# Theoretical Justification of Deep Learning

Generalization and Optimization

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### Agenda

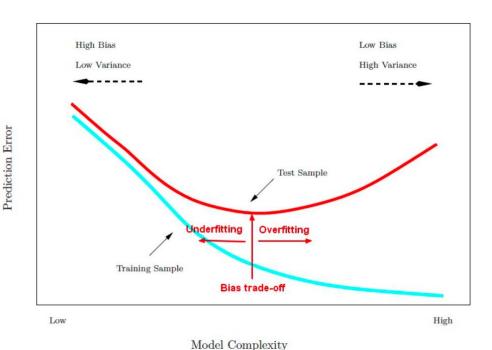
- Standard/classic statistical learning theory (SLT)
- Double Descent and limitations of classic SLT
- Benign Overfitting in interpolating shallow models
  - Linear Models
  - Kernel Methods
  - Boosting
  - Random Forests
- Theoretical Justifications of linear interpolators
- Linearization of Neural Networks
- Optimization: SGD Implicit Bias
- References

### Classic/Standard Statistical Learning Theory

	Finite ${\cal H}$	Infinite ${\cal H}$
Realizable	$\mathfrak{R}(g) \leq rac{1}{N}ig[\log  \mathcal{H}  + \log rac{1}{\delta}ig]$	$\mathfrak{R}(g) < rac{\mathit{VC}(\mathcal{H})\log 2\mathit{N} + \log rac{1}{\delta}}{\mathit{N}}$
Agnostic	$\mathfrak{R}(g) < \hat{\mathfrak{R}}(g) + \sqrt{rac{1}{2N}\lograc{2 \mathcal{H} }{\delta}}$	$\mathfrak{R}(g) < \hat{\mathfrak{R}}(g) + \sqrt{rac{VC(\mathcal{H})(\log rac{2N}{VC(\mathcal{H})}) + \log rac{4}{\delta}}{N}}$

where  $VC(\mathcal{H})$  is the Vapnik-Chervonenkis dimension of the hypothesis space  $\mathcal{H}$ . What is  $VC(\mathcal{H})$  dimension ?

- A dichotomy of a set S is a partition of S into two disjoint subsets
- A set of instances S is shattered by a hypothesis space  $\mathcal{H}$  if and only if for every dichotomy of S there exists some hypothesis in  $\mathcal{H}$  consistent with this dichotomy.
- The Vapnik-Chervonenkis dimension  $VC(\mathcal{H})$  of hypothesis space  $\mathcal{H}$  defined over instance space  $\mathcal{X}$  is the size of the largest finite subset of  $\mathcal{X}$  shattered by  $\mathcal{H}$ . If arbitrary large finite sets of  $\mathcal{X}$  can be shattered by  $\mathcal{H}$ , then  $VC(\mathcal{H}) = \infty$



### **VC** Dimension

Hypothesis class	VC dimension (approx/exact)
Half-line thresholds on $\mathbb{R}$ , $h_a(x)=1\{x\geq a\}$	1
Intervals on $\mathbb R$ (positive on $[a,b]$ )	2
Union of $k$ intervals on $\mathbb R$	2k
Homogeneous hyperplanes in $\mathbb{R}^d$ (through origin)	d
Affine hyperplanes in $\mathbb{R}^d$ (with bias)	d+1
Axis-aligned rectangles in $\mathbb{R}^d$	$2d$ ( $\Rightarrow$ in $\mathbb{R}^2$ : 4)
Euclidean balls in $\mathbb{R}^d$	d+1
Decision stumps on $d$ real features	d
Depth- $L$ axis-aligned decision trees	$O(2^L \log 2^L)$ (grows with #leaves)
1-Nearest Neighbor in $\mathbb{R}^d$ (prototypes unconstrained)	$\infty$
Degree- $p$ polynomial separators in $\mathbb{R}^d$	$\binom{d+p}{p}$
ReLU (piecewise-linear) NN with $oldsymbol{W}$ weights	$\Theta(W \log W)$ (tight up to constants)
Linear separators with margin $\gamma$ on radius- $R$ data	$\lesssim (R/\gamma)^2$ (dimension-free bound)

VC measures worst-case shattering power; higher VC implies a greater sample requirement or stronger regularization for good generalization.

### Double Descent (1)

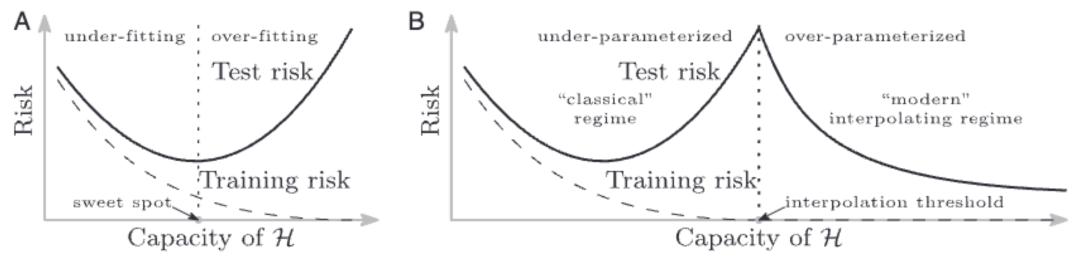


Fig. 1. Curves for training risk (dashed line) and test risk (solid line). (A) The classical U-shaped risk curve arising from the bias-variance trade-off. (B) The double-descent risk curve, which incorporates the U-shaped risk curve (i.e., the "classical" regime) together with the observed behavior from using high-capacity function classes (i.e., the "modern" interpolating regime), separated by the interpolation threshold. The predictors to the right of the interpolation threshold have zero training risk.

### Double Descent (2)

- As model capacity increases from small to moderate, test error first decreases and then increases.
- There is a point where the model exactly fits the training data while test error is near its highest.
- Beyond this point, further increasing capacity causes test error to decrease again.
- Training error keeps decreasing with capacity and remains near zero after exact fit.
- The behavior therefore is not a single U-shape but a "double descent" in test error as capacity grows.
- Models that are far larger than needed to interpolate the data can exhibit lower test error than smaller models.
- The location of the peak and the rate of the second descent depend on the task, data, and model family.

### Double Descent (3)

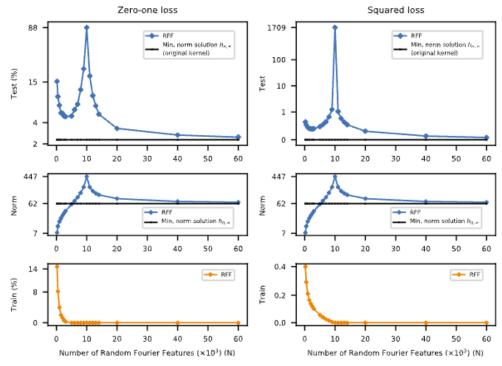


Fig. 2. Double-descent risk curve for the RFF model on MNIST. Shown are test risks (log scale), coefficient  $\ell_2$  norms (log scale), and training risks of the RFF model predictors  $h_{n,N}$  learned on a subset of MNIST ( $n = 10^4$ , 10 classes). The interpolation threshold is achieved at  $N = 10^4$ .

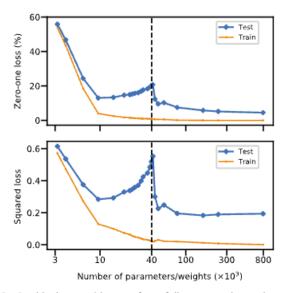


Fig. 3. Double-descent risk curve for a fully connected neural network on MNIST. Shown are training and test risks of a network with a single layer of H hidden units, learned on a subset of MNIST ( $n=4\cdot 10^3$ , d=784, K=10 classes). The number of parameters is  $(d+1)\cdot H+(H+1)\cdot K$ . The interpolation threshold (black dashed line) is observed at  $n\cdot K$ .

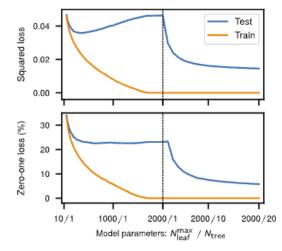


Fig. 4. Double-descent risk curve for random forests on MNIST. The double-descent risk curve is observed for random forests with increasing model complexity trained on a subset of MNIST ( $n=10^4$ , 10 classes). Its complexity is controlled by the number of trees  $N_{\rm tree}$  and the maximum number of leaves allowed for each tree  $N_{\rm leaf}^{\rm max}$ .

### Double Descent (4)

- The double-descent pattern shows up in all three: a linear model (with random features), a random forest, and a neural network.
- For each model family, **training error** falls to (near) zero as complexity increases and stays near zero afterward.
- For each model family, **test error peaks around the interpolation threshold**—the point where the model first exactly fits the training data.
- **Linear model:** increasing the number of random features moves from under- to over-parameterized and produces the same peak-then-drop in test error.
- **Neural network:** growing hidden units/parameters shows the same peak at interpolation followed by a second descent in test error.
- Random forest: increasing trees/allowed depth shows the same pattern—test error rises to a peak and then declines with more capacity.
- Across losses (zero-one and squared), the qualitative behavior is the same for all three model classes.
- The **peak location** and **descent rate** differ by model family and the chosen complexity knob, but the **overall pattern is consistent**.

### Justification of Double Descent in linear regression (1)

We consider a regression problem where the response y is equal to a linear function  $\boldsymbol{\beta} = (\beta_1, \dots, \beta_D) \in \mathbb{R}^D$  of D real-valued variables  $\boldsymbol{x} = (x_1, \dots, x_D)$  plus noise  $\sigma \epsilon$ :

$$y = x^*\beta + \sigma\epsilon = \sum_{j=1}^{D} x_j\beta_j + \sigma\epsilon.$$

Given n iid copies  $((\boldsymbol{x}^{(i)}, y^{(i)}))_{i=1}^n$  of  $(\boldsymbol{x}, y)$ , we fit a linear model to the data only using a subset  $T \subseteq [D] := \{1, \ldots, D\}$  of p := |T| variables.

Let  $\mathbf{X} := [\mathbf{x}^{(1)}|\cdots|\mathbf{x}^{(n)}]^*$  be the  $n \times D$  design matrix, and let  $\mathbf{y} := (y^{(1)}, \dots, y^{(n)})$  be the vector of responses. For a subset  $A \subseteq [D]$  and a D-dimensional vector  $\mathbf{v}$ , we use  $\mathbf{v}_A := (v_j : j \in A)$  to denote its |A|-dimensional subvector of entries from A; we also use  $\mathbf{X}_A := [\mathbf{x}_A^{(1)}|\cdots|\mathbf{x}_A^{(n)}]^*$  to denote the  $n \times |A|$  design matrix with variables from A. For  $A \subseteq [D]$ , we denote its complement by  $A^c := [D] \setminus A$ . Finally,  $\|\cdot\|$  denotes the Euclidean norm.

We fit regression coefficients  $\hat{\beta} = (\hat{\beta}_1, \dots, \hat{\beta}_D)$  with

$$\hat{\boldsymbol{\beta}}_T := \boldsymbol{X}_T^{\dagger} \boldsymbol{y}, \quad \hat{\boldsymbol{\beta}}_{T^c} := \boldsymbol{0}.$$

Above, the symbol  $^{\dagger}$  denotes the Moore-Penrose pseudoinverse. In other words, we use the solution to the normal equations  $\boldsymbol{X}_{T}^{*}\boldsymbol{X}_{T}\boldsymbol{v}=\boldsymbol{X}_{T}^{*}\boldsymbol{y}$  of least norm for  $\hat{\boldsymbol{\beta}}_{T}$  and force  $\hat{\boldsymbol{\beta}}_{T^{c}}$  to all-zeros.

**Theorem 1.** Assume the distribution of  $\mathbf{x}$  is the standard normal in  $\mathbb{R}^D$ ,  $\epsilon$  is a standard normal random variable independent of  $\mathbf{x}$ , and  $y = \mathbf{x}^*\boldsymbol{\beta} + \sigma\epsilon$  for some  $\boldsymbol{\beta} \in \mathbb{R}^D$  and  $\sigma > 0$ . Pick any  $p \in \{0, \ldots, D\}$  and  $T \subseteq [D]$  of cardinality p. The risk of  $\hat{\boldsymbol{\beta}}$ , where  $\hat{\boldsymbol{\beta}}_T = \mathbf{X}_T^T \mathbf{y}$  and  $\hat{\boldsymbol{\beta}}_{T^c} = \mathbf{0}$ , is

$$\mathbb{E}[(y - \boldsymbol{x}^* \hat{\boldsymbol{\beta}})^2] = \begin{cases} (\|\boldsymbol{\beta}_{T^c}\|^2 + \sigma^2) \cdot \left(1 + \frac{p}{n - p - 1}\right) & \text{if } p \leq n - 2; \\ +\infty & \text{if } n - 1 \leq p \leq n + 1; \\ \|\boldsymbol{\beta}_{T}\|^2 \cdot \left(1 - \frac{n}{p}\right) + (\|\boldsymbol{\beta}_{T^c}\|^2 + \sigma^2) \cdot \left(1 + \frac{n}{p - n - 1}\right) & \text{if } p \geq n + 2. \end{cases}$$

The proof of Theorem 1 is not hard, we give the details in Section 2.2. We now turn to the risk of  $\hat{\beta}$  under a random selection model for T.

### Justification Double Descent in linear regression (2)

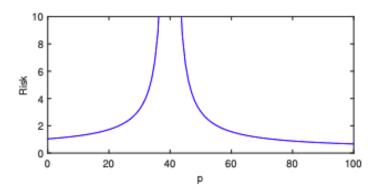


Figure 1: Plot of risk  $\mathbb{E}[(y - \boldsymbol{x}^*\hat{\boldsymbol{\beta}})^2]$  as a function of p, under the random selection model of T. Here,  $\|\boldsymbol{\beta}\|^2 = 1$ ,  $\sigma^2 = 1/25$ , D = 100, and n = 40.

In this setup with n=40n = 40n=40 and D=100D = 100D=100, least-squares regression uses ppp features, and as ppp increases up to nnn, the model overfits and risk worsens—reflecting the classical double descent without the initial descent. Interestingly, when p>np > np>n, the risk decreases again as the model benefits from overparameterization, perfectly fitting the data while becoming less volatile.

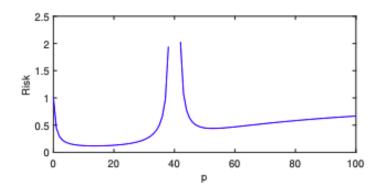


Figure 2: Plot of risk  $\mathbb{E}[(y-\boldsymbol{x}^*\dot{\boldsymbol{\beta}})^2]$  as a function of p, under the "prescient" selection model of T. Here,  $\|\boldsymbol{\beta}\|^2=1,\ \beta_j^2\propto 1/j^2,\ \sigma^2=1/25,\ D=100,\ \text{and}\ n=40.$ 

 The risk curve shows a "sweet spot" at low feature counts, resembling traditional descent, with higher risk in the interpolation regime. This happens because in a scientific feature selection setting, one cannot hand-pick informative features—so interpolation only helps when feature choice isn't biased. If features can be cherry-picked, good performance is achievable even with few variables.

### Minimum-norm solution in the underdetermined/overparameterized/interpolating regime

Setting. Linear regression with more parameters than samples:

- Data matrix  $X \in \mathbb{R}^{n \times d}$  with d > n; targets  $y \in \mathbb{R}^n$ .
- Interpolation means there exist  $w \in \mathbb{R}^d$  such that Xw = y (infinitely many solutions).

Main result (least-squares). Among all interpolating solutions,

$$w^\dagger = rg \min_{w: Xw = y} \|w\|_2 \ = \ X^+ y \ = \ X^ op (XX^ op)^{-1} y,$$

where  $X^+$  is the Moore–Penrose pseudoinverse.

Equivalently,

$$w^\dagger = \lim_{\lambda\downarrow 0}rg\min_w \|Xw-y\|_2^2 + \lambda\|w\|_2^2.$$

#### How it's selected in practice (implicit bias).

Gradient Descent on squared loss, initialized at  $w_0=0$  (and step size  $<2/\|X\|_2^2$ ), converges to  $w^\dagger$  among all exact-fit solutions.

#### Kernel/RKHS analogue (ridgeless).

For kernel matrix  $K = X_{\phi}X_{\phi}^{ op}$  :

$$f^\dagger(\cdot) = rg \min_{f: f(X) = y} \|f\|_{\mathcal{H}} \quad \Rightarrow \quad f^\dagger(x) = k(x,X)^ op K^+ y.$$

Why it matters. The minimum-norm (a.k.a. ridgeless) interpolant explains implicit regularization, connects to double-descent/benign overfitting, and often generalizes when signal aligns with the data span.

**Notation**. n: samples, d: features, X: design matrix, y: responses, w: parameters,  $\|\cdot\|_2$ : Euclidean norm,  $X^+$ : pseudoinverse, K: kernel Gram matrix,  $\mathcal{H}$ : RKP $^{\circ}$ 

#### References.

- Hastie, Montanari, Rosset, Tibshirani (2019), Surprises in High-Dimensional Ridgeless Least Squares, PNAS.
- Bartlett, Long, Lugosi, Tsigler (2020), Benign Overfitting in Linear Regression, PNAS.
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### Bridging to Neural Tangent Kernel (NTK)

- We've seen that linear models and kernel methods can interpolate the data, and gradient descent picks the minimum-norm (ridgeless) interpolant—matching the double-descent pattern we observed.
- Natural question: Can we approximate deep nets as linear or kernel machines so we can port these insights (interpolation, minimum-norm bias, double descent) to modern networks?
- Idea: look at very wide networks trained with small steps; their training can be linearized around initialization and viewed through a kernel induced by the network itself.
- This leads us to the Neural Tangent Kernel (NTK)—a framework where wide nets behave like kernel regressors with a learned, network-defined kernel.
- **Next:** define NTK, show the training dynamics, and connect it back to interpolation and generalization.

### Linearization of Neural Networks

For very wide nets trained with small steps, training behaves like fitting a linear model on fixed features determined at initialization.

#### Setup (scalar output for simplicity)

- Data:  $X=\{x_i\}_{i=1}^n$ , with  $x_i\in\mathbb{R}^d$ ; labels  $y\in\mathbb{R}^n$  where  $y_i\in\mathbb{R}$ .
- Network:  $f_{\theta}: \mathbb{R}^d \to \mathbb{R}$  with parameters  $\theta \in \mathbb{R}^p$ .
- Initialization:  $\theta_0 \in \mathbb{R}^p$ ; parameter change:  $\Delta \theta \coloneqq \theta \theta_0$ .

#### Fixed "tangent" features at initialization

•  $\phi(x)$ : gradient of the output w.r.t. parameters, evaluated at  $\theta_0$  (a length-p feature vector).

$$\phi(x) \coloneqq \nabla_{\theta} f_{\theta_0}(x) \in \mathbb{R}^p.$$

First-order (linear) approximation around  $\theta_0$ 

$$f_{ heta}(x) \ pprox \ f_{ heta_0}(x) \ + \ \phi(x)^ op \Delta heta$$

- f<sub>θ0</sub>(x): prediction at initialization.
- $\phi(x)^{\top}\Delta\theta$ : linear model in the fixed features  $\phi(x)$ .

Neural Tangent Kernel (NTK)

$$K(x,x') \ \coloneqq \ \phi(x)^ op \phi(x') \in \mathbb{R}.$$

- Kernel Gram matrix (train–train):  $K(X,X) \in \mathbb{R}^{n \times n}$  with entries  $K_{ij} = K(x_i,x_j)$ .
- Cross-kernel (test-train):  $K(x,X) \in \mathbb{R}^n$  with entries  $K(x,x_i)$ .

Training dynamics (squared loss, gradient flow)

Let  $f_t(X) \in \mathbb{R}^n$  collect predictions  $(f_{\theta(t)}(x_1), \dots, f_{\theta(t)}(x_n))$ . Under gradient flow (continuous-time limit of small-step GD) in the lazy regime:

$$rac{df_t(X)}{dt} \; = \; - \, K(X,X) \, ig( f_t(X) - y ig)$$

- t: (continuous) training time.
- Interpretation: predictions evolve by **kernel gradient descent** with kernel K.

Closed-form limit (interpolation, K(X, X) invertible)

$$f_{\infty}(x) = K(x,X) K(X,X)^{-1} y$$

- The trained wide NN behaves like kernel regression with the NTK.
- If K(X, X) is ill-conditioned, use ridge:

$$f_{\lambda}(x) = K(x,X) \left(K(X,X) + \lambda I 
ight)^{-1} y, \quad \lambda > 0.$$

Wide + small steps  $\Rightarrow$  NN  $\approx$  NTK kernel regression  $\Rightarrow$  double descent/benign overfitting results from linear & kernel models largely apply.

### Implicit Bias of Gradient Descent

- On linearly separable data, plain gradient descent on logistic/cross-entropy (and any smooth, strictly decreasing loss with an exponential tail) implicitly drives the predictor toward the hard-margin SVM solution. Concretely, the weights blow up in norm but their direction converges to the L2 max-margin separator.
- Under similar assumptions, **SGD also converges in direction to the L2 max-margin separator** on linearly separable data with exponentially-tailed losses (e.g., logistic).
- GD/SGD pick a specific interpolating classifier: on linearly separable data with an exponentially-tailed loss (e.g., logistic), both gradient descent and SGD drive the weight norm to infinity but their direction converges to the L2 hard-margin SVM. Practically, training longer after achieving 0 training error keeps increasing the margin, which often improves test accuracy.
  - If you're doing linear (or last-layer) classification with cross-entropy, **plain GD/SGD implicitly does SVM-like margin maximization**—so longer training can help margins without explicit regularization.

### Taxonomy of overfitting

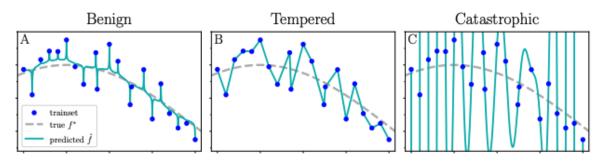


Figure 1: As  $n \to \infty$ , interpolating methods can exhibit three types of overfitting. (A) In benign overfitting, the predictor asymptotically approaches the ground-truth, Bayes-optimal function. Nadaraya-Watson kernel smoothing with a singular kernel, shown here, is asymptotically benign. (B) In tempered overfitting, the regime studied in this work, the predictor approaches a constant test risk greater than the Bayes-optimal risk. Piecewise-linear interpolation is asymptotically tempered. (C) In catastrophic overfitting, the predictor generalizes arbitrarily poorly. Rank-n polynomial interpolation is asymptotically catastrophic.

Benign	Tempered	Catastrophic	
Early-stopped DNNs	<ul> <li>Interpolating DNNs</li> </ul>	• Gaussian KR	
KR with ridge	• Laplace KR	<ul> <li>Critically-parameterized regression</li> </ul>	
• $k$ -NN ( $k \sim \log n$ )	• ReLU NTKs		
Nadaraya-Watson kernel	• k-NN (constant k)		
smoothing with Hilbert kernel	• Simplicial interpolation		

Table 2: A taxonomy of models under the three types of fitting identified in this work. **BOLD** are results from our work, others are known or folklore results.

	Regression	Classification
Benign	$\lim_{n\to\infty} \mathcal{R}_n = R^*$	$\lim_{n\to\infty} \mathcal{R}_n = R^*$
Tempered	$\lim_{n\to\infty} \mathcal{R}_n \in (R^*, \infty)$	$\lim_{n\to\infty} \mathcal{R}_n \in (R^*, 1-\frac{1}{K})$
Catastrophic	$\lim_{n\to\infty} \mathcal{R}_n = \infty$	$\lim_{n\to\infty} \mathcal{R}_n = 1 - \frac{1}{K}$

## Deep Neural Network Diagnostics via Random Matrix Theory

#### SETOL + WeightWatcher — one-slide takeaway

Main message. Deep nets that *generalize* have heavy-tailed layer spectra. You can score a model's quality without any data by fitting a power law to each layer's empirical spectral density (ESD) and reading off the  $\alpha$  (alpha) exponent; SETOL provides the theory, WeightWatcher gives you the button.

arXiv +1

Key result (paper). SETOL derives a layer-quality metric via a one-step Exact Renormalization Group (ERG) construction and shows it aligns closely with HTSR's  $\alpha/\alpha$  metrics computed from the ESDs of weight matrices—yielding conditions for "ideal learning" and flags for non-ideal regimes (e.g., overfit layers when  $\alpha < 2$ ).

#### What to remember.

- Good layers: α ≈ 2 (practically 2–6 acceptable).
- Red flags: α < 2 (over-trained/memorizing) or » 6 (under-trained/noisy); watch for correlation traps
  (spikes/outliers). (WeightWatcher +1)</li>
- Why it matters: Average α̂ (and related summaries) track test accuracy trends across model families—no train/test data needed.

### References

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   <u>Data: Exact Convergence with a Fixed Learning Rate</u>
- Neil Mallinar et al (2024), <u>Benign, Temered, or Catastrophic: A taxonomy of Overfitting</u>
- Charles Martin and Michael Mahoney (2021), <u>Implicit self-regularization in deep neural networks</u>