A Kronecker-factored approximate Fisher matrix for convolution layers

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Abstract

Second-order optimization methods such as natural gradient descent have the potential to speed up training of neural networks by correcting for the curvature of the loss function. Unfortunately, the exact natural gradient is impractical to compute for large models, and most approximations either require an expensive iterative procedure or make crude approximations to the curvature. We present Kronecker Factors for Convolution (KFC), a tractable approximation to the Fisher matrix for convolutional networks based on a structured probabilistic model for the distribution over backpropagated derivatives. Similarly to the recently proposed Kronecker-Factored Approximate Curvature (K-FAC), each block of the approximate Fisher matrix decomposes as the Kronecker product of small matrices, allowing for efficient inversion. KFC captures important curvature information while still yielding comparably efficient updates to stochastic gradient descent (SGD). We show that the updates are invariant to commonly used reparameterizations, such as centering of the activations. In our experiments, approximate natural gradient descent with KFC was able to train convolutional networks several times faster than carefully tuned SGD. Furthermore, it was able to train the networks in 10-20 times fewer iterations than SGD, suggesting its potential applicability in a distributed setting.

1. Introduction

Despite advances in optimization, most neural networks are still trained using variants of stochastic gradient descent (SGD) with momentum. It has been suggested that natural gradient descent (Amari, 1998) could greatly speed up optimization because it accounts for the geometry of the optimization landscape and has desirable invariance properties. (See Martens (2014) for a review.) Unfortunately, RGROSSE@CS.TORONTO.EDU JMARTENS@CS.TORONTO.EDU

computing the exact natural gradient is intractable for large networks, as it requires solving a large linear system involving the Fisher matrix, whose dimension is the number of parameters (potentially tens of millions for modern architectures). Approximations to the natural gradient typically either impose very restrictive structure on the Fisher matrix (e.g. LeCun et al., 1998; Le Roux et al., 2008) or require expensive iterative procedures to compute each update, analogously to approximate Newton methods (e.g. Martens, 2010). An ongoing challenge has been to develop a curvature matrix approximation which reflects enough structure to yield high-quality updates, while introducing minimal computational overhead beyond the standard gradient computations.

Much progress in machine learning has been driven by the development of structured probabilistic models whose independence structure allows for efficient computations, yet which still capture important dependencies between the variables of interest. In our case, since the Fisher matrix is the covariance of the backpropagated log-likelihood derivatives, we are interested in modeling the distribution over these derivatives. The model must support efficient computation of the inverse covariance, as this is what's required to compute the natural gradient. Recently, the Factorized Natural Gradient (FANG) (Grosse & Salakhutdinov, 2015) and Kronecker-Factored Approximate Curvature (K-FAC) (Martens & Grosse, 2015) methods exploited probabilistic models of the derivatives to efficiently compute approximate natural gradient updates. In its simplest version, K-FAC approximates each layer-wise block of the Fisher matrix as the Kronecker product of two much smaller matrices. These (very large) blocks can then be can be tractably inverted by inverting each of the two factors. K-FAC was shown to greatly speed up the training of deep autoencoders. However, its underlying probabilistic model assumed fully connected networks with no weight sharing, rendering the method inapplicable to two architectures which have recently revolutionized many applications of machine learning - convolutional networks (Le-Cun et al., 1989; Krizhevsky et al., 2012) and recurrent neural networks (Hochreiter & Schmidhuber, 1997; Sutskever et al., 2014).

We introduce Kronecker Factors for Convolution (KFC), an approximation to the Fisher matrix for convolutional net-

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works. Most modern convolutional networks have trainable parameters only in convolutional and fully connected layers. Standard K-FAC can be applied to the latter; our contribution is a factorization of the Fisher blocks corresponding to convolution layers. KFC is based on a structured probabilistic model of the backpropagated derivatives where the activations are independent of the derivatives, the activations and derivatives are spatially homogeneous, and the derivatives are spatially uncorrelated. Under these approximations, we show that the Fisher blocks for convolution layers decompose as a Kronecker product of smaller matrices (analogously to K-FAC), yielding tractable updates.

KFC yields a tractable approximation to the Fisher matrix of a conv net. It can be used directly to compute approximate natural gradient descent updates, as we do in our experiments. One could further combine it with the adaptive step size, momentum, and damping methods from the full K-FAC algorithm (Martens & Grosse, 2015). It could also potentially be used as a pre-conditioner for iterative second-order methods (Martens, 2010; Vinyals & Povey, 2012; Sohl-Dickstein et al., 2014). We show that the approximate natural gradient updates are invariant to widely used reparameterizations of a network, such as whitening or centering of the activations.

We have evaluated our method on training conv nets on object recognition benchmarks. In our experiments, KFC was able to optimize conv nets several times faster than carefully tuned SGD with momentum, in terms of both training and test error. Furthermore, it required 10-20 times fewer *iterations*, suggesting its usefulness in the context of highly distributed training algorithms.

2. Background

In this section, we outline the K-FAC method as previously formulated for standard fully-connected feed-forward networks without weight sharing (Martens & Grosse, 2015). Each layer of a fully connected network computes activations as:

$$\mathbf{s}_{\ell} = \mathbf{W}_{\ell} \bar{\mathbf{a}}_{\ell-1} \tag{1}$$

$$\mathbf{a}_{\ell} = \phi_{\ell}(\mathbf{s}_{\ell}),\tag{2}$$

where $\ell \in \{1, \ldots, L\}$ indexes the layer, \mathbf{s}_{ℓ} denotes the inputs to the layer, \mathbf{a}_{ℓ} denotes the activations, $\mathbf{\bar{W}}_{\ell} = (\mathbf{b}_{\ell} \mathbf{W}_{\ell})$ denotes the matrix of biases and weights, $\mathbf{\bar{a}}_{\ell} = (1 \ \mathbf{a}_{\ell}^{\top})^{\top}$ denotes the activations with a homogeneous dimension appended, and ϕ_{ℓ} denotes a nonlinear activation function (usually applied coordinate-wise). (Throughout this paper, we will use the index 0 for all homogeneous coordinates.) We will refer to the values \mathbf{s}_{ℓ} as *pre-activations*. By convention, \mathbf{a}_0 corresponds to the inputs \mathbf{x} and \mathbf{a}_L corresponds to the prediction \mathbf{z} made by the network. For convenience, we concatenate all of the parameters of the network into a vector $\boldsymbol{\theta} = (\operatorname{vec}(\mathbf{W}_1)^{\top}, \ldots, \operatorname{vec}(\mathbf{W}_L)^{\top})^{\top}$, where vec denotes the Kronecker vector operator which stacks the columns of a matrix into a vector. We denote the function computed by the network as $f(\mathbf{x}, \boldsymbol{\theta}) = \mathbf{a}_L$.

Typically, a network is trained to minimize an objective $h(\theta)$ given by $\mathcal{L}(\mathbf{y}, f(\mathbf{x}, \theta))$ as averaged over the training set, where $\mathcal{L}(\mathbf{y}, \mathbf{z})$ is a loss function. The gradient ∇h of $h(\theta)$, which is required by most optimization methods, is estimated stochastically using mini-batches of training examples. (We will often drop the explicit θ subscript when the meaning is unambiguous.)

For the remainder of this paper, we will assume the network's prediction $f(\mathbf{x}, \boldsymbol{\theta})$ determines the value of the parameter \mathbf{z} of a distribution $R_{\mathbf{y}|\mathbf{z}}$ over \mathbf{y} , and the loss function is the corresponding negative log-likelihood $\mathcal{L}(\mathbf{y}, \mathbf{z}) = -\log r(\mathbf{y}|\mathbf{z})$.

2.1. Second-order optimization of neural networks

Second-order optimization methods work by computing a parameter update v that minimizes (or approximately minimizes) a local quadratic approximation to the objective, given by $h(\boldsymbol{\theta}) = \nabla_{\boldsymbol{\theta}} h^{\top} \mathbf{v} + \frac{1}{2} \mathbf{v}^{\top} \mathbf{C} \mathbf{v}$, where **C** is a matrix which quantifies the curvature of the cost function hat θ . The exact solution to this minimization problem can be obtained by solving the linear system $\mathbf{C}\mathbf{v} = -\nabla_{\boldsymbol{\theta}}h$. The original and most well-known example is Newton's method, where C is chosen to be the Hessian matrix; this isn't appropriate in the non-convex setting because of the well-known problem that it searches for critical points rather than local optima (e.g. Pascanu et al., 2014). Therefore, it is more common to use natural gradient (Amari, 1998) or updates based on the generalized Gauss-Newton matrix (Schraudolph, 2002), which are guaranteed to produce descent directions because the curvature matrix C is positive semidefinite.

Natural gradient descent can be usefully interpreted as a second-order method (Martens, 2014) where C is the Fisher information matrix F, as given by

$$\mathbf{F} = \mathbb{E}_{\substack{\mathbf{y} \sim R_{\mathbf{y}|f(\mathbf{x},\boldsymbol{\theta})}}} \mathbb{E}_{\mathbf{y} \sim R_{\mathbf{y}|f(\mathbf{x},\boldsymbol{\theta})}} \left[\mathcal{D}\boldsymbol{\theta}(\mathcal{D}\boldsymbol{\theta})^{\top} \right],$$
(3)

where p_{data} denotes the training distribution, $R_{\mathbf{y}|f(\mathbf{x}, \theta)}$ denotes the model's predictive distribution, and $\mathcal{D}\theta = \nabla_{\theta} \mathcal{L}(\mathbf{y}, f(\mathbf{x}, \theta))$ is the log-likelihood gradient. For the remainder of this paper, all expectations are with respect to this distribution (which we term the *model's distribution*), so we will leave off the subscripts. (In this paper, we will use the \mathcal{D} notation for log-likelihood derivatives; derivatives of other functions will be written out explicitly.) In the case where $R_{\mathbf{y}|\mathbf{z}}$ corresponds to an exponential family model with "natural" parameters given by \mathbf{z} , \mathbf{F} is equivalent to the generalized Gauss-Newton matrix (Martens, 2014), which is an approximation of the Hessian which has also seen extensive use in various neural-network optimization methods (e.g. Martens, 2010; Vinyals & Povey, 2012).

F is an $n \times n$ matrix, where *n* is the number of parameters and can be in the tens of millions for modern deep architectures. Therefore, it is impractical to represent **F**

explicitly in memory, let alone solve the linear system exactly. There are two general strategies one typically takes to find a good search direction: either impose a structure on **F** enabling fast inversion (e.g. LeCun et al., 1998; Le Roux et al., 2008; Grosse & Salakhutdinov, 2015), or use an iterative procedure to approximately solve the linear system (e.g. Martens, 2010). These two strategies are not mutually exclusive: tractable curvature approximations can be used as preconditioners in second order optimization, and this has been observed to make a large difference (Martens, 2010; Chapelle & Erhan, 2011; Vinyals & Povey, 2012).

2.2. Kronecker-factored approximate curvature

Kronecker-factored approximate curvature (K-FAC; Martens & Grosse, 2015) is a recently proposed optimization method for neural networks which can be seen as a hybrid of the two approximation strategies: it uses a tractable approximation to the Fisher matrix \mathbf{F} , but also uses an optimization strategy which behaves locally like conjugate gradient. This section gives a conceptual summary of the aspects of K-FAC relevant to the contributions of this paper; a precise description of the full algorithm is given in Appendix B.2.

The block-diagonal version of K-FAC (which is the simpler of the two versions, and is what we will present here) is based on two approximations to \mathbf{F} which together make it tractable to invert. First, weight derivatives in different layers are assumed to be uncorrelated, which corresponds to \mathbf{F} being block diagonal, with one block per layer. Each block is given by $\mathbb{E}[\operatorname{vec}(\mathcal{D}\mathbf{W}_{\ell}) \operatorname{vec}(\mathcal{D}\mathbf{W}_{\ell})^{\top}]$. This approximation by itself is insufficient, because each of the blocks may still be very large. (E.g., if a network has 1,000 units in each layer, each block would be of size $10^6 \times 10^6$.) For the second approximation, observe that

$$\mathbb{E}\left[\mathcal{D}[\bar{\mathbf{W}}_{\ell}]_{ij}\mathcal{D}[\bar{\mathbf{W}}_{\ell}]_{i'j'}\right] = \mathbb{E}\left[\mathcal{D}[\mathbf{s}_{\ell}]_{i}[\bar{\mathbf{a}}_{\ell-1}]_{j}\mathcal{D}[\mathbf{s}_{\ell}]_{i'}[\bar{\mathbf{a}}_{\ell-1}]_{j'}\right]$$

If approximate activations we the and preindependent, this activation derivatives as can decomposed $\mathbb{E} \left| \mathcal{D}[\mathbf{W}_{\ell}]_{ij} \mathcal{D}[\mathbf{W}_{\ell}]_{i'j'} \right|$ be as \approx $\mathbb{E}[\mathcal{D}[\mathbf{s}_{\ell}]_i \mathcal{D}[\mathbf{s}_{\ell}]_{i'}] \mathbb{E}[[\bar{\mathbf{a}}_{\ell-1}]_j [\bar{\mathbf{a}}_{\ell-1}]_{j'}].$ This can be written algebraically as a decomposition into a Kronecker product of two smaller matrices:

$$\mathbb{E}[\operatorname{vec}(\bar{\mathbf{W}}_{\ell})\operatorname{vec}(\bar{\mathbf{W}}_{\ell})^{\top}] \approx \Psi_{\ell-1} \otimes \Gamma_{\ell} \triangleq \hat{\mathbf{F}}_{\ell}, \quad (4)$$

where $\Psi_{\ell-1} = \mathbb{E}[\bar{\mathbf{a}}_{\ell-1}\bar{\mathbf{a}}_{\ell-1}^{\top}]$ and $\Gamma_{\ell} = \mathbb{E}[\mathbf{s}_{\ell}\mathbf{s}_{\ell}^{\top}]$ denote the second moment matrices of the activations and preactivation derivatives, respectively. Call the block diagonal approximate Fisher matrix, with blocks given by Eqn. 4, $\hat{\mathbf{F}}$. The two factors are estimated online from the empirical moments of the model's distribution using exponential moving averages.

To invert $\hat{\mathbf{F}}$, we use the facts that (1) we can invert a block diagonal matrix by inverting each of the blocks, and (2) the Kronecker product satisfies the identity $(\mathbf{A} \otimes \mathbf{B})^{-1} =$

$$A^{-1} \otimes B^{-1}$$

$$\hat{\mathbf{F}}^{-1} = \begin{pmatrix} \boldsymbol{\Psi}_0^{-1} \otimes \boldsymbol{\Gamma}_1^{-1} & \mathbf{0} \\ & \ddots & \\ \mathbf{0} & \boldsymbol{\Psi}_{L-1}^{-1} \otimes \boldsymbol{\Gamma}_L^{-1} \end{pmatrix} \quad (5)$$

We do not represent $\hat{\mathbf{F}}^{-1}$ explicitly, as each of the blocks is quite large. Instead, we keep track of each of the Kronecker factors.

The approximate natural gradient $\hat{\mathbf{F}}^{-1}\nabla h$ can then be computed as follows:

$$\hat{\mathbf{F}}^{-1}\nabla h = \begin{pmatrix} \operatorname{vec}\left(\mathbf{\Gamma}_{1}^{-1}(\nabla_{\bar{\mathbf{W}}_{1}}h)\mathbf{\Psi}_{0}^{-1}\right) \\ \vdots \\ \operatorname{vec}\left(\mathbf{\Gamma}_{L}^{-1}(\nabla_{\bar{\mathbf{W}}_{L}}h)\mathbf{\Psi}_{L-1}^{-1}\right) \end{pmatrix}$$
(6)

We would often like to add a multiple of the identity matrix to F for two reasons. First, many networks are regularized with weight decay, which corresponds to a penalty of $\frac{1}{2}\lambda \theta^{\top} \theta$, for some parameter λ . Following the interpretation of \mathbf{F} as a quadratic approximation to the curvature, it would be appropriate to use $\mathbf{F} + \lambda \mathbf{I}$ to approximate the curvature of the regularized objective. The second reason is that the local quadratic approximation of h implicitly used when computing the natural gradient may be inaccurate over the region of interest, owing to the approximation of **F** by $\hat{\mathbf{F}}$, to the approximation of the Hessian by **F**, and finally to the error associated with approximating h as locally quadratic in the first place. A common way to address this issue is to damp the updates by adding $\gamma \mathbf{I}$ to the approximate curvature matrix, for some small value γ , before minimizing the local quadratic model. Therefore, we would ideally like to compute $\left[\hat{\mathbf{F}} + (\lambda + \gamma)\mathbf{I}\right]^{-1} \nabla h$.

Unfortunately, adding $(\lambda + \gamma)\mathbf{I}$ breaks the Kronecker factorization structure. While it is possible to exactly solve the damped system (see Appendix B.2), it is often preferable to approximate $\hat{\mathbf{F}} + (\lambda + \gamma)\mathbf{I}$ in a way that maintains the factorizaton structure. Martens & Grosse (2015) pointed out that

$$\hat{\mathbf{F}}_{\ell} + (\lambda + \gamma) \mathbf{I} \approx \left(\Psi_{\ell-1} + \pi_{\ell} \sqrt{\lambda + \gamma} \, \mathbf{I} \right) \otimes \left(\Gamma_{\ell} + \frac{1}{\pi_{\ell}} \sqrt{\lambda + \gamma} \, \mathbf{I} \right)$$
(7)

We will denote this damped approximation as $\hat{\mathbf{F}}_{\ell}^{(\gamma)} = \Psi_{\ell-1}^{(\gamma)} \otimes \Gamma_{\ell}^{(\gamma)}$. Mathematically, π_{ℓ} can be any positive scalar, but Martens & Grosse (2015) suggest the formula

$$\pi_{\ell} = \sqrt{\frac{\|\Psi_{\ell-1} \otimes \mathbf{I}\|}{\|\mathbf{I} \otimes \Gamma_{\ell}\|}},\tag{8}$$

where $\|\cdot\|$ denotes some matrix norm, as this value minimizes the norm of the residual in Eqn. 7. In this work, we use the trace norm $\|\mathbf{B}\| = \operatorname{tr} \mathbf{B}$. The approximate natural

gradient $\hat{\nabla}h$ is then computed as:

$$\hat{\nabla}h \triangleq [\hat{\mathbf{F}}^{(\gamma)}]^{-1} \nabla h = \begin{pmatrix} \operatorname{vec}\left([\boldsymbol{\Gamma}_{1}^{(\gamma)}]^{-1}(\nabla_{\bar{\mathbf{W}}_{1}}h)[\boldsymbol{\Psi}_{0}^{(\gamma)}]^{-1}\right) \\ \vdots \\ \operatorname{vec}\left([\boldsymbol{\Gamma}_{L}^{(\gamma)}]^{-1}(\nabla_{\bar{\mathbf{W}}_{L}}h)[\boldsymbol{\Psi}_{L-1}^{(\gamma)}]^{-1}\right) \end{pmatrix}$$
(9)

The algorithm as presented by Martens & Grosse (2015) has many additional elements which are orthogonal to the contributions of this paper. For concision, a full description of the algorithm is relegated to Appendix B.2.

2.3. Convolutional networks

Convolutional networks can require somewhat crufty notation when the computations are written out in full. In our case, we are interested in computing correlations of derivatives, which compounds the notational difficulties. In this section, we summarize the notation we use. (Table 1 lists all convolutional network notation used in this paper.) In sections which focus on a single layer of the network, we drop the explicit layer indices.

A convolution layer takes as input a layer of activations $\{a_{j,t}\}$, where $j \in \{1, \ldots, J\}$ indexes the input map and $t \in \mathcal{T}$ indexes the spatial location. (Here, \mathcal{T} is the set of spatial locations, which is typically a 2-D grid. For simplicity, we assume convolution is performed with a stride of 1 and padding equal to R, so that the set of spatial locations is shared between the input and output feature maps.) This layer is parameterized by a set of weights $w_{i,i,\delta}$ and biases b_i , where $i \in \{1, \ldots, I\}$ indexes the output map, j indexes the input map, and $\delta \in \Delta$ indexes the spatial offset (from the center of the filter). If the filters are of size $(2R + 1) \times (2R + 1)$, then we would have $\Delta = \{-R, \ldots, R\} \times \{-R, \ldots, R\}$. We denote the numbers of spatial locations and spatial offsets as $|\mathcal{T}|$ and $|\Delta|$, respectively. The convolution layer computes a set of preactivations $\{s_{i,t}\}$ as follows:

$$s_{i,t} = \sum_{\delta \in \Delta} w_{i,j,\delta} a_{j,t+\delta} + b_i, \tag{10}$$

where b_i denotes the bias parameter. The activations are defined to take the value 0 outside of \mathcal{T} . The pre-activations are passed through a nonlinearity such as ReLU to compute the output layer activations, but we have no need to refer to this explicitly when analyzing a single layer. (For simplicity, we assume operations such as pooling and response normalization are implemented as separate layers.)

Pre-activation derivatives $\mathcal{D}s_{i,t}$ are computed during backpropagation. One then computes weight derivatives as:

$$\mathcal{D}w_{i,j,\delta} = \sum_{t \in \mathcal{T}} a_{j,t+\delta} \mathcal{D}s_{i,t}.$$
 (11)

In some cases, it is useful to introduce vectorized notation for conv nets. We will represent the activations for a layer ℓ as a $|\mathcal{T}| \times J$ matrix \mathbf{A}_{ℓ} and the preactivations as a $|\mathcal{T}| \times I$ matrix \mathbf{S}_{ℓ} . The weights are represented as a $I \times |\Delta| J$ matrix \mathbf{W}_{ℓ} .

2.3.1. EFFICIENT IMPLEMENTATION AND VECTORIZED NOTATION

For modern large-scale vision applications, it's necessary to implement conv nets efficiently for a GPU (or some other massively parallel computing architecture). Since one contribution of our own work was to exploit the same underlying implementation to efficiently compute the statistics needed by our algorithm, we outline a typical GPU implementation of a conv net. As a bonus, discussing the implementation gives us a convenient high-level notation for analyzing conv nets mathematically. Due to space constrants, we relegate this material to Appendix A. This appendix also contains a table of all conv net notation used in this paper.

3. Kronecker factorization for convolution layers

We begin by assuming a block-diagonal approximation to the Fisher matrix like that of K-FAC, where each block contains all the parameters relevant to one layer (see Section 2.2). (Recall that these blocks are typically too large to invert exactly, or even represent explicitly, which is why the further Kronecker approximation is required.) The Kronecker factorization from K-FAC applies only to fully connected layers. Convolutional networks introduce several kinds of layers not found in fully connected feed-forward networks: convolution, pooling, and response normalization. Since pooling and response normalization layers don't have trainable weights, they are not included in the Fisher matrix. However, we must deal with convolution layers. In this section, we present our main contribution, an approximate Kronecker factorization for the blocks of \mathbf{F} corresponding to convolution layers. In the tradition of fast food puns (Ranzato & Hinton, 2010; Yang et al., 2014), we call our method Kronecker Factors for Convolution (KFC).

For this section, we focus on the Fisher block for a single layer, so we drop the layer indices. All conv net notation is summarized in Appendix A.

Recall that the Fisher matrix $\mathbf{F} = \mathbb{E} \left[\mathcal{D} \boldsymbol{\theta} (\mathcal{D} \boldsymbol{\theta})^{\top} \right]$ is the covariance of the log-likelihood gradient under the model's distribution. (In this paper, all expectations are with respect to the model's distribution unless otherwise specified.) By plugging in Eqn. 11, the entries corresponding to weight derivatives are given by:

$$\mathbb{E}[\mathcal{D}w_{i,j,\delta}\mathcal{D}w_{i',j',\delta'}] = \mathbb{E}\left[\left(\sum_{t\in\mathcal{T}} a_{j,t+\delta}\mathcal{D}s_{i,t}\right) \\ \left(\sum_{t'\in\mathcal{T}} a_{j',t'+\delta'}\mathcal{D}s_{i',t'}\right)\right]$$
(12)

To think about the computational complexity of computing

the entries directly, consider the second convolution layer of AlexNet (Krizhevsky et al., 2012), which has 48 input feature maps, 128 output feature maps, $27 \times 27 = 729$ spatial locations, and 5×5 filters. Since there are $128 \times 48 \times 5 \times 5 = 245760$ weights and 128 biases, the full block would require $245888^2 \approx 60.5$ billion entries to represent explicitly, and inversion is clearly impractical.

Recall that K-FAC approximation for classical fully connected networks can be derived by approximating activations and pre-activation derivatives as being statistically independent (this is the **IAD** approximation below). Deriving an analogous Fisher approximation for convolution layers will require some additional approximations.

Here are the approximations we will make in deriving our Fisher approximation:

- Independent activations and derivatives (IAD). The activations are independent of the pre-activation derivatives, *i.e.* $\{a_{j,t}\} \perp \{\mathcal{D}s_{i,t'}\}$.
- Spatial homogeneity (SH). The first-order statistics of the activations are independent of spatial location. The second-order statistics of the activations and preactivation derivatives at any two spatial locations t and t' depend only on t' - t. This implies there are functions M, Ω and Γ such that:

$$\mathbb{E}\left[a_{j,t}\right] = M(j) \tag{13}$$

$$\mathbb{E}\left[a_{j,t}a_{j',t'}\right] = \Omega(j,j',t'-t) \tag{14}$$

$$\mathbb{E}\left[\mathcal{D}s_{i,t}\mathcal{D}s_{i',t'}\right] = \Gamma(i,i',t'-t).$$
(15)

Note that $\mathbb{E}[\mathcal{D}s_{i,t}] = 0$ under the model's distribution, so $\operatorname{Cov}(\mathcal{D}s_{i,t}, \mathcal{D}s_{i',t'}) = \mathbb{E}[\mathcal{D}s_{i,t}\mathcal{D}s_{i',t'}].$

 Spatially uncorrelated derivatives (SUD). The preactivation derivatives at any two distinct spatial locations are uncorrelated, *i.e.* Γ(*i*, *i'*, δ) = 0 for δ ≠ 0.

We believe **SH** is fairly innocuous, as one is implicitly making a spatial homogeneity assumption when choosing to use convolution in the first place. **SUD** perhaps sounds like a more severe approximation, but in fact appeared to describe the model's distribution quite well in the networks we investigated; this is analyzed empirially in Section 5.1.

We now show that combining the above three approximations yields a Kronecker factorization of the Fisher blocks. For simplicity of notation, assume the data are twodimensional, so that the offsets can be parameterized with indices $\delta = (\delta_1, \delta_2)$ and $\delta' = (\delta'_1, \delta'_2)$, and denote the dimensions of the activations map as (T_1, T_2) . The formulas can be generalized to data dimensions higher than 2 in the obvious way. For clarity, we leave out the bias parameters in this section, but these are discussed in Appendix E.

Theorem 1. Combining approximations IAD, SH, and

SUD yields the following factorization:

$$\mathbb{E}\left[\mathcal{D}w_{i,j,\delta}\mathcal{D}w_{i',j',\delta'}\right] = \beta(\delta,\delta')\,\Omega(j,j',\delta'-\delta)\,\Gamma(i,i',0),$$
(16)

where

$$\beta(\delta, \delta') \triangleq (T_1 - \max(\delta_1, \delta'_1, 0) + \min(\delta_1, \delta'_1, 0)) \cdot (T_2 - \max(\delta_2, \delta'_2, 0) + \min(\delta_2, \delta'_2, 0)) \quad (17)$$

Proof. See Appendix E.
$$\Box$$

To talk about how this fits in to the block diagonal approximation to the Fisher matrix \mathbf{F} , we now restore the explicit layer indices and use the vectorized notation from Section 2.3.1. The above factorization yields a Kronecker factorization of each block, which will be useful for computing their inverses (and ultimately our approximate natural gradient). In particular, if $\hat{\mathbf{F}}_{\ell} \approx \mathbb{E}[\operatorname{vec}(\mathcal{D}\bar{\mathbf{W}}_{\ell}) \operatorname{vec}(\mathcal{D}\bar{\mathbf{W}}_{\ell})^{\top}]$ denotes the block of the approximate Fisher for layer ℓ , Eqn. 16 yields our KFC factorization of $\hat{\mathbf{F}}_{\ell}$ into a Kronecker product of smaller factors:

$$\hat{\mathbf{F}}_{\ell} = \mathbf{\Omega}_{\ell-1} \otimes \mathbf{\Gamma}_{\ell},\tag{18}$$

where

$$[\mathbf{\Omega}_{\ell-1}]_{j|\Delta|+\delta, \, j'|\Delta|+\delta'} \triangleq \beta(\delta,\delta') \, \Omega(j,j',\delta'-\delta)$$
$$[\mathbf{\Gamma}_{\ell}]_{i,i'} \triangleq \Gamma(i,i',0). \tag{19}$$

(We will derive much simpler formulas for $\Omega_{\ell-1}$ and Γ_{ℓ} in the next section.) Using this factorization, the rest of the K-FAC algorithm can be carried out without modification. For instance, we can compute the approximate natural gradient using a damped version of $\hat{\mathbf{F}}$ analogously to Eqns. 7 and 9 of Section 2.2:

$$\hat{\mathbf{F}}_{\ell}^{(\gamma)} = \mathbf{\Omega}_{\ell-1}^{(\gamma)} \otimes \mathbf{\Gamma}_{\ell}^{(\gamma)} \tag{20}$$

$$\triangleq \left(\mathbf{\Omega}_{\ell-1} + \pi_{\ell} \sqrt{\lambda + \gamma} \mathbf{I} \right) \otimes \\ \otimes \left(\mathbf{\Gamma}_{\ell} + \frac{1}{\pi_{\ell}} \sqrt{\lambda + \gamma} \mathbf{I} \right).$$
 (21)

$$\hat{\nabla}h = [\hat{\mathbf{F}}^{(\gamma)}]^{-1} \nabla h = \begin{pmatrix} \operatorname{vec}\left([\boldsymbol{\Gamma}_{1}^{(\gamma)}]^{-1}(\nabla_{\bar{\mathbf{W}}_{1}}h)[\boldsymbol{\Omega}_{0}^{(\gamma)}]^{-1}\right) \\ \vdots \\ \operatorname{vec}\left([\boldsymbol{\Gamma}_{L}^{(\gamma)}]^{-1}(\nabla_{\bar{\mathbf{W}}_{L}}h)[\boldsymbol{\Omega}_{L-1}^{(\gamma)}]^{-1}\right) \end{pmatrix}$$
(22)

Returning to our running example of AlexNet, $\bar{\mathbf{W}}_{\ell}$ is a $I \times (J|\Delta|+1) = 128 \times 1201$ matrix. Therefore the factors $\Omega_{\ell-1}$ and Γ_{ℓ} are 1201×1201 and 128×128 , respectively. These matrices are small enough that they can be represented exactly and inverted in a reasonable amount of time, allowing us to efficiently compute the approximate natural gradient direction using Eqn. 22.

3.1. Estimating the factors

Since the true covariance statistics are unknown, we estimate them empirically by sampling from the model's distribution, similarly to Martens & Grosse (2015). To sample derivatives from the model's distribution, we select a mini-batch, sample the outputs from the model's predictive distribution, and backpropagate the derivatives.

We need to estimate the Kronecker factors $\{\Omega_{\ell}\}_{\ell=0}^{L-1}$ and $\{\Gamma_{\ell}\}_{\ell=1}^{L}$. Since these matrices are defined in terms of the autocovariance functions Ω and Γ , it would appear natural to estimate these functions empirically. Unfortunately, if the empirical autocovariances are plugged into Eqn. 19, the resulting Ω_{ℓ} may not be positive semidefinite. This is a problem, since negative eigenvalues in the approximate Fisher could cause the optimization to diverge (a phenomenon we have observed in practice).

Instead, we estimate each Ω_{ℓ} directly using the following fact:

Theorem 2. Under assumption SH,

$$\boldsymbol{\Omega}_{\ell} = \mathbb{E}\left[\left[\mathbf{A}_{\ell} \right] \right]_{H}^{\top} \left[\mathbf{A}_{\ell} \right]_{H}^{\top} \right]$$
$$\boldsymbol{\Gamma}_{\ell} = \frac{1}{|\mathcal{T}|} \mathbb{E}\left[\mathcal{D} \mathbf{S}_{\ell}^{\top} \mathcal{D} \mathbf{S}_{\ell} \right].$$
(23)

(The $\llbracket \cdot \rrbracket$ notation is defined in Appendix A.)

We maintain exponential moving averages of the covariance statistics, where the empirical statistics are computed on each mini-batch using these formulas.

3.2. Using KFC in optimization

So far, we have defined an approximation $\hat{\mathbf{F}}^{(\gamma)}$ to the Fisher matrix \mathbf{F} which can be tractably inverted. This can be used in any number of ways in the context of optimization, most simply by using $\hat{\nabla}h = [\hat{\mathbf{F}}^{(\gamma)}]^{-1}\nabla h$ as an approximation to the natural gradient $\mathbf{F}^{-1}\nabla h$. Alternatively, we could use it in the context of the full K-FAC algorithm, or as a preconditioner for iterative second-order methods (Martens, 2010; Vinyals & Povey, 2012; Sohl-Dickstein et al., 2014).

In our experiments, we explored two particular instantiations of KFC in optimization algorithms. First, in order to provide as direct a comparison as possible to standard SGD-based optimization, we used $\hat{\nabla}h$ in the context of a generic approximate natural gradient descent procedure; this procedure is like SGD, except that $\hat{\nabla}h$ is substituted for the Euclidean gradient. Additionally, we used momentum, update clipping, and parameter averaging — all standard techniques in the context of stochastic optimization.¹ One can also view this as a preconditioned SGD method, where $\hat{\mathbf{F}}^{(\gamma)}$ is used as the preconditioner. Therefore, we refer to this method in our experiments as KFC-pre (to distinguish it from the KFC approximation itself). This method is spelled out in detail in Appendix B.1.

We also explored the use of $\hat{\mathbf{F}}^{(\gamma)}$ in the context of the full K-FAC training procedure (see Appendix B.2). Since this performed about the same as KFC-pre, we report results only for KFC-pre.

With the exception of inverting the Kronecker factors, all of the heavy computation for our methods was performed on the GPU. We based our implementation on CUDAMat (Mnih, 2009) and the convolution kernels provided by the Toronto Deep Learning ConvNet (TDLCN) package (Srivastava, 2015). Full details on our GPU implementation and other techniques for minimizing computational overhead are given in Appendix B.3.

4. Theoretical analysis

4.1. Invariance

Natural gradient descent is motivated partly by way of its invariance to reparameterization: regardless of how the model is parameterized, the updates are equivalent to the first order. Approximations to natural gradient don't satisfy full invariance to parameterization, but certain approximations have been shown to be invariant to more limited, but still fairly broad, classes of transformations (Ollivier, 2015; Martens & Grosse, 2015). For instance, K-FAC was shown to be invariant to affine transformations of the activations (Martens & Grosse, 2015).

For convolutional layers, we cannot expect an algorithm to be invariant to arbitrary affine transformations of a given layer's activations, as such transformations can change the set of functions which are representable. (Consider for instance, a transformation which permutes the spatial locations.) However, we show that the KFC updates are invariant to homogeneous, *pointwise* affine transformations of the activations, both before and after the nonlinearity. This is perhaps an overly limited statement, as it doesn't use the fact that the algorithm accounts for spatial correlations. However, it still accounts for a broad set of transformations, such as normalizing activations to be zero mean and unit variance either before or after the nonlinearity.

To formalize this, recall that a layer's activations are represented as a $|\mathcal{T}| \times J$ matrix and are computed from that layer's pre-activations by way of an elementwise nonlinearity, i.e. $\mathbf{A}_{\ell} = \phi_{\ell}(\mathbf{S}_{\ell})$. We replace this with an activation function ϕ_{ℓ}^{\dagger} which additionally computes affine transformations before and after the nonlinearity. Such transformations can be represented in matrix form:

$$\mathbf{A}_{\ell}^{\dagger} = \phi_{\ell}^{\dagger}(\mathbf{S}_{\ell}^{\dagger}) = \phi_{\ell}(\mathbf{S}_{\ell}^{\dagger}\mathbf{U}_{\ell} + \mathbf{1}\mathbf{c}_{\ell}^{\top})\mathbf{V}_{\ell} + \mathbf{1}\mathbf{d}_{\ell}^{\top}, \quad (24)$$

where \mathbf{U}_ℓ and \mathbf{V}_ℓ are invertible matrices, and \mathbf{c}_ℓ and \mathbf{d}_ℓ

¹Our SGD baseline used momentum and parameter averaging as well. Clipping was not needed for SGD, for reasons explained in Appendix B.1.

are vectors. For convenience, the inputs to the network can be treated as an activation function ϕ_0 which takes no arguments. We also assume the final layer outputs are not transformed, i.e. $\mathbf{V}_L = \mathbf{I}$ and $\mathbf{d}_L = \mathbf{0}$. KFC is invariant to this class of transformations:

Theorem 3. Let \mathcal{N} be a network with parameter vector $\boldsymbol{\theta}$ and activation functions $\{\phi_\ell\}_{\ell=0}^L$. Given activation functions $\{\phi_\ell^\dagger\}_{\ell=0}^L$ defined as in Eqn. 24, there exists a parameter vector $\boldsymbol{\theta}^\dagger$ such that a network \mathcal{N}^\dagger with parameters $\boldsymbol{\theta}^\dagger$ and activation functions $\{\phi_\ell^\dagger\}_{\ell=0}^L$ computes the same function as \mathcal{N} . The KFC updates on \mathcal{N} and \mathcal{N}^\dagger are equivalent, in that the resulting networks compute the same function.

Invariance to affine transformations also implies approximate invariance to smooth nonlinear transformations; see Martens (2014) for further discussion.

4.2. Relationship with other algorithms

It is possible to interpret many other neural net optimization methods as structured probabilistic approximations to natural gradient. This includes coordinatewise rescaling methods (e.g. LeCun et al., 1998; Duchi et al., 2011; Tieleman & Hinton, 2012; Zeiler, 2013; Kingma & Ba, 2015), centering of activations (Cho et al., 2013; Vatanen et al., 2013; Ioffe & Szegedy, 2015, e.g.), and the recently proposed Projected Natural Gradient (Desjardins et al., 2015). This allows us to compare the modeling assumptions implicitly made by different methods. See Appendix C for a full discussion.

5. Experiments

We have evaluated our method on two standard image recognition benchmark datasets: CIFAR-10 (Krizhevsky, 2009), and Street View Housing Numbers (SVHN; Netzer et al., 2011). Our aim is not to achieve state-of-the-art performance, but to evaluate KFC's ability to optimize previously published architectures. We first examine the probabilistic assumptions, and then present optimization results.

For CIFAR-10, we used the architecture from cuda-convnet² which achieved 18% error in 20 minutes. This network consists of three convolution layers and a fully connected layer. (While cuda-convnet provides some better-performing architectures, we could not use these, since these included locally connected layers, which KFC can't handle.) For SVHN, we used the architecture of Srivastava (2013). This architecture consists of three convolutional layers followed by three fully connected layers, and uses dropout for regularization. Both of these architectures were carefully tuned for their respective tasks. Furthermore, the TDLCN CUDA kernels

we used were carefully tuned at a low level to implement SGD updates efficiently for both of these architectures. Therefore, we believe our SGD baseline is quite strong.

5.1. Evaluating the probabilistic modeling assumptions

One of the benefits of using a structured probabilistic model to approximate the Fisher matrix is that we can analyze whether the modeling assumptions are satisfied. As discussed above, **IAD** is the standard approximation made by standard K-FAC, and was discussed in detail both theoretically and empirically by Martens & Grosse (2015). One implicitly assumes **SH** when choosing to use a convolutional architecture. However, **SUD** is perhaps less intuitive. Why should we suppose the derivatives are spatially uncorrelated? Conversely, why not go a step further and assume the *activations* are spatially uncorrelated (as do some methods; see Appendix C) or even drop all of the correlations (thereby obtaining a much simpler diagonal approximation to the Fisher matrix)?

Appendix D.1 analyzes empirically the validity of assumption **SUD** on conv nets trained to CIFAR-10 and SVHN. We conclude that **SUD** appears to describe the model distributions quite well for both networks. By contrast, the networks' activations have very strong spatial correlations, so it is significant that KFC does not assume spatially uncorrelated activations.

5.2. Optimization performance

We evaluated KFC-pre in the context of optimizing deep convolutional networks. We compared against stochastic gradient descent (SGD) with momentum, which is widely considered a strong baseline for training conv nets. All architectural choices (e.g. sizes of layers) were kept consistent with the previously published configurations. Since the focus of this work is optimization rather than generalization, metaparameters were tuned with respect to training error. This protocol was favorable to the SGD baseline, as the learning rates which performed the best on training error also performed the best on test error.³ We tuned the learning rates from the set $\{0.3, 0.1, 0.03, \dots, 0.0003\}$ separately for each experiment. For KFC-pre, we also chose several algorithmic parameters using the method of Appendix B.3, which considers only per-epoch running time and not final optimization performance.⁴

⁴For SGD, we used a momentum parameter of 0.9 and minibatches of size 128, which match the previously published config-

²https://code.google.com/p/cuda-convnet/

³For KFC-pre, we encountered a more significant tradeoff between training and test error, most notably in the choice of minibatch size, so the presented results do not reflect our best runs on the test set. For instance, as reported in Figure 1, the test error on CIFAR-10 leveled off at 18.5% after 5 minutes, after which the network started overfitting. When we reduced the mini-batch size from 512 to 128, the test error reached 17.5% after 5 minutes and 16% after 35 minutes. However, this run performed far worse on the training set. On the flip side, very large mini-batch sizes hurt generalization for both methods, as discussed in Section 5.3.

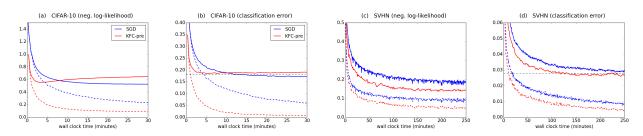


Figure 1. Optimization performance of KFC-pre and SGD. (a) CIFAR-10, negative log-likelihood. (b) CIFAR-10, classification error. (c) SVHN, negative log-likelihood. (d) SVHN, classification error. **Solid lines** represent test error and **dashed lines** represent training error. The **horizontal dashed line** represents the previously reported test error for the same architecture.

For both SGD and KFC-pre, we used an exponential moving average of the iterates (see Appendix B.1) with a timescale of 50,000 training examples (which corresponds to one epoch on CIFAR-10). This helped both SGD and KFC-pre substantially. All experiments for which wall clock time is reported were run on a single Nvidia GeForce GTX Titan Z GPU board.

As baselines, we also tried Adagrad (Duchi et al., 2011), RMSProp (Tieleman & Hinton, 2012), and Adam (Kingma & Ba, 2015), but none of these approaches outperformed carefully tuned SGD with momentum. This is consistent with the observations of Kingma & Ba (2015).

Figure 1(a,b) shows the optimization performance on the CIFAR-10 dataset, in terms of wall clock time. Both KFCpre and SGD reached approximately the previously published test error of 18% before they started overfitting. However, KFC-pre reached 19% test error in 3 minutes, compared with 9 minutes for SGD. The difference in training error was more significant: KFC-pre reaches a training error of 6% in 4 minutes, compared with 30 minutes for SGD. On SVHN, KFC-pre reached the previously published test error of 2.78% in 120 minutes, while SGD did not reach it within 250 minutes. (As discussed above, test error comparisons should be taken with a grain of salt.)

Appendix D.2 analyzes the performance of KFC-pre in relation to the recently proposed batch normalization method (Ioffe & Szegedy, 2015).

5.3. Potential for distributed implementation

Much work has been devoted recently to highly parallel or distributed implementations of neural network optimization (e.g. Dean et al. (2012)). Synchronous SGD effectively allows one to use very large mini-batches efficiently, which helps optimization by reducing the variance in the stochastic gradient estimates. However, the per-update performace levels off to that of batch SGD once the variance is no longer significant and curvature effects come to dominate. Asynchronous SGD partially alleviates this issue

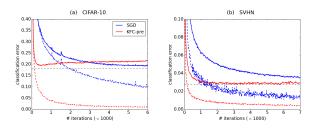


Figure 2. Classification error as a function of the number of iterations (weight updates). Heuristically, this is a rough measure of how the algorithms might perform in a highly distributed setting. (a) CIFAR-10. (b) SVHN. See Figure 1 caption for details.

by using new network parameters as soon as they become available, but needing to compute gradients with stale parameters limits the benefits of this approach.

As a proxy for how the algorithms are likely to perform in a highly distributed setting⁵, we measured the classification error as a function of the number of iterations (weight updates) for each algorithm. Both algorithms were run with large mini-batches of size 4096 (in place of 128 for SGD and 512 for KFC-pre). Figure 2 shows training curves for both algorithms on CIFAR-10 and SVHN, using the same architectures as above.⁶ KFC-pre required far fewer weight updates to achieve good training and test error compared with SGD. For instance, on CIFAR-10, KFC-pre obtained a training error of 10% after 300 updates, compared with 6000 updates for SGD, a 20-fold improvement. Similar speedups were obtained on test error and on the SVHN dataset. These results suggest that a distributed implementation of KFC-pre has the potential to obtain large speedups over distributed SGD-based algorithms.

urations. For KFC-pre, we used a momentum parameter of 0.9, mini-batches of size 512, and a damping parameter $\gamma = 10^{-3}$. In both cases, our informal explorations did not find other values which performed substantially better in terms of training error.

⁵The gradient computations can be farmed out to worker nodes, exactly as with SGD, and we expect the computations of Kronecker factors and their inverses can be performed asynchronously. Therefore, we would not expect additional sequential bottlenecks or communication overhead.

⁶Both SGD and KFC-pre reached a slightly worse test error before they started overfitting, compared with the smallminibatch experiments of the previous section. This is because large mini-batches lose the regularization benefit of stochastic gradients. One would need to adjust the regularizer in order to get good generalization performance in this setting.

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