

Mini-Lecture: Overview of the NIPALS-MCIA pipeline

Max Mattessich

Joaquin Reyna

Anna Konstorum

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Introduction

The Mini-Lecture, Brief outline of the math behind MCIA, covered the mathematical aspects of MCIA, this second Mini-Lecture will focus on the standard preprocessing of multi-block data and post-processing following MCIA. The target audience of this Mini-Lecture are bioinformaticians as well as other practitioners in data science.

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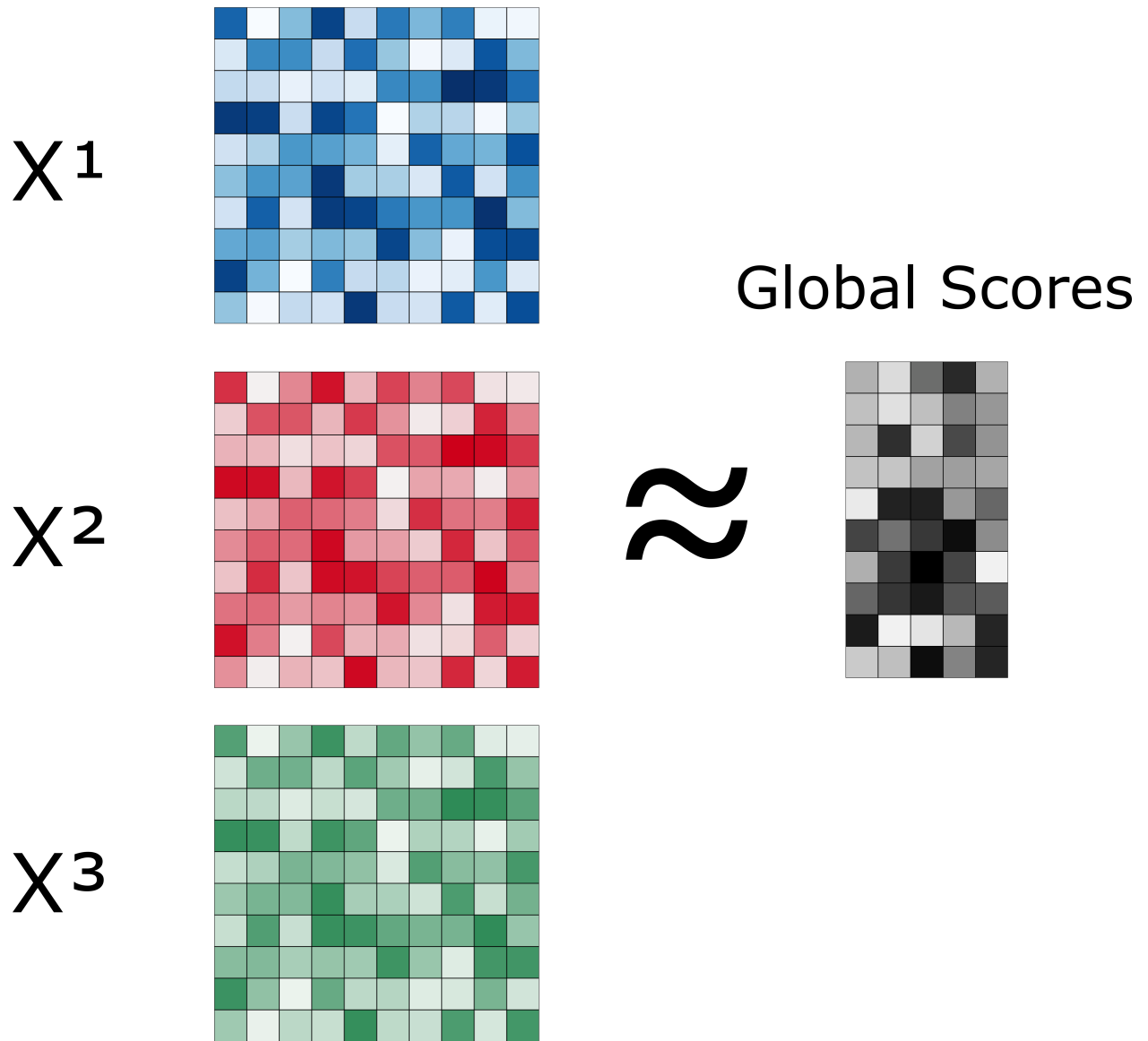


Figure 1: Generic MCIA pipeline. There are three data blocks X^1 , X^2 , and X^3 which are being decomposed via MCIA to produce a global scores matrix.