} and $S(\bar{A}_{11})$ reveal the singular values of A.

By applying the column and row permutations on the matrix A we obtain the desired factorization,

$$P_{r}AP_{c} = \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ \bar{A}_{21} & \bar{A}_{22} \end{pmatrix} = \begin{pmatrix} I \\ \bar{Q}_{21}\bar{Q}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{Q}_{11} & \bar{Q}_{12} \\ & S(\bar{Q}_{11}) \end{pmatrix} \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix}$$

$$= \begin{pmatrix} I \\ \bar{A}_{21}\bar{A}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ & S(\bar{A}_{11}) \end{pmatrix},$$
(23)

where

$$\bar{Q}_{21}\bar{Q}_{11}^{-1} = \bar{A}_{21}\bar{A}_{11}^{-1},$$

$$S(\bar{A}_{11}) = S(\bar{Q}_{11})R_{22} = \bar{Q}_{22}^{-T}R_{22}.$$

A similar derivation has been used by Pan in [34] to prove the existence of a rank revealing LU factorization. However, this derivation was not used in the rank revealing algorithms introduced in that paper. Indeed computing P_c by using a classic algorithm like QRCP would require computing R_{22} , and hence R_{22} could be used in the subsequent steps instead of computing an LU factorization and its Schur complement $S(\bar{A}_{11})$. In addition, the derivation in [34] does not bound $||A_{21}A_{11}^{-1}||_{max}$, which is important for the numerical stability of the LU factorization.

We present the LU factorization with column/row tournament pivoting obtained by the above procedure, which we refer to as LU_CRTP (A, k) in Algorithm 1. The first step of the algorithm (line 3) selects k columns by using QR with tournament pivoting on A. This factorization computes strong RRQR factorizations of subsets of 2k columns, but it never computes the complete QR factorization that was used to derive the algebra in (23). Once the QR factorization of the selected k columns from tournament pivoting is computed in step 4, k rows are selected by using tournament pivoting on $Q(:, 1:k)^T$ in step 5. The row and column permutations are applied to A and Q in step 6.

The computation of L_{21} in step 7 requires special attention. In infinite precision, $\bar{A}_{21}\bar{A}_{11}^{-1}=\bar{Q}_{21}\bar{Q}_{11}^{-1}$, however this might not be the case in finite precision. Due to round-off error, the computation of $\bar{Q}_{21}\bar{Q}_{11}^{-1}$ is numerically more stable than the computation of $\bar{A}_{21}\bar{A}_{11}^{-1}$. In the sparse case, not only the numerical values differ, but also $\bar{Q}_{21}\bar{Q}_{11}^{-1}$ has more nonzeros than $\bar{A}_{21}\bar{A}_{11}^{-1}$. This is because \bar{Q}_{11} and \bar{Q}_{21} are denser than \bar{A}_{11} and \bar{A}_{21} . The additional nonzeros correspond to exact cancellations and they are due to round-off errors. In our numerical experiments we observe that they have very small values. We do not investigate further this issue in this paper and we use a heuristic in which we first compute $L_{21} = \bar{A}_{21}\bar{A}_{11}^{-1}$ and we check if $||\bar{A}_{21}\bar{A}_{11}^{-1}||_{max}$ is smaller than $\sqrt{n}||A||_{max}$ (this corresponds to the growth factor observed experimentally for partial pivoting). If it is larger than $\sqrt{n}||A||_{max}$, we compute the L factor as $L_{21} = \bar{Q}_{21}\bar{Q}_{11}^{-1}$. We note that Algorithm 1 selects the columns and the rows that can be used in a CUR decomposition, in which case step 7 can be omitted.

Algorithm 1 LU_CRTP(A, k): rank-k truncated LU factorization with column/row tournament pivoting of a matrix A.

- 1: Input $A \in \mathbb{R}^{m \times n}$, target rank k
- 2: **Output** permutation matrices P_r , P_c , rank-k truncated factorization L_kU_k , factor R_k , such that $(AP_c)(1:k) = Q_kR_k$,

$$\begin{split} P_r A P_c &= \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ \bar{A}_{21} & \bar{A}_{22} \end{pmatrix} = \begin{pmatrix} I \\ \bar{A}_{21} \bar{A}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ & S(\bar{A}_{11}) \end{pmatrix}, \\ L_k &= \begin{pmatrix} I \\ \bar{A}_{21} \bar{A}_{11}^{-1} \end{pmatrix} = \begin{pmatrix} I \\ L_{21} \end{pmatrix}, \ U_k = \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \end{pmatrix}, \end{split}$$

where $L_k \in \mathbb{R}^{m \times k}$, $U_k \in \mathbb{R}^{k \times n}$, $R_k \in \mathbb{R}^{k \times k}$, and the remaining matrices have corresponding dimensions.

Note that $S(\bar{A}_{11})$ is not computed in the algorithm.

3: Select k columns by using QR with tournament pivoting on A,

$$P_c \leftarrow QR_TP(A, k)$$

- 4: Compute the thin QR factorization of the selected columns, $(AP_c)(:, 1:k) = Q_k R_k$, where $Q_k \in \mathbb{R}^{m \times k}$ and $R \in \mathbb{R}^{k \times k}$
- 5: Select k rows by using QR with tournament pivoting on Q_k^T . $P_r \leftarrow QR.TP(Q_k^T, k)$

$$ar{A} = P_r A P_c = \begin{pmatrix} ar{A}_{11} & ar{A}_{12} \\ ar{A}_{21} & ar{A}_{22} \end{pmatrix}, \quad P_r Q_k = \begin{pmatrix} ar{Q}_{11} \\ ar{Q}_{21} \end{pmatrix}$$

7: Compute

$$L_{21} = \bar{Q}_{21}\bar{Q}_{11}^{-1} = \bar{A}_{21}\bar{A}_{11}^{-1}$$
 (see discussion in the text)

In the following theorem we show that the LU_CRTP (A, k) factorization reveals the singular values of A and in addition also bounds element growth in the LU factorization.

THEOREM 3.1. Let A be an $m \times n$ matrix. The LU_CRTP(A, k) factorization obtained by using Algorithm 1,

$$(24) \qquad \bar{A} = P_r A P_c = \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ \bar{A}_{21} & \bar{A}_{22} \end{pmatrix} = \begin{pmatrix} I \\ \bar{Q}_{21} \bar{Q}_{11}^{-1} & I \end{pmatrix} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ & S(\bar{A}_{11}) \end{pmatrix},$$

where

(25)
$$S(\bar{A}_{11}) = \bar{A}_{22} - \bar{A}_{21}\bar{A}_{11}^{-1}\bar{A}_{12} = \bar{A}_{22} - \bar{Q}_{21}\bar{Q}_{11}^{-1}\bar{A}_{12},$$

satisfies the following properties.

(26)
$$\rho_{l}(\bar{A}_{21}\bar{A}_{11}^{-1}) = \rho_{l}(\bar{Q}_{21}\bar{Q}_{11}^{-1}) \leq F_{TP},$$

$$||S(\bar{A}_{11})||_{max} \leq \min((1 + F_{TP}\sqrt{k})||A||_{max}, q_{2}(m, k)F_{TP}\sigma_{k}(A),$$

(27)
$$q_1(n,k)q_2(m,k)\sigma_{k+1}(A)$$
,

(28)
$$1 \le \frac{\sigma_i(A)}{\sigma_i(\bar{A}_{11})}, \frac{\sigma_j(S(\bar{A}_{11}))}{\sigma_{k+j}(A)} \le q_1(n,k)q_2(m,k)$$

for any $1 \le l \le m-k$, $1 \le i \le k$, and $1 \le j \le \min(m,n)-k$. Here F_{TP} is the bound obtained from QR with tournament pivoting, as in (13), $q_1(n,k) = \sqrt{1+F_{TP}^2(n-k)}$, $q_2(m,k) = \sqrt{1+F_{TP}^2(m-k)}$.

Proof. The left part of (28) is satisfied for any permutation matrices P_r , P_c by the interlacing property of singular values. The factorization from (24) can be written as in (23), where

$$\bar{A}_{11} = \bar{Q}_{11}R_{11},$$

(30)
$$S(\bar{A}_{11}) = S(\bar{Q}_{11})R_{22} = \bar{Q}_{22}^{-T}R_{22}.$$

The permutation P_c and the R factor from (23) are obtained by using QR with tournament pivoting, and the singular values of R_{11} satisfy the bounds from (12). The row permutation P_r and \bar{Q}_{11} are obtained by using QR with tournament pivoting, and as discussed in section 2.3, the singular values of \bar{Q}_{11} satisfy the bounds from (18). We obtain

$$\sigma_i(\bar{A}_{11}) \ge \sigma_{min}(\bar{Q}_{11})\sigma_i(R_{11}) \ge \frac{1}{q_1(n,k)q_2(m,k)}\sigma_i(A).$$

Note that $\sigma_{min}(\bar{Q}_{11}) = \sigma_{min}(\bar{Q}_{22})$. We obtain

$$\sigma_j(S(\bar{A}_{11})) = \sigma_j(S(\bar{Q}_{11})R_{22}) \le ||S(\bar{Q}_{11})||_2 \sigma_j(R_{22}) \le q_1(n,k)q_2(m,k)\sigma_{k+j}(A),$$

the bounds on the singular values in (28).

To bound element growth in the L factor, (26), we note that $\bar{A}_{21}\bar{A}_{11}^{-1} = \bar{Q}_{21}\bar{Q}_{11}^{-1}$. As shown in section 2.3, we have that $\rho_l(\bar{A}_{21}\bar{A}_{11}^{-1}) = \rho_l(\bar{Q}_{21}\bar{Q}_{11}^{-1}) \leq F_{TP}$ for each row l of $\bar{A}_{21}\bar{A}_{11}^{-1}$. Element growth in $S(\bar{A}_{11})$ is bounded as follows:

$$\begin{split} |S(\bar{A}_{11})(i,j)| &= |\bar{A}_{22}(i,j) - (\bar{A}_{21}\bar{A}_{11}^{-1})(i,:)\bar{A}_{12}(:,j)| \\ &\leq ||A||_{max} + ||(\bar{A}_{21}\bar{A}_{11}^{-1})(i,:)||_{2}||\bar{A}_{12}(:,j)||_{2} \\ &\leq ||A||_{max} + \rho_{i}(\bar{A}_{21}\bar{A}_{11}^{-1})\sqrt{k}||A||_{max} \\ &\leq (1 + F_{TP}\sqrt{k})||A||_{max}. \end{split}$$

In addition, we use the fact that the QR factorization with tournament pivoting that selects k columns satisfies (10), and we obtain $\chi_j(R_{22}) = ||R_{22}(:,j)||_2 \le F_{TP}\sigma_{min}(R_{11}) \le F_{TP}\sigma_k(A)$. The absolute value of the element in position (i,j) of $S(\bar{A}_{11})$ can be bounded as follows:

$$|S(\bar{A}_{11})(i,j)| = |\bar{Q}_{22}^{-T}(i,:)R_{22}(:,j)| \le ||\bar{Q}_{22}^{-1}(:,i)||_2 ||R_{22}(:,j)||_2$$

$$\le ||\bar{Q}_{22}^{-1}||_2 ||R_{22}(:,j)||_2 = ||R_{22}(:,j)||_2 / \sigma_{min}(\bar{Q}_{22}) \le q_2(m,k)F_{TP}\sigma_k(A).$$

The last bound in (27) is obtained directly from (28).

The low rank approximation matrix obtained from Algorithm 1 can be written as

(31)
$$\tilde{A}_k = \begin{pmatrix} I \\ \bar{A}_{21}\bar{A}_{11}^{-1} \end{pmatrix} \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \end{pmatrix}$$

and

(32)
$$\bar{A} = \tilde{A}_k + \begin{pmatrix} 0_k & \\ & S(\bar{A}_{11}) \end{pmatrix},$$

where 0_k is a matrix of size $k \times k$ with all elements being zero. From (28) we obtain that

(33)
$$\|\bar{A} - \tilde{A}_k\|_2 \le q_1(n, k)q_2(m, k)\sigma_{k+1}(A).$$

Therefore \tilde{A}_k is within a factor of $q_1(n,k)q_2(m,k)$ of the best low rank approximation computed by the SVD. From (32) and (33), by applying Corollory 8.6.2 from [21], we obtain that $|\sigma_i(A) - \sigma_i(\tilde{A}_k)| \leq q_1(n,k)q_2(m,k)\sigma_{k+1}(A)$ for any $1 \leq i \leq k$. This leads to the following relation between the singular values of A, \tilde{A}_k , and $S(\bar{A}_{11})$:

(34)
$$1 - \frac{\sigma_{k+1}(A)}{\sigma_i(A)} q_1(n,k) q_2(m,k) \le \frac{\sigma_i(\tilde{A}_k)}{\sigma_i(A)} \le 1 + \frac{\sigma_{k+1}(A)}{\sigma_i(A)} q_1(n,k) q_2(m,k)$$

for any $1 \leq i \leq k$. Hence, the ratio $\sigma_i(\tilde{A}_k)/\sigma_i(A)$ is close to 1 when

$$\frac{\sigma_{k+1}(A)}{\sigma_i(A)}q_1(n,k)q_2(m,k)$$

is small. This is the case, for example, when there is a sufficiently large gap between $\sigma_k(A)$ and $\sigma_{k+1}(A)$.

3.2. Rank-K approximation, when K is to be determined. We consider now the case when the rank of the approximation needs to be determined during the factorization. Our factorization determines an overestimation K of the rank such that the approximated singular values are larger than a tolerance τ . The overestimation K is a multiple of k, i.e., K = tk and the rank has a value between (t-1)k and K = tk. Another option is to identify a gap in the singular values; we do not discuss this case here, but the algorithm that we present can be easily adapted.

Given an initial guess of the rank k, the factorization computes one step of a block LU factorization with column/row tournament pivoting, as described in the previous section, and obtains the factorization from (19). If the approximated singular values are all larger or equal than τ , then the factorization continues recursively on the trailing matrix $S(\bar{A}_{11})$. We refer to this factorization as $LU_{-}CRTP(A, k, \tau)$, which

is presented in Algorithm 2. After T steps, we obtain the factorization from (35). In the following theorem we give relations between the singular values of the diagonal blocks U_{ii} , $1 \leq i \leq T$ and the largest K singular values of A, and also between the singular values of $U_{T+1,T+1}$ and the smallest n-K singular values of A. We observe that with respect to the results in Theorem 3.1, the bounds here become exponential, where the exponent is the number of steps of tournament pivoting, K/k. An exponentially growing bound cannot be avoided without additional permutations, since in the special case k=1, the algorithm reduces to QRCP, where we know exponential growth is possible.

Theorem 3.2. Suppose we have computed T block steps of $LU_{-}CRTP(A, k, \tau)$ factorization by using Algorithm 2, where $A \in \mathbb{R}^{m \times n}$ and K = Tk. We obtain the following factorization:

$$(35) P_{r}AP_{c} = L_{K}U_{K}$$

$$= \begin{pmatrix} I & & & \\ L_{21} & I & & \\ \vdots & \vdots & \ddots & \\ L_{T1} & L_{T2} & \dots & I \\ L_{T+1,1} & L_{T+1,2} & \dots & L_{T+1,T} & I \end{pmatrix} \begin{pmatrix} U_{11} & U_{12} & \dots & U_{1T} & U_{1,T+1} \\ & U_{22} & \dots & U_{2T} & U_{2,T+1} \\ & & \ddots & \vdots & \vdots \\ & & & U_{TT} & U_{T,T+1} \\ & & & & & U_{T+1,T+1} \end{pmatrix},$$

where $L_{i+1,j}$ and U_{ij} are $k \times k$ for $1 \leq i,j \leq T$, and $U_{T+1,T+1}$ is $(m-Tk) \times (n-Tk)$. The following properties are satisfied:

(36)
$$\rho_l(L_{i+1,j}) \leq F_{TP},$$

$$||U_K||_{max} \leq \min((1 + F_{TP}\sqrt{k})^{K/k}||A||_{max}, q_2(m,k)q(m,n,k)^{K/k-1}\sigma_K(A),$$

$$K/k-1$$

(37)
$$\sigma_{K+1}(A) \prod_{n=0}^{K/k-1} q(m-vk, n-vk, k)),$$

(38)
$$\frac{1}{\prod_{v=0}^{t-2} q(m-vk, n-vk, k)} \le \frac{\sigma_{(t-1)k+i}(A)}{\sigma_i(U_{tt})} \le q(m-(t-1)k, n-(t-1)k, k),$$

(38)
$$\frac{1}{\prod_{v=0}^{t-2} q(m-vk, n-vk, k)} \le \frac{\sigma_{(t-1)k+i}(A)}{\sigma_i(U_{tt})} \le q(m-(t-1)k, n-(t-1)k, k),$$

$$1 \le \frac{\sigma_j(U_{T+1, T+1})}{\sigma_{K+j}(A)} \le \prod_{v=0}^{K/k-1} q(m-vk, n-vk, k)$$

for any $1 \le l \le k$, $1 \le i \le k$, $1 \le t \le T$, and $1 \le j \le \min(m, n) - K$. Here F_{TP} is the bound obtained from QR with tournament pivoting as given in (13), $q_2(m,k) =$ $\sqrt{1+F_{TP}^2(m-k)}$, and $q(m,n,k) = \sqrt{(1+F_{TP}^2(n-k))(1+F_{TP}^2(m-k))}$.

Proof. We consider the first and second step of the block factorization, written as

(40)
$$P_{r1}AP_{c1} = \begin{pmatrix} I \\ L_{2:T+1,1} & I \end{pmatrix} \begin{pmatrix} U_{11} & U_{1,2:T+1} \\ & S(U_{11}) \end{pmatrix},$$
(41)
$$P_{r2}S(U_{11})P_{c2} = \begin{pmatrix} I \\ L_{3:T+1,2} & I \end{pmatrix} \begin{pmatrix} U_{22} & U_{2,3:T+1} \\ & S(U_{22}) \end{pmatrix},$$

(41)
$$P_{r2}S(U_{11})P_{c2} = \begin{pmatrix} I \\ L_{3:T+1,2} & I \end{pmatrix} \begin{pmatrix} U_{22} & U_{2,3:T+1} \\ & S(U_{22}) \end{pmatrix}$$

where $L_{2:T+1,1}$ is the first block column formed by $[L_{21}; \ldots; L_{T+1,1}]$ from (35). Similar notation is used for the other block columns of L or block rows of U. We bound first the singular values of the obtained factorization. By applying Theorem 3.1 we have

(42)
$$1 \leq \frac{\sigma_{i}(A)}{\sigma_{i}(U_{11})}, \frac{\sigma_{j}(S(U_{11}))}{\sigma_{k+j}(A)} \leq q(m, n, k)$$
for any $1 \leq i \leq k, \ 1 \leq j \leq \min(m, n) - k,$

$$1 \leq \frac{\sigma_{i}(S(U_{11}))}{\sigma_{k+j}(U_{11})}, \frac{\sigma_{j}(S(U_{22}))}{\sigma_{j}(S(U_{22}))} \leq q(m - k, n - k, k)$$

(43)
$$1 \le \frac{\sigma_i(S(U_{11}))}{\sigma_i(U_{22})}, \frac{\sigma_j(S(U_{22}))}{\sigma_{k+j}(S(U_{11}))} \le q(m-k, n-k, k)$$
 for any $1 \le i \le k$, $1 \le j \le \min(m, n) - 2k$.

We have that

$$\frac{1}{q(m,n,k)} \le \frac{\sigma_{k+i}(A)}{\sigma_i(S(U_{11}))} \frac{\sigma_i(S(U_{11}))}{\sigma_i(U_{22})} \le q(m-k,n-k,k) \text{ for any } 1 \le i \le k,$$

and we obtain

(44)
$$\frac{1}{q(m,n,k)} \le \frac{\sigma_{k+i}(A)}{\sigma_i(U_{22})} \le q(m-k,n-k,k) \text{ for any } 1 \le i \le k.$$

We also have

$$1 \le \frac{\sigma_j(S(U_{22}))}{\sigma_{k+j}(S(U_{11}))} \frac{\sigma_{k+j}(S(U_{11}))}{\sigma_{2k+j}(A)} \le q(m,n,k)q(m-k,n-k,k)$$

for any $1 \le j \le \min(m, n) - 2k$, and we obtain

(45)
$$1 \le \frac{\sigma_j(S(U_{22}))}{\sigma_{2k+j}(A)} \le q(m,n,k)q(m-k,n-k,k).$$

We consider the steps t and t+1 of the factorization,

$$(46) P_{rt}S(U_{t-1,t-1})P_{ct} = \begin{pmatrix} I \\ L_{t+1:T+1,s} & I \end{pmatrix} \begin{pmatrix} U_{tt} & U_{t,t+1:T+1} \\ S(U_{tt}) \end{pmatrix},$$

$$(47) P_{r,t+1}S(U_{tt})P_{c,t+1} = \begin{pmatrix} I \\ L_{t+2:T+1,t+1} & I \end{pmatrix} \begin{pmatrix} U_{t+1,t+1} & U_{t+1,t+2:T+1} \\ S(U_{t+1,t+1}) \end{pmatrix}.$$

We suppose that the bounds on singular values at the tth step of the factorization satisfy the following relations:

$$\begin{split} \frac{1}{\prod_{v=0}^{t-2} q(m-vk,n-vk,k)} &\leq \frac{\sigma_{(t-1)k+i}(A)}{\sigma_{i}(U_{tt})} \leq q(m-(t-1)k,n-(t-1)k,k), \\ 1 &\leq \frac{\sigma_{j}(S(U_{tt}))}{\sigma_{tk+j}(A)} \leq \prod_{v=0}^{t-1} q(m-vk,n-vk,k) \end{split}$$

for any $1 \le i \le k$ and $1 \le j \le \min(m, n) - tk$. In addition by applying Theorem 3.1 to the factorization obtained at the (t+1)th step from (47), we have

(48)
$$1 \leq \frac{\sigma_i(S(U_{tt}))}{\sigma_i(U_{t+1,t+1})}, \frac{\sigma_j(S(U_{t+1,t+1}))}{\sigma_{k+j}(S(U_{tt}))} \leq q(m-tk, n-tk, k)$$
for any $1 < i < k, \ 1 < j < \min(m, n) - (t+1)k$.

We have that

$$(49) \qquad \frac{1}{\prod_{v=0}^{t-1} q(m-vk, n-vk, k)} \leq \frac{\sigma_{tk+i}(A)}{\sigma_i(S(U_{tt}))} \frac{\sigma_i(S(U_{tt}))}{\sigma_i(U_{t+1, t+1})} \leq q(m-tk, n-tk, k),$$

$$1 \leq \frac{\sigma_j(S(U_{t+1, t+1}))}{\sigma_{k+j}(S(U_{tt}))} \frac{\sigma_{k+j}(S(U_{tt}))}{\sigma_{(t+1)k+j}(A)} \leq q(m-tk, n-tk, k) \prod_{v=0}^{t-1} q(m-vk, n-vk, k)$$

for any $1 \le i \le k$ and $1 \le j \le min(m, n) - (t+1)k$. We obtain the following bounds for the (t+1) step of block factorization:

(50)
$$\frac{1}{\prod_{v=0}^{t-1} q(m-vk, n-vk, k)} \le \frac{\sigma_{tk+i}(A)}{\sigma_i(U_{t+1, t+1})} \le q(m-tk, n-tk, k),$$

(51)
$$1 \le \frac{\sigma_j(S(U_{t+1,t+1}))}{\sigma_{(t+1)k+j}(A)} \le \prod_{v=0}^t q(m-vk, n-vk, k)$$

for any $1 \le i \le k$ and $1 \le j \le min(m, n) - (t + 1)k$. This proves by induction the bounds from (38) and (39).

The bound on element growth in L_K from (36) follows from (26). The part of the bound from (37) that bounds $||U_K||_{max}$ with respect to $||A||_{max}$ can be easily obtained, and it has similarities with the bound obtained from LU with panel rank revealing factorization; see section 2.2.1 in [30]. The part of the bound of $||U_K||_{max}$ with respect to the singular values of A is obtained as follows. After the first iteration, by using the second part of the bound from (27), we have that $||S(U_{11})||_{max} \leq F_{TP}q_2(m,k)\sigma_k(A)$. After the second iteration, by using the second part of the bound from (27) and (42), we obtain

$$||S(U_{22})||_{max} \le F_{TP}q_2(m-k,k)\sigma_k(S(U_{11})) \le F_{TP}q_2(m-k,k)q(m,n,k)\sigma_{2k}(A).$$

After the third iteration, by using the second part of the bound from (27), we have that $||S(U_{33})||_{max} \leq F_{TP}q_2(m-2k,k)\sigma_k(S(U_{22}))$. By using (45), we obtain

$$||S(U_{33})||_{max} \le F_{TP}q_2(m-2k,k)q(m,n,k)q(m-k,n-k,k)\sigma_{3k}(A).$$

The second bound on pivot growth from (37) follows by induction. Note that a tighter bound can be obtained for each row block of U_K , but for simplicity we do not give it here. The third bound on pivot growth from (37) follows easily from (39).

Algorithm 2 presents the LU_CRTP (A,k,τ) factorization, which can be used when the rank of the low rank approximation needs to be determined. It can be easily seen from the bounds obtained in Theorems 3.1 and 3.2 that the R factor obtained from the QR factorization of every block of columns provides a slightly better estimation of the singular values of A than the diagonal blocks of U_K . This is why the algorithm uses the R factor to approximate the singular values of A and determine the rank K of the approximation.

3.3. A less expensive LU factorization with column tournament pivoting. In this section we present a less expensive LU factorization with column and row permutations which only satisfies some of the bounds from Theorem 3.1. We present only one step of the block factorization, in which the desired rank is k, and the extension to a larger rank K is straightforward.

Algorithm 2 LU_CRTP (A, k, τ) : rank-K truncated LU factorization with column/row tournament pivoting of a matrix A, using tolerance τ to identify singular values large enough to keep in the low rank approximation.

- 1: Input $A \in \mathbb{R}^{m \times n}$, k, tolerance τ
- 2: **Output** permutation matrices P_r, P_c , rank K, truncated factorization $B = \tilde{L}_K \tilde{U}_K$ such that $\tilde{L}_K \in \mathbb{R}^{m \times K}$, $\tilde{U}_K \in \mathbb{R}^{K \times n}$

 $\tilde{\sigma}_{K-k}(A) \ge \tau > \tilde{\sigma}_K(A)$, where $\tilde{\sigma}_k(A)$ is Kth singular value of A approximated by the algorithm,

$$\begin{split} P_{r}AP_{c} &= L_{K}U_{K} \\ &= \begin{pmatrix} I & & & \\ L_{21} & I & & \\ \vdots & \vdots & \ddots & \\ L_{T1} & L_{T2} & \dots & I \\ L_{T+1,1} & L_{T+1,2} & \dots & L_{T+1,T} \end{pmatrix} \begin{pmatrix} U_{11} & U_{12} & \dots & U_{1T} & U_{1,T+1} \\ & U_{22} & \dots & U_{2T} & U_{2,T+1} \\ & & \ddots & \vdots & \vdots \\ & & & U_{TT} & U_{T,T+1} \end{pmatrix}, \\ \tilde{L}_{K} &= L_{K}(:,1:K) & \tilde{U}_{K} &= (1:K,:) \end{split}$$

Note that $U_{T,T+1}$ is not updated during the last iteration.

```
3: A = A
 4: for T = 1 to n/k do
          j = (T-1)k+1, K = j+k-1
          Determine row/column permutations by using Algorithm 1,
                 [P_{r_k}, P_{c_k}, L_k, U_k, R_k] \leftarrow LU\_CRTP(\bar{A}(j:m, j:n), k)
         P_{r_T} = \begin{pmatrix} I & & \\ & P_{r_k} \end{pmatrix}, P_{c_T} = \begin{pmatrix} I & \\ & P_{c_k} \end{pmatrix}, \text{ where } I \text{ is } (j-1) \times (j-1)
\bar{A} = P_r A P_c, L_K = P_r L_K P_c, U_K = P_r U_K P_c, P_r = P_{r_T} P_r, P_c = P_c P_{c_T} P_r
 8:
          U_K(j:K,j:n) = U_k, L_K(j:m,j:K) = L_k
9:
          for i = 1 to k do
10:
               \tilde{\sigma}_{j+i-1}(A) = \sigma_i(R_k)
11:
          end for
12:
          if \tilde{\sigma}_K(A) < \tau then
13:
              Return \tilde{L}_K = L_K(:, 1:K), \tilde{U}_K = (1:K,:), K, P_r, P_c
14:
15:
               Update the trailing matrix,
16:
                   \bar{A}(K+1:m,K+1:n) = \bar{A}(K+1:m,K+1:n) - L_k U_k.
          end if
17:
18: end for
```

We refer to this factorization as LU_CTP, LU with column tournament pivoting. Given $A \in \mathbb{R}^{m \times n}$, the factorization selects k columns by using QR factorization with tournament pivoting on the matrix A. The factorization obtained is

$$AP_c = \begin{pmatrix} A_{11}^c & A_{12}^c \\ A_{21}^c & A_{22}^c \end{pmatrix} = Q \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix} = \begin{pmatrix} Q_{11} & Q_{12} \\ Q_{21} & Q_{22} \end{pmatrix} \begin{pmatrix} R_{11} & R_{12} \\ & R_{22} \end{pmatrix},$$

where $A_{11}^c \in \mathbb{R}^{k \times k}$, $Q \in \mathbb{R}^{m \times m}$, $Q_{11}, R_{11} \in \mathbb{R}^{k \times k}$. Note that the column permutation is the same as the one used in $LU_CRTP(A, k)$. However the row permutation is obtained by computing LU with partial pivoting of the first k columns of AP_c . To reduce communication, LU with tournament pivoting can be used to select the k rows

[24], or when necessary for more stability LU with tournament pivoting on the first k columns of Q [12],

$$P_r \begin{pmatrix} A_{11}^c \\ A_{21}^c \end{pmatrix} = \begin{pmatrix} L_{11} \\ L_{21} \end{pmatrix} U_{11},$$

where L_{11}, U_{11} are of dimension $k \times k$. The obtained LU factorization is

$$P_rAP_c = \begin{pmatrix} \bar{A}_{11} & \bar{A}_{12} \\ \bar{A}_{21} & \bar{A}_{22} \end{pmatrix} = \begin{pmatrix} L_{11} \\ L_{21} & I \end{pmatrix} = \begin{pmatrix} U_{11} & U_{12} \\ & S(\bar{A}_{11}) \end{pmatrix},$$

where $S(\bar{A}_{11}) = \bar{A}_{22} - L_{21}U_{12}$. Note that this factorization does not lower bound the singular values of Q_{11} , and this prevents us from bounding the singular values of \bar{A}_{11} and $S(\bar{A}_{11})$ with respect to the singular values of A. However, as we will see in the experimental results, in practice this factorization approximates well the singular values of A.

4. Complexity analysis of QR factorization with tournament pivoting. The selection of k columns by using QR factorization with tournament pivoting dominates the cost of the rank-k approximation factorization from Algorithm 1. We analyze in this section the cost of QR factorization with tournament pivoting for matrices with arbitrary sparsity structure, in terms of fill in the factors Q and R obtained during tournament, floating point operations, as well as interprocessor communication for the parallel version of the algorithm. Given the assumptions we make in our analysis, we expect that these bounds are loose. We then consider matrices whose column intersection graphs have small separators and obtain tighter bounds for the fill in the

Analyzing the cost of the low rank approximation factorization when the rank K needs to be determined (Algorithm 2) is a more difficult problem that we do not address in this paper. Tournament pivoting could potentially select the most dense k columns, and predicting an upper bound on the fill that occurs in the Schur complement can be very pessimistic.

factors Q and R. In both cases, our analysis is based on a column permutation of the

matrix which allows us to bound the fill.

4.1. Matrices with arbitrary sparsity structure. We discuss now the case of an arbitrary sparse matrix $A \in \mathbb{R}^{m \times n}$. Let d_i be the number of nonzeros in column i of A, $nnz(A) = \sum_{i=1}^{n} d_i$. We permute the matrix columns such that $d_1 \leq \ldots \leq d_n$.

Consider QR_TP with a flat tree. As explained in section 2.2, the matrix A is partitioned into n/k blocks of columns as $A = [A_{00}, \ldots, A_{n/k,0}]$. At the first step of tournament pivoting, the matrix $A_{01} = [A_{00}, A_{10}]$ is formed, and k columns are selected by computing the QR factorization of A_{01} , followed by a rank revealing factorization of the R factor. At the subsequent steps, a new matrix A_{ij} is formed with the previously selected k columns and a block A_{i0} of unvisited columns of A. From this matrix, k new columns are selected by using QR factorization followed by rank revealing factorization of the R factor. Given that k is small, we ignore in the following the cost of the rank revealing factorization of the R factors.

At the first step of reduction, the matrix A_{01} formed by the first 2k columns of A has at most $\sum_{i=1}^{2k} d_i$ rows (with equality when the sets of column indices are disjoint and each row has only one nonzero). By considering this matrix dense, $nnz(A_{01}) \leq 2k \sum_{i=1}^{2k} d_i$ and the number of flops required to compute its QR factorization is at most $8k^2 \sum_{i=1}^{2k} d_i$. The selected k columns have at most $\sum_{i=k+1}^{2k} d_i$ nonzeros and form a matrix with at most $\sum_{i=k+1}^{2k} d_i$ rows. The matrix formed at the second

step of tournament pivoting has at most $\sum_{i=k+1}^{3k} d_i$ rows and at most $2k \sum_{i=k+1}^{3k} d_i$ nonzeros, and its QR factorization requires at most $8k^2 \sum_{i=k+1}^{3k} d_i$. These bounds are asymptotically attainable if A has the following nonzero structure: the first $d_1 - 1$ rows have a nonzero only in the first column, while the d_1 th row has nonzeros in the first and the second column. The following $d_1 - 2$ rows have a nonzero in the second column, while the $(d_1 + d_2 - 1)$ th row has nonzeros in the second and the third column. The nonzero structure of the following rows follows the same pattern.

We consider the following costs associated with QR_TP: $nnz_{max}(QR_TP_{FT})$ refers to the maximum number of nonzeros of the Q and R factors over all the QR factorizations computed during tournament pivoting, $nnz_{total}(QR_TP_{FT})$ refers to the sum of the number of nonzeros of all the Q and R factors used during tournament pivoting, and $flops(QR_TP_{FT})$ refers to the total number of flops computed. Note that $nnz_{max}(QR_TP_{FT})$ reflects the memory needs of QR_TP, since once k columns are selected, the corresponding factors Q and R can be discarded before proceeding to the following step of tournament pivoting. We obtain

$$(52) \quad nnz_{max}(QR_TP_{FT}) \le 4d_nk^2,$$

(53)
$$nnz_{total}(QR_TP_{FT}) \le 2k \left(\sum_{i=1}^{2k} d_i + \sum_{i=k+1}^{3k} d_i + \ldots + \sum_{i=n-2k+1}^{n} d_i \right) \le 4nnz(A)k,$$

$$(54) flops(QR_TP_{FT}) \le 16nnz(A)k^2.$$

We discuss now the case when QR_TP is executed in parallel by using a binary tree of depth $\log(n/k)$, where $2k \leq n/P$. We consider a column cyclic distribution of the matrix on P processors, and we assume that n/P columns with a total of nnz(A)/P nonzeros are stored on each processor. We also make the reasonable assumption that k is small enough such that $\sum_{i=n-2k+1}^{n} d_i \leq nnz(A)/P$. At each step of the reduction, the matrix A_{ij} has 2k columns and at most $\sum_{i=n-2k+1}^{n} d_i$ rows, and each processor executes $\log(n/k)$ QR factorizations of such matrices. The cost of QR_TP per processor is

(55)
$$nnz_{max}(QR_TP_{BT}) \le 2k \sum_{i=n-2k+1}^{n} d_i \le 2\frac{nnz(A)}{P}k,$$

$$(56) nnz_{total}(QR_TP_{BT}) \le 2k \left(\sum_{i=n-2k+1}^{n} d_i\right) \log \frac{n}{k} \le 2 \frac{nnz(A)}{P} k \log \frac{n}{k},$$

(57)
$$flops(QR_TP_{BT}) \le 8k^2 \left(\sum_{i=n-2k+1}^n d_i\right) \log \frac{n}{k} \le 8\frac{nnz(A)}{P}k^2 \log \frac{n}{k}.$$

Given our assumption that the matrices used during tournament pivoting have only one nonzero per row, these bounds are loose.

In terms of interprocessor communication, it can be easily seen that during tournament pivoting, the processors exchange $\log P$ messages. Each message has at most $\sum_{i=n-2k+1}^n d_i$ words, for a total of $(\sum_{i=n-2k+1}^n d_i) \log P \le 2kd_n \log P$ words communicated on the critical path of parallel QR_TP.

4.2. Graphs with small separators. Given a matrix $A \in \mathbb{R}^{m \times n}$, the structure of its Q and R factors can be modeled by using the column intersection graph $G_{\cap}(A)$. We consider that at each iteration of the QR factorization, one Householder reflection is used to annihilate the elements below the diagonal. The Householder matrix

H which stores the Householder vectors can be used to implicitly represent Q. Its structure can also be modeled by $G_{\cap}(A)$. The graph $G_{\cap}(A)$ is the undirected graph of A^TA (we ignore numerical cancellations), has n vertices (one for each column of A), and has an edge between two vertices if the corresponding columns of A have a nonzero in a same row. The filled column intersection graph of A [17], $G_{\cap}^+(A)$, is the graph of the Cholesky factor L of A^TA and hence it is also the graph of the R factor of the QR factorization. Both the structure of R and R can be expressed in terms of paths in the column elimination tree of R, R, which is a depth-first spanning tree of R, R, which is an interpolation of R and R connected. The nonzero column indices of a row R of R correspond to vertices that belong to the path in R, and R similar definition holds for the last R and R rows of R.

We consider here the case when the column intersection graph belongs to a class S of graphs with small separators that are closed under the subgraph relation. (That is, if G is in S and G_1 is a subgraph of G, then G_1 is also in S.) We focus on the class S of graphs that are n^{λ} separable, $\lambda < 1$. These graphs have a separator S with cn^{λ} vertices, c > 0, whose removal disconnects the graph into two graphs A and B, each with less than 2/3n vertices. There are many example of graphs with good separators. For example, structured grids in d dimensions are $n^{d-1/d}$ -separable, and planar graphs are $n^{1/2}$ separable. For more detailed discussions see, e.g., [31].

We recall first results established in [19] on the number of nonzeros in the factors R, Q, as well as the Householder vectors H used during the QR factorization of A. These results are obtained by using a reordering of the matrix A based on nested dissection [16], which partitions the graph into two disjoint subgraphs and a separator, and continues recursively the same procedure on the two subgraphs. In the analysis, we consider that c and λ are fixed. We focus first on graphs which are \sqrt{n} -separable.

THEOREM 4.1 (Theorem 1 from [19]). Let A be an $m \times n$ matrix of full column rank, such that $G_{\cap}(A)$ is a member of a \sqrt{n} -separable class of graphs. Then there exists a column permutation P such that the matrices Q, R, and H in the thin QR factorization of AP satisfy the following bounds: $nnz(R) = O(n \log n)$, $nnz(Q) = O(m\sqrt{n})$, $nnz(H) = O(n \log n + (m-n)\sqrt{n})$.

These results are based on the fact that every row of Q has at most $O(\sqrt{n})$ nonzeros, the first n rows of H are a subset of the structure of the rows of R, while the last m-n rows of H have at most $O(\sqrt{n})$ nonzeros each.

Let $A \in \mathbb{R}^{m \times n}$ be a matrix whose column intersection graph $G_{\cap}(A)$ belongs to the class of \sqrt{n} -separable graphs, closed under the subgraph relation. As in the case of arbitrary matrices, we let d_i be the number of nonzeros in column i of A and we consider that A was permuted such that $d_1 \leq \ldots \leq d_n$. We make the same assumption as before that $kd_n \leq nnz(A)/P \leq nd_n/P$. In the case of QR with tournament pivoting based on a flat tree, the matrix formed at the first step of tournament A_{01} has 2k columns of A and has at most $\sum_{i=1}^{2k} d_i$ rows that have nonzero elements. Given that we consider a class of graphs closed under the subgraph relation, the graph of a submatrix A_{ij} formed by 2k columns, $G(A_{ij})$, has 2k vertices and satisfies the $(2k)^{1/2}$ separator theorem. We obtain that at the first step of tournament pivoting, the QR factorization of the current matrix A_{01} leads to factors that have $nnz(R) = O(k \log k)$, $nnz(Q) \leq O(\sqrt{k} \sum_{i=1}^{2k} d_i)$, and $nnz(H) \leq O(k \log k + (\sum_{i=1}^{2k} d_i - k)\sqrt{k})$. Since each row of H has at most $O(\sqrt{k})$ nonzeros, computing the QR factorization of A_{01} costs at most $O(k^{3/2} \sum_{i=1}^{2k} d_i)$ flops. By summing over the n/k QR factorizations used during tournament pivoting and by using the same approach as for matrices with arbitrary

sparsity structure, the cost of QR_TP is

(58)
$$nnz_{max}(QR_TP_{FT}) \le O(d_n k^{3/2}),$$

(59)
$$nnz_{total}(QR_TP_{FT}) \le O(nnz(A)\sqrt{k}),$$

(60)
$$flops(QR_TP_{FT}) \le O(nnz(A)k^{3/2}).$$

These bounds are smaller than the bounds for matrices with arbitrary sparsity structure from (52), (53), and (54) by a factor of $O(\sqrt{k})$. We discuss now the case when QR_TP is executed in parallel using a binary reduction tree as before. The reduction tree used during tournament pivoting has depth $\log(n/k)$, and each processor executes $\log(n/k)$ QR factorizations of matrices A_{ij} with 2k columns and at most $\sum_{i=n-2k+1}^{n} d_i \leq 2kd_n$ rows that have at least one nonzero. Each such QR factorization can be computed in at most $O(k^{3/2}\sum_{i=n-2k+1}^{n} d_i)$ flops. The cost of QR_TP per processor is

$$(61) \quad nnz_{max}(QR_TP_{BT}) \leq O\left(\sqrt{k}\sum_{i=n-2k+1}^n d_i\right) \leq O(d_nk^{3/2}) \leq O\left(\frac{nnz(A)}{P}\sqrt{k}\right),$$

$$(62) \quad nnz_{total}(QR_TP_{BT}) \le O\left(\sqrt{k}\left(\sum_{i=n-2k+1}^{n} d_i\right)\log\frac{n}{k}\right) \le O\left(\frac{nnz(A)}{P}\sqrt{k}\log\frac{n}{k}\right),$$

$$(63) \qquad flops(QR_TP_{BT}) \le O\left(k^{3/2} \left(\sum_{i=n-2k+1}^{n} d_i\right) \log \frac{n}{k}\right) \le O\left(\frac{nnz(A)}{P} k^{3/2} \log \frac{n}{k}\right).$$

These costs are $O(\sqrt{k})$ smaller than the corresponding costs from (55), (56), and (57) obtained for matrices with arbitrary sparsity structure.

By using the same reasoning, these results can be extended to graphs with larger separators. Consider that $G_{\cap}(A)$ belongs to the class of n^{λ} -separable graphs, with $1/2 < \lambda < 1$, closed under the subgraph relation. Nested dissection leads to the following bounds of the factors obtained from QR factorization [19], $nnz(R) = O(n^{2\lambda})$, $nnz(Q) = O(mn^{\lambda})$, $nnz(H) = O(n^{2\lambda} + (m-n)n^{\lambda})$. A path in the column elimination tree $G_{\cap}(A)$ going from its leaves to its root is formed by the vertices of a separator obtained at each level of nested dissection, and hence its length is $O(n^{\lambda})$. We deduce that the number of nonzeros in each row of Q and in the last m-n rows of H is at most $O(n^{\lambda})$. We obtain the following bounds for QR_TP:

$$(64) \quad nnz_{max}(QR_TP_{FT}) = nnz_{max}(QR_TP_{BT}) \le O(d_nk^{\lambda+1}) \le O\left(\frac{nnz(A)}{P}k^{\lambda}\right),$$

(65)
$$nnz_{total}(QR_TP_{FT}) \leq O(nnz(A)k^{\lambda}),$$

(66)
$$flops(QR_TP_{FT}) \leq O(nnz(A)k^{\lambda+1}),$$

(67)
$$nnz_{total}(QR_TP_{BT}) \le O\left(\frac{nnz(A)}{P}k^{\lambda}\log\frac{n}{k}\right),$$

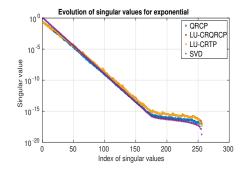
(68)
$$flops(QR_TP_{BT}) \le O\left(\frac{nnz(A)}{P}k^{\lambda+1}\log\frac{n}{k}\right).$$

We note again that these bounds are smaller than their counterparts in section 4.1 by factors $O(k^{1-\lambda})$.

5. Experimental results. In this section we present experimental results which show that LU_CRTP provides a good low rank approximation in terms of both accuracy and speed. We discuss first the accuracy of LU_CRTP by comparing the

Table 1
Test matrices generated in MATLAB.

No.	Matrix	Description
1	Baart	Discretization of the 1st kind Fredholm integral equation [27].
_		9 1 []
2	Break1	Break 1 distribution, matrix with prescribed singular values [3].
3	Break9	Break 9 distribution, matrix with prescribed singular values [3].
4	Deriv2	Computation of second derivative [27].
5	Devil	The devil's stairs, a matrix with gaps in
		its singular values; see [37] or [11].
6	Exponential	Exponential distribution, $\sigma_1 = 1$, $\sigma_i = \alpha^{i-1}$ $(i = 2,, n)$,
		$\alpha = 10^{-1/11}$ [3].
7	Foxgood	Severely ill-posed test problem of the 1st kind Fredholm
		integral equation used by Fox and Goodwin [27].
8	Gravity	One-dimensional gravity surveying problem [27].
9	Heat	Inverse heat equation [27].
10	PHILLIPS	Phillips test problem [27].
11	Random	Random matrix $A = 2* \text{rand(n)} - 1$ [25].
12	Shaw	One-dimensinal image restoration model [27].
13	Spikes	Test problem with a "spiky" solution [27].
14	Stewart	Matrix A = U Σ V ^T + 0.1 σ_m * rand(n), where σ_m
		is the smallest nonzero singular value [37].
15	Ursell	Integral equation with no square integrable solution [27].
16	Wing	Test problem with a discontinuous solution [27].



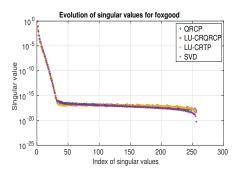


Fig. 1. Evolution of the singular values computed by SVD and approximated by QRCP, LU_CRQRCP (LU with column and row pivoting based on QRCP), and LU_CRTP (LU with column and row tournament pivoting).

approximation of the singular values obtained by our truncated LU factorization with the singular values obtained by SVD. In all the tests, the matrix is first permuted by using COLAMD (column approximate minimum degree) [9] followed by a postorder traversal of its column elimination tree.

5.1. Numerical accuracy. We use two sets of matrices in our numerical experiments, the first one is formed by 16 challenging matrices which are often used for testing rank revealing algorithms, and the second one is formed by matrices from the San Jose State University (SJSU) Singular Matrix Database [15]. We discuss first the results obtained on the first test set. The 16 matrices are generated in MATLAB, they are of size 256×256 , and their description can be found in Table 1. A more detailed description can be found in [11]. Experiments are carried out in double precision in MATLAB 2015a.

Figure 1 displays the evolution of singular values obtained by SVD and their approximation obtained by LU_CRTP for two matrices, EXPONENTIAL and FOXGOOD.

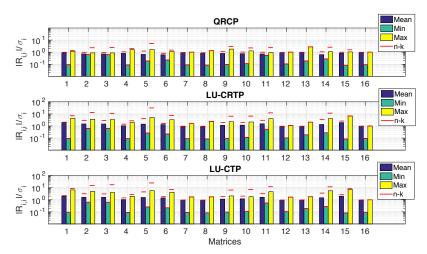


Fig. 2. Comparison of approximations of singular values obtained by LU_CRTP, LU_CTP, and QRCP for the matrices described in Table 1. Here k=16 and the factorization is truncated at K=128 (bars) or K=240 (red lines).

The value of k in LU_CRTP (A, k, τ) from Algorithm 2 is 16. However, the algorithm does not stop at k, but continues the factorization recursively until completion and hence approximates all the singular values of A. The approximated singular values correspond to the diagonal elements of the R factor of each block of k columns obtained from a tournament. In addition, the figure also displays the results obtained by QRCP and those obtained when the column and row permutations are determined by using QRCP instead of tournament pivoting, referred to as LU_CRQRCP. We note that the singular values are well approximated by the three algorithms, and the results obtained by LU_CRQRCP and LU_CRTP are almost superimposed. The usage of tournament pivoting instead of QRCP to select columns and rows in a block LU factorization does not lead to loss of accuracy in our experiments. The same behavior is observed for the remaining matrices in the set from Table 1, and the results can be found in the appendix of the technical report [23, Figure 3].

Figure 2 displays the ratios of the approximated singular values with respect to the singular values for the 16 matrices in our set, summarized by the minimum, maximum, and mean values of the ratios $|R_{i,i}|/\sigma_i(A)$. Three different methods are tested, QRCP, LU_CRTP, and LU_CTP. The last method, LU_CTP, corresponds to the cheaper factorization described in section 3.3 in which only the column permutation is selected by using tournament pivoting based on QR, while the row permutation is based on LU with partial pivoting. The bars display the results obtained when the algorithm is truncated at K = 128, and the red lines display the results obtained when the factorization runs almost to completion; it is truncated at K = n - k. In the results all singular values smaller than machine precision, ϵ , are replaced by ϵ . For the very last k columns, the ratios are slightly larger, and we do not include them in the results, since they are not relevant for a low rank approximation. These results show that the mean is very close to 1. On average, the approximated singular values are very close to the singular values computed by SVD. For the matrices in our set, except devil's stairs, the ratio is between 0.08 and 13.1 for LU_CRTP and between 0.08 and 17.5 for LU_CTP. For the devil's stairs, a more challenging problem for rank revealing factorizations, the maximum ratio for LU_CRTP is 27 and the maximum ratio for LU₋CTP is 26. More detailed results can be found in the appendix of [23],

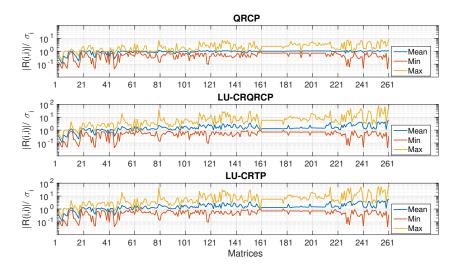


FIG. 3. Comparison of the ratios $|R(i,i)|/\sigma_i$ with $1 \le i \le \text{rank}$ for 261 matrices from the SJSU Singular Matrix Database [15] for QRCP (top), LU_CRQRCP (middle), and LU_CRTP (bottom). LU_CRQRCP and LU_CRTP use k = 16 and rank is the numerical rank of each matrix as given by the database. Each graph displays the mean, the minimum, and the maximum of the ratios $|R(i,i)|/\sigma_i$ for each matrix, and the index of the matrix in the collection is displayed on the x-axis.

where we also discuss the numerical accuracy of the factors L and U produced by LU_CRTP (A, k, τ) (Algorithm 2) on the set of matrices described in Table 3. The results show that the obtained factorization is very stable.

In the second set of experiments we use a set of 261 matrices from the SJSU Singular Matrix Database. These matrices have less than 1024 rows, and between 32 and 2048 columns, and they are obtained by using the following MATLAB code:

The singular values are approximated by using QRCP, LU_CRQRCP, and LU_CRTP until the numerical rank, provided by the database, is reached. For each matrix in the collection, Figure 3 presents the maximum, the minimum, and the mean of the ratios of the diagonal elements of the R factor, R(i,i), obtained by one of the three factorizations to the singular values. The matrices are sorted in increasing order of their numerical rank, and the index of each matrix is displayed on the x-axis.

These results show that the approximations of singular values obtained by LU_CRQRCP are very close to the approximations obtained by LU_CRTP. In other words, the selection of columns and rows during the LU factorization based on tournament pivoting leads to results very close to those obtained when the selection of columns uses QRCP, QR with column pivoting. The results obtained by LU_CRQRCP and LU_CRTP are slightly worse than those obtained by QRCP; these results outline the difference between the approximations of singular values obtained by a rank revealing LU factorization and a rank revealing QR factorization.

Table 2 gives the minimum and the maximum among all the computed ratios displayed in Figure 3. As also discussed in [11], QRCP has a minimum ratio of 4.17e - 2 and a maximum ratio of 8.96. While the minimum ratio is the same for

Table 2

The minimum and the maximum of all the ratios $|R(i,i)|/\sigma_i$, $1 \le i \le \text{rank}$, displayed in Figure 3, where the R factor is obtained by QRCP, LU_CRQRCP, and LU_CRTP. The numerical rank rank is provided by the database for all 261 matrices.

k	Method	mi	n	ma	max		
		matrix id	ratio	matrix id	ratio		
16	QRCP	260	4.17e-2	261	8.96e + 0		
	LU_CRQRCP	260	4.17e-2	253	$8.04e{+1}$		
	LU_CRTP	260	4.17e-2	261	8.49e + 1		

both LU_CRQRCP and LU_CRTP, the maximum ratio is 8.49e2, larger by two orders of magnitude. Typically, at the beginning of the factorization the diagonal elements of the R factor underestimate the singular values, and as the factorization proceeds, at some step they start overestimating the singular values. The largest ratios occur when the numerical rank is approached, and for several matrices, there are only a few outliers. We note also that our experiments use k=16, and for larger matrices we expect k to be larger, 128 or 256, and obtain better approximations of singular values.

As discussed in section 3.1, the computation of the L factor is performed by using the following heuristic. At each iteration of the block LU factorization when Algorithm 1 is called, step 7 computes first $L_{21} = \bar{A}_{21}\bar{A}_{11}^{-1}$. If $||\bar{A}_{21}\bar{A}_{11}^{-1}||_{max} \ge \sqrt{n}||A||_{max}$, the L factor is computed as $L_{21} = \bar{Q}_{21}\bar{Q}_{11}^{-1}$. For the first set of 16 matrices, $||\bar{A}_{21}\bar{A}_{11}^{-1}||_{max}$ is at most 1.49. However, for the second set of matrices from the SJSU Singular Matrix Database, there are 124 matrices for which at some steps $||\bar{A}_{21}\bar{A}_{11}^{-1}||_{max}$ is larger than $\sqrt{n}||A||_{max}$ and hence the L factor is computed as $L_{21} = \bar{Q}_{21}\bar{Q}_{11}^{-1}$.

5.2. Performance. We discuss first the performance of our algorithm by comparing the number of nonzeros in the factors of LU_CRTP with respect to the number of nonzeros in the factors of QRCP and LU with partial pivoting. As mentioned earlier, for all the factorizations, the columns of the matrix were first permuted by using COLAMD followed by a postorder traversal of its column elimination tree. The number of nonzeros in the factors gives not only the memory usage of these factorizations, but also a good indicator of their expected performance. We use several larger square sparse matrices obtained from the University of Florida Sparse Matrix Collection [10]. Table 3 displays the name of the matrices, their number of columns/rows (Size), their number of nonzeros (Nnz), and their field of application. Some of the matrix names are abbreviated. The matrix TSOPF_RS corresponds to the matrix TSOPF_RS_B39_C30 in the collection, PARAB_FEM corresponds to PARABOLIC_FEM, while MAC_ECON_FWD500.

Table 4 displays the results obtained for the first 10 matrices from Table 3 when a rank K approximation is computed, where K varies from 128 to 1024. The initial rank k=16. LU_CRTP uses a binary tree based tournament pivoting. MATLAB is used for these experiments, and there was not enough memory to obtain results for the last matrices in Table 3. The second column, $\operatorname{nnz} A(:,1:K)$, displays the number of nonzeros in the first K columns of A, once it was permuted as explained previously. The fourth column, $\frac{nnz\ QRCP}{nnz\ LU_CRTP}$, displays the ratio of the number of nonzeros of QRCP with respect to the number of nonzeros of LU_CRTP. The last column displays $\frac{nnz\ LU_CRTP}{nnz\ LU_PP}$, the ratio of the number of nonzeros of LU_CRTP with respect to LU with partial pivoting. For QRCP, we count the number of nonzeros in the first K columns of the Householder vectors H plus the number of nonzeros in the first K rows of the K factor. For LU_CRTP and LUPP we count the number of nonzeros in the first K columns of K and the first K rows of K. For LU_CRTP we

Table 3
Sparse matrices from University of Florida collection [10].

No.	Matrix	Size	Nnz	Problem description
17	ORANI678	2529	90158	Economic
18	GEMAT11	4929	33108	Power network sequence
19	raefsky3	21200	1488768	Computational fluid dynamics
20	WANG3	26064	177168	Semiconductor device
21	ONETONE2	36057	222596	Frequency-domain circuit simulation
22	TSOPF_RS	60098	1079986	Power network
23	RFDEVICE	74104	365580	Semiconductor device
24	NCVXQP3	75000	499964	Optimisation
25	MAC_ECON	206500	1273389	Economic
26	PARAB_FEM	525825	3674625	Computational fluid dynamics
27	ATMOSMODD	1270432	8814880	Fluid dynamics
28	CIRCUIT5M_DC	3523317	14865409	Circuit simulation

Table 4
Sparse matrices from University of Florida collection [10].

Name	$nnz\ A(:,1:K)$	Rank K	$\frac{nnz\ QRCP}{nnz\ LU_CRTP}$	nnz LU_CRTP nnz LUPP
ORANI678	7901	128	17.87	3.30
	55711	512	6.18	8.85
	71762	1024	4.86	11.01
GEMAT11	1232	128	2.1	2.2
	4895	512	3.3	2.6
	9583	1024	11.5	3.2
RAEFSKY3	7872	128	1.25	2.26
	31248	512	1.07	4.18
	63552	1024	1.06	6.58
WANG3	896	128	3.0	2.1
	3536	512	2.9	2.1
	7120	1024	2.9	1.2
ONETONE2	4328	128	36.0	2.8
	9700	512	73.5	1.1
	17150	1024	108.5	0.3
TSOPF_RS	4027	128	2.57	1.90
	5563	512	0.83	2.41
	7695	1024	0.61	2.13
RFDEVICE	633	128	10.0	1.1
	2255	512	82.6	0.9
	4681	1024	207.2	0.9
NCVXQP3	1263	128	2.87	1.21
	5067	512	3.50	1.01
	10137	1024	3.83	0.53
MAC_ECON	384	128	-	0.34
	1535	512	-	0.19
	5970	1024	-	0.11
PARAB_FEM	896	128	-	0.5
	3584	512	-	0.3
	7168	1024	-	0.2

ignore the number of nonzeros created during tournament pivoting since the memory requirements are small compared to the memory requirements displayed in Table 4.

We note that for smaller matrices, LU_CRTP leads to a factor of up to 17 times fewer nonzeros than QRCP for ORANI678. Larger improvement with respect to QRCP is observed for ONETONE2 and RFDEVICE, up to a factor of 207. As one can expect, LU_CRTP leads to up to 11 times more nonzeros than LUPP. We were not able to run QRCP for the last two matrices in Table 4 due to memory consumption. We

also observe that for the last four matrices, LU_CRTP has fewer nonzeros than LUPP. This means that the columns selected by tournament pivoting generate less fill-in than those selected before the factorization by using COLAMD and postorder traversal of the elimination tree as used in LUPP. This is something that we do not expect to happen in general.

We discuss now the parallel performance of LU_CRTP. There are routines for computing the QRCP factorization of a dense matrix as in LAPACK or ScaLAPACK, or for computing the QR factorization of a sparse matrix, such as the multifrontal SPQR software of Davis [8]. However there is no library available for computing in parallel a sparse rank revealing factorization. We present in this paper the performance of $LU_{-}CRTP(A, k)$ (Algorithm 1), which given a rank k, computes a rank-k approximation and produces a CUR factorization as presented in (3). The main step of this algorithm is to select k columns from the matrix A by using tournament pivoting based on QR. We do not have yet a parallel implementation of LU_CRTP (A, k, τ) which computes all the singular values larger than a parameter τ , as given in Algorithm 2; this remains future work. Table 6 reports runtimes for tournament pivoting based on QR, which selects k = 256 columns from the input matrix. We use in these tests the last larger matrices from our set in Table 3. The results are obtained on Edison, a Cray XC30 supercomputer at NERSC, formed by nodes with 2 x 12-core Intel "Ivy Bridge" processors. We run as many MPI processes per node as cores. Tournament pivoting uses a binary tree. At each step the selection of k columns from 2k columns is performed by first calling SPQR [8], and then calling the QRCP dGEQP3 routine from Lapack [1] (as implemented in MKL) on the R factor obtained from SPQR. SPQR reorders the matrix by using Metis [29].

Table 5 presents the breakdown of the times required for selecting k=256columns using P = 32 MPI processes. The second column displays the time required for reordering each matrix by using COLAMD followed by a postorder traversal of its column elimination tree. The third column displays the time required for selecting kcolumns from 2k columns by calling SPQR and DGEQP3. The runtimes are shown on three rows, the first row displays the minimum runtime, the second row displays the average runtime, while the last row displays the maximum runtime among processors. These statistics are computed over all calls to SPQR and DGEQP3 performed during tournament pivoting. The fourth column displays the time required for selecting k columns locally on each processor from the matrix of size $m \times n/P$ that it owns. Each processor uses a binary tree based tournament pivoting. As for the previous column, we display first the minimum runtime, then the average runtime, and finally the maximum runtime among processors. The last column displays the time required for selecting the final k columns from the sets of k columns selected locally on each of the P processors. The binary tree used by tournament pivoting among processors has depth log P. Our implementation uses a reduction like operation in which the result is available only on one processor, namely, processor 0. The number of processors involved at each level of the reduction is halved. Hence we display only the maximum runtime obtained on processor 0 that owns the result and is the root of the reduction tree.

These results show that selecting k columns from 2k columns can be performed efficiently. However, for a same matrix, the time spent in SPQR can vary considerably. For example, for MAC_ECOM, SPQR varies between 0.06 seconds and 22.71 seconds and takes on average 3.26 seconds. This results in load imbalance during tournament pivoting both in the first steps performed locally on each processor and in the last $\log P$ steps that involve communication. Essentially most of the time is spent in SPQR. We note that the time spent in the selection of k columns locally can be

Table 5

Runtime in seconds of tournament pivoting for selecting k=256 columns on P=32 MPI processes. The time is divided between the time required for reordering (COLAMD + etree), the time required for selecting k columns from n/P columns locally on each processor, and the time required for selecting k columns from the sets of columns selected by each processor (the last $\log P$ steps of tournament pivoting). For each matrix, the third and the fourth columns display on different rows minimum, average, and maximum runtimes obtained among different processors. The last column displays only the maximum runtime obtained among different processors.

		Tournament pivoting					
Matrix	COLAMD	Matrix $m \times 2k$		on each processor		among processors	
	+etree	SPQR	DGEQP3	SPQR	DGEQP3	SPQR	DGEQP3
		0.06	0.01	180.97	0.33		
MAC_ECON		3.26	0.02	246.65	0.36		
	0.52	22.71	0.02	361.84	0.39	3.37	0.08
		0.20	0.01	54.66	0.94		
PARAB_FEM		0.24	0.02	79.88	1.03		
	0.56	0.29	0.07	103.49	1.13	1.06	0.07
		0.48	0.01	183.36	2.17		
ATMOSMODD		0.92	0.02	236.69	2.39		
	3.35	4.96	0.02	483.61	2.63	2.50	0.05
		1.36	0.01	623.09	6.26		
circuit5M_dc		1.59	0.02	664.68	6.85		
	4.18	1.84	0.03	749.20	7.99	7.15	0.08

Table 6 Runtime in seconds for selecting k=256 columns using tournament pivoting based on QR.

Matrix	Number MPI processes						
	32	64	128	256	512	1024	2048
MAC_ECON	367	183	118	83	57	19	12
PARAB_FEM	106	65	36	22	15	5	4
ATMOSMODD	488	269	163	83	52	29	18
CIRCUIT5M_DC	771	367	196	109	69	44	37

further reduced by using a flat tree. Our future work will focus on this aspect as well as on designing a load balanced tournament pivoting.

Table 6 presents the runtime of binary tree based tournament pivoting when increasing the number of cores from 32 to 1024. For the matrices in our test set, the algorithm is scalable up to 1024 cores. For example, selecting 256 columns from PARAB_FEM requires only 5 seconds on 1024 cores. The runtime decreases only slightly when the number of processors is larger than 1024. This is because the number of columns at the leaves approaches 2k and most of the time is spent in the reduction tree. However more parallelism is possible by choosing a smaller k, updating the trailing matrix, and performing more applications of tournament pivoting until the desired low rank approximation is obtained. In this case, k becomes a tuning parameter.

6. Conclusions and future work. In this paper we have introduced LU_CRTP, a block LU factorization based on column and row permutations for computing a low rank approximation of a sparse matrix. The selection of columns and rows at each step of the block factorization uses tournament pivoting based on QR. The experimental results show that LU_CRTP provides a good trade-off between accuracy and speed. The approximated singular values are on avergage very close to the singular values computed by SVD, and usually within a factor of 10. On 1024 cores of a Cray XC30 computer, the algorithm computes a rank-256 approximation in 5 seconds for a matrix of size 525825 × 525825 (PARAB_FEM).

Acknowledgments. The authors thank the anonymous reviewers and the editor for their very useful comments that helped us improve the paper. In particular the authors would like to thank the referee who suggested the derivation which leads to (34).

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