

On Space Folds by Neural Networks

Michał Lewandowski

16.10.2025

Co-authors



Hamid
(Meta)



Raphael
(SCCH)



Bernhard
(SCCH)

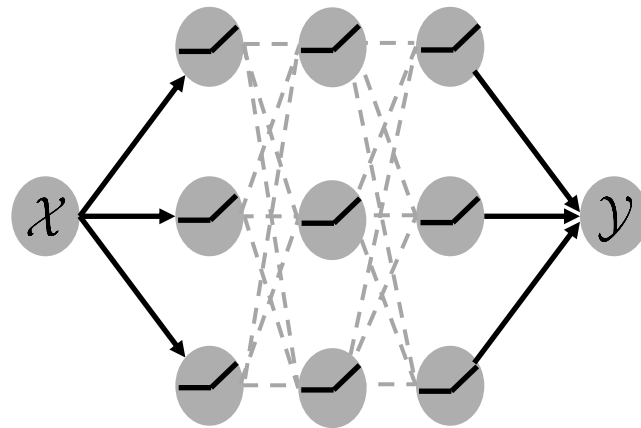


Bernhard
(JKU & SCCH)

- **SCCH**: a Research Center in Hagenberg, Austria, focused on applied AI and software sciences
- **S3AI**: a research project focused on geometry of neural networks (2020-2024)

Introduction

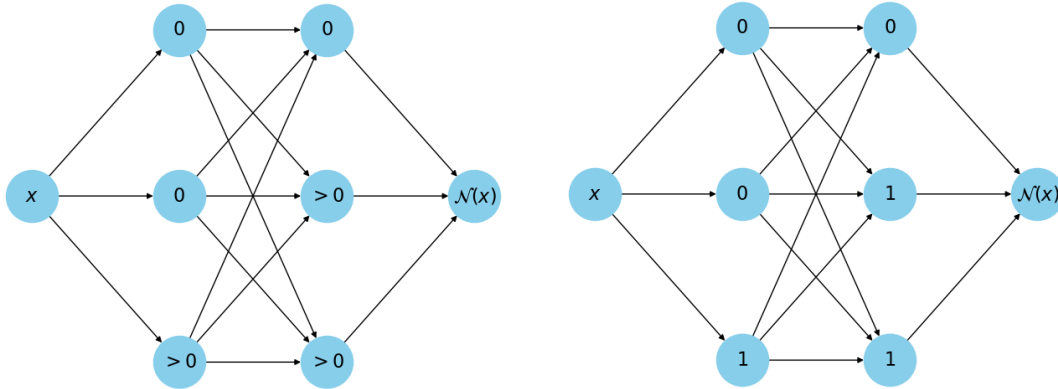
- **Focus:** Analysis of the **phenomena of folding** through a novel measure
- **How:** We focus on fully connected neural networks (MLPs)
- **Goal:** Gaining theoretical insights – not optimizing real-world performance
 - ✓ Universal approximation and exponential expressiveness
 - ✓ Piece-wise linear functions - inherit a geometrical interpretation



(Reproduced from [1])

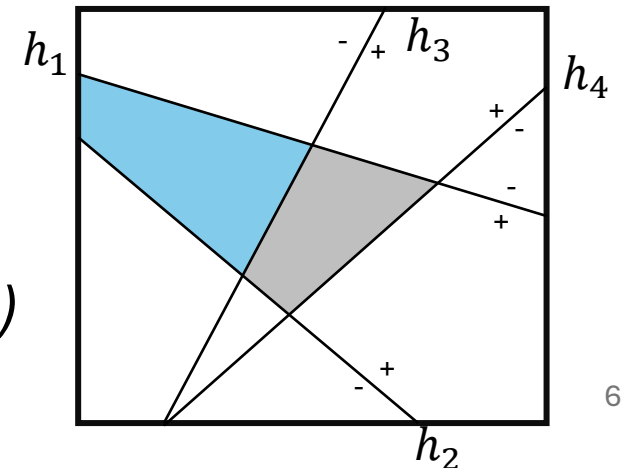
Hamming Activation Space

- A ReLU NN (denoted \mathcal{N}) is an alternating composition of the ReLU ($\sigma(x) := \max(x, 0)$) and affine functions $g_i(\mathbf{x}) := W_i \mathbf{x} + b_i$
- Fix $(W_i, b_i)_{i=1}^L$; push an input \mathbf{x} through \mathcal{N} ; at every layer we obtain a vector of activation values



- Activation pattern: $\pi_1 = (001011)$
- For any $\mathbf{x}_i \in \mathcal{X}$, there is an associated activation pattern π_i
- By construction, each (observable) π_i is a solution $\{\mathbf{x}: h_1(\mathbf{x}) \geq 0 \ \& \ \dots \ \& \ h_4(\mathbf{x}) \geq 0\}$
- In geometry: a polytope [1], in ML: a linear region [2]

- Hamming distance d_H quantifies difference between $\pi_i, \pi_j \in \{0,1\}^N : d_H(\pi_i, \pi_j) := |\{i: \pi_{i,k} \neq \pi_{j,k}\}|$
- $(\{0,1\}^N, d_H)$ is a metric space (*The Hamming Activation Space*)



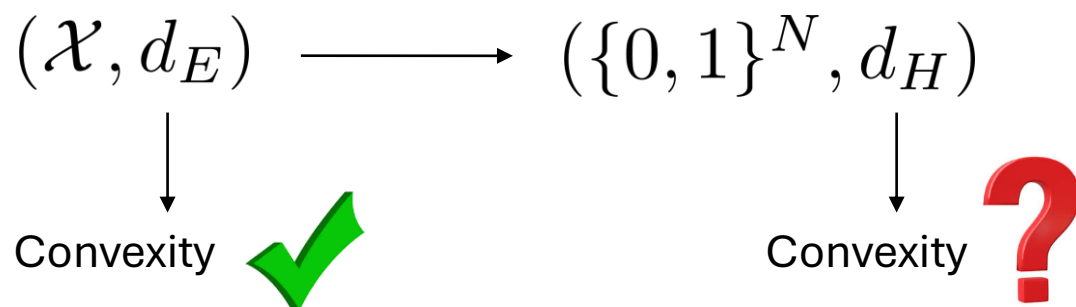
[1] Ziegler, G. M. (2000). *Lectures on 0/1-polytopes*. Polytopes—combinatorics and computation.

[2] Montufar et al. (2014). *On the Number of Linear Regions of Deep Neural Networks*. NeurIPS.

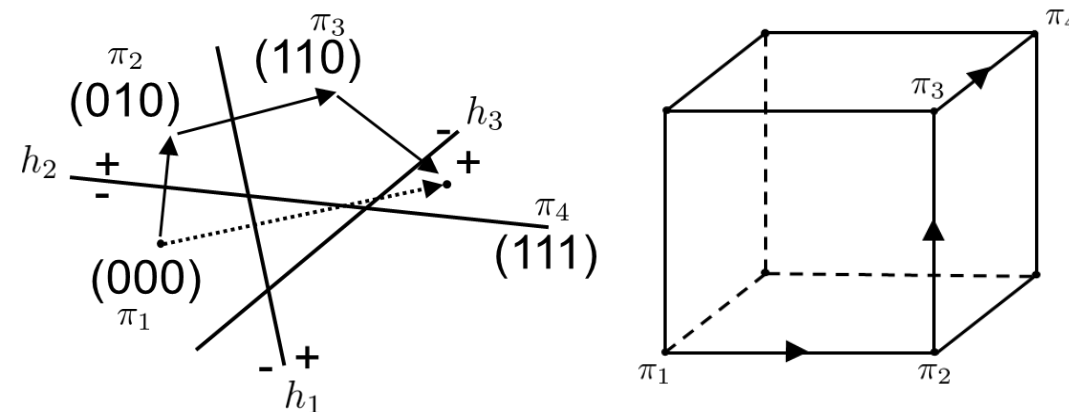
Analysis of ReLU-Induced Tessellations

The Hamming activation space contains every $\pi \in \{0,1\}^N$.
BUT not every π is a pre-image of a linear region!

Impacts the definition of convexity
in the Hamming space [1, 2]

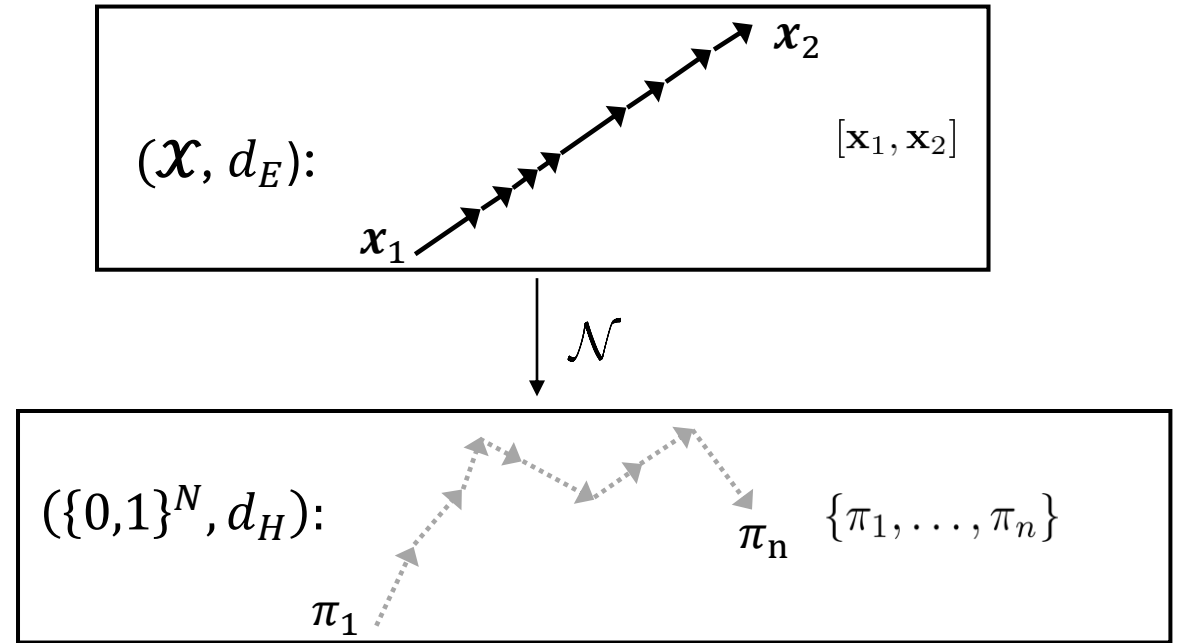
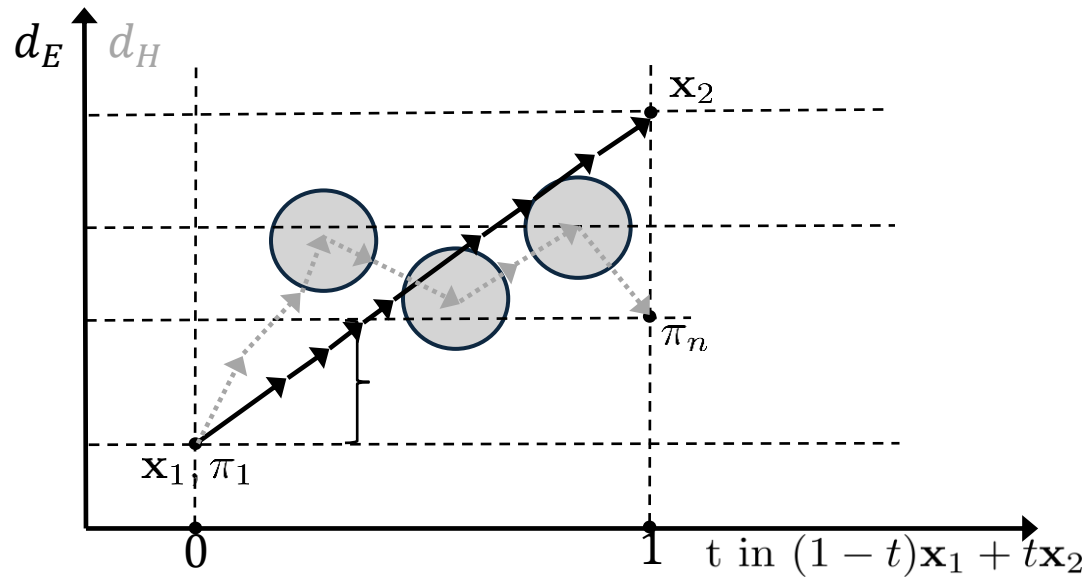


Definition 2. A subset S of the Hamming cube H^N is convex if, for every pair of points $\pi_i, \pi_j \in S$, all (observable) points on every shortest path between π_i, π_j are also in S .



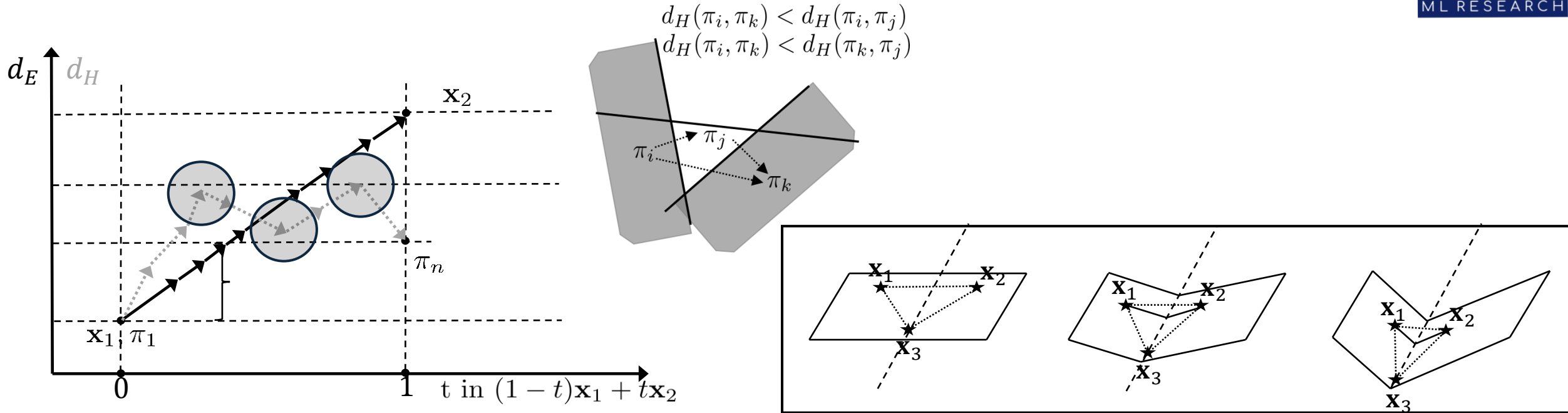
Lemma 1: Given a space partition into regions $R_{\pi_1}, \dots, R_{\pi_r}$ labelled by $A = \{\pi_1, \dots, \pi_r\}$, a union $R = \bigcup_{\pi \in S \subset A} R_{\pi}$ is convex in R^n iff A is convex in H^N .

Mappings of Paths: Connection to Convexity



- $[\mathbf{x}_1, \mathbf{x}_2]$ - straight line in (\mathcal{X}, d_E) – the smallest convex “set”
- We *map* $[\mathbf{x}_1, \mathbf{x}_2]$ from (\mathcal{X}, d_E) to $(\{0,1\}^N, d_H)$ obtaining $\{\pi_1, \dots, \pi_n\}$
- But in $(\{0,1\}^N, d_H)$ not every shortest path γ (under d_H) between π_1 and π_n contains all $\{\pi_1, \dots, \pi_n\}$, i. e., $\gamma \neq \{\pi_1, \dots, \pi_n\} \Rightarrow \{\pi_1, \dots, \pi_n\}$ is not convex (cf. Def 2)
- By investigating γ we can measure *deviations* from convexity between \mathbf{x}_1 and \mathbf{x}_2

Mappings of Paths: Connection to Folding



- On a path $\Gamma = \{\pi_1, \dots, \pi_n\}$, we monitor two *range measures*:
 - r_1 : Max change in d_H at each step i wrt π_1 : $r_1(\Gamma) := \max_i d_H(\pi_1, \pi_i)$
 - r_2 : Total travelled distance on the hypercube: $r_2(\Gamma) := \sum_{i=1}^{n-1} d_H(\pi_i, \pi_{i+1})$

Local Folding

$$\chi(\Gamma) := 1 - \frac{r_1(\Gamma)}{r_2(\Gamma)} \in [0,1]$$

Global Folding

$$\Phi_{\mathcal{N}} := \text{median}(\{\chi(\Gamma) : \chi(\Gamma) > 0\})$$

Alternatively, map $\sum_{i,j} \alpha_j \mathbf{x}_i$ through \mathcal{N} for $\forall i, j$

Space Folding Measure – Properties

Edge cases:

- $\chi(\Gamma) = 0$ if $d_H(\pi_1, \pi_i)$ increases monotonically.
- $\chi(\Gamma) = 1$ for a looped path: $\max_i d_H(\pi_1, \pi_i) = c$, while $\sum_{i=1}^{n-1} d_H(\pi_i, \pi_{i+1}) \rightarrow \infty$.

The space folding measure has the following properties:

1. **[Stability]** Multiple steps in the same linear region do not influence its values.
2. **[Asymmetry]** The folding measure is sensitive to the direction of path traversal, i.e., $\chi(\Gamma) \neq \chi(-\Gamma)$, where $-\Gamma = \{\pi_n, \dots, \pi_1\}$.
3. **[Flatness Invariance]** $\chi(\Gamma) = 0$ if and only if $\chi(-\Gamma) = 0$.
4. **[Non-additivity]** Is neither sub- nor super-additive.

Let $\Gamma = \Gamma(\mathbf{x}_1, \mathbf{x}_2)$ be a path spanned between the edge points $\mathbf{x}_1, \mathbf{x}_2$.

Proposition. Let d_χ be a symmetrized space folding measure,

$$d_\chi(\mathbf{x}_1, \mathbf{x}_2) = \frac{1}{2} \left(\chi(\Gamma(\mathbf{x}_1, \mathbf{x}_2)) + \chi(\Gamma(\mathbf{x}_2, \mathbf{x}_1)) \right).$$

Then, d_χ is a pseudo-metric: **(1) Positivity:** from bounds on χ ; **(2) Symmetry:** from construction; **(3) Triangle inequality:** from the triangle inequality of d_H

Algorithm & Complexity

Algorithm 1: Computation of the Space Folding Measure

Input: Two input samples $\mathbf{x}_1, \mathbf{x}_2$, the number of intermediate points n , the total number of hidden neurons N , cost of running the network in the inference mode $O(C)$

Output: Space Folding $\chi(\Gamma)$

Step 1: Linearly interpolate \mathbf{x}_1 and \mathbf{x}_2 , sampling n points; // Sampling Complexity: $O(n)$

Step 2: For each sampled point:

begin

 Compute the binarisation; // Binarization Complexity Per Point: $O(C)$

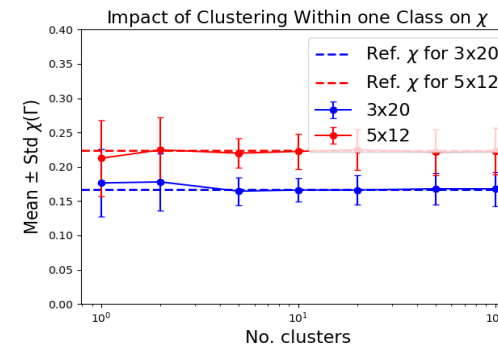
// Total Binarization Complexity: $O(n \cdot C)$

Step 3: Compute the maximal (from the starting point) and total Hamming distances between intermediate points; // Computation of Range Measures Complexity: $O(n \cdot N)$

return Space Folding $\chi(\Gamma)$; // Total Algorithm Complexity: $O(n \cdot (N + C))$

Computational complexity: $O(n \cdot (N + \text{Cost of Sample Propagation}) \cdot |C_1| \cdot |C_2|)$

1. For a given the dataset, $D = \{(\mathbf{x}_i, y_i)\}_{i=1}^m$
2. Cluster data samples \mathbf{x}_i of the same label
3. Use clusters' centroids for computation of χ



Space Folding - Caveats

- For an activation function f , we considered its pre-image thresholded at 0, i.e., $f^{(-1)}((-\infty, 0]) \rightarrow 0$ and $f^{(-1)}((0, \infty)) \rightarrow 1$
- Why thresholding at 0?
 - consider thresholds $a, b \in (-\infty, \infty)$ in the range of the activation function f such that $|a| < |b|$,
 - $\#a$ -induced regions $\geq \#b$ -induced regions $\Rightarrow a$ -induced folding measure $\geq b$ -induced folding measure;
 - for ReLU $a = 0$ is "more informative" than any $b > 0$

Space Folding - Beyond ReLU

- Space Folding computation is based on a walk traversing linear regions
- The computation can be extended to a walk traversing equivalence classes [1,2]:
$$\mathbf{x}_1 \sim_{\mathcal{N}} \mathbf{x}_2 \Leftrightarrow d_H(\pi_1, \pi_2) = 0$$
- For ReLU NN, $[\mathbf{x}]_{\mathcal{N}} := \{\mathbf{z} : \mathbf{z} \sim_{\mathcal{N}} \mathbf{x}\}$ correspond to linear regions
- For non-monotonous f equivalence classes may be topologically disconnected, but the construction applies as is

[1] N. Shepeleva, et al. (2020). *Relu code space: a basis for rating network quality besides accuracy*. ICLR W.

[2] M. Lewandowski, et al. (2025). *The Space Between: On Folding, Symmetries and Sampling*. ICLR W.

Summary

- Same output function yet different architectures?
 - Studied in [1] based on CantorNet [2]; higher folding values for less Kolmogorov-complex representation

The question:

Given a space folding value $\chi(\Gamma) = \tau \in [0, 1]$ for some path Γ , what can we learn?

- χ can be seen as a feature of the network
- χ is upper- and lower-bounded - a reference point across neural networks
- **Interpretation:**
 - Higher $\chi \rightarrow$ indicates a more compact representation
 - Lower $\chi \rightarrow$ indicates potential architecture improvements for the task

[1] M.Lewandowski et al. (2025). On Space Folds of Neural Networks. TMLR.

[2] M.Lewandowski et al. (2024). *CantorNet: A Sandbox For Testing Topological and Geometrical Complexity Measures*. NeurIPS₁₄ Workshop.

Space Folding and Generalization

- Folding values increase with the depth of the network *conditioned on* high validation accuracy
- On CIFAR100, we observed a steady increase in global folding during the training process

Beyond fully connected networks

- Consider a **subset of layers** – different underlying tessellation
- **Skip Connections** – applies as is
- **Transformer** – fcnn after the attention head
- **Convolutional layers** – after convolution and pooling

Low sensitivity to batch-norm, dropout

Convexity in the Hamming Activation Space

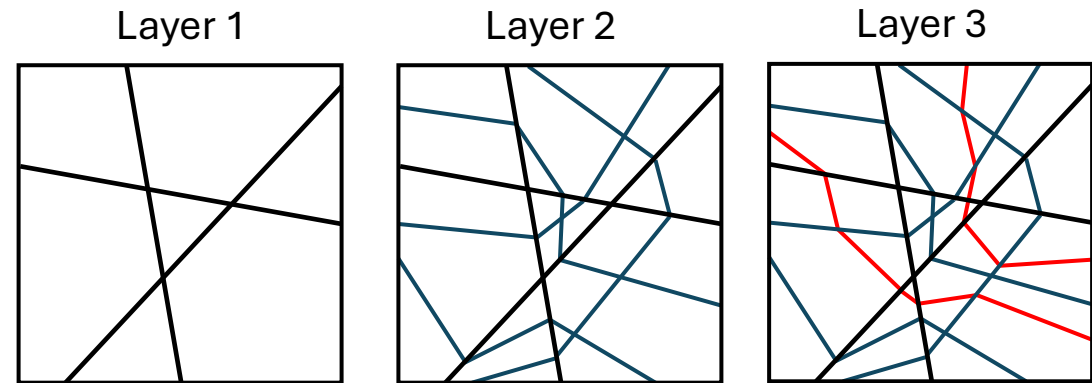
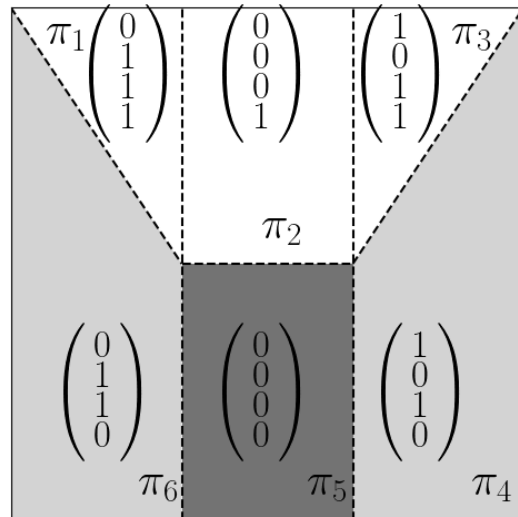
Example 1. Consider activation patterns $\pi_1 = (0111), \pi_2 = (0001), \pi_3 = (1011)$. For a walk (1) $\pi_1 \rightarrow \pi_2$, (2) $\pi_2 \rightarrow \pi_3$, (3) $\pi_1 \rightarrow \pi_3$, there are intermediate, non-observable activation patterns π_{inter} that we traverse:

(1) $\pi_{\text{inter}} = \{(0011), (0101)\}$

(2) $\pi_{\text{inter}} = \{(0011), (1001)\}$

(3) $\pi_{\text{inter}} = \{(0011), (1111)\}$

Thus, the activation patterns $\{\pi_1, \pi_2, \pi_3\}$ form a convex set in the Hamming cube sense.



(Adapted from [1])

[1] M. Raighu et al. (2017). *On the expressive power of deep neural networks*. ICML.

[2] M. Lewandowski et al. (2024). *CantorNet: A Sandbox for Testing Topological and Geometrical Complexity Measures*. NeurIPS Workshop.

Interaction Coefficient and Adversarial Examples

χ is neither sub- nor super-additive - define *interaction coefficient*

$$I(k) := |\chi(\Gamma_1 \oplus \Gamma_2) - \chi(\Gamma_1) - \chi(\Gamma_2)|,$$

where $k \in \mathbb{N}$ is the connecting index of paths Γ_1 and Γ_2 defined as

$$\Gamma_1 = \{\pi_1, \dots, \pi_k\}, \Gamma_2 = \{\pi_k, \dots, \pi_n\}, \Gamma_1 \oplus \Gamma_2 = \{\pi_1, \dots, \pi_n\}.$$

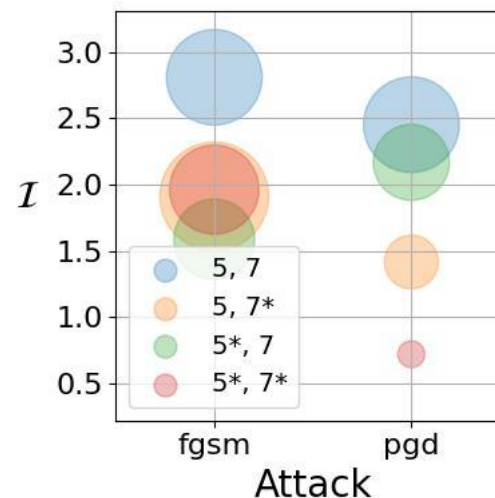
- I computed using digits 5 and 7 from the MNIST test set; * denotes adversarial perturbation
- I is consistently higher for unperturbed samples

Algorithm 1: \mathcal{I} and Adversarial Attacks

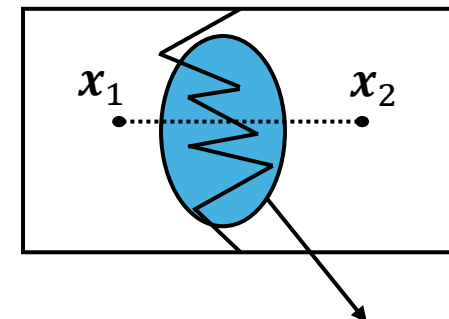
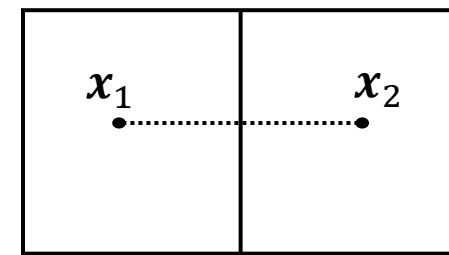
Input: Dataset $\{(\mathbf{x}_i, y_i)\}_{i \in I}$

Output: Mean non-zero \mathcal{I}

- 1: **Step 1:** Adversarially perturb the input \mathbf{x} obtaining \mathbf{x}^* .
- 2: **Step 2:** Assert that $\mathcal{N}(\mathbf{x}) \neq \mathcal{N}(\mathbf{x}^*)$.
- 3: **Step 3:** Compute \mathcal{I} on a path spanned between $[\mathbf{x}, \mathbf{x}^*]$
- 4: (a) check the linear combination of \mathbf{x} and \mathbf{x}^* for varying connecting index k .
- 5: (b) store k s.t. \mathcal{I} increases step-wise
- 6: **return** $\frac{1}{n-k+1} \sum_{i>k} \mathcal{I}_i$



Intuition: folding and adversarial geometry



We can measure this!

Equivalence of Convexity Notions: Sketch of the Proof

Convexity in $R_n \Rightarrow$ Convexity in the Hamming space

- Consider convex $R = \bigcup_{\pi \in A} R_\pi$ in R^n .
- Connectivity of A : take π_i, π_j and some points $P \in R_{\pi_i}; Q \in R_{\pi_j}$
- $R = \bigcup_{\pi \in A} R_\pi$ is convex $\Rightarrow [P, Q]$ lies entirely within R .
 - Along $[P, Q]$ we cross h_1, \dots, h_l flipping 1-bit in activation patterns at a time.
 - Sequence of flips forms a shortest path in the Hamming space from π_i to π_j .
 - If A wasn't convex, there would exist γ connecting π_i with π_j and leaving A .
 - But $[P, Q]$ is a straight line - it corresponds to the minimal sequence of bit flips.

