

# Other Decompositions

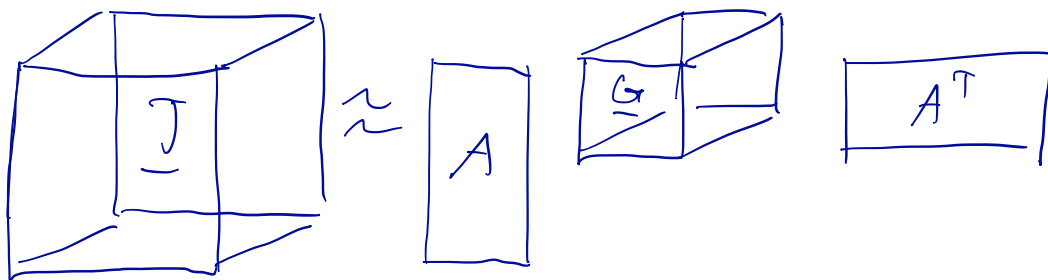
## RESICAL

RESICAL decomposition is a combination of Tucker2 and INDSCAL, that is, it is a Tucker2 decomposition with symmetric frontal slices.

Given  $\underline{I} \in \mathbb{R}^{I \times I \times K}$  and  $R \in \mathbb{N}$ , RESICAL is

$$\underline{I} \approx \llbracket \underline{G}; A, A, I \rrbracket \Leftrightarrow \underline{I}_R \approx A G_R A^T,$$

where  $A \in \mathbb{R}^{I \times R}$  and  $\underline{G} \in \mathbb{R}^{R \times R \times K}$ .



We do not necessarily require that the frontal slices of  $\underline{I}$  are symmetric.

To compute RESCAL, consider the mode-1 matricization:

$$T_{(1)} = A G_{(1)} (I \otimes A)^T.$$

Like INDSCAL, we have  $A$  twice. One solution here is to "fix" the right-hand  $A$  and only update the left-hand  $A$ .

If  $T_k$  are not symmetric, this alone can yield sub-optimal results. To make the updated  $A$  to work also with

$T_{(2)}$ , we replace  $T_{(1)}$  with  $[T_1 \ T_1^T \ T_2 \ T_2^T \ \dots \ T_k \ T_k^T]$ .

This gives

$$\|Y - AH(I_{2k} \otimes A)^T\|_F$$

with  $Y = [T_1 \ T_1^T \ T_2 \ T_2^T \ \dots \ T_k \ T_k^T]$  and

$H = [G_1 \ G_1^T \ G_2 \ G_2^T \ \dots \ G_k \ G_k^T]$ , with update

rule

$$A = Y(H(I_{2k} \otimes A)^T)^+$$

$$= \left( \sum_{k=1}^k (T_k \ G_k^T + T_k^T \ A G_k) \right) \left( \sum_{k=1}^k (B_k + C_k) \right)^+$$

with  $B_k = G_k A^T A G_k^T$  and  $C_k = G_k^T A^T A G_k$

To update the core, we update each frontal slice thereof

$$G_k = \arg \min_{G_k} \|\text{vec}(T_k) - (A \otimes A) \text{vec}(G_k)\|$$

giving  $\text{vec}(G_k) = (A \otimes A)^{\dagger} \text{vec}(T_k)$ . This still requires computing a pseudo-inverse of a big matrix  $(A \otimes A)$ . We can reduce the computation with a skinny QR decomposition of  $A$ . Let  $A = QU$ , where  $Q \in \mathbb{R}^{I \times R}$  is column-orthogonal and  $U \in \mathbb{R}^{R \times R}$  is upper-triangular. It should be noted that every matrix has a QR decomposition. We now have

$$\begin{aligned} \|T_k - A G_k A^T\| &= \|T_k - Q U G_k U^T Q^T\| \\ &= \|Q^T T_k Q - U G_k U^T\| \end{aligned}$$

which is optimized by

$$\text{vec}(G_k) = (U \otimes U)^{\dagger} \text{vec}(Q^T T_k Q).$$

RESICAL can be used when we want to model non-symmetric data with DEDICOM type decomposition. Using the  $A \underline{G} A^T$  model provides an "information flow" from mode 1 to mode 2 and vice versa. For example, if we have  $(s, p, o)$  data, we can assume that subjects and objects come from the same set of entities, and there should be just one factor matrix for them. But  $A A^T$  is symmetric, while most predicates are not (is President of, is Son of, ...). If we have resolved the predicates in a surface  $(np, up, np)$  tensor (so that we have a  $(np, p, np)$  tensor), we can decompose it with RESICAL to obtain  $(s, p, o)$  core  $\underline{G}$  and  $A: np \rightarrow s \cup o$ .



# DEDICOM

DEDICOM decomposition is a matrix decomposition that is equivalent to a frontal slice of RESCAL:

$$T = AGA^T.$$

Like RESCAL, DEDICOM can be used to model asymmetric relations between entities.

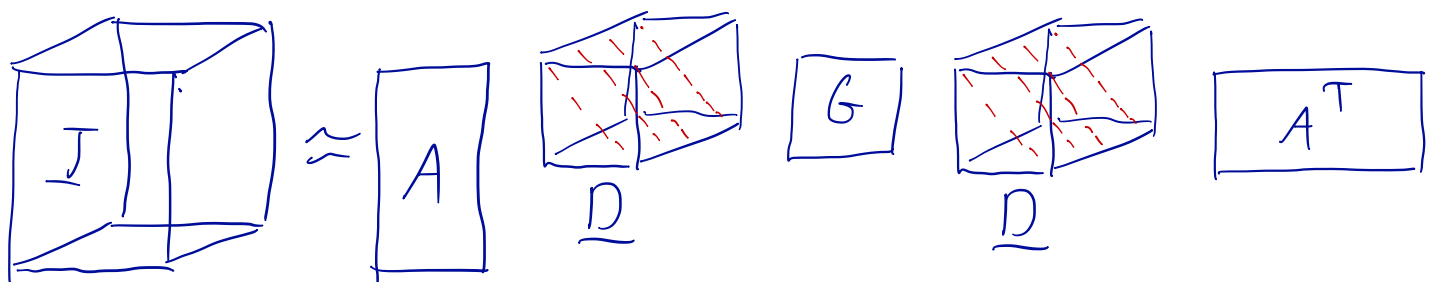
The three-way DEDICOM adds weights for each entity factors participation in each position in the third mode. For example, if we have countries-by-countries-by-time tensor  $T \in \mathbb{R}^{I \times I \times K}$ , where  $t_{ijk}$  has the value of trade from country  $i$  to country  $j$

at time point  $k$ , 3-way DEDICOM adds information on how much does a country factor  $\vec{a}_i$  act as a seller or buyer at time  $k$ .

Each frontal slice of a 3-way DEDICOM is

$$T_k \approx A D_k G D_k A^T,$$

where  $A$  and  $G$  are as in the matrix case, and  $\underline{D}$  is an  $R \times R \times K$  tensor such that each frontal slice  $D_k$  is diagonal.  $(D_k)_{rr} = d_{rrk}$  is the weight of a factor  $r$  at time  $k$ .



Where RESCAL decomposes each relation separately using  $G_k$ , DEDICOM assumes there is only one (potentially asym-

metric) relation encoded in  $G$ , but that the participation to that relation varies over time.

### Computing DEDICOM: ASALSAN

Given  $\underline{T} \in \mathbb{R}^{I \times I \times K}$  and  $R \in \mathbb{N}$ , in DEDICOM we want to find  $A \in \mathbb{R}^{I \times R}$ ,  $\underline{D} \in \mathbb{R}^{R \times R \times K}$ , and  $G \in \mathbb{R}^{R \times R}$  that minimize

$$\sum_{k=1}^K \|T_k - A D_k G D_k A^T\|.$$

ASALSAN (alternating simultaneous approximation, least squares, and Newton) uses the methods in the name for optimizing different factors.

As with RESCAL, DEDICOM has matrix  $A$  on both sides, and ASALSAN uses

similar approach handle it: it stacks pairs  $T_k T_k^T$  to get

$$Y = [T_1 T_1^T \dots T_k T_k^T]$$

The error function becomes

$$\|Y - AH (I_{2k} \otimes A^T)\|$$

with

$$H = [D_1 G D_1 \quad D_1 G^T D_1 \quad \dots \quad D_k G D_k \quad D_k G^T D_k]$$

Similarly to RESCAL, we fix the right  $A$  and update

$$A \leftarrow \left( \sum_{k=1}^K (T_k A D_k G^T D_k + T_k^T A D_k G D_k) \right) \left( \sum_{k=1}^K (B_k + C_k) \right)^{-1}$$

where

$$B_k = D_k G D_k (A^T A) D_k G^T D_k$$

$$C_k = D_k G^T D_k (A^T A) D_k G D_k$$

Matrix  $G$  we can update using the vectorized representation

$$G \leftarrow \arg \min_G \left\| \begin{pmatrix} \text{vec}(T_1) \\ \text{vec}(T_2) \\ \vdots \\ \text{vec}(T_R) \end{pmatrix} - \begin{pmatrix} AD_1 \otimes AD_1 \\ \vdots \\ AD_R \otimes AD_R \end{pmatrix} \text{vec}(G) \right\|$$

Finally, to update  $\underline{D}$ , we can update each frontal slice separately, having only  $R$  unknowns. As there doesn't seem to be any easy closed-form solution, ALSAN uses the Newton's method.

## PARATUCK2

PARATUCK2 (portmanteau of PARAFAC and Tucker2) generalizes DEDCOM to allow different factors on the right-hand side:

$$T_k = AD_k G D_k B^T,$$

where  $A$ ,  $\underline{D}$ , and  $G$  are as in DEDCOM.

# Tensor train (TT) decomposition

Given a tensor  $\underline{T} \in \mathbb{R}^{I_1 \times I_2 \times \dots \times I_N}$ , the Tensor train (or TT) decomposition has  $N$  3-way factor tensors  $\underline{G}^{(n)} \in \mathbb{R}^{R_{n-1} \times R_n \times I_n}$ , where  $R_0 = R_N = 1$ , and the rest are model parameters ("TT rank")  $R_1, R_2, \dots, R_{N-1}$ . TT expresses an element of  $\underline{T}$  as a product of the frontal slices of  $\underline{G}^{(n)}$ 's:

$$t_{i_1 i_2 \dots i_N} = G_{i_1}^{(1)} G_{i_2}^{(2)} \dots G_{i_N}^{(N)}$$

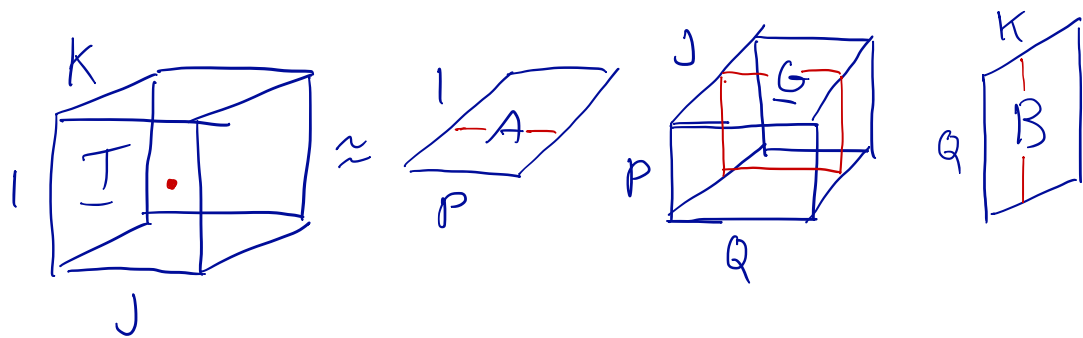
Notice that this is a scalar as  $G_{i_1}^{(1)} \in \mathbb{R}^{1 \times R_1}$  and  $G_{i_N}^{(N)} \in \mathbb{R}^{R_{N-1} \times 1}$ . Writing the products open, we have

$$t_{i_1 i_2 \dots i_N} = \sum_{r_1=1}^{R_1} \sum_{r_2=1}^{R_2} \dots \sum_{r_{N-1}=1}^{R_{N-1}} \left( G^{(1)}(1, r_1, i_1) G^{(2)}(r_1, r_2, i_2) \dots \right. \\ \left. \times G^{(N-1)}(r_{N-2}, r_{N-1}, i_{N-1}) G^{(N)}(r_{N-1}, 1, i_N) \right)$$

For a 3-way tensor, this simplifies to

$$t_{ijk} = \sum_{p=1}^P \sum_{q=1}^Q A(p, i) G(p, q, j) B(q, k)$$

where  $A \in \mathbb{R}^{P \times I}$ ,  $G \in \mathbb{R}^{P \times Q \times J}$ , and  $B \in \mathbb{R}^{Q \times K}$



Let's consider an example: Let  $\underline{T} \in \mathbb{R}^{3 \times 4 \times 5}$  be such that  $t_{ijk} = i + j + k$  for all  $i, j$ , and  $k$ . Define  $A$ ,  $G$ , and  $B$  so that

$$A(i, i) = (i \ 1) \quad G_j = \begin{pmatrix} 1 & 0 \\ j & 1 \end{pmatrix} \quad B(i, k) = \begin{pmatrix} 1 \\ k \end{pmatrix}.$$

Now

$$t_{ijk} = (i \ 1) \begin{pmatrix} 1 & 0 \\ j & 1 \end{pmatrix} \begin{pmatrix} 1 \\ k \end{pmatrix} = (i + j \ 1) \begin{pmatrix} 1 \\ k \end{pmatrix} = i + j + k.$$

Tensor  $\underline{T}$  has 60 elements, while the tensor train has only  $3 \cdot 2 + 4 \cdot 2 \cdot 2 + 5 \cdot 2 = 32$ .

In general, TT can express an  $\underbrace{1 \times 1 \times \dots \times 1}_N$  tensor with  $1^N$  elements with

$$1 \cdot N \cdot R^2$$

elements, where  $R = \max_{n=1}^N R_n$ .

This number  $R = \max_{n=1}^N R_n$  is called the (maximal) TT-rank.

TT allows certain operations to be effective for tensors stored in the TT-format. If  $\underline{T}$  and  $\underline{S}$  are such that

$$T_{i_1 \dots i_N} = G_{i_1}^{(1)} G_{i_2}^{(2)} \dots G_{i_N}^{(N)} \quad \text{and} \quad S_{i_1 \dots i_N} = H_{i_1}^{(1)} \dots H_{i_N}^{(N)},$$

then  $\underline{U} = \underline{T} + \underline{S}$  has TT decomposition to

$$F_{i_n}^{(n)} = \begin{pmatrix} G_{i_n}^{(n)} & H_{i_n}^{(n)} \end{pmatrix} \quad (n=2, \dots, N-1)$$

$$F_{i_1}^{(1)} = \begin{pmatrix} G_{i_1}^{(1)} & H_{i_1}^{(1)} \end{pmatrix}$$

$$F_{i_N}^{(N)} = \begin{pmatrix} G_{i_N}^{(N)} \\ H_{i_N}^{(N)} \end{pmatrix}$$

If the TT-ranks of  $\underline{T}$  are  $R_1^T, R_2^T, \dots$ , and of  $\underline{S}$  are  $R_1^S, R_2^S, \dots$ , then for  $\underline{U}$  they are  $R_1^T + R_1^S, R_2^T + R_2^S, \dots$ .

Similarly for the Hadamard (element-wise) product we have that



$$\underline{U} = \underline{I} * \underline{S} \iff F_{i_n}^{(n)} = G_{i_n}^{(n)} \otimes H_{i_n}^{(n)}, n=1, \dots, N$$

To see this, notice that

$$\begin{aligned} u_{i_1 i_2 \dots i_N} &= (G_{i_1}^{(1)} \dots G_{i_N}^{(N)}) (H_{i_1}^{(1)} \dots H_{i_N}^{(N)}) \\ &= (G_{i_1}^{(1)} \dots G_{i_N}^{(N)}) \otimes (H_{i_1}^{(1)} \dots H_{i_N}^{(N)}) \\ &= (G_{i_1}^{(1)} \otimes H_{i_1}^{(1)}) (G_{i_2}^{(2)} \otimes H_{i_2}^{(2)}) \dots (G_{i_N}^{(N)} \otimes H_{i_N}^{(N)}). \end{aligned}$$

Hence,  $R_n^U = R_n^I R_n^S$ .

## Computing the TT

The basic algorithm for computing the TT decomposition within any accuracy  $\epsilon$  is the TT-SVD. Given  $\underline{I}$ , it finds  $\underline{S}$  that has the smallest possible ranks  $R_n^S$  such that

$$\|\underline{I} - \underline{S}\| \leq \epsilon \|\underline{I}\|.$$

The algorithm is based on computing SVD to obtain unfolded  $\underline{G}$ 's.

TT-SVD (T,  $\epsilon$ )

$$\delta \leftarrow \epsilon(N-1)^{-1/2} \| \underline{T} \|$$

$$\underline{S} \leftarrow \underline{T}; \quad J \leftarrow 1:2:\dots:l_n; \quad R_0 = 1$$

for  $n=1, \dots, N$  do

$$S \leftarrow \text{reshape}(\underline{S}, R_{n-1} \times l_n, J / (R_{n-1} \times l_n))$$

$$(U, \Sigma, V) \leftarrow \text{SVD}(S)$$

$$R_n \leftarrow \text{arg min}_r \left\{ \sum_{i=r+1}^{J/(R_{n-1} \times l_n)} \sigma_i^2 \leq \delta \right\}$$

$$\underline{G}^{(n)} \in \mathbb{R}^{R_{n-1} \times l_n \times R_n}$$

↓

$$\underline{G}^{(n)} \leftarrow \text{reshape}(U(:, 1:R_n), R_{n-1}, l_n, R_n)$$

$$\underline{S} \leftarrow \Sigma V^T$$

$$J \leftarrow J R_n / (l_n R_{n-1})$$

end

$$\underline{G}^{(N)} \leftarrow \underline{S}$$

$$\underline{\text{return}} (\underline{G}^{(1)}, \underline{G}^{(2)}, \dots, \underline{G}^{(N)})$$

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N.B. Here  $\underline{G}^{(n)} \in \mathbb{R}^{R_{n-1} \times l_n \times R_n}$ , not  $\mathbb{R}^{R_{n-1} \times R_n \times l_n}$ .

Applications of TT

TT is not (often?) used for data analysis as such, as it lacks easy

interpretation of the "cars"  $\underline{G}^{(1)}, \underline{G}^{(2)}, \dots$   
It is, however, commonly used to  
reduce the number of free parameters  
in different machine learning applications.  
The parameters can also be stored in a  
matrix: we can "fold" it into a tensor  
and then apply TT.

TT can also be applied to the core tensor  
of the Tucker decomposition, if we want  
to compress it more.

# CORCONDIA

CORCONDIA (Core consistency diagnostic) is not a tensor factorization per se. Rather, it's a method for selecting the rank in a CP decomposition, or, alternatively, for deciding between CP and Tucker.

The idea of CORCONDIA is to use the fact that CP is a special case of Tucker. Given  $\underline{T}$ , we can first compute CP  $[[A, B, C]]$  and then use these as the factor matrices for Tucker, and find the core tensor  $\underline{G}$  as

$$\underline{G} \leftarrow (A \otimes B \otimes C)^+ \text{vec}(\underline{T})$$

Now CORCONDIA statistic measures how diagonal  $\underline{G}$  is

$$\text{CORCONDIA} = 100 \left( 1 - \frac{\sum_{i,j,k} (g_{ijk} - \mathbb{1}(i=j=k))^2}{R} \right)$$

Here

$$\mathbb{1}(i=j=k) = \begin{cases} 1 & \text{if } i=j=k \\ 0 & \text{otherwise} \end{cases}$$

CORCONDIA statistic gives us an indication how good a model rank-R CP is for the data. The statistic assumes values from  $(-\infty, 100]$ , and small values indicate a bad match. Usually, values less than 50 indicate some problems.

That's all  
folks!