# Deep Learning - Parameters and Functions Mildly Overparametrized ReLU Nets

Guido Montúfar montufar@math.ucla.edu

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#### Kedar Karhadkar



### Michael Murray



#### Hanna Tseran



• "Mildly Overparameterized ReLU Networks Have a Favorable Loss Landscape"



Parametric model



Parametric model and input data set



Parametric model over the input data set



Function space over the input data set

 Neural networks have a non-convex loss landscape with local minima and plateaus<sup>1</sup>.

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- We can avoid excessive overparameterization by emphasizing qualitative aspects of the loss landscape, using only the rank of the Jacobian rather than e.g. the smallest eigenvalue of the NTK.
- We obtain theorems under more realistic mild overparameterization  $d_1 = \Omega(n \log n)$  or even  $d_1 = \Omega(1)$  for high-dimensional inputs.

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- Theorem 2: If  $d_0 d_1 \ge n$  and  $d_1 = \Omega(\log(\frac{n}{\epsilon d_0}))$ , then all activation regions, except for an  $\epsilon$  fraction, have no bad local minima.
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- Theorem 7: If d<sub>0</sub> = 1 and d<sub>1</sub> = Ω(n log(<sup>n</sup>/<sub>ε</sub>)), all but at most an ε fraction of *non-empty* activation regions have no bad local minima.

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- Theorem 12 provide bounds on the fraction of regions with bad local minima by volume.

### Setup

• We consider input and output data

$$X = (x^{(1)}, \ldots, x^{(n)}) \in \mathbb{R}^{d \times n}, \quad y = (y^{(1)}, \ldots, y^{(n)}) \in \mathbb{R}^{1 \times n}.$$

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• We consider a parameterized model

F	:	$\mathbb{R}^{m}$	X	$\mathbb{R}^{d}$	$\rightarrow$	$\mathbb R$
		parameter		input		prediction

and the vector of predictions on input data X,

$$F(\theta, X) := (F(\theta, x^{(1)}), F(\theta, x^{(2)}), \dots, F(\theta, x^{(n)})).$$

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• The mean squared error loss  $L: \mathbb{R}^m_{parameter} \times \mathbb{R}^{d \times n} \times \mathbb{R}^{1 \times n} \to \mathbb{R}^1$ ,

$$L(\theta, X, y) := \frac{1}{2} \sum_{i=1}^{n} (F(\theta, x^{(i)}) - y^{(i)})^{2}.$$
(1)

Lemma 1 (Full rank Jacobian implies critical point is global min) Fix a dataset  $(X, y) \in \mathbb{R}^{d \times n} \times \mathbb{R}^{1 \times n}$ , a parametrized model F, and a differentiable critical point  $\theta \in \mathbb{R}^m$  of the squared error loss (1).

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Proof.

$$0 = \nabla_{\theta} L(\theta, X, y) = \underbrace{\nabla_{\theta} F(\theta, X)}_{\mathsf{rank} = n} \cdot \underbrace{(F(\theta, X) - y)}_{= 0}$$

### ReLU network

• E.g., a two-layer ReLU network  $F : \mathbb{R}^{d_1 \times d_0}_{parameter} \times \mathbb{R}^{d_0} \to \mathbb{R}_{prediction}$ 

$$F(W, x) = v^T \sigma(W x),$$

where  $\sigma: s \mapsto \max\{0, s\}$  componentwise, and  $v \in \mathbb{R}^{d_1}$ .

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- To accommodate a bias, we can add a 1 component to x.
- This map is piecewise polynomial in W, v and piecewise linear in x.

### Activation regions and Jacobian

- For data X, the smooth pieces are separated by  $\langle w^{(i)}, x^{(j)} \rangle = 0$ .
- For each  $A = [a^{(1)}, \dots, a^{(n)}] \in \{0, 1\}^{d_1 \times n}$  define activation region

$$\mathcal{S}_X^A := \left\{ W \in \mathbb{R}^{d_1 \times d_0} : (2A_{ij} - 1) \langle w^{(i)}, x^{(j)} \rangle > 0 \ \forall i \in [d_1], j \in [n] \right\}.$$

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The Jacobian of the vector of predictions is

$$abla_ heta F(W,X) = [(v \odot a^{(j)}) \otimes x^{(j)}]_j, \quad orall W \in \mathcal{S}_X^A, \quad orall A.$$

• Similar definitions for deep ReLU nets.

## Subdivision of parameter space



Figure 1: Fan of activation regions; activation patterns indicate the input data points on which each unit is active.

## Activation regions with no bad local minima

Theorem 2 (Most activation regions are good) Let  $\epsilon > 0$ . If  $d_1 \ge \max\left(\frac{n}{d_0}, \Omega\left(\log\left(\frac{n}{\epsilon d_0}\right)\right)\right)$ ,

then for generic datasets (X, y), the following holds.

In all but at most an  $\epsilon$  fraction of all activation regions (i.e. at most  $\lceil \epsilon 2^{d_1} \rceil$ ), every differentiable critical point of L is a global minimum.

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Uses an upper bound on probability that a binary random matrix is singular. Caveat: This refers to all activation regions, empty or non-empty.

## Non-empty activation regions

## Subdivision of parameter space

### Proposition 3 (Number of non-empty regions)

Consider a network with one layer of  $d_1$  ReLUs. If the columns of X are in general position in a d-dimensional linear space, then the number of non-empty activation regions in the parameter space is  $(2\sum_{k=0}^{d-1} \binom{n-1}{k})^{d_1}$ .

Regions of a product central hyperplane arrangement.

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Regions of a product central hyperplane arrangement.

Proposition 4 (Identity of non-empty regions) Let  $A \in \{0, 1\}^{d_1 \times n}$ . The corresponding activation region is non-empty if and only if  $\sum_{j:A_{ij}=1} x^{(j)}$  is a vertex of  $\sum_{j \in [n]} \operatorname{conv}\{0, x^{(j)}\}$  for all  $i \in [d_1]$ .

Combination of covectors of the oriented matroid of the input data.



Figure 2: The polytope *P* of a ReLU on data points  $x^{(1)}, x^{(2)}, x^{(3)}$  is the Minkowski sum of the line segments  $P_i = \text{conv}\{0, x^{(i)}\}$ . The activation regions are the normal cones of *P*. The vertices of *P* correspond to the non-empty activation regions.

## High-dimensional input

For high-dimensional inputs, most activation regions are non-empty, thus:

Corollary 5 (Most non-empty activation regions are good)

Under the same assumptions as Theorem 2, if  $d \ge n$ , then for X in general position and arbitrary y:

In all but at most an  $\epsilon$  fraction of all non-empty activation regions, every differentiable critical point of L is a zero loss global minimum.

Non-empty activation regions with no bad local minima

For 1D input, we can explicitly list the non-empty activation regions. Lemma 6 (Non-empty activation regions for 1D data) Fix a dataset (X, y) with  $x^{(1)} < x^{(2)} < \cdots < x^{(n)}$ . Let  $A \in \{0, 1\}^{d_1 \times n}$ . Then  $S_X^A$  is non-empty if and only if the rows of A are step vectors. In particular, there are exactly  $(2n)^{d_1}$  non-empty activation regions. Theorem 7 (Most non-empty activation regions are good) Let  $\epsilon \in (0, 1)$ . Suppose that X consists of distinct data points, and

$$d_1 \ge 2n \log\left(\frac{n}{\epsilon}\right).$$

Then in all but at most an  $\epsilon$  fraction of non-empty activation regions,  $\nabla_{\theta} F$  is full rank and every differentiable critical point of L is a global minimum.

Theorem 8 (Fraction of regions with global minima) Let  $\epsilon \in (0, 1)$ . Suppose that X consists of distinct data points, and

$$|\{i \in [d_1]: v^{(i)} > 0\}| \ge 2n \log\left(\frac{2n}{\epsilon}\right),$$

and

$$|\{i \in [d_1] : v^{(i)} < 0\}| \ge 2n \log\left(\frac{2n}{\epsilon}\right).$$

Then in all but at most an  $\epsilon$  fraction of non-empty activation regions  $S_X^A$ , the subset of global minimizers  $\mathcal{G}_{X,y} \cap \mathcal{S}_X^A$  is a non-empty affine set of codimension n. Moreover, all global minima of L have zero loss.

## Nonsmooth critical points

We can extend the analysis to handle points on the boundaries between regions, where the loss is non-differentiable.

Theorem 9 Let  $\epsilon \in (0, 1)$ . If  $d_1 \ge 2n \log\left(\frac{n}{\epsilon}\right)$ ,

then in all but at most a fraction  $\epsilon$  of non-empty activation regions A, every local minimum of L in  $\mathcal{S}_{X}^{A} \times \mathbb{R}^{d_{1}}$  is a global minimum.

### Extension to deep nets

• For  $l \in \{0, \dots, L\}$ , we define the *l*-th layer  $f_l \colon \mathbb{R}^m \times \mathbb{R}^{d_{l-1}} \to \mathbb{R}^{d_l}$  by

$$\begin{split} & f_0(W, x) := x, \\ & f_l(W, x) := \sigma(W_l f_{l-1}(\theta, x)) & \text{if } l \in [L-1], \text{ and} \\ & f_L(W, x) := v^T f_{l-1}(\theta, x), \end{split}$$

where  $v \in \mathbb{R}^{d_{l-1}}$  is a fixed vector whose entries are nonzero.

• The activation patterns of a deep network are given by tuples

$$A=(A_1,A_2,\ldots,A_{L-1}),$$

where for each  $l \in [L-1]$ ,  $A_l \in \{0,1\}^{d_l \times n}$ .

• Denote  $\mathcal{S}_X^A$  subset of parameters with activation pattern A.

Theorem 10 Let  $X \in \mathbb{R}^{d_0 \times n}$  be an input dataset with distinct points. Suppose that for all  $l \in [L-2]$ ,

$$d_l = \Omega\left(\log\frac{n}{\epsilon L}\right),\,$$

and that

$$d_{L-1} = n + \Omega\left(\log \frac{1}{\epsilon}\right).$$

Then for at least a  $(1 - \epsilon)$  fraction of all activation patterns A, the following holds. For all  $W \in S_X^A$ ,  $\nabla_W F(W, X)$  has rank n.

## Volumes of activation regions

## One-dimensional input data

We bound the volume of activation regions with full rank Jacobian in terms of the amount of separation between the data points.

#### Proposition 11

Let  $n \ge 2$ . Suppose the entries of v are nonzero. Suppose that for all  $j, k \in [n]$  with  $j \ne k$ , we have  $|x^{(j)}| \le 1$  and  $|x^{(j)} - x^{(k)}| \ge \phi$ . If

$$d_1 \geq rac{4}{\phi} \log\left(rac{n}{\epsilon}
ight),$$

then, writing  $\mu$  for the Lebesgue measure,

 $\mu(\cup\{\mathcal{S}_X^{\mathcal{A}}\cap [-1,1]^{d_1}\times [-1,1]^{d_1}:\nabla_{w,b}\mathsf{F}\text{ has full rank on }\mathcal{S}_X^{\mathcal{A}}\})\geq (1-\epsilon)2^{2d_1}.$ 

### Arbitrary dimension input data

We say that an input dataset  $X \in \mathbb{R}^{d_0 \times n}$  is  $\gamma$ -anticoncentrated if for all nonzero  $u \in \mathbb{R}^n$ ,  $\mathbb{P}_{a \sim \mathcal{D}_X}(u^T a = 0) \leq 1 - \gamma$ . We can interpret this as a condition on the amount of separation between data points.

Theorem 12 Let  $\epsilon, \gamma \in (0, 1)$ . Suppose that  $X \in \mathbb{R}^{d_0 \times n}$  is generic and  $\gamma$ -anticoncentrated. If

$$d_1 \geq rac{8}{\gamma^2} \log\left(rac{d_0}{\epsilon}
ight) + rac{2}{\gamma} \left(rac{n}{d_0} + 1
ight),$$

then with probability at least  $1 - \epsilon$ ,  $\nabla_{(W,v)}F(W, v, X)$  has rank n.

# Experiments

## Probability of full rank Jacobian for random init



Figure 3: Input dimension  $d_0$  is left fixed. Minimum  $d_1$  to achieve full rank linear in *n*, slope decreases as  $d_0$  increases, as predicted by Theorem 2.

## Probability of full rank Jacobian for random init



Figure 3: Input dimension  $d_0$  scales linearly in the number of samples *n*. Minimum  $d_1$  to achieve full rank remains constant in *n*, consistent with Theorem 2.

## Percentage of regions with global min, $d_0 = 1$



Figure 4: Percentage of randomly sampled activation regions that contain a global minimum of the loss for networks with  $d_0 = 1$ . Black line is Theorem 8.

## Percentage of regions with global min, $d_0 = 2, 5$



Figure 5: Percentage of randomly sampled activation regions that contain a global minimum for networks with input dimension  $d_0 = 2$  (top) and  $d_0 = 5$  (bottom). Consistent with Theorem 2 and Corollary 5.

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# Function space on 1D data

Proposition 13 (Function space on one-dimensional data) Let X be a list of n distinct points in  $1 \times \mathbb{R}$  with  $x^{(1)} < x^{(2)} < \cdots < x^{(n)}$ . Let  $\overline{x}^{(i)} = [x_2^{(i)}, -1]$  and  $X_{\geq i} = [0, \dots, 0, x^{(i)}, \dots, x^{(n)}]$ .

• Then the functions a ReLU represents on X form a polyhedral cone,  $\alpha f \in \mathbb{R}^n$  with  $\alpha \ge 0$  and f in the polyline with vertices

$$ar{x}^{(i)}X_{\leq i}, \ i=1,\ldots,n$$
 and  $-ar{x}^{(i)}X_{\geq i}, \ i=1,\ldots,n.$  (2)

- A sum of m ReLUs represents non-negative scalar multiples of convex combinations of any m points on this polyline.
- Arbitrary linear combinations of m ReLUs represent scalar multiples of affine combinations of any m points on this polyline.

### Function space



Figure 6: Function space of a ReLU on *n* data points in  $1 \times \mathbb{R}$ , for n = 3, 4.

#### Summary

- We studied the loss landscape of two-layer ReLU networks in the mildly overparameterized regime.
- Most activation regions have no bad differentiable local minima.
- In the univariate case, most non-empty activation regions contain a high-dimensional set of global minimizers.

#### Further topics

Gradient descent.

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