

Matrix and Tensor Factorization with Scientific Constraints

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Acknowledgements

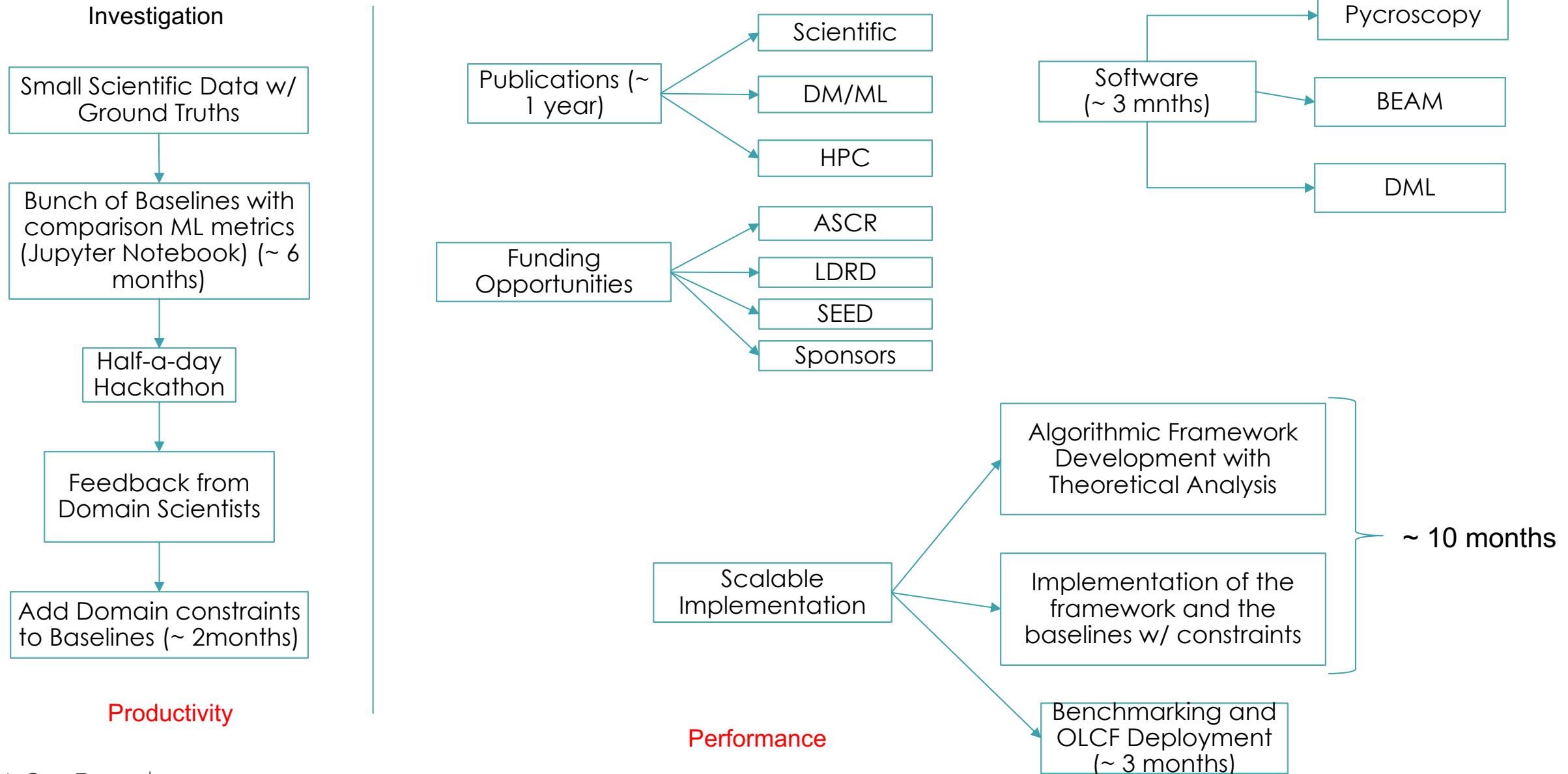
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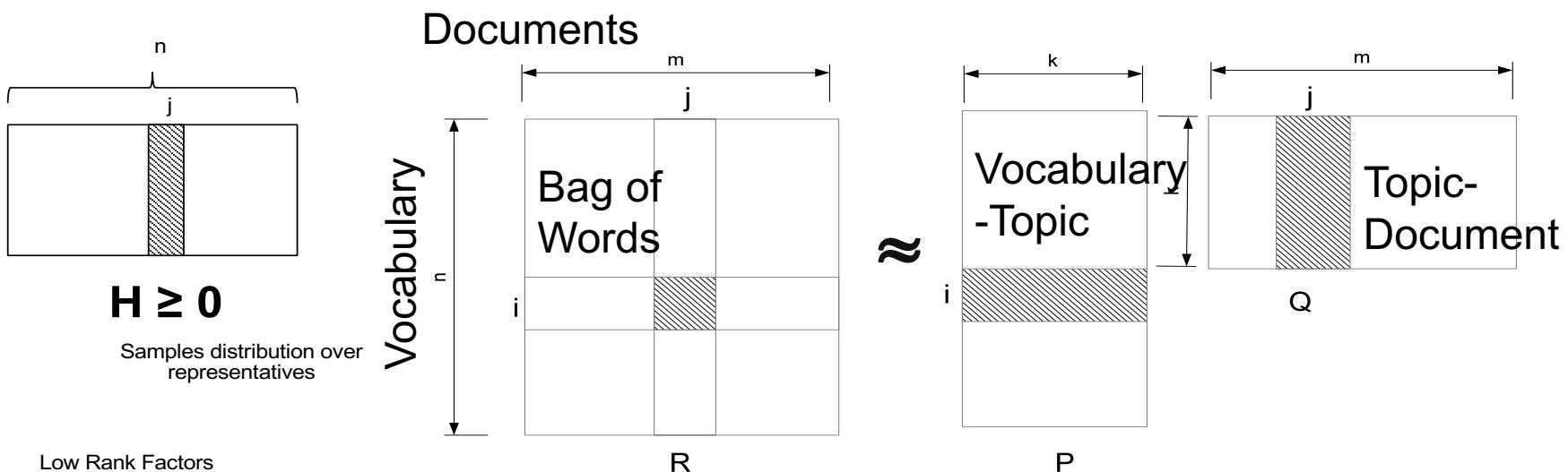
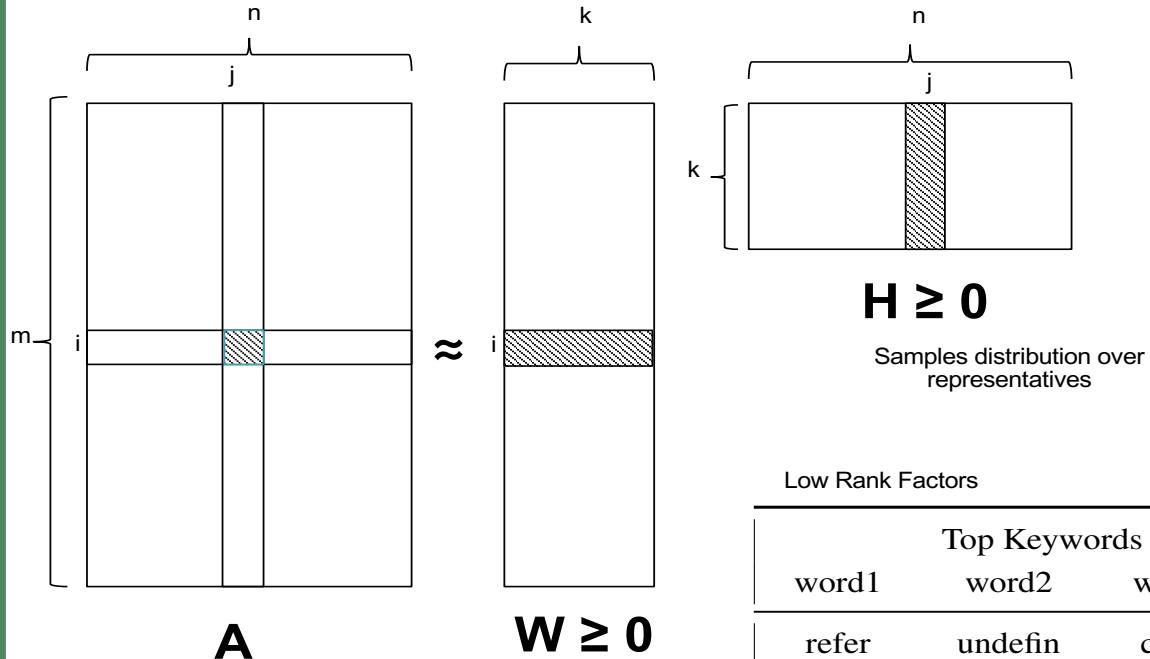
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Scientific Engagement Model



NMF and Applications

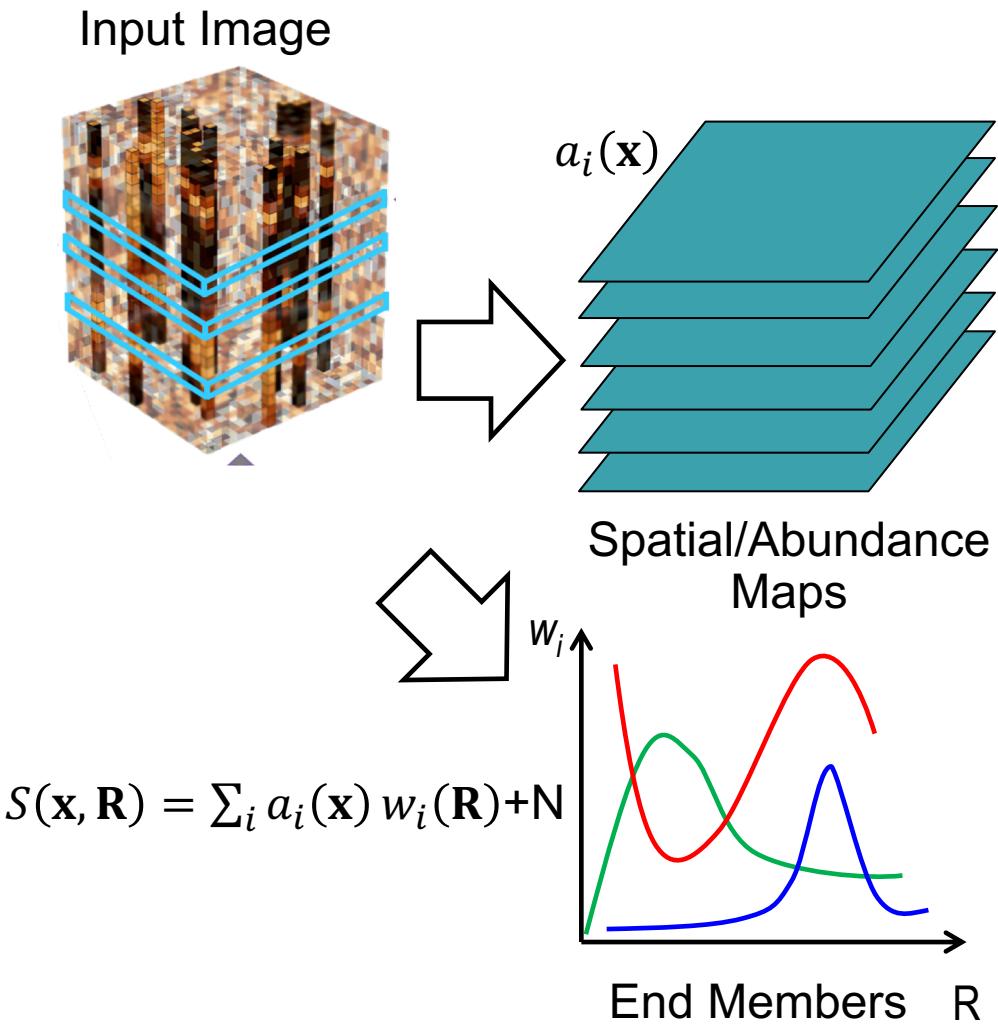


| Top Keywords from Topics 1-25 | | | | | Top Keywords from Topics 26-50 | | | | |
|-------------------------------|-----------|----------|---------|---------|--------------------------------|-----------|---------|---------|---------|
| word1 | word2 | word3 | word4 | word5 | word1 | word2 | word3 | word4 | word5 |
| refer | undefin | const | key | compil | echo | type=text | php | form | result |
| text | field | box | word | static | test | perform | fail | unit | result |
| imag | src | descript | alt=ent | size | tabl | key | queri | databas | insert |
| button | click | event | form | add | user | email | usernam | login | log |
| creat | bean | add | databas | except | data | json | store | read | databas |
| string | static | final | catch | url | page | load | content | url | link |
| width | height | color | left | display | privat | static | final | import | float |
| app | applic | servic | thread | work | row | column | date | cell | valu |
| ipsum | lorem | dolor | sit | amet | line | import | command | print | recent |
| node | list | root | err | element | var | map | marker | match | url |
| 0x00 | 0xff | byte | 0x01 | 0xc0 | server | connect | client | messag | request |
| file | directori | read | open | upload | number | byte | size | print | input |

Ramakrishnan Kannan, Grey Ballard, Haesun Park: MPI-FAUN: An MPI-Based Framework for Alternating-Updating Nonnegative Matrix Factorization. *IEEE Trans. Knowl. Data Enq.* 30(3): 544-558 (2018)

Motivation

- Understanding terrestrial information in an unknown place from satellite images
- Identifying presence of hidden unknown/foreign bodies in a scanned image - Eg., contamination in food articles, camouflaged explosives etc.
- Biological application - spectral karyotyping, immunofluorescence, live-cell imaging, drug discovery, and tissue pathology – Eg., Unmixing on Spectral imaging of the stained tissues using multiple dyes.
- Physics and Material Sciences – Mapping properties to end-members. Comparing different materials



- Distributed Communication avoiding NMF Algorithms
- <https://github.com/ramkikannan/nmflibrary>
- <http://dx.doi.org/10.1109/TKDE.2017.2767592>

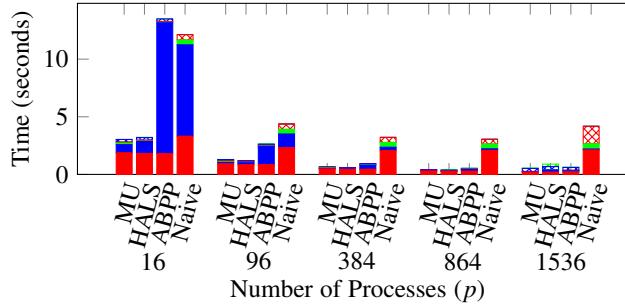
Titan – Dense Matrix, Low Rank
 50, 100 Iterations, 12650 Nodes,
 202500 Cores,

| Matrix Size | Algos | NMF Time (in Secs) |
|-----------------------------|----------|--------------------|
| 3.03 million x 3.03 million | MU | 554 |
| | HALS | 197.75 |
| | ANLS/BPP | 219.8 |

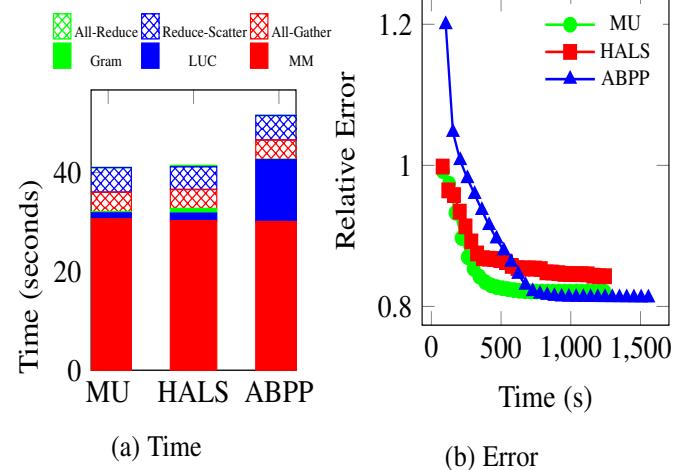
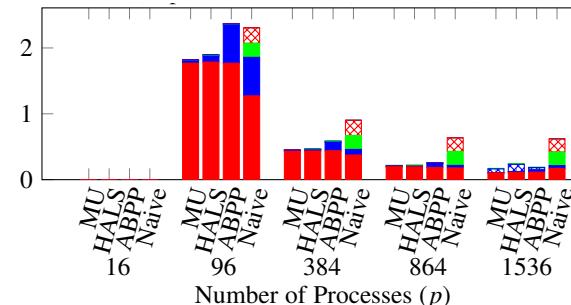
Rhea, 100 nodes, 1600 cores, Low Rank 50,

| Dataset | Type | Matrix size | NMF Time |
|----------------|--------|---------------------------|--------------|
| Video | Dense | 1 Million x 13,824 | 5.73 seconds |
| Stack Exchange | Sparse | 627,047 x 12 Million | 67 seconds |
| Webbase-2001 | Sparse | 118 Million x 118 Million | 25 minutes |

Sparse Webbase –
 1 Million Vertex Graph



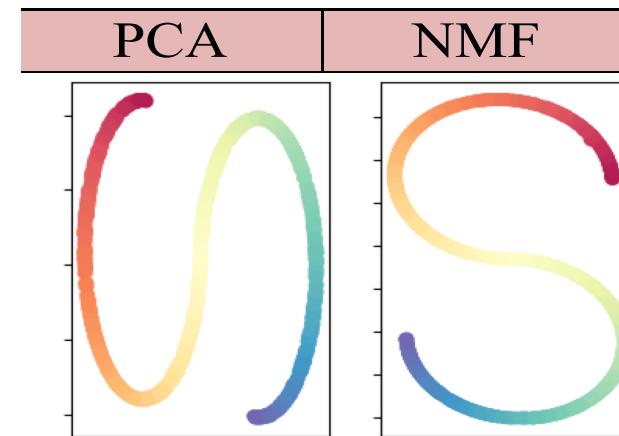
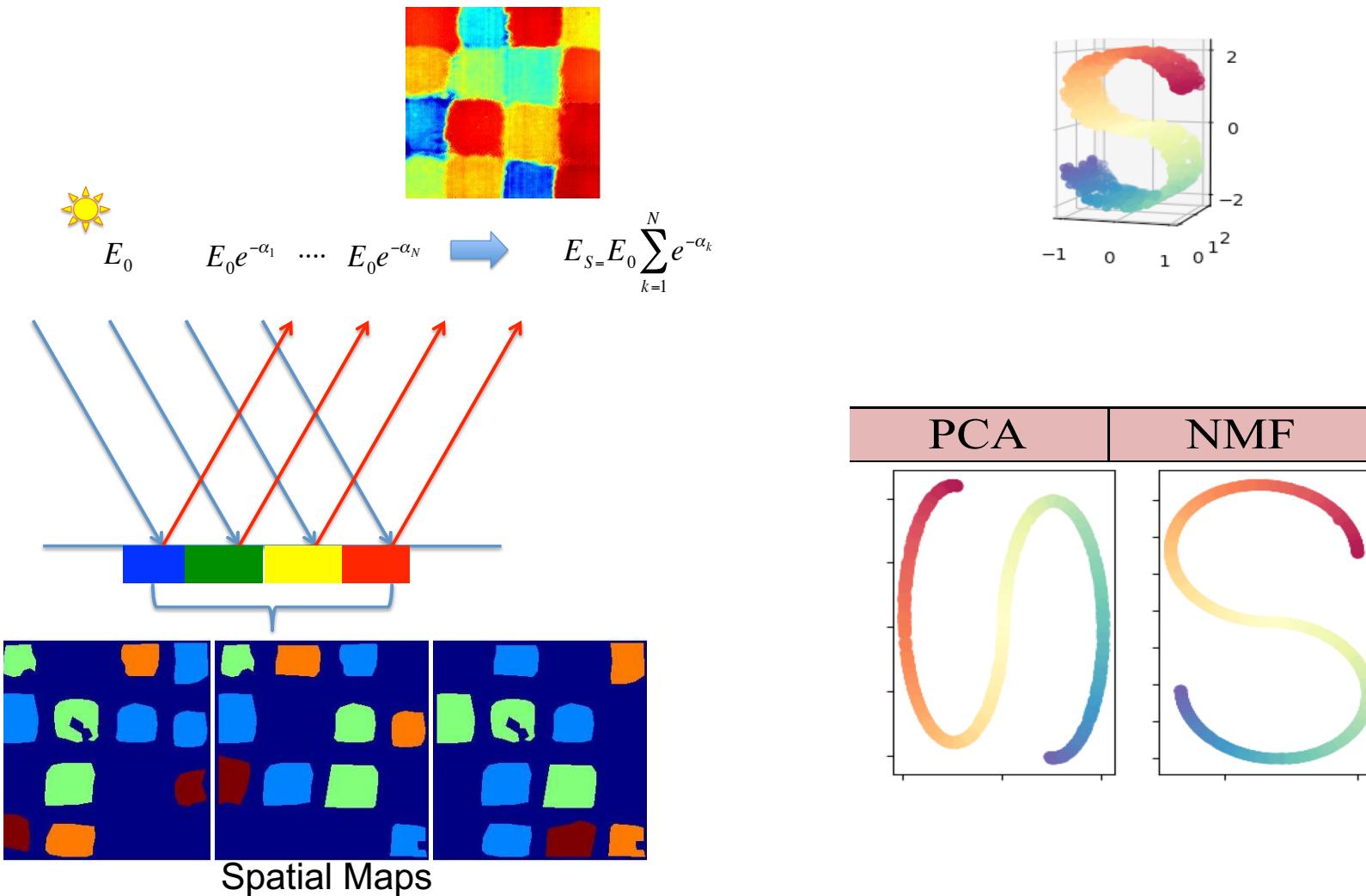
Dense Real world – Video



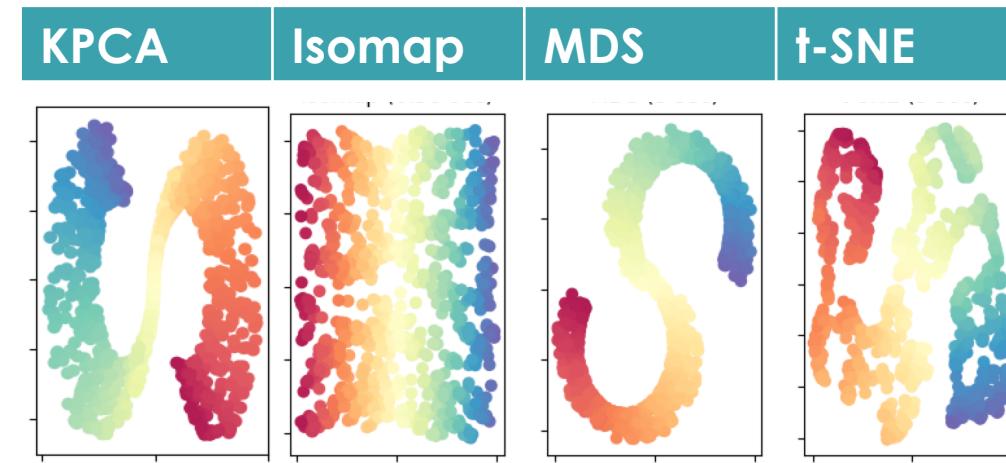
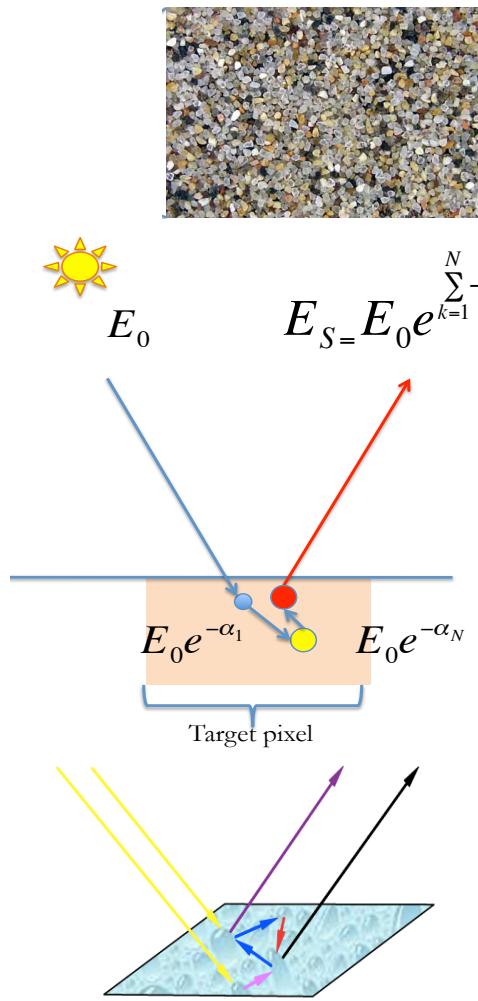
NMF on 118 million Web-graph

Existing Approach : Linear Unmixing

1. Good at Capturing Macroscopic Information
2. Spatially segregated patterns



Existing Non-linear Unmixing (NLUM)



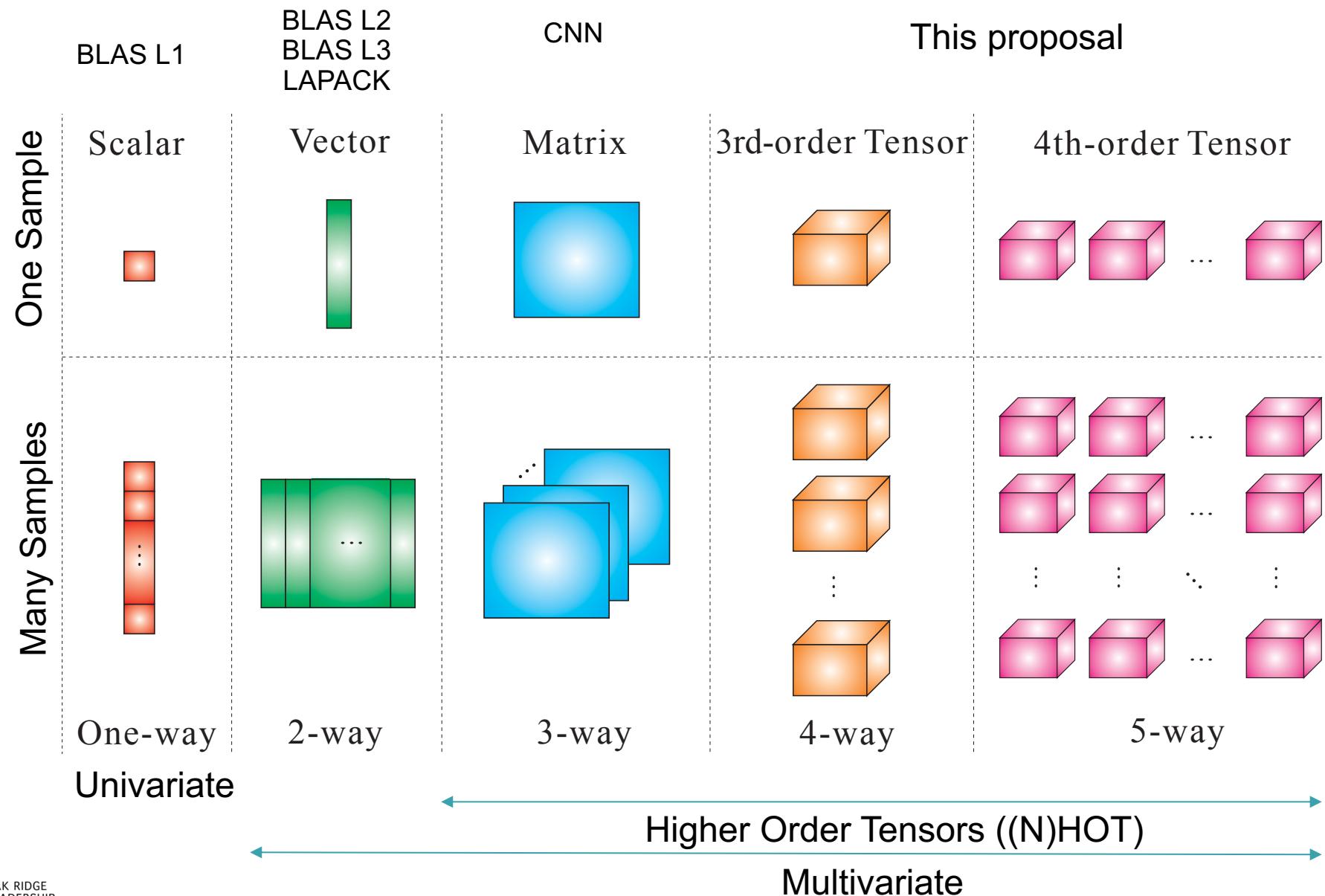
Non-Meaningful results for following reasons:

1. End-members and abundance maps are negative
2. Too many end-members participate in a particular location
3. Similar end-members and not distinctive enough
4. Ratio of end-members are not correct
5. Rotated end-members

Solution:

NLUM w/ Physical Constraints such as non-negativity, sparsity, spatial smoothness, sum to 1, orthogonal etc.

Higher Order Tensors



Dimensionality Reduction in Scientific Data

- Multimodal characterization of materials –
comprehensive characterization from chemical composition to functional properties on the nanoscale

