Supplementary material to GmGM: a fast Gaussian graphical model for multi-modal data

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¹ Contents

18 1 Notation

 In addition to the notation used in the main paper, we also introduce further notation to aid in the proofs. For working with tensors, Kolda and Bader [\[8\]](#page-15-0) proved to be an invaluable resource; we have borrowed their notation in most cases. The only exception is that we have chosen to denote the ℓ -mode matricization of a tensor $\mathcal T$ as $\text{mat}_{\ell}[\mathcal T]$ rather than $\mathcal T_{(\ell)}$, to highlight its similarity to vec $[\mathcal T]$ and free up the subscript for other purposes.

- ²⁴ To keep track of lengths of axes, we define the following notation:
- 25 d_{ℓ}^{γ} is the length of axis ℓ
- \bullet $d_{\gt\ell}^{\gamma}$ is the product of lengths of all axes after ℓ
- ²⁷ $d_{\langle\ell\,\rangle}^{\gamma}$ is the product of lengths of all axes before ℓ
- d^{γ}_{λ} 28 • $d_{\sqrt{\ell}}^{\gamma}$ is the product of lengths of all axes except for ℓ
- \bullet d_{\forall}^{γ} is the product of lengths of all axes (i.e. the number of elements in \mathcal{D}^{γ})
- ³⁰ $d_{\forall} = \sum_{\gamma} d_{\forall}^{\gamma}$ is the total number of elements across all datasets

In prior work, d_ℓ has been used to represent the lengths of axes but m_ℓ was used where we write d_ℓ 31 32 (such as in [\[5\]](#page-15-1)). As prior work also used ℓ to represent leaving out the ℓ th axis in other contexts 33 (such as in [\[6\]](#page-15-2)), and the analogous definitions of $d_{\geq \ell}$ and $d_{\leq \ell}$ were convenient for use in proofs, we 34 chose to introduce $d_{\setminus \ell}$ as the variable to represent leave-one-out length products. By representing all ³⁵ of these related concepts with similar symbols, we hope the maths will be easier to parse.

36 We will let I_a be the $a \times a$ identity matrix, which allows a concise definition of the Kronecker sum: з7 $\quad \bigoplus_\ell \Psi_\ell = \sum_\ell \mathbf{I}_{d_{<\ell}} \otimes \Psi_\ell \otimes \mathbf{I}_{d_{>\ell}}.$

38 We make frequent use of the vectorization vec $[M]$ of a matrix M, and more generally of a tensor α vec $[T]$. We adopt the rows-first convention of vectorization, such that

$$
29 \text{ VEC} \left[T \right].
$$
 We adopt the rows-first contribution of vectorization, such that

$$
\text{vec}\begin{bmatrix} 1 & 2 \\ 3 & 4 \end{bmatrix} = \begin{bmatrix} 1 & 2 & 3 & 4 \end{bmatrix} \tag{1}
$$

⁴⁰ While columns-first is more common, rows-first is more natural when we adopt the convention that ⁴¹ rows are the first axis of tensor; this is the convention that matricization uses, and matricization ⁴² is much more important for our work due to its role in defining the Gram matrices. Note that, for 43 matrices, a rows-first vectorization of M is equivalent to a columns-first vectorization of \mathbf{M}^T , so there ⁴⁴ is no fundamental difference between the two. For vectorizing a tensor, we proceed by stacking the the axes in order, such that an element $(i_1, ..., i_K)$ in $\mathcal T$ gets mapped to the element $\sum_{\ell} i_{\ell} d_{\leq \ell}$ 46 in vec $[\mathcal{T}]$.

- 47 We define the Gram matrices as $S_\ell^\gamma = \text{mat}_\ell \left[\mathcal{D}^\gamma \right] \text{mat}_\ell \left[\mathcal{D}^\gamma \right]^T$. Typically we consider only the one-
- 48 sample case but if you have multiple samples, indexed by a subscript i , then the Gram matrix becomes
- 49 an average: $\mathbf{S}_{\ell}^{\gamma} = \frac{1}{n} \sum_{i}^{n} \text{mat}_{\ell} \left[\mathcal{D}_{i}^{\gamma} \right] \text{mat}_{\ell} \left[\mathcal{D}_{i}^{\gamma} \right]^{T}$.
- ⁵⁰ An essential concept is that of the "stridewise-blockwise trace", defined as:

$$
\operatorname{tr}_{b}^{a}[\mathbf{M}] = \left[\operatorname{tr}\left[\mathbf{M}\left(\mathbf{I}_{a} \otimes \mathbf{J}^{ij} \otimes \mathbf{I}_{b}\right)\right]\right]_{ij} \tag{2}
$$

51 Where J^{ij} is the matrix of zeros except at (i, j) where it has a 1. It is a generalization of the 52 blockwise trace considered by Kalaitzis et al. [\[6\]](#page-15-2), and is related to the $proj_{\mathcal{K}}$ operation defined by 53 Greenewald, Zhou, and Hero III [\[5\]](#page-15-1). Specifically, $proj_{\mathcal{K}}[\mathbf{M}]$ is equivalent to $\bigoplus_{\ell} \text{tr}_{d_{>\ell}}^{d_{<\ell}}[\mathbf{M}]$ up to 54 an additive diagonal factor (Lemma 33 from Greenewald, Zhou, and Hero III [\[5\]](#page-15-1)). $proj_K$ [M] was 55 defined to be the matrix that best approximates M (in terms of the Frobenius norm) while being ⁵⁶ Kronecker-sum-decomposable. This matrix is not unique; the choice by Greenewald, Zhou, and 57 Hero III [\[5\]](#page-15-1) to include an additive factor was to enforce $tr[proj_K[\mathbf{M}]]=0$. We do not wish to ⁵⁸ enforce this constraint as it would be impossible to preserve in the multi-tensor case. 59 The parameter b of the stridewise-blockwise trace partitions the $m \times m$ matrix M into a block matrix

- 60 with $b \times b$ blocks of size $(\frac{m}{b} \times \frac{m}{b})$. The parameter a then partitions these blocks into a "strided" 61 matrix with $a \times a$ strides containing $\frac{m}{ab} \times \frac{m}{ab}$ blocks. We take the trace of each stride, and the final
- ⁶² matrix is the matrix of these traces. As this is conceptually complicated, we provide an example.

$$
tr_{2}^{2} = \begin{bmatrix} 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \\ 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \\ 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \\ 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \\ 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \\ 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \\ 1 & 2 & 3 & 4 & 5 & 6 & 7 & 8 \end{bmatrix}
$$
\n
$$
= tr_{2}^{2} \begin{bmatrix} tr \begin{bmatrix} 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1 & 2 \\ 1
$$

⁶³ Notice the construction of the "strides" in Line [6](#page-2-2) - the parameter of 2 told us to grab every second ⁶⁴ element from each row and each column.

65 2 Proofs

 We will assume that no dataset contains repeated axes (i.e. no single tensor has two axes represented by the same graph), as this greatly affects the derived gradients. Shared axes - two tensors having one or more axes in common - are allowed. The case of shared axes is, after all, the whole point of developing this extension to prior work.

⁷⁰ 2.1 Permutations

- 71 Note that both vec $[\text{mat}_1 [\mathcal{D}^\gamma]]$ and vec $[\text{mat}_\ell [\mathcal{D}^\gamma]]$ are row vectors containing the same ele-72 ments, just in a different order. This means that there is a permutation matrix $P_{\ell \to 1}$ such that 73 $\text{vec}\left[\text{mat}_1\left[\mathcal{D}^{\gamma}\right]\right]^T \mathbf{P}_{\ell \rightarrow 1} = \text{vec}\left[\text{mat}_{\ell}\left[\mathcal{D}^{\gamma}\right]\right]^T.$
- **Lemma 1** (Rearrangement lemma). $P_{\ell \to 1} \left(I_{d_{<\ell}} \otimes \Psi_\ell \otimes I_{d_{> \ell}} \right) P_{\ell \to 1}^T = \Psi_\ell \otimes I_{d_{\setminus \ell}}$ 74

Proof. While vec, mat_ℓ and \otimes are defined as operations on matrices, for the purposes of permuta-

⁷⁶ tions we can consider them as operations on indices. We can express them as follows:

vec:
$$
(i_1, ..., i_K) \rightarrow \left(\sum_{\ell} i_{\ell} d_{\leq \ell}\right)
$$
 (8)

$$
\text{mat}_{\ell}: (i_1, ..., i_K) \to \left(i_{\ell}, \sum_{\ell' < \ell} i_{\ell'} d_{\ell'} + \sum_{\ell' > \ell} i_{\ell'} \frac{d_{\ell'}}{d_{\ell}} \right) \tag{9}
$$

$$
\bigotimes : ((i_1^1, i_1^2), ..., (i_K^1, i_K^2)) \to \left(\sum_{\ell} i_{\ell}^1 d_{<\ell}, \sum_{\ell} i_{\ell}^2 d_{<\ell}\right)
$$
(10)

77 We'll consider just the rows of \otimes , \otimes _{rows} - although the same argument applies with columns:

$$
\bigotimes_{rows} : (i_1^1, ..., i_K^1) \to \left(\sum_{\ell} i_{\ell}^1 d_{<\ell}\right)
$$
 (11)

78 Finally, we'll introduce the permutation operation $\sigma_{\ell \to 1}$ that will change the order of our Kronecker ⁷⁹ product:

$$
\sigma_{\ell \to 1} : \left((i_1^1, i_1^2), \dots, (i_K^1, i_K^2) \right) \to \left(((i_\ell^1, i_\ell^2), (i_1^1, i_1^2), \dots, (i_{\ell-1}^1, i_{\ell-1}^2), (i_{\ell+1}^1, i_{\ell+1}^2), \dots, (i_K^1, i_K^2) \right)
$$
\n(12)

so And again without loss of generality we restrict ourself to $\sigma_{\ell \to 1}^{rows}$:

$$
\sigma_{\ell \to 1}^{rows} : (i_1^1, ..., i_K^1) \to (i_\ell^1, i_1^1, ..., i_{\ell-1}^1, i_{\ell+1}^1, ..., i_K^1)
$$
\n(13)

81 After a Kronecker product our indices are in the form $\sum_{\ell} i_{\ell} d_{\leq \ell}$, and if we were to reorder it with 82 $\sigma_{\ell \to 1}$ they would be in the form $i_{\ell} + \sum_{\ell' < \ell} i_{\ell'} d_{\ell'} d_{\ell'} d_{\ell} + \sum_{\ell' > \ell} i_{\ell'} d_{\ell'} d_{\ell'}$. Likewise, if we had matricized as it we would have $(i_\ell, \sum_{\ell' < \ell} i_{\ell'} d_{\ell'} + \sum_{\ell' > \ell} i_{\ell'} \frac{d_{\ell'} \ell'}{d_\ell}$, which is vectorized to $i_\ell + \sum_{\ell' < \ell} i_{\ell'} d_{\ell'} d_\ell +$ it we would have $\left(i_\ell, \sum_{\ell' < \ell} i_{\ell'} d_{<\ell'} + \sum_{\ell' > \ell} i_{\ell'} \frac{d_{<\ell'}}{d_\ell}\right)$ $e^{i\psi} \sum_{\ell' > \ell} i_{\ell'} d_{\leq \ell'}$. These reorderings are the same, and hence the matrix that represents it is $P_{\ell \to 1}$. \Box 85

86 2.2 Derivation of the probability density function

⁸⁷ Recall that the Kronecker-sum-structured normal distribution for a single tensor is defined as follows:

vec
$$
[\mathcal{D}^{\gamma}] \sim \mathcal{N}\left(\mathbf{0}, \left(\bigoplus_{\ell \in \gamma} \Psi_{\ell}\right)^{-1}\right) \iff \mathcal{D}^{\gamma} \sim \mathcal{N}_{KS}\left(\{\Psi_{\ell}\}_{\ell \in \gamma}\right)
$$
 (14)

⁸⁸ The log-likelihood for this distribution is given in [\[6\]](#page-15-2) for the matrix case and [\[5\]](#page-15-1) for the general tensor

⁸⁹ case. However, neither of these papers provide a derivation. As the full derivation will motivated the

⁹⁰ construction of lemmas useful for the proofs of Theorems [1](#page-7-1) and [2,](#page-9-4) we will give it here. First, observe ⁹¹ that the density function is that of a normal distribution.

$$
p\left(\mathcal{D}^{\gamma}\right) = \frac{\sqrt{\left|\bigoplus_{\ell \in \gamma} \Psi_{\ell}\right|}}{\left(2\pi\right)^{\frac{d_{\gamma}^{\gamma}}{2}}} e^{\frac{-1}{2} \text{vec}[\mathcal{D}^{\gamma}]^{T} \left(\bigoplus_{\ell} \Psi_{\ell}\right) \text{vec}[\mathcal{D}^{\gamma}]}\tag{15}
$$

92 Lemma 2 (⊕-vec lemma). $\text{vec} \left[\mathcal{D}^{\gamma}\right]^T (\bigoplus_{\ell} \Psi_{\ell}) \text{vec} \left[\mathcal{D}^{\gamma}\right] = \sum_{\ell} \text{tr} \left[\mathbf{S}_{\ell}^{\gamma} \Psi_{\ell}\right]$

Proof. This proof relies on the following two properties of vec: $(A \otimes B)$ vec $[C] = \text{vec} [\mathbf{B} \mathbf{C}^T \mathbf{A}^T]$ 93

94 and $\text{tr}[\mathbf{A}^T \mathbf{B}] = \text{vec}[\mathbf{A}]^T \text{vec}[\mathbf{B}]$. The C term picks up a transpose due to our use of the rows-first 95 vectorization; when using columns-first notation the right hand side becomes vec $[\text{BCA}^T]$.

vec
$$
[\mathcal{D}^{\gamma}]
$$
 $\left(\bigoplus_{\ell} \Psi_{\ell}\right)$ vec $[\mathcal{D}^{\gamma}] = \sum_{\ell} \text{vec} [\mathcal{D}^{\gamma}]^{T} \left(\mathbf{I}_{d_{\leq \ell}} \otimes \Psi_{\ell} \otimes \mathbf{I}_{d_{>\ell}}\right) \text{vec } [\mathcal{D}^{\gamma}]$ (Definition of \bigoplus)
\n
$$
= \sum_{\ell} \text{vec} [\text{mat}_{1} [\mathcal{D}^{\gamma}]]^{T} \left(\mathbf{I}_{d_{\leq \ell}} \otimes \Psi_{\ell} \otimes \mathbf{I}_{d_{>\ell}}\right) \text{vec} [\text{mat}_{1} [\mathcal{D}^{\gamma}]] \quad (16)
$$
\n
$$
= \sum_{\ell} \text{vec} [\text{mat}_{\ell} [\mathcal{D}^{\gamma}]]^{T} \mathbf{P}_{\ell \to 1}^{T} \left(\mathbf{I}_{d_{\leq \ell}} \otimes \Psi_{\ell} \otimes \mathbf{I}_{d_{>\ell}}\right) \mathbf{P}_{\ell \to 1} \text{vec} [\text{mat}_{\ell} [\mathcal{D}^{\gamma}]] \quad (17)
$$

$$
= \sum_{\ell} \text{vec} \left[\text{mat}_{\ell} \left[\mathcal{D}^{\gamma} \right] \right]^{T} \left(\mathbf{\Psi}_{\ell} \otimes \mathbf{I}_{d_{\setminus \ell}} \right) \text{vec} \left[\text{mat}_{\ell} \left[\mathcal{D}^{\gamma} \right] \right]
$$
\n(Rearrangement Lemma)

$$
= \sum_{\ell} \text{vec} \left[\text{mat}_{\ell} \left[\mathcal{D}^{\gamma} \right] \right]^T \text{vec} \left[\text{mat}_{\ell} \left[\mathcal{D}^{\gamma} \right] \Psi_{\ell}^T \right] \tag{18}
$$

$$
=\sum_{\ell} \operatorname{tr} \left[\mathbf{S}_{\ell}^{\gamma} \boldsymbol{\Psi}_{\ell} \right] \tag{19}
$$

 \Box

96

⁹⁷ With this lemma, the probability density function in the single-tensor case can be expressed in the ⁹⁸ form:

$$
p\left(\mathcal{D}^{\gamma}\right) = \frac{\sqrt{\left|\bigoplus_{\ell \in \gamma} \Psi_{\ell}\right|}}{\left(2\pi\right)^{\frac{d\gamma}{2}}} e^{\frac{-1}{2}\sum_{\ell} \text{tr}\left[\mathbf{S}_{\ell}^{\gamma} \Psi_{\ell}\right]}
$$
(20)

⁹⁹ Leading to the probability density function for the multi-tensor case as:

$$
p(\{\mathcal{D}^{\gamma}\}) = \prod_{\gamma} \frac{\sqrt{\left|\bigoplus_{\ell \in \gamma} \Psi_{\ell}\right|}}{(2\pi)^{\frac{d\sqrt{\gamma}}{2}}} e^{\frac{-1}{2}\sum_{\ell} \text{tr}\left[\mathbf{S}_{\ell}^{\gamma} \Psi_{\ell}\right]}
$$
(21)

$$
= \frac{\prod_{\gamma} \sqrt{\left|\bigoplus_{\ell \in \gamma} \Psi_{\ell}\right|}}{\left(2\pi\right)^{\frac{d_{\mathbf{y}}}{2}}} e^{\frac{-1}{2}\sum_{\gamma} \sum_{\ell} \text{tr}\left[\mathbf{S}_{\ell}^{\gamma} \Psi_{\ell}\right]}
$$
(22)

$$
= \frac{\prod_{\gamma} \sqrt{\left|\bigoplus_{\ell \in \gamma} \Psi_{\ell}\right|}}{\left(2\pi\right)^{\frac{d_{\mathbf{y}}}{2}}} e^{\frac{-1}{2} \sum_{\ell} \text{tr}[\mathbf{S}_{\ell} \Psi_{\ell}]} \tag{23}
$$

¹⁰⁰ The negative log-likelihood is thus:

$$
NLL (\{\mathcal{D}^{\gamma}\}) = \frac{d_{\forall}}{2} \log (2\pi) + \frac{1}{2} \sum_{\ell} \text{tr} \left[\mathbf{S}_{\ell} \boldsymbol{\Psi}_{\ell}\right] - \frac{1}{2} \sum_{\gamma} \log \left|\bigoplus_{\ell \in \gamma} \boldsymbol{\Psi}_{\ell}\right| \tag{24}
$$

¹⁰¹ 2.3 Gradient

 The derivation of the gradient of the negative log-likelihood is essentially the same as the derivation given by Kalaitzis et al. [\[6\]](#page-15-2) for the original Bi-Graphical Lasso. Our derivation is complicated by the fact that we are considering general tensors rather than matrices. We'll let sym be the symmetricizing operator that must be applied as we are taking the derivative with respect to a symmetric matrix: 106 sym $[M] = K \circ M$, where K is a matrix with 1s on the diagonal and 2s everywhere else. We'll also 107 define \mathbf{J}^{ij} to be the matrix of zeros except for a 1 at position (i, j) .

$$
\frac{d}{d\mathbf{\Psi}_{\ell}} \text{NLL} \left(\{ \mathcal{D}^{\gamma} \} \right) = \frac{1}{2} \text{sym} \left[\mathbf{S}_{\ell} \right] - \frac{1}{2} \sum_{\gamma} \frac{d}{d\mathbf{\Psi}_{\ell}} \log \left| \bigoplus_{\ell' \in \gamma} \mathbf{\Psi}_{\ell'} \right| \tag{25}
$$

$$
= \frac{1}{2} \text{sym} \left[\mathbf{S}_{\ell}\right] - \frac{1}{2} \sum_{\gamma} \text{tr}\left[\left(\bigoplus_{\ell' \in \gamma} \mathbf{\Psi}_{\ell'}\right)^{-1} \frac{d}{d\psi_{\ell}^{ij}} \bigoplus_{\ell' \in \gamma} \mathbf{\Psi}_{\ell'}\right]_{ij} \tag{26}
$$

$$
= \frac{1}{2} \text{sym} \left[\mathbf{S}_{\ell}\right] - \frac{1}{2} \sum_{\gamma} \text{tr}\left[\left(\bigoplus_{\ell' \in \gamma} \boldsymbol{\Psi}_{\ell'}\right)^{-1} \left(\mathbf{I}_{d_{<\ell}} \otimes \frac{d}{d\psi_{\ell}^{ij}} \boldsymbol{\Psi}_{\ell} \otimes \mathbf{I}_{d_{>\ell}}\right)\right]_{ij} (27)
$$

$$
= \frac{1}{2} \text{sym} \left[\mathbf{S}_{\ell}\right] - \frac{1}{2} \sum_{\gamma} \text{tr}\left[\left(\bigoplus_{\ell' \in \gamma} \boldsymbol{\Psi}_{\ell'}\right)^{-1} \left(\mathbf{I}_{d_{<\ell}} \otimes (\mathbf{J}^{ij} + \mathbf{J}^{ji} - \delta_{ij} \mathbf{J}^{ij}\right) \otimes \mathbf{I}_{d_{>\ell}}\right]_{ij} (28)
$$

$$
= \frac{1}{2} \text{sym} \left[\mathbf{S}_{\ell}\right] - \frac{1}{2} \sum_{\gamma} \left[(2 - \delta_{ij}) \text{tr} \left[\left(\bigoplus_{\ell' \in \gamma} \Psi_{\ell'} \right)^{-1} \left(\mathbf{I}_{d_{<\ell}} \otimes \mathbf{J}^{ij} \otimes \mathbf{I}_{d_{> \ell}} \right) \right] \right]_{ij}
$$
\n
$$
= \frac{1}{2} \text{sym} \left[\mathbf{S}_{\ell}\right] - \frac{1}{2} \sum_{\gamma} (2\mathbf{J} - \mathbf{I}) \circ \text{tr} \left[\left(\bigoplus_{\ell' \in \gamma} \Psi_{\ell'} \right)^{-1} \left(\mathbf{I}_{d_{<\ell}} \otimes \mathbf{J}^{ij} \otimes \mathbf{I}_{d_{> \ell}} \right) \right]_{ij}
$$
\n(29)\n(30)

$$
= \frac{1}{2} \text{sym} \left[\mathbf{S}_{\ell}\right] - \frac{1}{2} \sum_{\gamma} \text{sym} \left[\text{tr}_{d>\ell}^{d<\ell} \left[\left(\bigoplus_{\ell' \in \gamma} \boldsymbol{\Psi}_{\ell'} \right)^{-1} \right] \right] \tag{31}
$$

¹⁰⁸ The MLE occurs when this gradient is zero, i.e. when the following equation is satisfied:

$$
\mathbf{S}_{\ell} = \sum_{\gamma} \operatorname{tr}_{d > \ell}^{d < \ell} \left[\left(\bigoplus_{\ell' \in \gamma} \Psi_{\ell} \right)^{-1} \right]
$$
(32)

 In other words, our effective Gram matrices are the best Kronecker-sum decomposition of the covariance matrix of the maximum likelihood estimate. Unfortunately, Kronecker-sum decomposition does not interact well with matrix inverses, so this does not directly yield an analytic solution. It does, however, yield a solution for the eigenvectors.

¹¹³ 2.4 Maximum Likelihood Estimate for the Eigenvectors

¹¹⁴ We first produce two lemmas to aid in the derivation.

115 **Lemma 3** (Cyclic property of the stridewise-blockwise trace). *For any matrices* $M, A_{a \times a}, B_{b \times b}$, 116 *we have that* $\mathbf{tr}_{b}^{a} \, [(\mathbf{\tilde{A}} \otimes \mathbf{I} \otimes \mathbf{B}) \, \mathbf{M}] = \mathbf{tr}_{b}^{a} \, [\mathbf{M} \, (\mathbf{A} \otimes \mathbf{I} \otimes \mathbf{B})]$

- ¹¹⁷ *Proof.* This follows directly from the cyclic property of the (normal) trace operator and the definition ¹¹⁸ of the stridewise-blockwise trace. \Box
- ¹¹⁹ Lemma 4 (Conjugacy extraction of the stridewise-blockwise trace). *For any matrices* M *and* V*, we* 120 *have that* $\mathrm{tr}_b^a\left[\left(\mathbf{I}_a\otimes\mathbf{V}\otimes\mathbf{I}_b\right)\mathbf{M}\left(\mathbf{I}_a\otimes\mathbf{V}\otimes\mathbf{I}_b\right)^T\right]=\mathbf{V}\mathrm{tr}_b^a\left[\mathbf{M}\right]\mathbf{V}^T.$

$$
\begin{aligned} \textit{Proof.} \\ \textit{tr}_b^a\left[\left(\mathbf{I}_a\otimes\mathbf{V}\otimes\mathbf{I}_b\right)\mathbf{M}\left(\mathbf{I}_a\otimes\mathbf{V}\otimes\mathbf{I}_b\right)^T\right] &= \left[\textit{tr}\left[\left(\mathbf{I}_a\otimes\mathbf{V}\otimes\mathbf{I}_b\right)\mathbf{M}\left(\mathbf{I}_a\otimes\mathbf{V}\otimes\mathbf{I}_b\right)^T\left(\mathbf{I}_a\otimes\mathbf{J}^{ij}\otimes\mathbf{I}_b\right)\right]\right]_{ij} \\ \textit{(Definition of }\textit{tr}_b^a) \end{aligned}
$$

¹²¹ Thanks to the Rearrangement Lemma, we can get this just in terms of the standard blockwise trace,

- ¹²² for which there exists a convenient lemma from Dahl et al. [\[3\]](#page-15-3) that does the heavy lifting for us.
- ¹²³ Unfortunately, this requires inserting permutation matrices into every nook and cranny.

$$
= \left[\text{tr} \left[\mathbf{P} \left(\mathbf{I}_a \otimes \mathbf{V} \otimes \mathbf{I}_b \right) \mathbf{P}^T \mathbf{P} \mathbf{M} \mathbf{P}^T \mathbf{P} \left(\mathbf{I}_a \otimes \mathbf{V} \otimes \mathbf{I}_b \right)^T \mathbf{P}^T \mathbf{P} \left(\mathbf{I}_a \otimes \mathbf{J}^{ij} \otimes \mathbf{I}_b \right) \mathbf{P}^T \right] \right]_{ij} \tag{33}
$$

$$
= \left[\text{tr} \left[(\mathbf{V} \otimes \mathbf{I}_{ab})^T \mathbf{P} \mathbf{M} \mathbf{P}^T (\mathbf{V} \otimes \mathbf{I}_{ab}) (\mathbf{J}^{ij} \otimes \mathbf{I}_{ab}) \right] \right]_{ij}
$$
(34)

$$
= tr_{ab} \left[(\mathbf{V} \otimes \mathbf{I}_{ab}) \mathbf{P} \mathbf{M} \mathbf{P}^T (\mathbf{V} \otimes \mathbf{I}_{ab})^T \right]
$$
\n(Definition of tr_{ab})
\n
$$
= \mathbf{V}^{tr} \left[\mathbf{P} \mathbf{M} \mathbf{P}^T \mathbf{I} \mathbf{V}^T \right]
$$
\n(Definition of tr_{ab})

$$
= \mathbf{V} \text{tr}_{ab} \left[\mathbf{P} \mathbf{M} \mathbf{P}^T \right] \mathbf{V}^T
$$
 (Lemma 2 of Dahl et al. [3])

 \Box

124 We then can see analogously that tr_{ab} $\left[\mathbf{PMP}^T \right] = tr_b^a$ [M], completing the proof.

125

126 **Theorem 1.** Let $V_{\ell}e_{\ell}V_{\ell}^{T}$ be the eigendecomposition of S_{ℓ} . Then V_{ℓ} are the eigenvectors of the ¹²⁷ *maximum likelihood estimate of* Ψℓ*.*

Proof.

$$
\mathbf{S}_{\ell} = \sum_{\gamma} \operatorname{tr}_{d > \ell}^{d_{\leq \ell}} \left[\left(\bigoplus_{\ell' \in \gamma} \Psi_{\ell} \right)^{-1} \right]
$$
(35)

$$
= \sum_{\gamma} \text{tr}_{d>\ell}^{d} \left[\left(\bigoplus_{\ell' \in \gamma} \mathbf{V}_{\ell} \Lambda_{\ell} \mathbf{V}_{\ell}^{T} \right)^{-1} \right]
$$
(36)

$$
= \sum_{\gamma} \operatorname{tr}_{d>\ell}^{d<\ell} \left[\left(\bigotimes_{\ell'} \mathbf{V}_{\ell'} \right) \left(\bigoplus_{\ell' \in \gamma} \mathbf{\Lambda}_{\ell} \right)^{-1} \left(\bigotimes_{\ell \ell'} \mathbf{V}_{\ell'} \right)^{T} \right]
$$
(37)

$$
= \sum_{\gamma} \operatorname{tr}_{d_{> \ell}}^{d_{< \ell}} \left[\left(\mathbf{I}_{d_{< \ell}} \otimes \mathbf{V}_{\ell} \otimes \mathbf{I}_{d_{> \ell}} \right) \left(\bigoplus_{\ell' \in \gamma} \mathbf{\Lambda}_{\ell} \right)^{-1} \left(\mathbf{I}_{d_{< \ell}} \otimes \mathbf{V}_{\ell} \otimes \mathbf{I}_{d_{> \ell}} \right)^{T} \right] \quad \text{(Cyclic Property)}
$$

$$
= \sum_{\gamma} \mathbf{V} \mathrm{tr}_{d_{>\ell}}^{d_{<\ell}} \left[\left(\bigoplus_{\ell' \in \gamma} \Lambda_{\ell} \right)^{-1} \right] \mathbf{V}^{T}
$$
\n(Conjugacy Extraction)\n
$$
= \mathbf{V} \left(\sum_{\gamma} \mathrm{tr}_{d_{>\ell}}^{d_{<\ell}} \left[\left(\bigoplus_{\ell' \in \gamma} \Lambda_{\ell} \right)^{-1} \right] \right) \mathbf{V}^{T}
$$
\n(38)

¹²⁸ We conclude the proof by observing that the central matrix is diagonal, and thus the right hand side 129 constitutes an eigendecomposition of \mathbf{S}_{ℓ} . Thus \mathbf{S}_{ℓ} and Ψ_{ℓ} share eigenvectors. \Box

¹³⁰ 2.5 Maximum Likelihood Estimate for the Eigenvalues

¹³¹ In the previous section, we derived the eigenvectors of the maximum likelihood estimate. While ¹³² interesting (they correspond to the principal components of our data), we need the eigenvalues 133 to reconstruct Ψ_{ℓ} . Our strategy for this is to transform our data such that the precision matrices ¹³⁴ are diagonal, and estimate these diagonals. This transformation is stated in terms of the Tucker 135 operator $([\mathcal{D}^\gamma; \{\mathbf{V}_{\ell}^T\}_{\ell \in \gamma}])$. In the case where \mathcal{D} is a matrix, we have that $[\mathbf{D}]; \mathbf{V}_{rows}^T, \mathbf{V}_{cols}^T] =$ 136 $V_{rows} DV_{cols}^T$. While the definition of the Tucker operator can be given in terms of "n-mode prod-¹³⁷ ucts"[\[8\]](#page-15-0), we will only use the following property relating the Tucker operator to matricizationKolda ¹³⁸ [\[7\]](#page-15-4):

$$
\mathcal{Y} = [\![\mathcal{X}]; \{\mathbf{M}_{\ell}\}]
$$

\n
$$
\implies \text{mat}_{\ell}[\mathcal{Y}] \quad = \mathbf{M}_{\ell} \text{mat}_{\ell}[\mathcal{X}] \left(\mathbf{M}_{K} \otimes ... \otimes \mathbf{M}_{\ell+1} \otimes \mathbf{M}_{\ell-1} \otimes ... \otimes \mathbf{M}_{1}\right)^{T} \quad (\text{Kolda [7]})
$$

¹³⁹ The Tucker operator is an important concept for our calculation of the eigenvalues, but it is only the ¹⁴⁰ existence of such an operator that is important for our work; we never need to calculate it.

 Lemma 5 (Eigendecompositions of the Kronecker-sum-structured normal distribution). *Suppose* $\{D^{\gamma}\}\sim \mathcal{N}_{KS}(\{\Psi_{\ell}\})$. Then $\{[\![D^{\gamma};\{\mathbf{V}_{\ell}^{T}\}]\!]\}\sim \mathcal{N}_{KS}(\{\mathbf{\Lambda}_{\ell}\})$ and the effective Gram matrices of *this distribution are given by the eigenvalues* e^ℓ *of the effective Gram matrices* S^ℓ *of the original distribution.*

¹⁴⁵ *Proof.* We will prove this by showing that the probability density function is that of a Kronecker-¹⁴⁶ sum-structured normal distribution with the given parameters.

¹⁴⁷ In the first part of the proof, we will massage the probability density function into a convenient form -

¹⁴⁸ this does not depend on the Tucker decomposition, and holds for our original dataset as well.

$$
p([\mathcal{D}^{\gamma}; \{\mathbf{V}_{\ell}^{T}\}_{\ell \in \gamma}]) = p(\{\mathcal{D}^{\gamma}\})
$$
\n(39)

$$
= \frac{\prod_{\gamma} \sqrt{|\bigoplus_{\ell \in \mathcal{D}^{\gamma}} \Psi_{\ell}|}}{(2\pi)^{\frac{d_{\mathbf{y}}}{2}}} e^{\frac{-1}{2} \sum_{\ell} \text{tr}[\Psi_{\ell} \mathbf{S}_{\ell}]} \tag{40}
$$

$$
= \frac{\prod_{\gamma} \sqrt{|\bigoplus_{\ell \in \mathcal{D}^{\gamma}} \Lambda_{\ell}|}}{(2\pi)^{\frac{d_{\mathbf{y}}}{2}}} e^{\frac{-1}{2} \sum_{\ell} \text{tr} \left[\mathbf{V}_{\ell} \Lambda_{\ell} \mathbf{V}_{\ell}^{T} \mathbf{S}_{\ell}\right]}
$$
(41)

$$
= \frac{\prod_{\gamma} \sqrt{|\bigoplus_{\ell \in \mathcal{D}^{\gamma}} \Lambda_{\ell}|}}{(2\pi)^{\frac{d_{\mathbf{y}}}{2}}} e^{\frac{-1}{2} \sum_{\ell} \text{tr} \left[\mathbf{\Lambda}_{\ell} \mathbf{V}_{\ell}^{T} \mathbf{S}_{\ell} \mathbf{V}_{\ell} \right]}
$$
(42)

$$
=\frac{\prod_{\gamma}\sqrt{|\bigoplus_{\ell\in\mathcal{D}^{\gamma}}\mathbf{\Lambda}_{\ell}|}}{(2\pi)^{\frac{d_{\mathbf{v}}}{2}}}e^{\frac{-1}{2}\sum_{\ell}\text{tr}[\mathbf{\Lambda}_{\ell}\mathbf{e}_{\ell}]}\tag{43}
$$

149 To complete the proof, we must show that e_{ℓ} are the effective Gram matrices for $[\mathcal{D}_j; \{\mathbf{V}_{\ell}^T\}_{\ell \in \mathcal{D}_j}]$. 150 For brevity, let $\mathbf{V}_{\setminus \ell} = (\mathbf{V}_K \otimes ... \otimes \mathbf{V}_{\ell+1} \otimes \mathbf{V}_{\ell-1} \otimes ... \otimes \mathbf{V}_1).$

$$
\mathbf{e}_{\ell} = \mathbf{V}_{\ell}^T \mathbf{S}_{\ell} \mathbf{V}_{\ell} \tag{44}
$$

$$
= \sum_{\ell' \in \gamma} \frac{1}{n} \sum_{i}^{n} \mathbf{V}_{\ell}^{T} \text{mat}_{\ell} \left[\mathcal{D}_{i}^{\gamma} \right] \text{mat}_{\ell} \left[\mathcal{D}_{i}^{\gamma} \right]^{T} \mathbf{V}_{\ell}
$$
 (Definition of \mathbf{S}_{ℓ})

$$
= \sum_{\ell' \in \gamma} \frac{1}{n} \sum_{i}^{n} \mathbf{V}_{\ell}^{T} \text{mat}_{\ell} \left[\mathcal{D}_{i}^{\gamma} \right] \mathbf{V}_{\backslash \ell}^{T} \mathbf{V}_{\backslash \ell} \text{mat}_{\ell} \left[\mathcal{D}_{i}^{\gamma} \right]^{T} \mathbf{V}_{\ell}
$$
\n(45)

$$
= \sum_{\ell' \in \gamma} \frac{1}{n} \sum_{i}^{n} \text{mat}_{\ell} \left[\left[\mathcal{D}_{j}; \left\{ \mathbf{V}_{\ell}^{T} \right\}_{\ell \in \mathcal{D}_{j}} \right] \right] \text{mat}_{\ell} \left[\left[\mathcal{D}_{j}; \left\{ \mathbf{V}_{\ell}^{T} \right\}_{\ell \in \mathcal{D}_{j}} \right] \right]^{T} \tag{46}
$$

¹⁵¹ This completes the proof.

152

¹⁵³ Since this transformed data is still normally distributed with Kronecker-sum structure, we can use the ¹⁵⁴ previously derived gradient (Line [32\)](#page-6-1):

$$
\frac{d}{d\Lambda_{\ell}}\text{NLL}\left(\{\mathcal{D}^{\gamma}\}\right) = \mathbf{e}_{\ell} - \sum_{\gamma} \text{tr}_{d>\ell}^{d<\ell} \left[\left(\bigoplus_{\ell' \in \gamma} \Lambda_{\ell}\right)^{-1}\right]
$$
(47)

 \Box

- ¹⁵⁵ This yields Theorem [2:](#page-9-4)
- 156 **Theorem 2.** Let $\{G_\ell^\gamma\}$ be matrices such that the expression $\bigoplus_{\ell \in \gamma} G_\ell^\gamma$ is the best Frobenius-norm
- 157 *approximation of* $\left(\bigoplus_{\ell \in \gamma} \Lambda_\ell^t \right)^{-1}$. Then, for a learning rate μ_t , gradient descent can be performed

158 *with the update equation* $\Lambda_{\ell}^{t+1} = \Lambda_{\ell}^{t} - \mu_{t} \left[e_{\ell} - \sum_{\gamma|\ell \in \gamma} \mathbf{G}_{\ell}^{\gamma} \right]$. As Ψ_{ℓ} is positive definite, μ_{t} must

159 *be chosen to prevent* Λ^t_ℓ *from becoming negative.*

160 This is convenient because we have reduced our optimization task from one with $\sum_{\ell} d_{\ell}^2$ parameters 161 to one with $\sum_{\ell} d_{\ell}$ parameters.

¹⁶² 3 Dependences

 All tests and figures were generated on a Linux (Ubuntu 20.04) with an Intel Core i7 chip and 8GB of RAM. Along with our code, we provide an environment file (environment.yml) that contains full [d](https://github.com/NeurIPS-GmGM-Paper/GmGM)etails of all the dependencies used. In our GitHub repository [\(https://github.com/NeurIPS-GmGM-](https://github.com/NeurIPS-GmGM-Paper/GmGM) [Paper/GmGM\)](https://github.com/NeurIPS-GmGM-Paper/GmGM), we give precise and simple instructions on how to create a conda environment with the same setup as ours. Most of the packages used were specific to the experiments we ran. The dependencies necessary for our algorithm were Python 3.9 and NumPy 1.23.5.

¹⁶⁹ 4 Experiments

¹⁷⁰ 4.1 Synthetic data

 We generated random graphs by modelling each edge's probability of existing as being drawn from independent Bernoulli distributions. When estimating the runtimes, we ran all models five times and averaged out the results. When creating precision-recall curves, we averaged the results of fifty runs of the models. Due to space reasons, we omitted the precision-recall curves for the tensor-variate case in our main paper, so we provide this here in Figure [1.](#page-10-2)

¹⁷⁶ 4.2 COIL video

 We downloaded the processed COIL-20[\[10\]](#page-16-0) dataset, and tested our model on it. We wanted to see if our model could understand the structure of a video, which we expected to consist of two linear graphs (for the rows and columns, i.e. each row is connected only to its neighbor rows) and a circular graph (for the frames, because the video is of a 360° rotation). To generate these graphs, we ran our algorithm on the duck video from the dataset, and then greedily kept the largest edge from each vertex such that vertices in the final graph had at most two edges. If we shuffled our data (shuffle rows, columns, and frames) and try to reconstruct it with these graphs, we get mixed results (Figure ¹⁸⁴ [2\)](#page-10-3).

 We can put a numeric value to the reconstruction, by measuring the percentage of the time that our reconstructed edges connect two adjacent rows/columns/frames. We get an accuracy of 80% for the rows, 91% for the columns, and 99% for the frames. This hints that it is quite good at reconstructing frames of videos, but rows and columns are a harder task. This could be due to the specific characteristics of this video, in which there are a lot of rows that spend most of their time being mostly blue.

Figure 1: PR curves for the graphs generated from a 3-axis tensor. TeraLasso does almost perfectly; it can be hard to see as it is hugging the top right corner.

Figure 2: A reconstruction of the COIL-20 duck video after shuffling the rows, columns, and frames, using GmGM. While portions of the duck are well-reconstructed, it is clearly imperfect. Notably, the duck kisses itself.

4.3 EchoNet-Dynamic ECGs

 We downloaded all of the EchoNet-Dynamic[\[11\]](#page-16-1) data. This dataset did not have heartbeats labeled, so we picked a few videos at random and labeled them ourselves as a proof of concept. Specifically, we labeled every frame in which the mitral valve opened. Our goal was to see if the graphs produced by our algorithm could predict this opening. Table [1](#page-13-0) contains the videos we picked, the labels we gave, and the labels we predicted.

 Mitral valve predictions were done by taking GmGM's output frames graph in precision matrix form, and measuring the mass along the diagonals. We treated this as a time series (since each diagonal corresponds to an increasing time offset from all frames simultaneously). We applied gaussian blur and then a continuous wavelet transform peak detection algorithm[\[4\]](#page-15-5) to find which diagonals had the most mass (Figure [3\)](#page-11-2). These represent the offsets corresponding with a heartbeat. Given the first mitral valve opening and these offsets, we predict the remaining openings.

4.4 Mouse embryo stem cell transcriptomics

 We used the mouse embryo stem cell dataset E-MTAB-2805[\[2\]](#page-15-6). This dataset had already been labeled by what stage of the cell cycle each cell was in. The data was log-transformed, and we restricted the

Figure 3: An example heartbeat offset detection, from EchoNet-Dynamic video 0XFE6E32991136338. The blue curve represents our Gaussian-blurred diagonal mass (if x=10, it represents the blurred mass of the 10th diagonal to the right of the main diagonal). The red lines represent the predicted peaks via a continuous wavelet transform peak detection algorithm. These represent offsets from the first mitral valve opening. For this video, the mitral valve opened on frame 17 and our first offset was on the 30th diagonal. Hence, we would predict the second mitral valve opening to occur at frame 47 (which, in this case, was correct).

 gene set to the same mitosis-related genes used for Li et al. [\[9\]](#page-16-2)'s analysis of this same dataset. We kept the top 100 edges in our output graphs for each vertex, and set the rest to zero.

4.5 10x Genomics flash frozen lymph node

 For this experiment, we looked at a single-cell RNA-sequencing+ATAC-sequencing dataset from 10x Genomics[\[1\]](#page-15-7). We wanted to know whether clusters in UMAP-space make sense when viewed on GmGM's predicted graphs, whether clusters on the graphs made sense in UMAP-space, and whether these clusters had any meaning. Before performing the experiment, we removed cells whose library size was three median absolute deviations from the median, and similarly removed genes and peaks if the the total amount of times they were expressed was three median absolute deviations from the median. In our output graphs, we kept the top 5 edges per vertex.

 From Figures [4](#page-12-0) and [5,](#page-12-1) we can see that the clusters indeed seem to make sense in both UMAP-space and on the GmGM graph, as they all form coherent regions in both spaces.

 To validate that these clusters are meaningful, we performed a GO term enrichment analysis; the full results of this analysis are saved on our GitHub repository, but we summarize them here.

 Clusters 3 and 7 are clearly distinct in both spaces, and this is reflected in their GO terms. Cluster 3 was strongly associated with the CCKR signalling map and apoptosis, which none of the other clusters were. Cluster 7 was the most distinct, associated with the integrin signalling pathway, blood coagulation, and insulin. The other clusters all related to B and T cell-specific pathways. GmGM always grouped clusters 4 and 6 together, whereas UMAP would sometimes prefer to group cluster 6 with the rest of the clusters (compare Figures [5a](#page-12-1) and [6\)](#page-14-1).

4.6 LifeLines-DEEP metagenomics + metabolomics

 We used the LifeLines-DEEP metagenomics and metabolomics datasets[\[13\]](#page-16-3). We did not do any pre- processing to the metabolomics, and we used the already pre-processed version of the metagenomics data from Prost, Gazut, and Brüls [\[12\]](#page-16-4). We kept only patients that appeared in both datasets, and

Figure 4: (a) UMAP of the cells in the 10x Genomics dataset. Colored by kmeans (k=3). (b) GmGM's predicted graph over those cells, colored using the same clusters as on UMAP and plotted using igraph without reference to the outputs of UMAP.

Figure 5: (a) UMAP of the cells in the 10x Genomics dataset. Colored using same clusters as GmGM. (b) GmGM's predicted graph over those cells, colored using Louvain clustering.

0XF46CF63A2A1FA90 [25, 79, 134, 188] [25, 80, 133, 184] Table 1: Mitral valve labellings and precision matrices for the EchoNet-Dynamic dataset. The precision matrices, for the most part, seem to have clear off-diagonal structures, as expected, and the mitral valve prediction is generally quite good; it is only significantly off for the last opening in 0XF072F7A9791B060.

²³¹ log-transformed the data. We compared our model's results to the model given by Prost, Gazut, and ²³² Brüls [\[12\]](#page-16-4) in the main paper.

Figure 6: Another UMAP plot showing the same concept as Figure [5a,](#page-12-1) with the clusters labeled

Figure 7: (a) Runtimes of our algorithm and prior work on matrix-variate data. Our regularized algorithm is denoted "GmGM L1", and takes about the same time as the unregularized "GmGM". (b) Precision-recall curves for tensor-variate data. TeraLasso and our regularized "GmGM L1" perform almost perfectly.

²³³ 5 Regularization

 As remarked in the main paper, our algorithm by default includes no regularization. This is because our algorithm leverages the fact that we have a closed-form expression for the eigenvectors of the maximum likelihood estimate to avoid costly eigendecompositions every iteration. We do not have a closed-form expression for the eigenvectors in the regularized case.

 Nevertheless, we can add regularization to the eigenvalue estimation by performing an eigenrecom- position and regularizing that. Eigenrecomposition requires a matrix multiplication, which is quite costly compared to the cost of an unregularized iteration - both in practice, and asymptotically in 241 the matrix-variate case (matrix multiplication is $O(\sum_{\ell} d_{\ell}^3)$ whereas an unregularized iteration is

242 $O(\prod_{\ell} d_{\ell})$). Thus, to regularize we first let our algorithm converge to the MLE before considering the ²⁴³ penalty term. This allows us to avoid a major increase in runtime; our regularized algorithm runs in ²⁴⁴ roughly the same time as the unregularized one (Figure [7a\)](#page-14-2).

²⁴⁵ It is important to note that this estimator is slightly different than the standard Lasso estimator, as the 246 g standard estimator would minimize $\|\Psi_\ell\|_1$ and our estimator minimizes $\|\hat{V}_\ell\Lambda_\ell\hat{V}_\ell^T\|_1$ (where only

247 the eigenvalues Λ_{ℓ} are free to vary). It can be derived as follows:

$$
\frac{\partial}{\partial \lambda_i} \|\mathbf{V}\mathbf{\Lambda}\mathbf{V}^T\|_1 = \frac{\partial}{\partial \lambda_i} \|\sum_j \lambda_j v_{ja} v_{bj}\|_1
$$
\n(48)

= \lceil $\overline{}$ ∂ $\partial \lambda_i$ \sum .
j $\lambda_jv_{ja}v_{bj}$ 1 $\overline{1}$ ab (49)

$$
= \left[\frac{\partial}{\partial \lambda_i} \text{sign}\left[\sum_j \lambda_j v_{ja} v_{bj}\right] v_{ia} v_{bi}\right]_{ab} \tag{50}
$$

$$
= \left[\text{sign} \left[\mathbf{V} \mathbf{\Lambda} \mathbf{V}^T \right]_{ab} v_{ia} v_{bi} \right]_{ab} \tag{51}
$$

$$
= \mathbf{v}_i^T \text{sign} \left[\mathbf{V} \boldsymbol{\Lambda} \mathbf{V}^T \right] \mathbf{v}_i \tag{52}
$$

²⁴⁸ Despite this difference, it performs comparably to prior work. We show in Figure [7b](#page-14-2) the precision-²⁴⁹ recall curves for the 3-axis case, and observe that it performs almost perfectly. This is notable as it

²⁵⁰ was the case that the unregularized algorithm performed worse than prior work.

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