
Randomized Two-Sided Gram-Schmidt Process With Applications

Laura Grigori¹, Lorenzo Piccinini^{*2}, and Igor Simunec¹

¹EPFL – Switzerland

²Università di Bologna – Italy

Abstract

Given two matrices X, Y of dimension n times m , with $m < n$ and full rank, the Two-Sided Gram-Schmidt process aims to find two bases Q, P of the same dimension such that $\text{range}(X) = \text{range}(Q)$, $\text{range}(Y) = \text{range}(P)$ and $Q^T P = D$, with D diagonal, i.e. Q and P are biorthogonal. It is widely known that this algorithm frequently suffers from numerical instability, and the bases Q and P are often ill-conditioned.

In this talk, we present a randomized version of the algorithm, which computes two matrices Q and P that satisfy the sketched biorthogonality condition $(SQ)^T SP = D$, where S is a sketching matrix of dimension s times n satisfying an oblivious eps-embedding property, such as a subsampled randomized Hadamard transform or a sparse sign matrix. We show how this approach can improve the stability of the algorithm and the condition number of the computed bases Q and P .

As an application, we consider the computation of approximate eigenvalues and both right and left eigenvectors, where our randomized two-sided Gram-Schmidt orthogonalization process can be implemented within the non-symmetric Lanczos algorithm.

*Speaker