

Random matrix theory analysis of neural network weight matrices

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Abstract

As neural network weight matrices are initialized randomly, they conform precisely to random matrix theory (RMT) predictions before training. Post-training, deviations from RMT predictions indicate task-specific information encoded in the weights. We analyze feedforward and convolutional neural network weights trained on image recognition tasks. We demonstrate that most of the weights' singular values follow universal RMT predictions even after training, suggesting that major parts of weights remain random. By comparing singular value spectra with the Marchenko-Pastur distribution and singular vector entries with the Porter-Thomas distribution, we identify significant deviations only in the parts associated with the largest singular values. We argue that a comparison to RMT predictions allows locating learned information in the weights. In addition, the RMT analysis enables us to differentiate between networks trained within various learning regimes.

1. Introduction

Neural networks are often highly over-parametrized [1–12] and capable of memorizing large amounts of random training data [13, 14]. Traditional "bias-variance tradeoff" [15] suggests that such networks should overfit and fail with unseen data. However, they exhibit a double descent behavior [16–18] relative to the number of parameters, performing well even in the over-parametrized limit. This apparent contradiction is addressed by evidence showing that ultra-wide neural networks are biased towards simple functions [19–22].

To analyze these networks, we employ random matrix theory (RMT) [23–27] as a zero-information hypothesis, where deviations from RMT indicate system-specific information. RMT has been useful for studying systems with inherent randomness, such as nuclear spectra [24, 26–28], stock market correlations [29–31, 31–35], and biological networks [36, 37].

RMT has previously been applied to estimate the asymptotic performance of single-layer networks [38, 39] and to analyze the generalization dynamics of linear networks [40]. Outliers and the random part of pre-activation covariance matrices were examined in [41], while other studies focused on the spectra of Jacobians at initialization [42] and the eigenvalue distribution of the Hessian of the loss matrix [43, 44]. The spectral evolution of weight matrices during training was analyzed in [45], assessing the quality of pretrained DNNs [45, 46] by computing spectral norms of weight matrices and fitting the exponent of a power law tail of the singular value spectrum.

Here, we use various RMT tools to demonstrate that the weight matrices of deep, overparameterized neural networks are predominantly random. By comparing the singular values of several DNNs with the Wigner nearest neighbor spacing distribution, we find that the bulk of the spectrum aligns with RMT predictions. This is further corroborated by analyzing the number variance of the singular

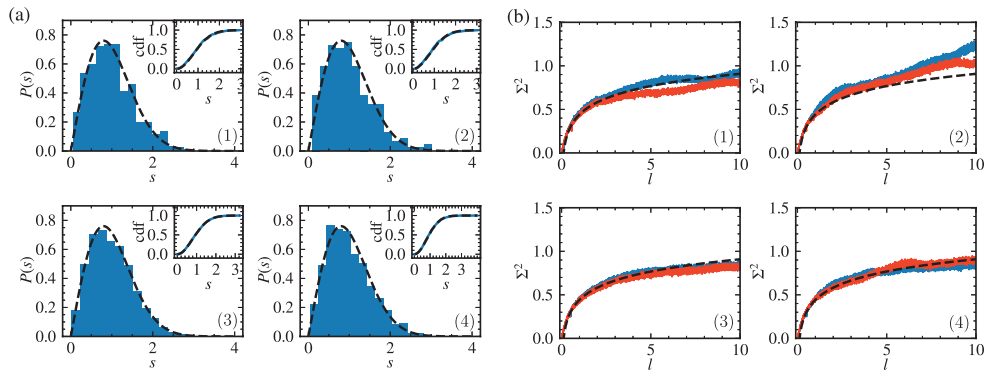


Figure 1. Nearest neighbor spacing distributions (a) and level number variance (b) of unfolded singular values of weights for various neural networks. The RMT predictions are depicted by dashed, black lines. The insets in (a) depict the cumulative distributions. Subpanels (1) show results for the second hidden layer weight matrix of MLP1024, (2) the second convolutional layer in the CNN miniAlexNet, (3) the second fully connected layer in AlexNet, and (4) for the third dense layer in VGG19. In all cases there is excellent visual agreement with the RMT predictions. For the level spacings this is further supported by Kolmogorov-Smirnov tests which cannot reject the null hypothesis at a significance level of (a1) 81%, (a2) 85%, (a3) 31%, and (a4) 96%.

value spectra of weight matrices. We investigate the hypothesis that a large fraction of singular values does not encode information by comparing the distribution of eigenvector components to the Porter-Thomas distribution. Significant deviations from the Porter-Thomas distribution are found only in a small fraction of eigenvectors with large singular values, indicating that learned information is encoded in them.

Additionally, we train networks across different learning regimes, from lazy networks where weights barely change during training to rich networks where final weights significantly differ from initial ones [47–50]. Networks trained in the lazy regime adhere closely to RMT predictions, unlike those trained in rich and intermediate regimes. Consequently, the weight spectrum and the comparison of singular vector entries to the Porter-Thomas distribution can distinguish between learning regimes, with the best generalization performance found between the two extremes.

The results presented here are a concise summary of Ref. [51], which includes additional findings and detailed information.

2. Experimental setup

In this study, we consider the singular value decomposition of a weight matrix W defined via $W = USV^T$, where U and V are orthogonal matrices, and S is a diagonal matrix of non-negative entries ν_i on its diagonal, the so-called singular values. While this is straightforward for dense layers, in the case of convolutional layers we first reshape the four dimensional weight tensors to a rectangular shape and then compute their singular values and vectors.

We consider several networks for image recognition with different architectures and sizes: a) a fully connected feedforward network with layers of size [3072, 1024, 512, 512, 10] denoted as MLP1024 and b) a convolutional network called miniAlexNet consisting of two convolutional layers followed by three dense layers, both trained on the CIFAR-10 [10] dataset. In addition, we analyze

the two larger networks c) AlexNet [52] and d) VGG19 [11], whose models trained on the ImageNet [53] dataset are available via `pytorch` [54] and `tensorflow` [55], respectively. The network weights are initialized using a Glorot uniform distribution [56] and then trained with stochastic gradient descent and cross-entropy loss. More details on the networks can be found in Appendix A.

3. Universal random matrix theory properties

We first consider universal properties that do only depend on the class of random matrix, not on a specific realization. These properties can be formulated for the unfolded singular value spectrum. Here, unfolding refers to normalizing the mean density of states of the singular values ν_i to unity, yielding the unfolded spectrum ξ_i [23–27]. For real random matrices in the universality class of the Gaussian orthogonal ensemble (GOE), the level spacings $s_k = \xi_{k+1} - \xi_k$, i.e. the differences between neighboring unfolded singular values, are distributed according to the Wigner surmise [23–28]

$$P_{\text{GOE}}(s) = \frac{\pi s}{2} \exp\left(-\frac{\pi}{4} s^2\right). \quad (1)$$

The nearest neighbor spacings of the weight matrix singular values are in excellent agreement with the RMT prediction Eq. (1) before and after training the networks (Fig. 1a). This is supported by Kolmogorov-Smirnov tests of the empirical data against Eq. (1) that cannot reject the null hypothesis even at a significance level as high as $\alpha = 0.30$ (for specific p -values, see Appendix B).

Another prediction of RMT that allows to test the random nature of weight matrices is the level number variance, which is sensitive to long range correlations in the spectrum. The number variance describes fluctuations in the number of unfolded singular values $N_{\xi_i}(l)$ in intervals of length l around each singular value ξ_i : $\Sigma^2(l) = \langle (N_{\xi_i}(l) - l)^2 \rangle_{\xi}$. For random matrices from the GOE universality class, the level number variance depends on the interval width l according to $\Sigma^2(l) \propto \ln(2\pi l)$ in the regime $l \gtrsim 1$ [24–27] in excellent agreement with the data (see Fig. 1b).

4. Distribution of singular values and singular vector entries

To locate the learned information in the weights, we compare the agreement with non-universal RMT predictions between initial and trained matrices. For a random $n \times m$ matrix of zero mean and variance σ , the singular values follow the Marchenko-Pastur distribution [57–59, 59],

$$P(\nu) = \begin{cases} \frac{n/m}{\pi \tilde{\sigma}^2 \nu} \sqrt{(\nu_{\max}^2 - \nu^2)(\nu^2 - \nu_{\min}^2)} & \nu \in [\nu_{\min}, \nu_{\max}] \\ 0 & \text{else} \end{cases} \quad (2)$$

where $\nu_{\min}^{\max} = \tilde{\sigma}(1 \pm \sqrt{m/n})$ and $\tilde{\sigma} = \sigma\sqrt{n}$. We assume without loss of generality that $m \leq n$. While the distribution Eq. (2) describes the spectrum of the weights of untrained networks, the trained weight deviates from the Marchenko-Pastur law (see [45]). In the absence of a microscopic theory for the spectrum of a trained weight matrix, we estimate its random part by fitting the empirical spectra with a modified Marchenko-Pastur law: setting ν_{\min} to the smallest empirical eigenvalue, and using ν_{\max} and σ^2 as fit parameters. This introduces an additional free parameter compared to the strict Marchenko-Pastur distribution, representing the percentage of the spectrum still following the Marchenko-Pastur law (see Fig. 2a). We also consider the normalized eigenvectors of the $m \times m$ matrix $W^\dagger W$ (right singular vectors of W), whose components for a random matrix follow the

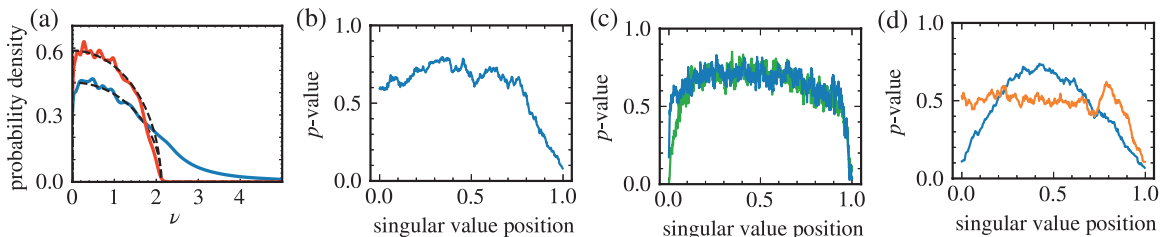


Figure 2. (a) Spectral distributions for initial weight (red), trained weight of the second hidden layer of a MLP512 network (blue) and RMT prediction (black). (b-d) Analysis of the eigenvectors of $W^T W$. Here, we show p -values of Kolmogorov-Smirnov tests of the eigenvector entries versus a Gaussian distribution. All results are averaged over neighboring eigenvectors with a window size of 15. The x -direction describes the position of rank ordered singular values, such that 0 corresponds to the smallest and 1 to the largest singular value of each weight matrix. We show results for (b) the second hidden layer of MLP1024, (c) the first dense layer of the large pretrained DNNs AlexNet (blue) and VGG19 (green), and (d) the second convolutional layer (blue) and first dense layer (orange) of the CNN miniAlexNet.

Porter-Thomas distribution [24, 26, 27], i.e., a Gaussian distribution with mean zero and standard deviation $1/\sqrt{m}$. To check if the observation that most singular values of trained networks are random extends to the eigenvectors, we test the empirical distribution of each eigenvector’s entries against this Gaussian using a Kolmogorov-Smirnov test. A large p -value means the Gaussian hypothesis cannot be rejected, indicating the vector contains only noise, while a small p -value suggests stored information. To reveal trends, we average the p -values over neighboring singular values with a window size of 15. We find that most eigenvectors are random, especially those corresponding to small singular values (Fig. 2b-d). For large singular values, the p -values decrease, indicating stored information, consistent with [45].

5. RMT analysis of different learning regimes

It was shown [14, 60–64] that neural networks can achieve good generalization accuracies even when their weights change only by very small amounts during training. The opposite to this *lazy learning* is denoted as *rich learning*, where the final weights \mathbf{W} after training deviate significantly from the initial ones \mathbf{W}_0 . We train several MLP512 networks, where laziness is controlled by introducing a hyperparameter α that modifies the output activations via [48] $a_L = \text{softmax}(\alpha(W_{L-1}\mathbf{a}_{L-1} + \mathbf{b}_L))$ and the cost function as

$$l(\mathbf{W}, \mathbf{b}) = -\frac{1}{N\alpha^2} \sum_{k=1}^N \mathbf{y}^{(k)} \cdot \ln(\mathbf{a}_{\text{out}}^{(k)}). \quad (3)$$

Here, a large $\alpha > 1$ scales down the gradient updates and therefore encourages lazy learning, while small $\alpha < 1$ steers training towards the rich learning regime [48]. We denote $\alpha = 1$ as *typical learning*. A comparison of the RMT analysis in the three regimes—rich ($\alpha = 0.5$), typical ($\alpha = 1$), and lazy ($\alpha = 5$)—is illustrated in Fig. 3. For all networks, the bulk spectra exhibit random characteristics, with level spacings (panel b) and level number variance (panel c) closely aligning with RMT predictions. Notably, the level number variances do not indicate slower growth for networks with larger α . In the rich network, there are more large singular values compared to the

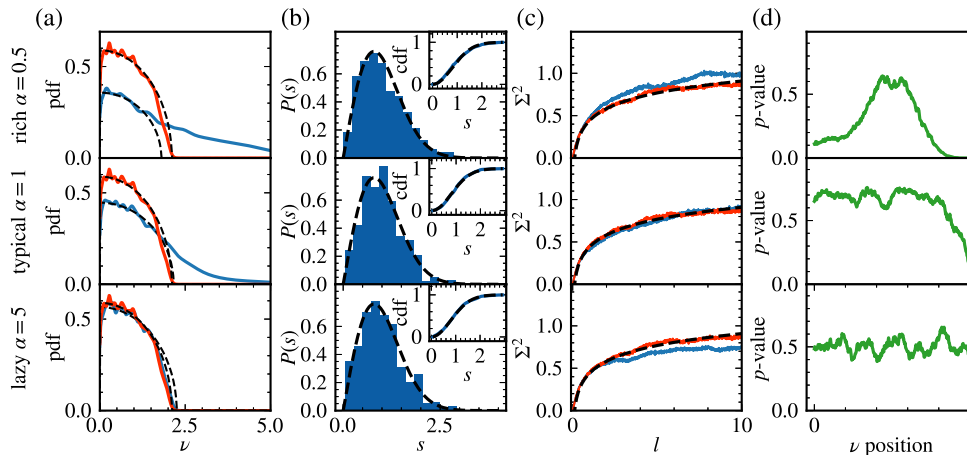


Figure 3. Random matrix theory analysis of second layer weights of MLP512 networks trained in different learning regimes: rich learning ($\alpha = 0.5$, top panel), typical learning ($\alpha = 1$, middle panel), and lazy learning ($\alpha = 5$, bottom panel). We show (a) the spectra for trained (blue) and randomly initialized networks (red) with fits of modified Marchenko-Pastur laws (dashed, black), (b) unfolded level spacing distributions (main panel, blue, window size 15) and corresponding cumulative distributions (insets) with the Wigner surmise (dashed, black), (c) unfolded level number variance (trained: blue, initialized: red), and (d) p -values for comparing singular vector entries to a Porter-Thomas distribution. Trained networks in all cases follow universal RMT predictions (b and c), indicating a random bulk. Lazy networks can be distinguished from typical and rich networks by the spectral distributions in (a) and p -values in (d) [51].

typical network, while the lazy network’s Marchenko-Pastur spectrum remains almost unchanged (panel a). Despite this, the lazy network achieves a test accuracy of 50.4% on CIFAR-10, compared to 52.7% for the rich network and 55.2% for the typical network.

When examining the p -values for Kolmogorov-Smirnov tests of eigenvector entries against a Porter-Thomas distribution (panel d), it is observed that in the typical case, small p -values occur only for large singular values. In contrast, the rich network shows small p -values for vectors corresponding to the smallest singular values as well. For the lazy network, all p -values fluctuate around 0.5, consistent with the behavior expected for random weights.

6. Conclusion

We employed RMT as a zero-information hypothesis to distinguish randomness from learned information during training. At initialization, weight matrices perfectly align with RMT predictions, with singular value spectra following the Marchenko-Pastur distribution, singular vector entries obeying the Porter-Thomas distribution, and level spacing adhering to the Wigner surmise. A comparison between initialized and trained networks reveals where information is stored in the weight matrices. Even after training, much of the eigenvalue spectrum remains random, and the spectral statistics continues to match RMT predictions. Singular vectors are mostly random, except for those associated with the largest singular values, indicating that learned information is concentrated in these vectors. Separating random parts from information may allow improved network compression algorithms to be designed.

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Appendix A. Network architectures and performance

Table 1. Neural network architectures and performance of trained networks. We use d to indicate a dense layer, c for a convolutional layer, p for max pooling, f for flattening, rc for random crop layer, and r for response normalization layer (with a depth radius of 5, a bias of 1, $\alpha = 1$, and $\beta = 0.5$).

	NETWORK	DATASET	TEST ACC
I)	MLP512, SEED 1 (d 3072, d 512, d 512, d 512, d 10) [14]	CIFAR-10	54.7%
	MLP512, SEED 2 (d 3072, d 512, d 512, d 512, d 10)	CIFAR-10	55.1%
	MLP512, SEED 3 (d 3072, d 512, d 512, d 512, d 10)	CIFAR-10	55.2%
II)	MLP1024 (d 3072, d 1024, d 512, d 512, d 10)	CIFAR-10	55.4%
III)	MINIALEXNET (c 300 5×5, p 3×3, r, c 150 5×5, p 3×3, r, f, d 384, d 192, d 10) [14]	CIFAR-10	78.5%
IV)	ALEXNET [52]	IMAGENET	56.5%
V)	VGG16 [11]	IMAGENET	67.6%
VI)	VGG19 [11]	IMAGENET	72.4%

We consider a variety of different networks to show that our results are valid for a wide range of architectures. Tab. 1 lists the network architectures, training datasets, and accuracies achieved on each dataset. We downloaded the large pre-trained networks iv) AlexNet [52] via `pytorch` [54], v) VGG16 [11] via `tensorflow` [55], and vi) VGG19 [11] via `pytorch` [54]. Networks i)-iii) are trained using mini-batch stochastic gradient descent for 100 epochs. The weights are initialized using the Glorot uniform distribution [56] and the biases are initialized with zeros. We standardize each image of the CIFAR-10 dataset by subtracting the mean and dividing by the standard deviation. We set the learning rate to 0.001 at the beginning and use an exponential learning rate schedule with decay constant 0.95. For all networks, we choose 0.95 as momentum and the mini-batch size is 32. Network architectures i)-ii) in Tab. 1 are trained without L_2 regularization, while we use an L_2 regularization strength of 10^{-4} for training miniAlexNet networks iii).

Appendix B. Kolmogorov-Smirnov tests against the Wigner surmise

The p -values of Kolmogorov-Smirnov tests of the unfolded singular value spacings against the Wigner surmise are shown in Tab. 2.

Appendix C. Details on unfolding the spectrum

We perform a Gaussian broadening [25, 65] by approximating the probability density as a sum of Gaussian functions centered around each of the m singular values ν_k with widths $\sigma_k = (\nu_{k+a} - \nu_{k-a})/2$, where $2a$ is the window size of the broadening [32, 66]

$$P(\nu) \approx \frac{1}{m} \sum_{k=1}^m \frac{1}{\sqrt{2\pi\sigma_k^2}} \exp\left(-\frac{(\nu - \nu_k)^2}{2\sigma_k^2}\right). \quad (4)$$

Unfolding is a transformation that maps the singular values ν_i to uniformly distributed singular values ξ_i [23–25, 27, 32]. For this purpose, we first determine the probability density $P(\nu)$ using

Table 2. Kolmogorov-Smirnov test results of the distribution of unfolded singular value spacings of the weight matrices against the Wigner surmise. Rejection of the null hypothesis is based on the $\alpha = 0.05$ significance level. The p-value indicates how likely it is to obtain a distribution with at least as much cumulative density function deviation as the one tested for drawing random numbers from a Wigner surmise distribution. We find excellent agreement with the Wigner surmise for a variety of network architectures.

NETWORK	REJECT NULL HYPOTHESIS?			KS-TEST P-VALUE		
	LAYER 1	LAYER 2	LAYER 3	LAYER 1	LAYER 2	LAYER 3
MLP512 (SEED 1)	NO	NO	NO	0.347	0.401	0.812
MLP512 (SEED 2)	NO	NO	NO	0.993	0.421	0.844
MLP512 (SEED 3)	NO	NO	NO	0.768	0.784	0.863
MLP1024	NO	NO	NO	0.799	0.812	0.792
MINIALEXNET (SECOND CONV. LAYER)		NO			0.859	
ALEXNET (DENSE LAYERS)	NO	NO	NO	0.670	0.229	0.160
VGG16 (DENSE LAYERS)	NO	NO	NO	0.923	0.312	0.309
VGG19 (DENSE LAYERS)	NO	NO	NO	0.376	0.652	0.557

Eq. (4) and calculate the corresponding cumulative distribution

$$F(\nu) = m \int_{-\infty}^{\nu} P(x) dx . \quad (5)$$

The unfolded singular values are defined as $\xi_i = F(\nu_i)$.

Appendix D. Controlling the learning regime

A criterion for estimating the learning regime was proposed by Chizat et al. [48]: For a neural network $f_{\mathbf{W}}$ that maps an input \mathbf{x} to an output, and an accuracy function $\mathcal{A}(f_{\mathbf{W}}, \{\mathbf{x}\}, \{\mathbf{y}\})$, where $\{\mathbf{x}\}$ is a dataset with labels $\{\mathbf{y}\}$, one computes the network’s linearization around the initial weights \mathbf{W}_0

$$\tilde{f}_{\mathbf{W}}(\mathbf{x}) = f_{\mathbf{W}_0}(\mathbf{x}) + (\mathbf{W} - \mathbf{W}_0) \cdot \nabla_{\mathbf{W}} f_{\mathbf{W}}|_{\mathbf{W}_0}(\mathbf{x}) . \quad (6)$$

In the lazy learning regime, where $\mathbf{W} \approx \mathbf{W}_0$, linearization is a good approximation such that the accuracies are barely different, i.e.

$$\mathcal{A}(f_{\mathbf{W}}, \{\mathbf{x}\}, \{\mathbf{y}\}) \approx \mathcal{A}(\tilde{f}_{\mathbf{W}}, \{\mathbf{x}\}, \{\mathbf{y}\}) . \quad (7)$$

On the contrary, in the rich learning regime, one expects significant deviations such that

$$\mathcal{A}(f_{\mathbf{W}}, \{\mathbf{x}\}, \{\mathbf{y}\}) \gg \mathcal{A}(\tilde{f}_{\mathbf{W}}, \{\mathbf{x}\}, \{\mathbf{y}\}) . \quad (8)$$

This criterion has the advantage that it can also be studied on a layer-wise basis by linearizing around a single weight matrix only, and as accuracies are in the range $[0, 1]$, it gives a scale for laziness comparable between different network architectures. A disadvantage is that it requires to compute the linearization which can be resource intensive for large networks. For obtaining $\tilde{f}_{\mathbf{W}}$, we use

the `autodiff` implementation in the `jax` python package together with the `neural_tangents` package [67].

Accuracies for linearized and full MLP512 networks as a function of α are depicted in Fig. 4. As expected, the networks are in the rich regime for $\alpha < 1$, where the full networks (blue crosses) perform significantly better than the linearized networks (green circles), while we observe lazy learning for $\alpha > 1$. The network with $\alpha = 1$ (black symbols), lies about in the middle between the two regimes, where we also find the best test accuracy. We therefore denote $\alpha = 1$ as *typical learning*.

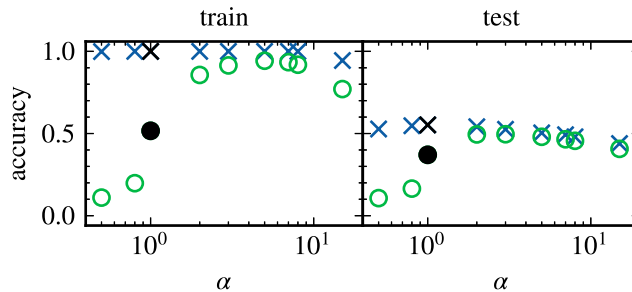


Figure 4. Comparison of training and test accuracies for full MLP512 networks (blue crosses) and linearized networks (green circles) around the initial weights of the second layer as a function of the laziness hyperparameter α . The black symbols indicate accuracies for $\alpha = 1$. For small $\alpha < 1$ accuracies of linearized and full networks deviate significantly which indicates rich learning, while for large $\alpha > 1$ performance differences are small indicating lazy learning [51].