Low-Dimensional Lipschitz Embeddings Invariant to Permutations

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Norbert Wiener Center for Harmonic Analysis and Applications

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Joint work with:

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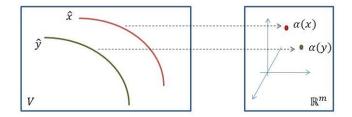
Maneesh Singh (Verisk)

arXiv preprint: 2203.07546 [math.FA], [cs.LG]



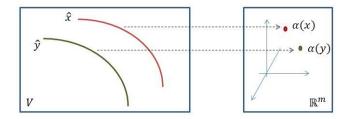
High-Level View

In this talk, we discuss Euclidean embeddings of metric spaces induced by representations of permutation (sub)groups \mathcal{S}_n on linear spaces V. Problem: Construct bi-Lipschitz embeddings of the metric space $\hat{V} = V / \sim$ of orbits, $\alpha: \hat{V} \to \mathbb{R}^m$, where $d(\hat{x}, \hat{y}) = \min_{u \in \hat{x}, v \in \hat{y}} \|u - v\|_V$.



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Today we focus on the case $V = \mathbb{R}^{n \times d}$, $X \sim Y \Leftrightarrow Y = PX$ for some $P \in \mathcal{S}_n$.

4 D > 4 B > 4 E > 4 E > 9 Q Q

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- **2** Embeddings of \hat{V} for $V = \mathbb{R}^{n \times d}$
- 3 Sorting based Embeddings
- Numerical Examples

Motivation

Motivation

- 2 Embeddings of \hat{V} for $V = \mathbb{R}^{n \times d}$
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Graph Learning Problems

Given a data graph (e.g., social network, transportation network, citation network, chemical network, protein network, biological networks):

- Graph adjacency or weight matrix, $A \in \mathbb{R}^{n \times n}$:
- Data matrix, $X \in \mathbb{R}^{n \times r}$, where each row corresponds to a feature vector per node.

Contruct a map $f:(A,X)\to f(A,X)$ that performs:

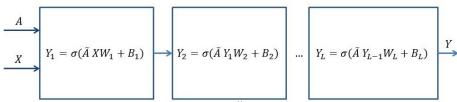
- classification: $f(A, X) \in \{1, 2, \dots, c\}$
- 2 regression/prediction: $f(A, X) \in \mathbb{R}$.

Key observation: The outcome should be invariant to vertex permutation: $f(PAP^T, PX) = f(A, X)$, for every $P \in \mathcal{S}_n$.



Graph Convolution Networks (GCN), Graph Neural Networks (GNN)

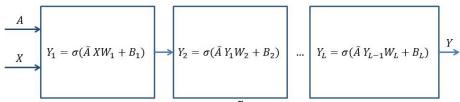
General architecture of a GCN/GNN



GCN (Kipf and Welling ('16)) choses $\tilde{A} = I + A$; GNN (Scarselli et.al. ('08), Bronstein et.al. ('16)) choses $\tilde{A} = p_I(A)$, a polynomial in adjacency matrix. *L*-layer GNN has parameters $(p_1, W_1, B_1, \dots, p_L, W_L, B_L)$.

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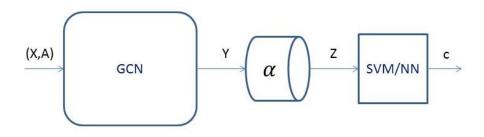


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Note the covariance (or, equivariance) property: for any $P \in O(n)$ (including S_n), if $(A, X) \mapsto (PAP^T, PX)$ and $B_i \mapsto PB_i$ then $Y \mapsto PY$.

Deep Learning with GCN/GNN

The approach for the two learning tasks (classification or regression) is based on the following scheme (see also Maron et.al. ('19)):



where α is a permutation invariant map (embedding), and SVM/NN is a single-layer or a deep neural network (Support Vector Machine or a Fully Connected Neural Network) trained on invariant representations.

The purpose of this talk is to analyze the α component.

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- Embeddings of \hat{V} for $V = \mathbb{R}^{n \times d}$



Motivation

The metric space \hat{V} when $V = \mathbb{R}^{n \times d}$

Recall the equivalence relation \sim on $V=\mathbb{R}^{n\times d}$ induced by the group of permutation matrices S_n acting on V by left multiplication: for any $X, X' \in \mathbb{R}^{n \times d}$.

$$X \sim X' \Leftrightarrow X' = PX$$
, for some $P \in \mathcal{S}_n$

Let $\mathbb{R}^{n \times d} = \mathbb{R}^{n \times d} / \sim$ be the quotient space endowed with the natural distance induced by Frobenius norm $\|\cdot\|_F$

$$d(\hat{X}_1, \hat{X}_2) = \min_{P \in S_n} \|X_1 - PX_2\|_F \ , \ \hat{X}_1, \hat{X}_2 \in \widehat{\mathbb{R}^{n \times d}}.$$

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The computation of the minimum distance is performed by solving the Linear Assignment Problem (LAP) whose convex relaxation is exact:

$$\max_{P \in \mathcal{S}_n} trace(PX_2X_1^T) = \max_{W \in DS(n)} trace(WX_2X_1^T)$$

where $DS(n) = \{W \in [0,1]^{n \times n} : W1 = 1, W^T1 = 1\}$ is the convex set of doubly stochastic matrices.

The embedding problem

Problem: Construct a bi-Lipschitz embedding $\hat{\alpha}: \mathbb{R}^{n \times d} \to \mathbb{R}^m$, i.e., an integer m = m(n,d), a map $\alpha: \mathbb{R}^{n \times d} \to \mathbb{R}^m$ with constants $0 < a \le b < \infty$ so that for any $X, X' \in \mathbb{R}^{n \times d}$,

- If $X \sim X'$ then $\alpha(X) = \alpha(X')$.
- 2 If $\alpha(X) = \alpha(X')$ then $X \sim X'$.

where $d(\hat{X}, \hat{X}') = \min_{P \in \mathcal{S}_n} \|X - PX'\|_F$.

Consider the map

$$\mu: \widehat{\mathbb{R}^{n \times d}} \to \mathcal{P}(\mathbb{R}^d) \ , \ \mu(X)(x) = \frac{1}{n} \sum_{k=1}^n \delta(x - x_k)$$

where $\mathcal{P}(\mathbb{R}^d)$ denotes the convex set of probability measures over \mathbb{R}^d , and δ denotes the Dirac measure. x_k is the k^{th} row of X.

Clearly $\mu(X') = \mu(X)$ iff X' = PX for some $P \in \mathcal{S}_n$.

The Wasserstein-2 distance is equivalent to the natural metric:

$$W_2(\mu(X), \mu(Y))^2 := \inf_{q \in J(\mu(X), \mu(Y))} \mathbb{E}_q[\|x - y\|_2^2] = \min_{P \in \mathcal{S}_n} \|Y - PX\|^2$$

By Kantorovich-Rubinstein theorem, the Wasserstein-1 distance (the Earth moving distance) extends to a norm on the space of signed Borel measures.

Main drawback: $\mathcal{P}(\mathbb{R}^d)$ is infinite dimensional!

Motivation

Finite Dimensional Embeddings

Idea: "Project" the measure onto a finite dimensional space. This is accomplished by *kernel methods*:

Fix a family of functions f_1, \dots, f_m and consider:

$$\mu(X) \mapsto \int_{\mathbb{R}^d} f_j(x) d\mu(X) = \frac{1}{n} \sum_{k=1}^n f_j(x_k) \quad , \quad j \in [m]$$

Finite Dimensional Embeddings

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Possible choices:

- Polynomial embeddings: $\mathbb{R}[X]^{S_n}$, ring of invariant polynomials; [Lipman&al.], [Peyré&al.], [Sanay&al.], [Kemper book] ...
- ② Gaussian kernels: $f_j(x) = exp(-\|x a_j\|^2/\sigma_j^2)$; [Gilmer&al.],[Zaheer&al.], [Vinyals&al.],...
- **3** Fourier kernels (cmplx embd): $f_j(x) = exp(2\pi i \langle x, \omega_j \rangle)$; related to Prony method; [Li&Liao] for bi-Lipschitz estimates.

Main drawback: No global bi-Lipschitz embeddings [Cahill&al.]. Ok on (some) compacts.

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The Max Pool approach

Motivation

The idea is provided by the following observation.

Let
$$\downarrow$$
: $\mathbb{R}^n \to \mathbb{R}^n$ denote the *sorting map* $x \mapsto \downarrow x = \Pi x$, $\Pi \in \mathcal{S}_n$, so that $(\Pi x)_1 \geq (\Pi x)_2 \geq \cdots \geq (\Pi x)_n$.

The Max Pool approach

The idea is provided by the following observation.

Let $\downarrow: \mathbb{R}^n \to \mathbb{R}^n$ denote the sorting map $x \mapsto \downarrow x = \Pi x$, $\Pi \in \mathcal{S}_n$, so that $(\Pi x)_1 \geq (\Pi x)_2 \geq \cdots \geq (\Pi x)_n$

Lemma

 $\downarrow: \widehat{\mathbb{R}^n} \to \mathbb{R}^n$ is an isometry (hence bi-Lipschitz):

$$\|\downarrow(x)-\downarrow(y)\|=\min_{P\in\mathcal{S}_n}\|x-Py\|$$
, for all $x,y\in\mathbb{R}^n$.

Proof is based on the rearrangement inequality (see Wikipedia, or Hardy-Littlewood-Pólya "Inequalities" §10.2).

Our main goal is to extend this construction from \mathbb{R}^n to $\mathbb{R}^{n\times d}$

The Encoder β_A

Notations

Motivation

Recall the equivalence relation, for $X, Y \in \mathbb{R}^{n \times d}$,

$$X \sim Y \Leftrightarrow \exists \Pi \in \mathcal{S}_n, Y = \Pi X$$

that induces a quotient space $\mathbb{R}^{n \times d} = \mathbb{R}^{n \times d} / \sim$ and the natural distance

$$d:\widehat{\mathbb{R}^{n\times d}}\times\widehat{\mathbb{R}^{n\times d}}\to\mathbb{R}$$
 , $d(X,Y)=\min_{\Pi\in\mathcal{S}_n}\|X-\Pi Y\|_F$

In the following we look for an Euclidean embedding of the form

$$\beta_A: \mathbb{R}^{n \times d} \to \mathbb{R}^{n \times D}$$
 , $\beta_A(X) = \downarrow (XA)$

where \downarrow (·) sorts decreasingly each column of ·, independently. The matrix $A \in \mathbb{R}^{d \times D}$ is called the *key* of encoder β_A .

The key is called *universal* if $\widehat{\beta_A} : \widehat{\mathbb{R}^{n \times d}} \to \mathbb{R}^{n \times D}$ is injective

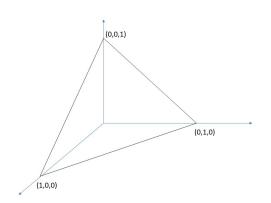
Sorting

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Consider the case

n = 2, d = 3

$$X = \left[\begin{array}{ccc} X_{11} & X_{12} & X_{13} \\ X_{21} & X_{22} & X_{23} \end{array} \right]$$



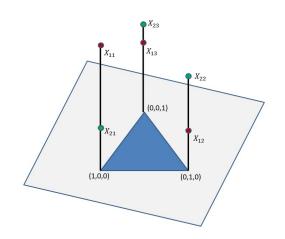
Intuition behind universality of keys

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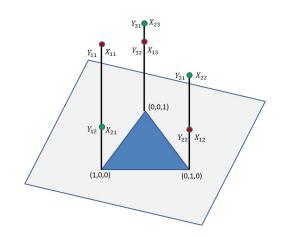
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$$X = \left[\begin{array}{ccc} X_{11} & X_{12} & X_{13} \\ X_{21} & X_{22} & X_{23} \end{array} \right]$$

$$Y = \downarrow X$$

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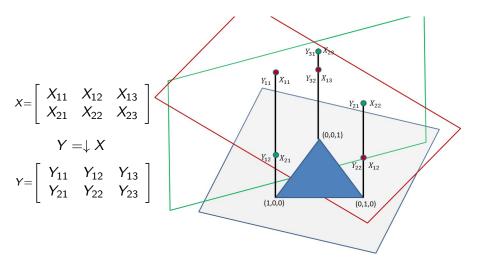
Information lost!



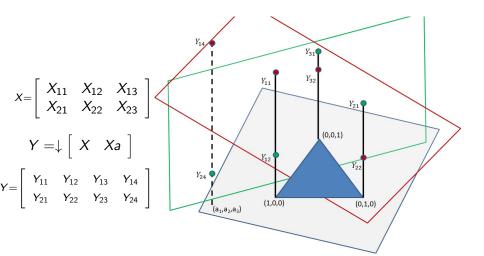
Intuition behind universality of keys

Sorting

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Intuition for this encoder



Sorting

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Theorem

Motivation

Consider the metric space $(\widehat{\mathbb{R}^{n\times d}},d)$. Set D=1+(d-1)n! and let $A\in\mathbb{R}^{d\times D}$ be a matrix whose columns form a full spark frame. Then the key A is universal and the induced map $\hat{\beta}_A:\widehat{\mathbb{R}^{n\times d}}\to\mathbb{R}^{n\times D}$, $\hat{\beta}_A(\hat{X})=\downarrow (XA)$ is injective. Furthermore, $\hat{\beta}_A$ is bi-Lipschitz with constants $a_0=\min_{J\subset [D],|J|=d}s_d(A[J])$ and $b_0=s_1(A)$, where $s_1(A)$ denotes the largest singular value of A, A[J] denotes the submatrix of A formed by columns indexed by J, and $s_d(A[J])$ denotes the d^{th} singular value (in this case, the smallest) of A[J]. Specifically, for any $X,Y\in\mathbb{R}^{n\times d}$,

$$a_0 d(\hat{X}, \hat{Y}) \le \|\beta_A(X) - \beta_A(Y)\| \le b_0 d(\hat{X}, \hat{Y})$$
 (3.1)

where all norms are Frobenius norms.

Three results (2)

Bi-Lipschitz Property of Universal Keys

Theorem

Assume the key $A \in \mathbb{R}^{d \times D}$ is universal, i.e., the induced map $\hat{\beta}_A : \widehat{\mathbb{R}^{n \times d}} \to \mathbb{R}^{n \times D}$, $X \mapsto \beta_A(X) = \downarrow (XA)$ is injective. Then $\hat{\beta}_A$ is bi-Lipschitz, that is, there are constants $a_0 > 0$ and $b_0 > 0$ so that for all $X, Y \in \mathbb{R}^{n \times d}$.

$$a_0 d(\hat{X}, \hat{Y}) \le \|\beta_A(X) - \beta_A(Y)\| \le b_0 d(\hat{X}, \hat{Y})$$
 (3.2)

where all are Frobenius norms. Furthermore, an estimate for b_0 is provided by the largest singular value of A, $b_0 = s_1(A)$.



Dimension Reduction

Theorem

Assume $A \in \mathbb{R}^{d \times D}$ is a universal key for $\mathbb{R}^{n \times d}$ with $D \geq 2d$. Then, for $m \geq 2nd$, a generic linear operator $B : \mathbb{R}^{n \times D} \to \mathbb{R}^m$ with respect to Zariski topology on $\mathbb{R}^{n \times D \times m}$, the map

$$\hat{\beta}_{A,B}: \widehat{\mathbb{R}^{n \times d}} \to \mathbb{R}^{2nd} , \ \hat{\beta}_{A,B}(\hat{X}) = B\left(\hat{\beta}_A(\hat{X})\right)$$
 (3.3)

is bi-Lipschitz. In particular, almost every full-rank linear operator $B: \mathbb{R}^{n \times D} \to \mathbb{R}^{2nd}$ produces such a bi-Lipschitz map.

This result is compatible with a Whitney embedding theorem with the important caveat that the Whitney embedding result applies to smooth manifolds, whereas $\widehat{\mathbb{R}^{n\times d}}$ is not a manifold.

Highlights of proofs

Universal keys

Motivation

The upper bound is imediate. For lower bound, fix $X, Y \in \mathbb{R}^{n \times d}$:

$$\|\beta_{A}(X) - \beta_{A}(Y)\|_{2}^{2} = \sum_{k=1}^{D} \|\downarrow (Xa_{k}) - \downarrow (Ya_{k})\|_{2}^{2} = \sum_{k=1}^{D} \|P_{k}Xa_{k} - Q_{k}Ya_{k}\|_{2}^{2}$$

$$\stackrel{\Pi_{k}:=Q_{k}^{T}P_{k}}{=} \sum_{k=1}^{D} \|(\Pi_{k}X - Y)a_{k}\|_{2}^{2}$$

Highlights of proofs

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$$= \sum_{k=1}^{\Pi_{k}:=Q_{k}^{T}P_{k}} \sum_{k=1}^{D} \|(\Pi_{k}X - Y)a_{k}\|_{2}^{2} > \sum_{k=1}^{D} \|(\Pi_{k}X - Y)a_{k}\|_{2}^{2}$$

$$\stackrel{\Pi_{k:=Q_{k}^{T}P_{k}}}{=} \sum_{k=1}^{D} \|(\Pi_{k}X - Y)a_{k}\|_{2}^{2} \ge \sum_{j=1}^{d} \|(\Pi_{k_{j}}X - Y)a_{k_{j}}\|_{2}^{2}$$

so that $\Pi_{k_1} = \cdots = \Pi_{k_d} = \Pi_0$ (pigeonhole principle: needs D > (d-1)n!).

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$$\stackrel{\prod_{k:=Q_k^T P_k}}{=} \sum_{k=1}^D \|(\prod_k X - Y)a_k\|_2^2 \ge \sum_{j=1}^d \|(\prod_{k_j} X - Y)a_{k_j}\|_2^2$$

so that $\Pi_{k_1} = \cdots = \Pi_{k_d} = \Pi_0$ (pigeonhole principle: needs D > (d-1)n!). Then:

$$\|\beta_{A}(X) - \beta_{A}(Y)\|_{2}^{2} \geq \sum_{i=1}^{d} \|(\Pi_{0}X - Y)a_{k_{j}}\|_{2}^{2} \stackrel{\text{full spark}}{\geq} s_{d}(A[J])^{2} \|\Pi_{0}X - Y\|^{2}$$

$$\geq s_d(A[J])^2 \min_{\Pi \in S_n} \|\Pi X - Y\|^2 = s_d(A[J])^2 d(\hat{X}, \hat{Y})^2$$

Highlights of proofs

Bi-Lipschitz Property

The proof resembles the treatment of phase retrieval problem:

- Homogeneity and compactness reduce the problem to local analysis.
- The encoder is "locally" linearized. The failure of local lower Lipschitz bound implies a certain behavior for a Quadratically Constrained Ratio of Quadratics (QCRQ).
- QCRQ has a minimizer:inf ⇒ min. [Teboulle&al.] This step took most of time and lots of (self)convincing!
- Contradiction to injectivity assumption.

Highlights of proofs

Dimension Reduction

Motivation

The proof follows the approach in [Cahill&al.], [Dufresne]:

$$0 = B(\beta_A(X)) - B(\beta_A(Y)) \Rightarrow \beta_A(X) - \beta_A(Y) \in \ker(B)$$

Need to show: $\beta_A(X) - \beta_A(Y) = 0$, or, $Ran(\Delta) \cap \ker(B) = \{0\}$, where

$$\Delta: \mathbb{R}^{n\times d} \times \mathbb{R}^{n\times d} \to \mathbb{R}^{n\times D}, \ \Delta(X,Y) = \beta_A(X) - \beta_A(Y).$$

In the polynomial case, [Cahill&al.] exploit arguments from algebraic geometry. Here the problem is simpler since $Ran(\Delta)$ is included in a finite union of linear subspaces of dimension at most 2nd.

By a dimension argument it follows that the target space for B must be of dimension at least 2nd to obtain an injective embedding. In this case, generically, $Ran(\Delta)$ and ker(B) intersect transversally.



Towards universal keys

The arXiv preprint provides necessary and sufficient conditions for a key to be universal.

Open Problem: Given (n, d) find the smallest dimension D so that there exists a universal key $A \in \mathbb{R}^{d \times D}$ for $\mathbb{R}^{n \times d}$.

So far we obtained (joint with Daniel Levy (UMD)):

n	d	D-d
2	2	1
3	2	2
4	2	2
5	2	3
6	2	≥ 4

Open Problem: If a universal key exists for a triple (n, d, D) then is it true that universal keys are generic in $\mathbb{R}^{d \times D}$?

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The Protein Dataset

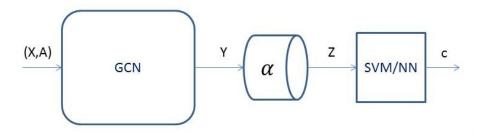
Protein Dataset: 663 non-enzymes and 450 enzymes out of 1113 proteins. Each graph associated to one protein: nodes represent amino acids and edges represent the bonds between them. Number of nodes (aminoacids): varying between 20 and 620 with average of 39. Input feature vectors os size r=29.

Task: the task is classification of each protein into enzyme or non-enzyme.

The Deep Network Architecture

Architecture: ReLU activation and

- GCN with L=3 layers and 29 input feature vectors, and 50 hidden nodes in each layer; no dropouts, no batch normalization. output of GCN: d=1,10,50,100.
- Mid-layer component: α
- Fully connected NN with dense 3-layers and 150 internal units; no dropouts, with batch normalization.



The Network

Training has been done over 300 epochs with a batch size of 128. Loss function: binary cross-entropy.

The following 7 α modules have been tested:

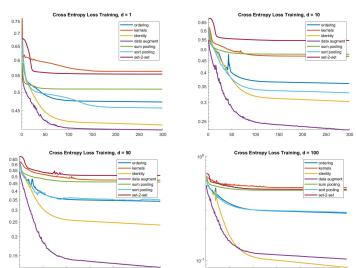
- **1** identity: $\alpha(X) = X$; no permutation invariance.
- ② data augmentation: $\alpha(X) = X$ BUT the training data set has been augmented with 4 random permutations of each graph.
- **3** ordering: $\alpha(X) = \downarrow (XA)$, $A = [I \ 1]$
- kernels: $\alpha(X) = (\sum_{k=1}^{n} exp(-\|x_k a_j\|^2))_{1 \le j \le m = 5nd}$
- **3** sumpooling: $\alpha(X) = 1^T X$
- o sort-pooling: sorted by last column
- set-to-set: introduced in [Vinyals&al.]



Enzyme Classification Example

Training Loss: X Entropy

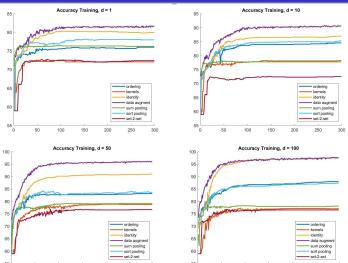
Motivation





Enzyme Classification Example

Accuracy on Training set

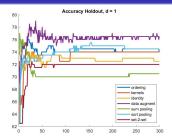


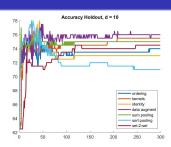
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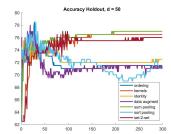
100 150

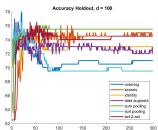
200 250

Accuracy on Holdout data



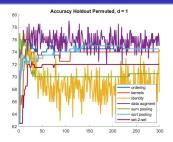


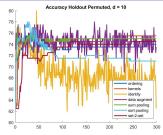


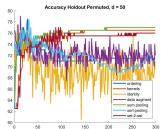


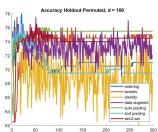
Enzyme Classification Example

Accuracy on Holdout data with nodes randomly permuted









Performance Results: Accuracy

d = 50	ordering	kernels	identity	data	sum-	sort-	set-2-
				augment	pooling	pooling	set
Training	83.1	78.8	91	96	79.2	83.7	76.7
Holdout	71.5	76.5	72.5	71	77	71	76
Holdout Perm	71.5	76.5	69.5	72	77	71	76

Table: Accuracy ACC(%) for enzyme/non-enzyme classification of the seven algorithms on PROTEINS_FULL dataset after 300 epochs for embedding dimension d = 50

For comparison: [Dobson&al.] obtain an accuracy of 77-80% using an SVM based classifier.

The QM9 Dataset

Dataset: Consists of about 134,000 isomers of organic molecules made up of CHONF, each containing 10-29 atoms. see

http://quantum-machine.org/datasets/ Nodes corresponds to atoms; each feature vector containins geometry (x,y,z coordinates), partial charge per atom (Mulliken charge), and atom type.

Task: the task is regression: predict a physical feature (electron energy gap $\Delta \varepsilon$) computed for each molecule.

Architecture: ReLU activation and

- GCN with L=3 layers and 50 hidden nodes in each layer; no dropouts, no batch normalization; zero padding to m = 29 number of rows. output of GCN: d = 1, 10, 50, 100.
- Mid-layer component: α
- Fully connected NN with dense 3-layers and 150 internal units in each of the two hidden layers; no dropouts, with batch normalization.

The Network

Training has been done over 300 epochs with a batch size of 128. Loss function: Mean-Square Error (MSE).

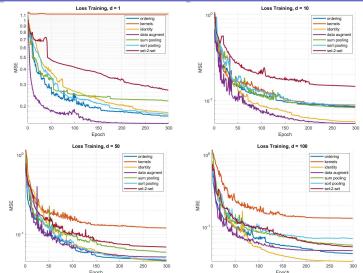
The same 7 α modules have been tested:

- **1** identity: $\alpha(X) = X$; no permutation invariance.
- ② data augmentation: $\alpha(X) = X$ BUT the training data set has been augmented with 4 random permutations of each graph.
- ordering: $\alpha(X) = \downarrow (XA)$, $A = [I \ 1]$
- kernels: $\alpha(X) = (\sum_{k=1}^{n} exp(-\|x_k a_j\|^2))_{1 \le j \le m = 5nd}$
- **3** sumpooling: $\alpha(X) = 1^T X$
- sort-pooling: sorted by last column
- set-to-set: introduced in [Vinyals&al.]



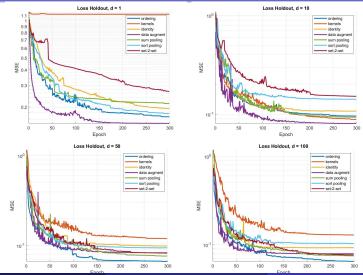
QM9 Regression Example

Training MSE



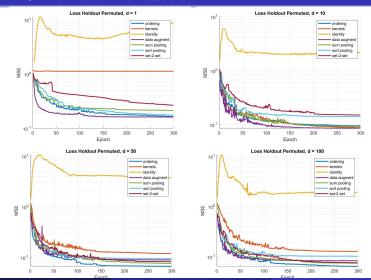
QM9 Regression Example

Validation MSE



QM9 Regression Example

Validation MSE with Random Permutations



d = 100	ordering	kernels	identity	data	sum-	sort-	set-2-
				augment	pooling	pooling	set
Training	0.155	0.269	0.139	0.164	0.178	0.199	0.173
Holdout	0.187	0.267	0.227	0.206	0.201	0.239	0.201
Holdout Perm	0.187	0.267	1.086	0.213	0.201	0.239	0.201

Table: Mean Absolute Error (MAE) for regression of the electron energy gap $\Delta \varepsilon = LUMO - HOMO$ (eV) of the seven algorithms on QM9 dataset after 300 epochs for embedding dimension d=100

For comparison:

- chemical accuracy is 0.043eV
- the best ML method [Gilmer&al.] achieves MAE of 0.053eV
- Coulomb method [Rupp&al.] achieves MAE of 0.229eV



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Thank you! Questions?

The Embedding Problem

Notations (2)

Definition

Fix $X \in \mathbb{R}^{n \times d}$. A matrix $A \in \mathbb{R}^{d \times D}$ is called admissible for X if $\beta_A^{-1}(\beta_A(X)) = \hat{X}$. In other words, if $Y \in \mathbb{R}^{n \times d}$ so that $\downarrow (XA) = \downarrow (YA)$ then there is $\Pi \in \mathcal{S}_n$ sot that $Y = \Pi X$.

We denote by $A_{d,D}(X)$ (or A(X)) the set of admissible keys for X.

Definition

Fix $A \in \mathbb{R}^{d \times D}$. A data matrix $X \in \mathbb{R}^{n \times d}$ is said separated by A if $A \in \mathcal{A}(X)$.

We let S(A) denote the set of data matrices separated by A. The key A is universal iff $S(A) = \mathbb{R}^{n \times d}$.



Genericity Results for d > 2

Admissible keys

Theorem

Let $X \in \mathbb{R}^{n \times d}$. For any $D \geq d+1$ the set $\mathcal{A}_{d,D}(X)$ of admissible keys for X is dense in $\mathbb{R}^{d \times D}$ with respect to Euclidean topology, and it is generic with respect to Zariski topology. In particular, $\mathbb{R}^{d \times D} \setminus \mathcal{A}_{d,D}(X)$ has Lebesgue measure 0, i.e., almost every key is admissible for X.

Proof

It is sufficient to consider the case D = d + 1. Also, it is sufficient to analyze the case $A = [I_d \ b]$ and to show that a generic $b \in \mathbb{R}^d$ defines an admissible key. The vector $b \in \mathbb{R}^d$ does **not** define an admissible key if there are $\Xi, \Pi_1, \dots, \Pi_d \in S_n$ so that for $Y = [\Pi_1 x_1, \dots, \Pi_d x_d]$,

$$Yb = \Xi Xb$$
 but $Y - \Pi X \neq 0$, $\forall \Pi \in S_n$

Define the linear operator

Admissible keys

Motivation

Proof - cont'd

Let

$$\mathcal{P} = \left\{ (\Pi_1, \cdots, \Pi_d) \in (\mathcal{S}_n)^d \ \forall \Pi \in \mathcal{S}_n, \exists k \in [d] \ s.t. \ (\Pi - \Pi_k) x_k \neq 0 \right\}$$

Then

$$\{b \in \mathbb{R}^d : [I_d \ b] \text{ not admissible for } X\} = \bigcup_{(\Xi; \Pi_1, \cdots, \Pi_d) \in \mathcal{S}_n \times \mathcal{P}} \ker(B(\Xi; \Pi_1, \cdots, \Pi_d)) \in \mathcal{S}_n \times \mathcal{P}$$

It is now sufficient to show that each null space has dimension less than d. Indeed, the alternative would mean $B(\Xi;\Pi_1,\cdots,\Pi_d)=0$ but this would imply $(\Pi_1,\cdots,\Pi_d)\not\in\mathcal{P}$. \square



Non-Universality of vector keys

Insufficiency of a single vector key

The following is a no-go result, which shows that there is no universal single vector key for data matrices tall enough.

Proposition

If d > 2 and n > 3.

$$\bigcup_{X \in \mathbb{R}^{n \times d}} \{b \in \mathbb{R}^d: \ A = [I_d \ b] \ \text{not admissible for} X\} = \mathbb{R}^d.$$

Consequently,

$$\bigcap_{X\in\mathbb{R}^{n\times d}}\mathcal{A}_{d,d+1}(X)=\emptyset.$$

On the other hand, for n = 2, d = 2, any vector $b \in \mathbb{R}^2$ with $b_1 b_2 \neq 0$ defines a universal key $A = [I_2 \ b]$.

Insufficiency of a single vector key - cont'd

Proof

To show the result, it is sufficient to consider a counterexample for n = 3, d = 2, with key $b = [1, 1]^T$.

$$X = \begin{bmatrix} 1 & -1 \\ -1 & 0 \\ 0 & 1 \end{bmatrix} , Y = \begin{bmatrix} 1 & 0 \\ -1 & 1 \\ 0 & -1 \end{bmatrix}$$

Then $Xb = [0, -1, 1]^T$ and $Yb = [1, 0, -1]^T$, yet $X \not\sim Y$. Thus $[I_2 \ b]$ is not admissible for X.

Then note if $a \in \mathbb{R}^d$ so that $[I_d \ a]$ is admissible for X then for any $P \in S_d$ and L an invertible $d \times d$ diagonal matrix, $L^{-1}P^TA \in \mathcal{A}_{d,1}(XPL)$. This shows how for any $b \in \mathbb{R}^2$, one can construct $X \in \mathbb{R}^{3 \times 2}$ so that $b \notin \mathcal{A}_{2,1}(X)$.

For n > 3 or d > 2, proof follows by embedding this example.

Genericity Results for d > 2

Admissible Data Matrices

Theorem

Assume $a \in \mathbb{R}^d$ is a vector with non-vanishing entries, i.e., $a_1 a_2 \cdots a_d \neq 0$. Then for any n > 1, $S([I_d \ a])$ is dense in $\mathbb{R}^{n \times d}$ and includes an open dense set with respect to Zariski topology. In particular, $\mathbb{R}^{n\times d}\setminus\mathcal{S}([I_d\ a])$ has Lebesgue measure 0, i.e., almost every data matrix X is separated by the vector key a.

Genericity Results for $d \ge 2$

Admissible Data Matrices

Theorem

Assume $a \in \mathbb{R}^d$ is a vector with non-vanishing entries, i.e., $a_1a_2\cdots a_d \neq 0$. Then for any $n \geq 1$, $\mathcal{S}([I_d\ a])$ is dense in $\mathbb{R}^{n\times d}$ and includes an open dense set with respect to Zariski topology. In particular, $\mathbb{R}^{n\times d}\setminus\mathcal{S}([I_d\ a])$ has Lebesgue measure 0, i.e., almost every data matrix X is separated by the vector key a.

Corollary

Assume $A \in \mathbb{R}^{d \times (D-d)}$ is a matrix such that at least one column has non-vanishing entries. Then for any $n \geq 1$, $\mathcal{S}([I_d \ A])$ is dense in $\mathbb{R}^{n \times d}$ and is generic with respect to Zariski topology. In particular, $\mathbb{R}^{n \times d} \setminus \mathcal{S}([I_d \ A])$ has Lebesgue measure 0, i.e., almost every data matrix X is separated by the matrix key $[I_d \ A]$.

Proof that $\mathcal{S}([I_d A])$ is generic

The case D > d

Motivation

Assume $A \in \mathbb{R}^{d \times (D-d)}$ satisfies $A_{1,k}A_{2,k}\cdots A_{d,k} \neq 0$ for some $k \in [D-d]$. The set of non-separated data matrices $X \in \mathbb{R}^{n \times d}$ (i.e., the complement of $\mathcal{S}([I_d A])$ factors as follows:

$$\mathbb{R}^{n\times d}\setminus\mathcal{S}([I_d\ A])=\bigcup_{(\Xi_1,\cdots,\Xi_{D-d};\Pi_1,\cdots,\Pi_d)\in(\mathcal{S}_n)^D}(\ker L(\Xi_1,\cdots,\Xi_{D-d};\Pi_1,\cdots,\Pi_d;A))$$

$$\setminus \bigcup_{\Pi \in \mathcal{S}_n} \ker M(\Pi, \Pi_1, \cdots, \Pi_d)$$
 (*)

where, with $A = [a_1, \dots, a_{D-d}], X = [x_1, \dots, x_d]$:

$$L(\Xi_1,\cdots,\Xi_{D-d};\Pi_1,\cdots,\Pi_d;A):\mathbb{R}^{n\times d}\to\mathbb{R}^{n\times D-d}\ ,\ (L((\ldots)X)_k=[(\Xi_k-\Pi_1)x_1,\cdots,(\Xi_k-\Pi_d)x_d]a_k\ ,\ k\in[D-d]$$

$$M(\Pi,\Pi_1,\cdots,\Pi_d):\mathbb{R}^{n\times d}\to\mathbb{R}^{n\times d}\quad,\quad M(\Pi,\Pi_1,\cdots,\Pi_d)X=[(\Pi-\Pi_1)\underline{x_1},\cdots,(\underline{\Pi}-\Pi_d)\underline{x_d}]$$

Proof that S(A) is generic

cont'd

1. The outer union can be reduced by noting that on the "diagonal" Δ ,

$$\Delta = \{ (\Xi_1, \cdots, \Xi_{D-d}; \Pi_1, \cdots, \Pi_d) \in (\mathcal{S}_n)^D \ , \ \Pi_1 = \Pi_2 = \cdots = \Pi_d \}$$
$$M(\Pi_1, \Pi_1, \cdots, \Pi_d) = 0 \to \bigcup_{\Pi \in \mathcal{S}_n} \ker M(\Pi, \Pi_1, \cdots, \Pi_d) = \mathbb{R}^{n \times d}$$

2. If $(\Xi_1,\cdots,\Xi_{D-d};\Pi_1,\cdots,\Pi_d)\in (\mathcal{S}_n)^D\setminus \Delta$ then for every $k\in [D-d]$ there is $j\in [d]$ such that $\Xi_k-\Pi_j\neq 0$. In particular choose the k column of A that is non-vanishing. Let $x_j\in \mathbb{R}^n$ so that $(\Xi_k-\Pi_j)x_j\neq 0$. Consider the matrix $X=[0,\cdots,0,x_j,0,\cdots,0]$ where x_j is the only non identically 0 column. Claim: $X\not\in\ker L(\Xi_1,...,\Pi_d;A)$. Indeed, the resulting k column of L()X is $A_{j,k}(\Xi_k-\Pi_j)x_j\neq 0$. It follows that $\dim\ker L(\Xi_1,\cdots,\Xi_{D-d};\Pi_1,\cdots,\Pi_d;A)< nd$

Hence $\mathbb{R}^{n\times d}\setminus\mathcal{S}([I_d\ A])$ is a finite union of subsets of closed linear spaces properly included in $\mathbb{R}^{n\times d}$. This proves the theorem.

Additional Relations

Note the following relationship and matrix representation of X when matrices are column-stacked:

$$M(\Pi, \Pi_1, \dots, \Pi_d) = L(\Pi, \dots, \Pi; \Pi_1, \dots, \Pi_d; I)$$

$$L \equiv \begin{bmatrix} A_{1,1}(\Xi_1 - \Pi_1) & A_{2,1}(\Xi_1 - \Pi_2) & \cdots & A_{d,1}(\Xi_1 - \Pi_d) \\ A_{1,2}(\Xi_2 - \Pi_1) & A_{2,2}(\Xi_2 - \Pi_2) & \cdots & A_{d,2}(\Xi_2 - \Pi_d) \\ \vdots & \vdots & \ddots & \vdots \\ A_{1,D-d}(\Xi_{D-d} - \Pi_1) & A_{2,D-d}(\Xi_{D-d} - \Pi_2) & \cdots & A_{d,D-d}(\Xi_{D-d} - \Pi_d) \end{bmatrix}$$

a $n(D-d) \times nd$ matrix.

