

G-Invariant Representations using Coorbits

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Preprints

Preprints:

1. R.B., Naveed Haghani, Maneesh Singh, “Permutation Invariant Representations with Applications to Graph Deep Learning”, arXiv: 2203.07546 [math.FA] , [cs.LG]
2. R.B., Efstratios Tsoukanis, “Relationships between the Phase Retrieval Problem and Permutation Invariant Embeddings”, arXiv:2306.13111 [math.FA] , [cs.IT] , [math.IT]
3. R.B., Efstratios Tsoukanis, “G-Invariant Representations using Coorbits: Bi-Lipschitz Properties”, arXiv:2308.11784 [math.RT]
4. R.B., Efstratios Tsoukanis, “G-Invariant Representations using Coorbits: Injectivity Properties”, arXiv:2310.16365 [math.RT]
5. R.B, Efstratios Tsoukanis, Matthias Wellershoff, “Stability of sorting based embeddings”, arXiv:2410.05446 [math.FA]

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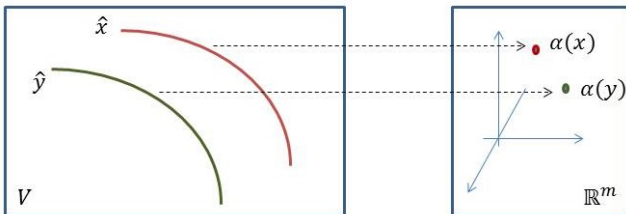
6 Extra

High-Level View

In this talk, we discuss Euclidean embeddings of metric spaces induced by orthogonal representations of finite groups G acting on a linear space V with inner product.

Problem: Construct bi-Lipschitz embeddings of the metric space $\hat{V} = V / \sim$ of orbits, $\alpha : \hat{V} \rightarrow \mathbb{R}^m$, where $\mathbf{d}([x], [y]) = \inf_{u \in [x], v \in [y]} \|u - v\|$

$$a_0 \mathbf{d}([x], [y]) \leq \|\alpha([x]) - \alpha([y])\|_2 \leq b_0 \mathbf{d}([x], [y]).$$



The Program

Given a discrete group G acting unitarily on a normed real space V , we formulate four general problems

- ① Construct injective embeddings of the quotient space V/G , $\alpha : \hat{V} \rightarrow \mathbb{R}^m$. [The injectivity problem.](#)
- ② Construct/Obtain bi-Lipschitz properties for the Euclidean embedding $\alpha : \hat{V} \rightarrow \mathbb{R}^m$. [The stability problem.](#)
- ③ Develop algorithms for inversion $\alpha^{-1} : \mathbb{R}^m \rightarrow \hat{V}$. [The recovery problem.](#)
- ④ Analyze specific cases. [Applications.](#)

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Today we discuss results about the first two problems: **injectivity**, **bi-Lipschitz stability**.

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I. 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.

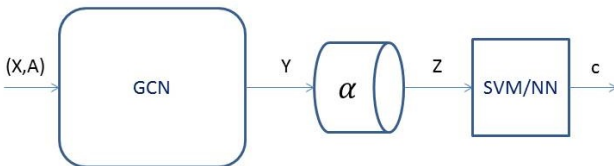
Construct a map $f : (A, X) \rightarrow f(A, X)$ that performs:

- 1 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 Deep Learning with GCN/GNN

Our approach for these learning tasks (classification or regression) is based on the following scheme (see GCN¹ and equivariance²):



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.

[Our focus is on the \$\alpha\$ component.](#)

¹Kipf, T. N. and Welling, M., Semi-Supervised Classification with Graph Convolutional Networks, arXiv e-prints , arXiv:1609.02907 (Sep 2016).

²H. Maron, E. Fetaya, N. Segol, Y. Lipman, On the Universality of Invariant Networks, arXiv:1901.09342 [cs.LG] (May 2019).

II. Assignment Problems

The Graph Isomorphism Problem

Consider two graphs $G = (\mathcal{V}, \mathcal{E})$ and $\tilde{G} = (\tilde{\mathcal{V}}, \tilde{\mathcal{E}})$ with n nodes. The graph isomorphism problem is the computational problem of determining whether these graphs are identical after a relabeling of nodes.

If A and \tilde{A} denote their adjacency matrices, **these graphs are isomorphic if and only if $\tilde{A} = \Pi A \Pi^T$ for some permutation matrix $\Pi \in \mathcal{S}_n$.**

Current state-of-the-art (Wikipedia): Babai (2015,2017) presented a quasi-polynomial algorithm with running time $2^{O((\log n)^c)}$, for some fixed $c > 0$. Helfgott (2017) claims that one can take $c = 3$.

Similar problem can be stated for weighted graphs: $A, \tilde{A} \in \text{Sym}(n)$ with nonnegative entries, isomorphic if and only if $\tilde{A} = \Pi A \Pi^T$ for some $\Pi \in \mathcal{S}_n$.

Graph Alignment Problems

Consider two $n \times n$ symmetric matrices A, B . The “vanilla” alignment problem for quadratic forms asks for the orthogonal matrix $U \in O(n)$ that minimizes

$$\|UAU^T - B\|_F^2 := \text{trace}((UAU^T - B)^2) = \|A\|_F^2 + \|B\|_F^2 - 2\text{trace}(UAU^T B).$$

The solution is well-known and depends on the eigendecomposition of matrices A, B : if $A = U_1 D_1 U_1^T$, $B = U_2 D_2 U_2^T$ then

$$U_{opt} = U_2 U_1^T, \quad \|U_{opt} A U_{opt}^T - B\|_F^2 = \sum_{k=1}^n |\lambda_k - \mu_k|^2,$$

where $D_1 = \text{diag}(\lambda_k)$ and $D_2 = \text{diag}(\mu_k)$ are diagonal matrices with eigenvalues ordered monotonically.

Quadratic Assignment Problem (QAP)

The challenging case is when U is constrained to the permutation group as is the case in the *graph matching problem*. In this case, the optimization problem becomes

$$\min_{U \in \mathcal{S}_n} \|UAU^T - B\|_F$$

which turns into a QAP: $\max_{U \in \mathcal{S}_n} \text{trace}(UAU^T B)$.

This is equivalent to computing the natural distance

$d(\hat{A}, \hat{B}) = \min_{P, Q \in \mathcal{S}_n} \|PAP^T - QBQ^T\|_F$ between the equivalence classes $\hat{A}, \hat{B} \in \widehat{\text{Sym}(n)}$ induced by action $(\Pi, A) \mapsto \Pi A \Pi^T$.

How is this connected to the embedding problem? If one can design an *efficient* nearly isometric map $\Phi : \text{Sym}(n) \rightarrow \mathbb{R}^m$ so that

(1) $\Phi(PAP^T) = \Phi(A)$ for all $P \in \mathcal{S}_n$ and $A \in \text{Sym}(n)$, and

(2) $(1-\delta) \min_{P \in \mathcal{S}_n} \|PAP^T - B\| \leq \|\Phi(A) - \Phi(B)\| \leq (1+\delta) \min_{P \in \mathcal{S}_n} \|PAP^T - B\|$,

then the QAP solved efficiently up to a multiplicative factor.

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Problem Setup

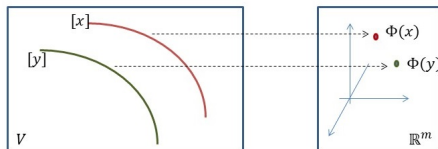
Consider a group $G \subset O(d)$ acting on the Euclidean space $V = \mathbb{R}^d$.

General problem

Construct an embedding map $\Phi : V \rightarrow \mathbb{R}^m$

- 1 Invariance: $\Phi(U_g x) = \phi(x) \quad \forall g \in G, x \in V$
- 2 Injectivity: if $\Phi(x) = \Phi(y)$ then there exists $g \in G$ so that $y = U_g x$.
- 3 Φ is bi-Lipschitz on $(\hat{V} = V/G, \mathbf{d})$:

$$a_0 \inf_{u \in [x], v \in [y]} \|u - v\| \leq \|\Phi(x) - \Phi(y)\| \leq b_0 \inf_{u \in [x], v \in [y]} \|u - v\|.$$



Approaches

Over the past many years, several constructions have been proposed:

- ① Invariant Polynomials: Hilbert, Noether, ..., Cahill³, Bandeira⁴
- ② Kernels: replace monomials by other kernels, e.g. $e^{i\omega x}$, e^{-x^2} , $\sigma(\langle x, a \rangle)$ ⁵
- ③ Sorting: extends the 1-D sorting, $x \mapsto \downarrow x$ ^{6,7}

~~1+2: *sum pooling layer*; 3: extension of *max pooling layer* in deep nets^{8,9}.~~

³J. Cahill, A. Contreras, A.C. Hip, Complete Set of translation Invariant Measurements with Lipschitz Bounds, Appl. Comput. Harm. Anal. 49 (2020), 521–539.

⁴A. Bandeira, B. Blum-Smith, J. Kileel, J. Nilés-Weed, A. Perry, A.S. Wein, Estimation under group actions: Recovering orbits from invariants, ACHA 66 (2023)

⁵D. Yarotsky, Universal approximations of invariant maps by neural networks, Constructive Approximation (2021)

⁶R. Balan, N. Haghani, M. Singh, Permutation Invariant Representations with Applications to Graph Deep Learning, arXiv:2203.07546

⁷J. Cahill, J.W. Iverson, D.G. Mixon, D. Packer, Group-invariant max filtering, arXiv:2205.14039.

⁸O. Vinyals, S. Bengio, M. Kudlur, Order Matters: Sequence to sequence for sets, Proc. ICLR 2016

Sorting based Representations and G-invariance

Assume V is a real d -dimensional Hilbert space and G a finite orthogonal group of size $N = |G|$, acting on V , $\{U_g, g \in G\}$.

Fix a generator $w \in V$ (call it, *window*, or *template*, or *wavelet*) and consider the nonlinear map induced by sorting its coorbit:

$$\phi_w : V \rightarrow \mathbb{R}^N, \quad \phi_w(x) = \downarrow ((\langle x, U_g w \rangle)_{g \in G}).$$

where $\downarrow(y) = (y_{\pi(i)})_{i \in [N]}$ is the non-increasing sorting operator:

$$y_{\pi(1)} \geq \dots \geq y_{\pi(N)}.$$

Key observations:

- 1 $\phi_w(U_g x) = \phi_w(x)$, ϕ is G -invariant.
- 2 ϕ_w is piecewise linear (in fact, $\phi_w(x) = \phi_x(w)$, and $(w, x) \mapsto \phi_w(x)$ is piecewise bilinear).

G-Invariant Coorbit Representations

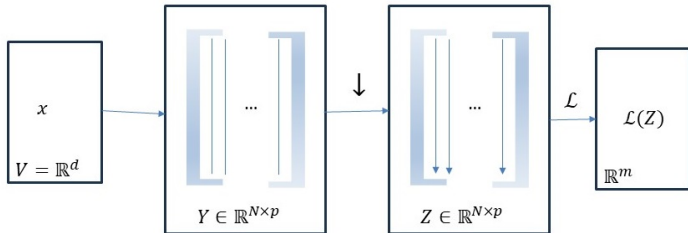
For a collection $\mathbf{w} = (w_1, \dots, w_p) \in V^p$ the sorted coorbit representation:

$$\Phi_{\mathbf{w}} : V \rightarrow \mathbb{R}^{N \times p}, \quad \Phi_{\mathbf{w}}(x) = [\phi_{w_1}(x) | \dots | \phi_{w_p}(x)].$$

Pass through a linear operator $\mathcal{L} : \mathbb{R}^{N \times p} \rightarrow \mathbb{R}^m$, the G-invariant coorbit representation:

$$\Psi_{\mathbf{w}, \mathcal{L}} : V \rightarrow \mathbb{R}^m, \quad \Psi_{\mathbf{w}, \mathcal{L}}(x) = \mathcal{L}(\Phi_{\mathbf{w}}(x))$$

$$x \mapsto Y := (\langle x, U_g w \rangle)_{g \in G} \times p \quad Y \mapsto Z := \downarrow (\langle x, U_g w \rangle)_{g \in G} \times p \quad Z \mapsto \mathcal{L}(Z)$$



In particular, if $S \subset [N] \times [p]$, $\Phi_{\mathbf{w}, S} := \Psi_{\mathbf{w}, 1_S} = \Phi_{\mathbf{w}}|_S$.

G-Invariant Coorbit Representations

Special cases:

1. For $G = S_n$ and $V = \mathbb{R}^{n \times d}$ with action $(P, X) \mapsto PX$ ¹⁰ introduced the embedding $\beta_A(X) = \downarrow (XA)$, for key $A \in \mathbb{R}^{d \times D}$ and sorting operator acting independently in each column. This is of the type $\Psi_{w, \mathcal{L}}$ for $w_1 = \delta_1 \cdot a_1^T, \dots, w_D = \delta_1 \cdot a_D^T$, where $\delta_1 = (1, 0, \dots, 0)^T$ and $A = [a_1 | \dots | a_D]$, and \mathcal{L} a restriction operator to an appropriate subset $S \subset [n!] \times [D]$ of size nD .

2. The *max filter* introduced in ¹¹ for some template $w \in V$ is defined by $\langle \langle \cdot, w \rangle \rangle : V \rightarrow \mathbb{R}$, $\langle \langle x, w \rangle \rangle = \max_{g \in G} \langle x, U_g w \rangle$. Equivalent recasting: $\langle \langle x, w \rangle \rangle = \mathcal{L}(\Phi_w(X))$, for a restriction operator \mathcal{L} to the subset $S = \{1\}$.

3. The operator $\Psi_{w, \mathcal{L}}$, $\Psi_{w, \mathcal{L}}(X) = \mathcal{L}(\Phi_w(X))$ has been introduced in ¹²

¹⁰R. Balan, N. Haghani, M. Singh, Permutation Invariant Representations with Applications to Graph Deep Learning, arXiv:2203.07546 (2022)

¹¹J. Cahill, J. W. Iverson, D. G. Mixon, D. Packer, Group-invariant max filtering, arXiv:2205.14039 (2022)

¹²R.B, Efstratios Tsoukanis, Matthias Wellershoff, "Stability of sorting based embeddings", arXiv:2410.05446 (2024)

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Main Results

Injectivity

Let $V_G = \{x \in V : U_g x = x, \forall g \in G\}$, $d_G = \dim(V_G)$, $q \geq 0$ and for $g = (g_1, \dots, g_n), h = (h_1, \dots, h_n) \in G^n$ distinct, $\rho_n(q) = \max_{g,h} \gamma_{g,h}^q$ where

$$\gamma_{g,h}^q = \text{semi.alg.dim.} \{(x, y) \in V \times V : \dim(\text{span}\{U_{g_k} x - U_{h_k} y, k \in [n]\}) = q\}$$

Theorem (R.B., E. Tsoukanis '23-'25)

In any of the following cases

- 1 Assume $p \geq 2 \dim(V) - d_G$ and set $\mathbf{n} = (1, \dots, 1) \in [N]^p$.
- 2 Fix $n \in [N]$ and choose $p > \max_{q \in [n]} \frac{1}{q} (\rho_n(q) - d_G - 1)$. Set $\mathbf{n} = (n, \dots, n) \in [N]^p$.
- 3 Choose $p \geq 1$ and $\mathbf{n} = (n_1, \dots, n_p) \in [N]^p$ so that $\max_{q_1 \in [n_1], \dots, q_p \in [n_p]} (\min_{i \in [p]} \rho_{n_i}(q_i) - (q_1 + \dots + q_p)) \leq d_G$.

For a generic (w.r.t. Zariski topology) \mathbf{w} and for any $S \subset [N] \times [p]$ with $|\{k : (k, i) \in S\}| \geq n_i$, the map $\Phi_{\mathbf{w}, S} : (\widehat{V}, \mathbf{d}) \rightarrow (\mathbb{R}^{|S|}, \|\cdot\|_2)$ is injective.

Main Results (2)

Theorem (R.B, E.T., M. Wellershoff '24)

Consider the same setup as before. Assume $\mathbf{w} \in V^p$ and $\mathcal{L} : \mathbb{R}^{N \times p} \rightarrow \mathbb{R}^m$ so that $\Psi_{\mathbf{w}, \mathcal{L}} : (\widehat{V}, \mathbf{d}) \rightarrow (\mathbb{R}^m, \|\cdot\|_2)$ is **injective**.

- 1 The map $\Psi_{\mathbf{w}, \mathcal{L}} : (\widehat{V}, \mathbf{d}) \rightarrow (\mathbb{R}^m, \|\cdot\|_2)$ is bi-Lipschitz. Let a_0, b_0 denote its bi-Lipschitz constants.
- 2 If $f : V \rightarrow H$ is a Lipschitz continuous function so that $f(U_g x) = f(x)$ for all g, x , where H is a Hilbert space, then there exists a Lipschitz continuous function $g : \mathbb{R}^m \rightarrow H$ so that $f = g \circ \Psi_{\mathbf{w}, \mathcal{L}}$, i.e. $f(x) = g(\Psi_{\mathbf{w}, \mathcal{L}}(x))$. Furthermore, $Lip(g) \leq Lip(f)/a_0$.
- 3 Assume $g : \mathbb{R}^m \rightarrow H$ is a Lipschitz function with Lipschitz constant $Lip(g)$. Then $f = g \circ \Psi_{\mathbf{w}, \mathcal{L}} : V \rightarrow H$ is G -invariant and Lipschitz, with Lipschitz constant $Lip(f) \leq b_0 Lip(g)$.

Its proof is based on Kirszbraun's extension theorem.

Existing Results

Injectivity problem

Over the past 15 years or so, there have been works that recognized the difference between *generating polynomials* and *separating invariants*¹³. A seminal paper that resurfaces results on semi-algebraic sets is ¹⁴. The method goes back to earlier works in phase retrieval¹⁵.

More recently, in the context of G-invariance, ^{16, 17}, or permutation invariance¹⁸

¹³Emilie Dufresne, Separating invariants and finite reflection groups, *Advances in Mathematics* 221 (2009), no. 6, 1979–1989.

¹⁴Dym Nadav, Steven J. Gortler. "Low dimensional invariant embeddings for universal geometric learning." arXiv preprint arXiv:2205.02956.

¹⁵R. Balan, P. Casazza, D. Edidin, On signal reconstruction without phase, *ACHA* 20(2006)

¹⁶D. G. Mixon, D. Packer, Max filtering with reflection groups, arXiv:2212.05104

¹⁷R. Balan, E. Tsoukanis, G-invariant representations using coorbits: Injectivity properties, arXiv:2310.16365

¹⁸On the equivalence between graph isomorphism testing and function approximation with GNNs, Z. Chen, S. Villar, L. Chen, I. Bruna, *NeurIPS* 2019

Existing Results (2)

Lipschitz and Bi-Lipschitz properties

Earlier results obtain Lipschitz/bi-Lipschitz properties on compacts, or certain classes of functions.

Global L/bi-L are harder to establish and typically rule out polynomial based embeddings.

So far only sorting based embeddings showed such global properties^{19, 20, 21}

¹⁹R. Balan, E. Tsoukanis, G-invariant representations using coorbits: Bi-lipschitz properties, arXiv:2308.11784

²⁰J. Cahill, J. W. Iverson, D. G. Mixon, Bilipschitz group invariants, arXiv:2305.17241

²¹D. G. Mixon, Y. Qaddura, Injectivity, stability, and positive definiteness of max filtering, arXiv:2212.11156

Sketch of Proof: Injectivity Result

Define the “bad” set of \mathbf{w} 's that fail to separate all distinct classes:

$$\mathcal{F} = \{ \mathbf{w} \in V^p, \exists x \neq y \Phi_{\mathbf{w}}(x) = \Phi_{\mathbf{w}}(y) \}.$$

The work is to embed \mathcal{F} into a semi-algebraic set of semi-algebraic dimension strictly less than $pd = p \dim(V)$.

This technique is called “lift-and-project”²²: we construct a semi-algebraic vector bundle embedded into a certain Grassmanian vector bundle $\gamma_{n,k}^\perp$. The bad set \mathcal{F} is then identified with a subset of the projection of this vector bundle into its second component.

The full result for $\Psi_{\mathbf{w},\mathcal{L}}$ follows from analyzing the semi-algebraic dimension of difference set $\{ \Phi_{\mathbf{w}}(x) - \Phi_{\mathbf{w}}(y) \}$ and of the kernel of \mathcal{L} .

²²R. Balan, P. Casazza, D. Edidin, On signal reconstruction without phase, ACHA 20(2006)

Sketch of Proof: Lower Lipschitz bound

The proof is by contradiction. Consider the simpler case when \mathcal{L} is given by restriction to a subset $S \subset [M] \times [p]$.

1. If lower Lipschitz constant vanishes, then it must vanish locally: there are $(x_n)_n, (y_n)_n$ such that

$$\lim_{n \rightarrow \infty} \frac{\|\Phi_{\mathbf{w}, S}(x_n) - \Phi_{\mathbf{w}, S}(y_n)\|^2}{\mathbf{d}([x_n], [y_n])^2} = 0$$

and

$$\lim_{n \rightarrow \infty} x_n = \lim_{n \rightarrow \infty} y_n = z_1, \quad \|x_n\| = 1, \quad \|y_n\| \leq 1, \quad \|z_1\| = 1$$

and they are aligned with one another:

$$\|x_n - y_n\| = \min_{g \in G} \|x_n - U_g y_n\| \tag{4.1}$$

$$\|x_n - z_1\| = \min_{g \in G} \|x_n - U_g z_1\| \tag{4.2}$$

$$\|y_n - z_1\| = \min_{g \in G} \|y_n - U_g z_1\| \tag{4.3}$$

Lower Lipschitz bound

2. We construct inductively z_2, z_3, \dots, z_d such that for all $1 \leq k \leq d - 1$:

$$\|z_{k+1}\| \ll \|z_k\|, \quad \dim(\text{span}(z_1, \dots, z_k)) = k$$

and the local lower Lipschitz constant vanishes in a convex set

$$\left\{ \sum_{r=1}^k a_r z_r, \quad |a_r - 1| < \epsilon \right\}.$$

3. For $k = d$ this construction defines a non-empty open set

$\left\{ \sum_{r=1}^d a_r z_r, \quad |a_r - 1| < \epsilon \right\}$ where the local lower Lipschitz constant vanishes.

4. Finally, we can construct $u, v \neq 0$, so that $x = u + \sum_{r=1}^d z_r$ and $y = v + \sum_{r=1}^d z_r$ satisfy $x \neq y$ and yet

$$\Phi_{w,S}(x) = \Phi_{w,S}(y).$$

This contradicts the injectivity hypothesis.

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The Protein Dataset

Protein Dataset: PROTEINS_FULLL²³ consists of 1113 proteins: 663 non-enzymes and 450 enzymes. 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 of size $r = 29$.

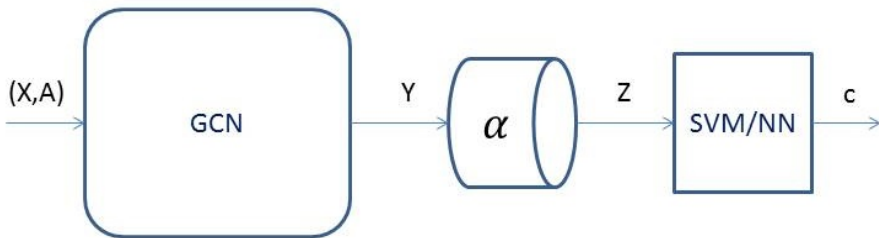
Task: the task is classification of each protein into *enzyme* or *non-enzyme*.

²³P.D. Dobson, A.J. Doig, "Distinguishing Enzyme Structures from Non-enzymes without Alignments", J. Mol. Biol. 330, 771-783, 2003.

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.

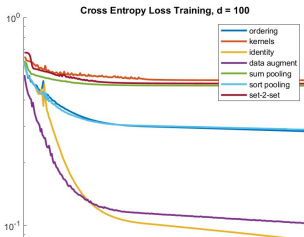
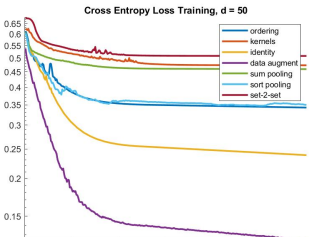
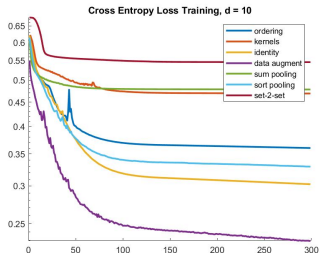
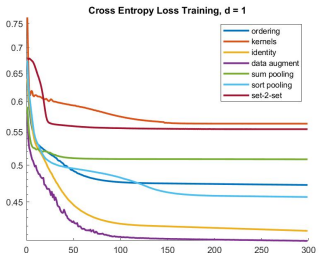
The following 7 α modules have been tested:

- 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 permutatons of each graph.
- ③ ordering: $\alpha(X) = \downarrow (XA)$, $A = [I \ 1]$
- ④ kernels: $\alpha(X) = (\sum_{k=1}^n \exp(-\|x_k - a_j\|^2))_{1 \leq j \leq m=5nd}$
- ⑤ sumpooling: $\alpha(X) = 1^T X$
- ⑥ sort-pooling: sorted by last column
- ⑦ set-to-set: introduced in [Vinyals&al.]²⁴

²⁴Vinyals, O., Bengio, S. Kudlur, M., Order Matters: Sequence to sequence for sets, ICLR 2016.

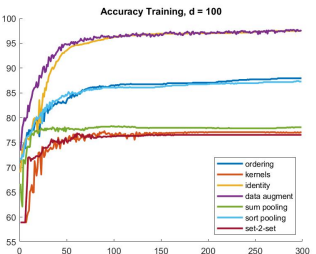
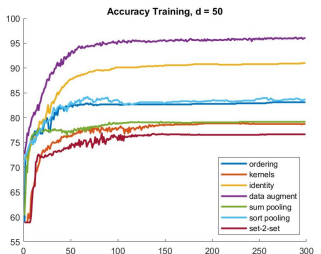
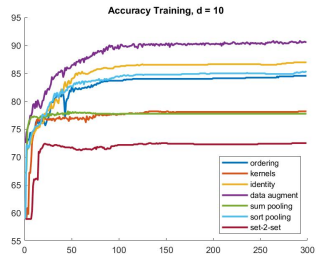
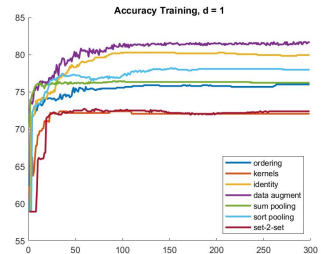
Enzyme Classification Example

Training Loss: X Entropy



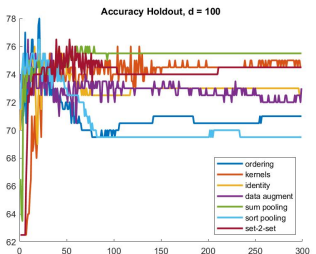
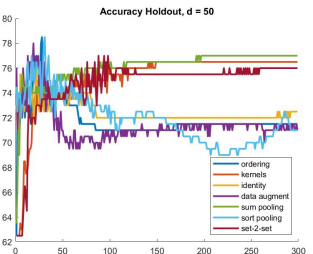
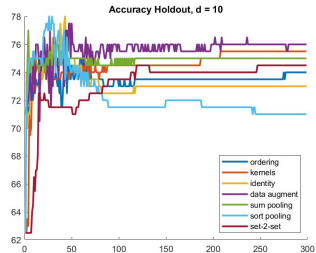
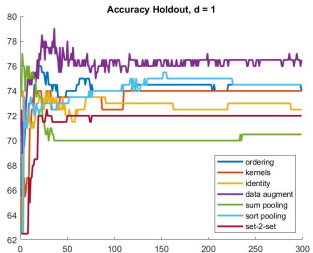
Enzyme Classification Example

Accuracy on Training set



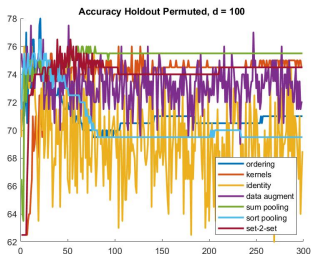
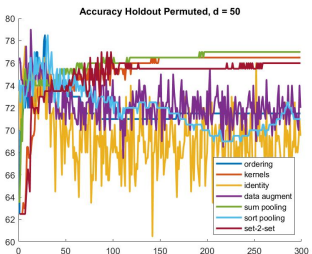
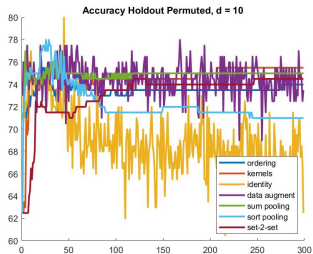
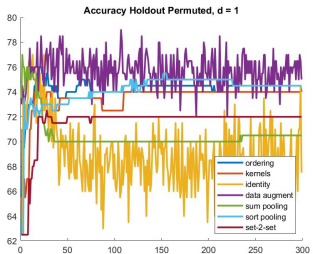
Enzyme Classification Example

Accuracy on Holdout data



Enzyme Classification Example

Accuracy on Holdout data with nodes randomly permuted



Performance Results: Accuracy

$d = 50$	ordering	kernels	identity	data augment	sum- pooling	sort- pooling	set-2- 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.]²⁵ obtains an accuracy of 77-80% using an SVM based classifier.

²⁵P.D. Dobson, A.J. Doig, "Distinguishing Enzyme Structures from Non-enzymes without Alignments", J. Mol. Biol. 330, 771-783, 2003.

The QM9 Dataset

Dataset: QM9²⁶ 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 contains 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 $\Delta\epsilon$) 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.

²⁶R. Ramakrishnan, P.O. Dral, M. Rupp, and O.A. von Lilienfeld. Quantum chemistry structures and properties of 134 kilo molecules. *Scientific data*, 1(1):1-7, 2014.

The Network

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

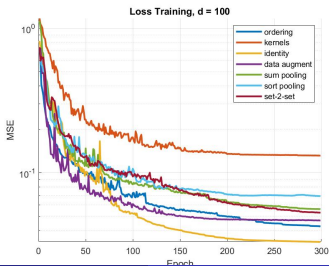
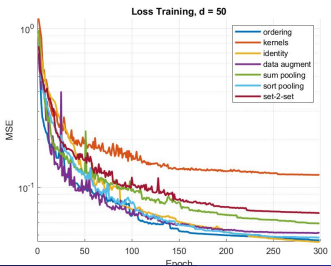
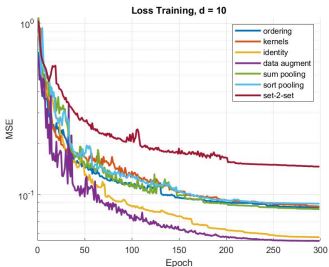
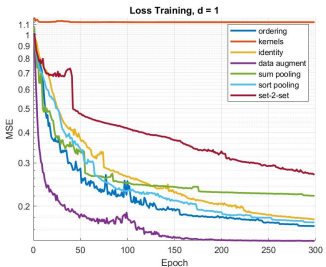
The same 7 α modules have been tested:

- 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 permutatons of each graph.
- ③ ordering: $\alpha(X) = \downarrow (XA)$, $A = [I \ 1]$
- ④ kernels: $\alpha(X) = (\sum_{k=1}^n \exp(-\|x_k - a_j\|^2))_{1 \leq j \leq m=5nd}$
- ⑤ sumpooling: $\alpha(X) = 1^T X$
- ⑥ sort-pooling: sorted by last column
- ⑦ set-to-set: introduced in [Vinyals&al.]²⁷

²⁷Vinyals, O., Bengio, S. Kudlur, M., Order Matters: Sequence to sequence for sets, ICLR 2016

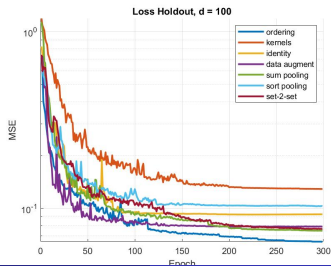
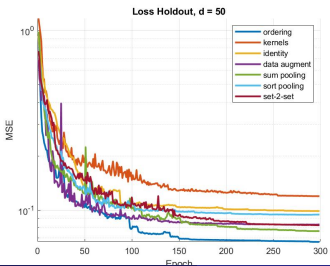
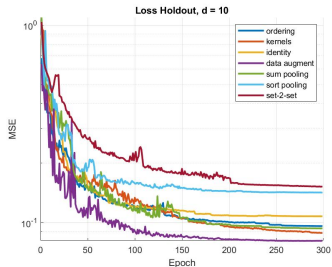
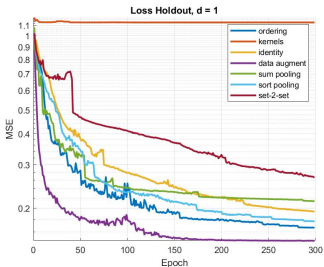
QM9 Regression Example

Training MSE



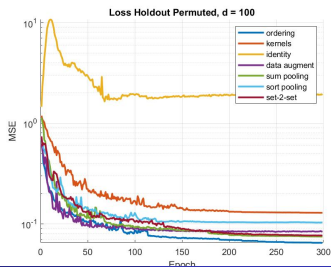
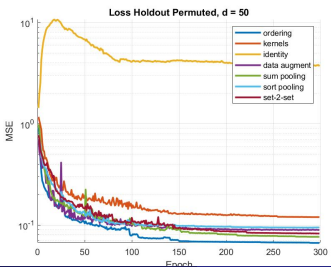
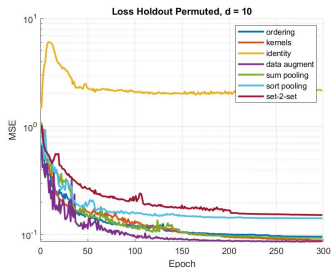
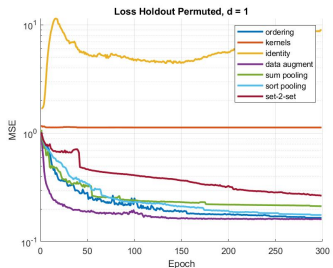
QM9 Regression Example

Validation MSE



QM9 Regression Example

Validation MSE with Random Permutations



Performance Results: MAE

$d = 100$	ordering	kernels	identity	data augment	sum- pooling	sort- pooling	set-2- 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

Thank you!
Questions?

Table of Contents

- 1 Problem Formulation
- 2 Motivation
- 3 Approach
- 4 Analysis Results
- 5 Numerical Examples
- 6 Extra**

A Universal Embedding

Consider the map

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

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!

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:

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

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Possible choices:

- ① 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.

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 $\Pi \in \mathcal{S}_n$ so that $Y = \Pi X$.

We denote by $\mathcal{A}_{d,D}(X)$ (or $\mathcal{A}(X)$) 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 $\mathcal{S}(A)$ denote the set of data matrices separated by A .

The key A is universal iff $\mathcal{S}(A) = \mathbb{R}^{n \times d}$.

Genericity Results for $d \geq 2$

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 $Y = [\Pi_1 x_1, \dots, \Pi_d x_d]$,

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

Define the linear operator

Genericity Results for $d \geq 2$

Admissible keys

Proof - cont'd

Let

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

Then

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

It is now sufficient to show that each null space has dimension less than d .
Indeed, the alternative would mean $B(\Xi; \Pi_1, \dots, \Pi_d) = 0$ but this would imply $(\Pi_1, \dots, \Pi_d) \notin \mathcal{P}$. \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 $d \geq 2$ and $n \geq 3$,

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

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 $A = [I_2 \ b]$.

Non-Universality of vector keys

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 $b = [1, 1]^T$.

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

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, $L^{-1}P^T A \in \mathcal{A}_{d,1}(XPL)$. 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 $d \geq 2$

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 $n \geq 1$, $\mathcal{S}([I_d \ a])$ 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 .

Genericity Results for $d \geq 2$

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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 $D > d$

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:

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

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

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

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

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

cont'd

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

$$\Delta = \{(\Xi_1, \dots, \Xi_{D-d}; \Pi_1, \dots, \Pi_d) \in (\mathcal{S}_n)^D, \Pi_1 = \Pi_2 = \dots = \Pi_d\}$$

$$M(\Pi_1, \Pi_1, \dots, \Pi_d) = 0 \rightarrow \bigcup_{\Pi \in \mathcal{S}_n} \ker M(\Pi, \Pi_1, \dots, \Pi_d) = \mathbb{R}^{n \times d}$$

2. If $(\Xi_1, \dots, \Xi_{D-d}; \Pi_1, \dots, \Pi_d) \in (\mathcal{S}_n)^D \setminus \Delta$ 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 $X = [0, \dots, 0, x_j, 0, \dots, 0]$ where x_j is the only non identically 0 column. Claim: $X \notin \ker L(\Xi_1, \dots, \Pi_d; A)$. Indeed, the resulting k column of $L()X$ is $A_{j,k}(\Xi_k - \Pi_j)x_j \neq 0$. It follows that

$$\dim \ker L(\Xi_1, \dots, \Xi_{D-d}; \Pi_1, \dots, \Pi_d; A) < nd$$

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. \square

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.

Planar Rotations

Consider the action on \mathbb{R}^2 of the cyclic group $\langle U \rangle \simeq \mathbb{Z}_N$ generated by a planar rotation by $a = \frac{2\pi}{N}$, $U = [\cos(a) \quad -\sin(a); \sin(a) \quad \cos(a)]$. The quotient space \mathbb{R}^2 / \sim is topologically equivalent to a 2 dim. cone in \mathbb{R}^3 , by identifying the positive x-semiaxis with the half-line of angle a .

Recall $\rho_n(q) = \max_{g,h} \gamma_{g,h}^q$ where

$\gamma_{g,h}^q = \text{semi.alg.dim.} \{(x,y) \in V \times V : \dim(\text{span}\{U_{gk}x - U_{hk}y, k \in [n]\}) = q\}$

Explicit computation:

$$\rho_1(q) = \begin{cases} 2, & q = 0, \\ 4, & q = 1, \\ -1, & q \geq 2. \end{cases} \quad \rho_2(q) = \begin{cases} 2, & q = 0, \\ 3, & q = 1 \text{ \& } N \text{ odd,} \\ 4, & q = 1 \text{ \& } N \text{ even,} \\ 4, & q = 2 \\ -1, & q \geq 3. \end{cases}$$

Planar Rotations (2)

Theorem

- 1 For any $w \in \mathbb{R}^2$, the map $\Phi_w : \widehat{\mathbb{R}^2} \rightarrow \mathbb{R}^N$ is never injective.
- 2 For any $w_1, w_2 \in \mathbb{R}^2$ and $S = \{(1, 1), (1, 2)\}$ (the max filter case), the map $\Phi_{w,S} : \widehat{\mathbb{R}^2} \rightarrow \mathbb{R}^2$ is never injective.
- 3 If $w_1, w_2, w_3 \in \mathbb{R}^2$ are linearly independent and $\text{angle}(w_i, w_j) < \frac{2\pi}{N}$ then for either $S = \{(1, 1), (1, 2), (1, 3)\}$ (the max filter, or $\mathbf{n} = (1, 1, 1)$ configuration), or $S = \{(1, 1), (2, 1), (1, 2)\}$ (a $\mathbf{n} = (2, 1)$ configuration), generically, the map $\Phi_{w,S} : \widehat{\mathbb{R}^2} \rightarrow \mathbb{R}^3$ is injective and bi-Lipschitz.

A careful analysis of our main theorem would guarantee the embedding $\Phi_{w,S} : \widehat{\mathbb{R}^2} \rightarrow \mathbb{R}^4$ is injective (and hence bi-Lipschitz) for certain $\mathbf{n} = (2, 1, 1)$ configurations, or any $\mathbf{n} = (1, 1, 1, 1)$ configuration.

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