

Improving Predictive Uncertainty Estimation using Dropout - Hamiltonian Monte Carlo

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Abstract Estimating predictive uncertainty is crucial for many computer vision tasks, from image classification to autonomous driving systems. Hamiltonian Monte Carlo (HMC) is an sampling method for performing Bayesian inference. On the other hand, Dropout regularization has been proposed as an approximate model averaging technique that tends to improve generalization in large scale models such as deep neural networks. Although, HMC provides convergence guarantees for most standard Bayesian models, it does not handle discrete parameters arising from Dropout regularization. In this paper, we present a robust methodology for improving predictive uncertainty in classification problems, based on Dropout and Hamiltonian Monte Carlo. Even though Dropout induces a non-smooth energy function with no such convergence guarantees, the resulting discretization of the Hamiltonian proves empirical success. The proposed method allows to effectively estimate the predictive accuracy and to provide better generalization for difficult test examples.

Keywords Bayesian learning · Hamiltonian Monte Carlo · Dropout · Transfer Learning · Classification.

1 Introduction

Artificial Intelligence systems have a wide variety of applications and in some cases are part of complex systems whose operation involves making delicate

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and dangerous decisions[20]. Uncertainty in knowledge representation and reasoning has been studied since the fall of symbolic expert systems [29]. Therefore, several research efforts focused on efficient methods for estimating model uncertainty and capturing the variability inherent to real world situations [6].

Inference using uncertainty estimates is crucial for many important tasks, such as image classification, detecting noisy examples (adversarial examples) and analyzing failure cases in decision-making systems [31]. These problems can benefit on predictive uncertainty for achieving good performance, requiring well calibrated probabilities. In such cases, Bayesian inference provides posterior predictive distributions that can be used to reduce over-confidence of the model outputs [12].

Bayesian neural networks consider model weights as random variables and have been proposed as a method to estimate predictive uncertainty [24], however inference is computationally intractable for deep neural networks. More recently, [11] proposed Monte Carlo Dropout as a method to obtain uncertainty estimates from deep learning architectures. Dropout has been previously proposed as a regularization technique for deep neural networks (see [34]) and the relationship with Bayesian model uncertainty being justified as a variational approximation. The key idea is to train a deep learning model and perform inference using Monte Carlo and dropout in order to obtain an approximate posterior distribution.

Hamiltonian Monte Carlo (HMC) is a Markov Chain Monte Carlo (MCMC) method for obtaining a sequence of random samples while maintaining asymptotic consistency with respect to a target distribution [25]. HMC provides a mechanism for defining proposals with high acceptance rate, enabling more efficient exploration of the state space than standard random-walk proposals. In addition, another property of HMC is the feasibility to support high dimensional models arising from deep neural architectures [5].

Although HMC was originally proposed for training Bayesian neural networks, the method is not well suited for discrete parameters arising from dropout layers. In this work, we propose a methodology named Dropout - Hamiltonian Monte Carlo (D-HMC). The proposed approach is tested with the MNIST digit recognition [19] and predictive uncertainty is compared to other sampling approaches such as Stochastic Gradient HMC (SGHMC) [9] and Stochastic Gradient Langevin Dynamics (SGLD) [39].

Transfer learning is a popular approach for domain adaption, when a large amount of data is used to pre-train a deep neural network and only a smaller amount of data is available for the new task [22]. The new model is prone to over-fitting, therefore a careful choice of the hyper-parameters used for transfer learning must be carried out in order to preserve the performance of the original task [23]. Conversely, we develop a novel methodology to represent predictive uncertainty when using convolutional neural networks for transfer learning for age recognition [10].

2 Related Work

HMC relies on a numerical approximation (an integrator) of a continuous dynamical system. The method is designed in a way that the main properties of the dynamical system are preserved. However, the approximation error is highly sensitive to critical user-specified hyper-parameters, such as the number of steps and the step size. Hoffman *et. al.* introduced the No-U-Turn sampler (NUTS) [17], as a method to determine the appropriate number of steps that the algorithm needs in order to converge to the target distribution. This method uses a recursive algorithm to build a set of likely candidate points, stopping automatically when it starts to double back and retrace its steps. In practice, simple heuristics can also be used to establish the number of steps and the step size, based on the assumption that the posterior distribution is Gaussian with diagonal covariance [16].

Other highly successful implementations focus on tackling large data sets using ideas from stochastic optimization such as Stochastic Gradient Descent (SGD). The SGHMC and SGLD techniques support data batches and introduces a novel integrator using Langevin dynamics which takes into account the extra-noise induced in the gradient [9, 39]. Another technique, the Metropolis Adjusted Langevin Algorithm (MALA) [32, 33], uses a Metropolis-Hastings (Metropolis) correction scheme to accept or reject proposals. Also, the Riemann Manifold Hamiltonian Monte Carlo method provides better adaption to the problem geometry for strongly correlated densities (RMHMC) [13, 37]. Finally, proximal algorithms and convex optimization techniques have also been studied in the context of MCMC [30], and HMC with log-concave or non-differentiable (non-smooth) energy functions [8].

3 Methods

Different architectures have been proposed for image classification using deep neural networks. However, most models have a fully connected layer that compares the network output \mathbf{x} with the sample training label y . Conversely, an image dataset $\mathbf{D} = \{\mathbf{d}_1, \dots, \mathbf{d}_N\}$ can be represented by a set of tuples $d = (\mathbf{x}, y)$ that contains the image features $\mathbf{x} \in \mathbb{R}^D$ and the labels $y = \{1, \dots, K\}$. The fully connected layer consists of an activation function such as the softmax, which encodes the image features into a vector of class probabilities $[\phi_1(\theta), \dots, \phi_k(\theta)]$, whose elements can be written as:

$$\phi_i(\theta) = \frac{\exp(\mathbf{x}^T \mathbf{w}_i + b_i)}{\sum_{k=1}^K \exp(\mathbf{x}^T \mathbf{w}_k + b_k)} \quad (1)$$

where $\theta = \{(\mathbf{w}_1, \mathbf{b}_1), \dots, (\mathbf{w}_K, \mathbf{b}_K)\}$, $\mathbf{w}_i \in \mathbb{R}^D$ is a vector, b_i is a scalar, D represents the dimensionality of the feature space \mathbf{x} and K the number of classes.

3.1 Hamiltonian Monte Carlo

Now, we consider the unknown parameter θ as a random variable. In a Bayesian setting, we want to sample from the posterior distribution $p(\theta|\mathbf{D})$ given by:

$$p(\theta|\mathbf{D}) = \frac{p(\mathbf{D}|\theta)p(\theta)}{p(\mathbf{D})} = \frac{p(\mathbf{D}|\theta)p(\theta)}{\int p(\mathbf{D}|\theta)p(\theta) d\theta} \quad (2)$$

$$\propto p(\mathbf{D}|\theta)p(\theta) \quad (3)$$

The target distribution $p(\theta|\mathbf{D})$ is known up to a normalization constant. Standard Markov Chain Monte Carlo methods such as the Metropolis-Hastings algorithm can be used to obtain the required posterior probabilities [12]. However due to the complexity of the target distribution (e.g. high-dimensional and possibly correlated image features), the algorithm would perform an inefficient exploration of the posterior. Instead, HMC improves the exploration by simulating Hamiltonian dynamics. A Hamiltonian function H is composed as a potential energy U and a kinetic energy K . These terms are constructed as follows:

$$H(\theta, r) = U(\theta) + K(r) \quad (4)$$

where r is called the momentum and is considered as an auxiliary variable. A positive-definite mass matrix M is also introduced as follow:

$$U(\theta) = - \sum_{d \in \mathbf{D}} \log p(d|\theta) - \log p(\theta) \quad (5)$$

$$K(r) = \frac{1}{2} r^T M^{-1} r \quad (6)$$

For image classification problems, the density $p(d|\theta) = \prod_{i=1}^K \phi_i(\theta)^{[y=k]}$ takes the form of a categorical distribution over the K different classes and the prior $p(\theta) \sim \mathcal{N}(0, \alpha \mathbf{I}_D)$ takes the form of a multivariate Gaussian distribution. Now, in order to sample from $p(\theta|\mathbf{D})$, the method simulates Hamiltonian dynamics while leaving the joint distribution (θ, r) invariant, such that:

$$p(\theta|\mathbf{D}) \propto \exp(-U(\theta)) \quad (7)$$

$$\pi(\theta, r) \propto \exp(-H(\theta, r)) \quad (8)$$

The first step proposes a new value for the momentum variable from a Gaussian distribution. Then, a Metropolis update using Hamiltonian dynamics is used to propose a new state. The state evolves in a fictitious continuous time t , and the partial derivatives of the Hamiltonian can be seen in Equation 10.

$$d\theta = M^{-1}r dt \quad (9)$$

$$dr = -\nabla U(\theta) dt \quad (10)$$

In order to simulate continuous dynamics, the leapfrog integrator can be used to discretize time. Using a small step size ϵ , Equation 10 becomes:

$$r_i^{(t+\frac{\epsilon}{2})} = r_i^{(t)} - \frac{\epsilon}{2}\nabla U(\theta^{(t)}) \quad (11)$$

$$\theta_i^{(t+\epsilon)} = \theta_i^{(t)} + \epsilon r^{(t+\frac{\epsilon}{2})} M_i^{-1} \quad (12)$$

$$r_i^{(t+\epsilon)} = r_i^{(t+\frac{\epsilon}{2})} - \frac{\epsilon}{2}\nabla U(\theta^{(t+\epsilon)}) \quad (13)$$

After $i = 1, \dots, m$ iterations of the leapfrog integrator with finite ϵ , the joint proposal becomes $(\theta^{(t+1)}, r^{(t+1)}) = (\theta_m, r_m)$. Subsequently, a Metropolis update is used to accept the proposal with probability $\rho > \text{unif}(0, 1)$, such that:

$$\rho = \min\{1, \exp(-H(\theta^{(t+1)}, r^{(t+1)})) + H(\theta^{(t)}, r^{(t)})\} \quad (14)$$

3.1.1 Properties of Hamiltonian dynamics

An integrator is an algorithm that numerically approximates an evolution of the exact solution of a differential equation. Some important properties of Hamiltonian dynamics are important for building MCMC updates [25]:

- Reversibility: The dynamics are time-reversible.
- Volume Preservation: Hamiltonian dynamics are volume preserving.
- Conservation of the Hamiltonian: H is constant as θ and r vary.

The leapfrog method in HMC satisfies the criteria of volume conservation and reversibility over time. However, the total energy is only conserved approximately, in this way a bias is introduced in the joint density $\pi(\theta, r)$. Conversely, the Metropolis update is used to satisfy the detailed balance condition.

3.1.2 Limitations of HMC

One of the main limitations of HMC is the lack of support for discrete parameters. The difficulty in extending HMC to a discrete parameter space stems from the fact that the construction of proposals relies on the numerical solution of a differential equation. In other hand, approximating the likelihood of a discrete parameter by a continuous density is not always possible [26]. Moreover, when any discontinuity is introduced into the energy function ($U(\theta)$), the first-order discretization does not ensure that a Metropolis correction maintains the stationary distribution invariant. Therefore, the standard implementation of HMC does not guarantee convergence when the parameter of interest has a discontinuous density. This is because integrators are designed for differential equations with smooth derivatives over time.

3.2 Dropout / DropConnect

Dropout and DropConnect arise in the context regularization of deep neural networks and provide a way to combine exponentially many different architectures [34, 36]. Dropout/DropConnect can be described as follows:

- Regular (binary) DropConnect adds noise to global network weights, by setting a randomly selected subset of weights to zero computed by element-wise multiplication of a binary mask Γ :

$$\phi_{c,i}(\theta, \Gamma) = \frac{\exp(\mathbf{x}^T(\Gamma \odot \mathbf{w}_i) + b_i)}{\sum_{k=1}^K \exp(\mathbf{x}^T(\Gamma \odot \mathbf{w}_k) + b_k)} \quad (15)$$

- Instead, binary Dropout randomly selects local inputs:

$$\phi_{d,i}(\theta, \Gamma) = \frac{\exp((\Gamma \odot \mathbf{x}^T)\mathbf{w}_i + b_i)}{\sum_{k=1}^K \exp((\Gamma \odot \mathbf{x}^T)\mathbf{w}_k + b_k)} \quad (16)$$

where $\Gamma \sim \mathcal{B}(1 - p)$ is a Bernoulli distributed random mask and p is the probability that the weight/input being dropped.

Due to its great effectiveness in various types of neural networks, the Dropout learning scheme has been the subject of research in recent years. Thus, it has been shown that Dropout is similar to bagging and other related ensemble methods [38]. Since all models are averaged in a efficient and fast approximation with weights scaling, Dropout can be seen as an approximation to the geometric mean in the space of all possible models, this approximation is less expensive than the arithmetic mean in bagging methods [3].

4 Dropout - Hamiltonian Monte Carlo

Stochastic gradient descent optimization has been extensively used for training neural networks. Instead of using the gradient of the full likelihood, stochastic gradients are used to update model parameters. In the context of Bayesian models, SGLD combines noisy gradient updates with Langevin dynamics in order to generate proposals from data subsets $B_i \subset \mathbf{D}$ such that:

$$\tilde{U}(\theta) = -\frac{N}{n} \sum_{d \in B_i} \log p(d|\theta, \Gamma) - \log p(\theta) \quad (17)$$

where $i = 1, \dots, N$, N is the number of data subsets and $n = |B_i|$ the size of the mini-batch and p the dropout rate.

Chen *et.al.* introduce a new momentum variable ν as an alternative discretization for HMC [9]. Similar to SGLD, the choice of the step size must

balance between efficient sampling and high acceptance rates (speed and accuracy). Equation 18 shows the SGHMC update.

$$\theta^{(t+1)} = \theta^{(t)} + \nu^{(t)} \quad (18a)$$

$$\nu^{(t+1)} = (1 - \alpha)\nu^{(t)} + \epsilon \nabla \tilde{U}(\theta^{(t+1)}) + \xi^{(t)} \quad (18b)$$

$$\xi^{(t)} = \mathcal{N}(0, 2[\alpha - \beta]\epsilon) \quad (18c)$$

where \mathcal{N} denotes a multivariate normal density, ν denotes the momentum variable in SGHMC, ϵ denotes the step size, while α and β are tuning constants.

Integrating Dropout into SGHMC can be seen as adding a regularization term. Since HMC cannot perform inference on the discrete parameters, the gradient can be computed by either marginalization or by means of a local reparameterization [18]. Therefore, on each iteration a multivariate Bernoulli mask $\Gamma \sim \mathcal{B}(1 - p)$ with probability $1 - p$ is applied to the θ parameter and then to the gradient of the energy function $\tilde{U}(\theta)$. Conversely, the density $p(d|\theta, \Gamma) = \prod_{i=1}^K \phi_{d,i}(\theta, \Gamma)^{[y=k]}$ takes the form of a modified categorical distribution. Each one of the input components is randomly deleted and the remaining elements are scaled up by $1/p$. The noisy gradient updates generate proposals from a perturbed target distribution [4]. The proposed method is described in algorithm 1.

Algorithm 1 D-SGHMC

Require: $\theta_*, M, p, \alpha, \beta, \epsilon$.

$v_* \sim \mathcal{N}(0, M)$

$(\theta_0, v_0) = (\theta_*, v_*)$

for $t = 1, 2, \dots$ **do**

Sample dropout mask Γ

$\Gamma \sim \mathcal{B}(1 - p)$

Transform mini-batch B_t using binary mask

$B_t^i \leftarrow 1/p (\Gamma \odot B_t)$

$(\theta^0, v^0) = (\theta_{t-1}, v_{t-1})$

for $i = 1, 2, \dots, L$ **do**

$\theta^i \leftarrow \theta^{i-1} + v^{i-1}$

$v^i \leftarrow (1 - \alpha)v^{i-1} + \epsilon \nabla \tilde{U}(\theta^i) + \mathcal{N}(0, 2[\alpha - \beta]\epsilon)$

end for

$(\theta_t, v_t) = (\theta^L, v^L)$

end for

return *Fully constructed Markov Chain*

SGHMC introduces a second-order term that reduces the discrepancy between the discretization noise and the stationary distribution, so it is no longer necessary to make a Metropolis correction. However, when dropout is incorporated, the energy function is no longer differentiable. In this context, the energy function of HMC requires a specially tailored discretization. Nishimura [26] finds a solution that preserves the critical properties of the Hamiltonian dynamics, through soft approximations, where the dynamics can be analytically integrated near the discontinuity in a way that preserves the total energy.

[27] have also shown that it is possible to build a discretization where the integrator maintains the irreversibility of the Markov chain and preserves energy, but volume preservation is no longer guaranteed [1]. Although dropout involves a discontinuous gradient, it is still possible to evaluate the gradient using automatic differentiation [2, 14], where the discontinuous parameters are taken to a continuous space [7].

5 Experiments

The SGHMC, SGLD and D-SGHMC algorithms were implemented in Edward [35], a Python library for posterior probabilistic modeling, inference, and criticism. We perform inference in two highly cited computer vision problems. These data sets were not selected with the goal of improving state-of-the-art results, but to evaluate whether we can incorporate uncertainty estimates on difficult examples.

In order to compare predictive uncertainty estimates obtained with the state-of-the-art sampling methods with the proposed method, the settings for the hyper-parameters are shown in Table 1.

Method	SGHMC	D-SGHMC	SGLD
Step Size (ϵ)	0.0001		
Friction Constant (α)	1.0		–
Mini-Batch Size	100		
Epochs	100		
Warmup	500		
Iterations	Epochs \times Num. Batches + Warmup		
Prediction Samples	30		
Dropout probability (p)	–	0.1 to 0.9	–

Table 1 Hyper-Parameters Setting

Training data is split in mini-batches of size $n = 100$ and whitening is performed on each one of these batches. On the other hand, predictive distribution is estimated using 30 Monte Carlo samples over the Test dataset

5.1 Digit Recognition on MNIST

In the first experiment, the predictive uncertainty on the the MNIST dataset is evaluated [19]. The database contains 60.000 training images and 10.000 test images, normalized to 28×28 pixels (784 features) and stored in gray scale. Figure 1 shows some examples of the database.

A total number of 5 independent chains are used for each method on MNIST. The Dropout rate is varied as $p \in [0.1, 0.5, 0.9]$, and the predictive results are compared with SGHMC and SGLD using the hyper-parameter setting established in the Table 1. The results can be seen in Table 2.



Fig. 1 Examples from the MNIST database.

Method	Total Accuracy (%)
SGHMC	90.94 ± 0.28
SGLD	88.06 ± 0.18
D-SGHMC ($p = 0.9$)	91.66 ± 0.10
D-SGHMC ($p = 0.5$)	91.72 ± 0.11
D-SGHMC ($p = 0.1$)	88.26 ± 0.08

Table 2 Test Accuracy for MNIST data Set. 5 independent chains.

Compared to other sampling techniques, the results show that D-SGHMC obtains a lower error when the dropout probability is set to 0.5, which is consistent to the state-of-the-art results in terms of linear classification methods [19].

The role of the Dropout probability (p) for D-SGHMC is also studied. Figure 2 show accuracy results with the execution of 5 independent chains. It is possible to establish that for the studied data set a dropout probability between on 0.6 and 0.8 generates lower error rates. On the other hand, it can also be seen that probabilities lower than 0.4 rapidly decreases the variable's influence on the classification results.

Figure 3 shows the correlation matrix between the true class and the predicted class. Higher class uncertainty is achieved when classifying digits 9 and 8 of MNIST, which could be classified as digits 4 and 5 respectively.

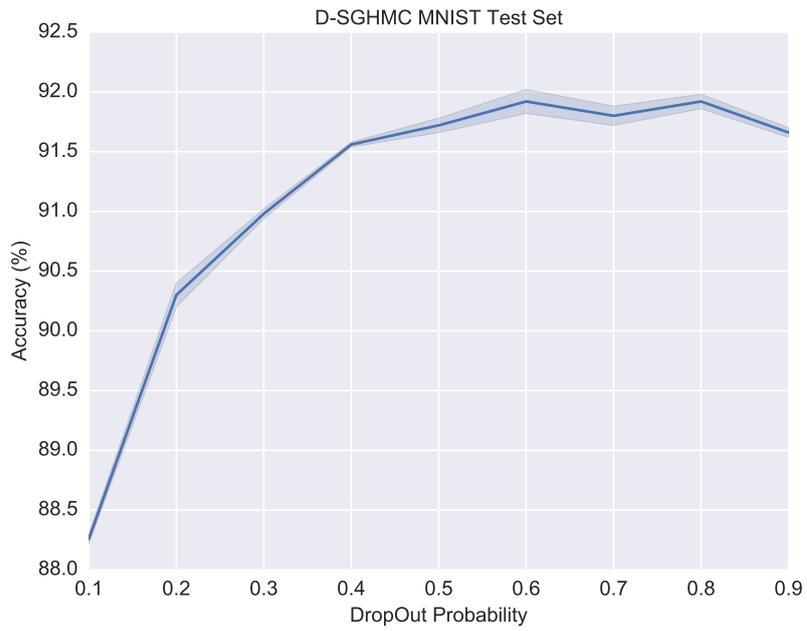


Fig. 2 D-SGHMC Sensitivity analysis for p hyper-parameter. 5 independent chains on MNIST.

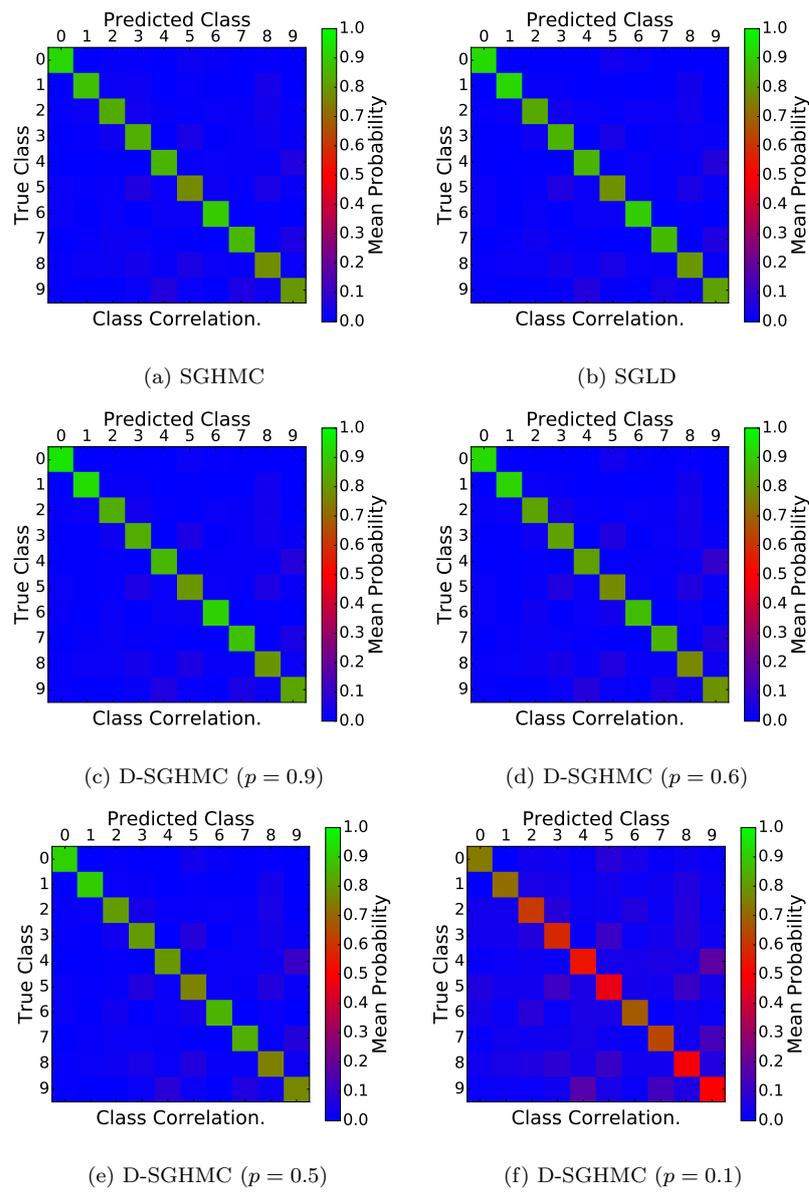


Fig. 3 Matrix Correlation for error rates on MNIST.

Predictive accuracy is now evaluated by comparing the different methods, where the expected predictive accuracy is computed using a Monte Carlo approximation. D-SGHMC (Fig. 4.a and 4.b) achieves higher predictive accuracy when compared to SGHMC (Fig. 4.c) and SGLD (Fig. 4.d).

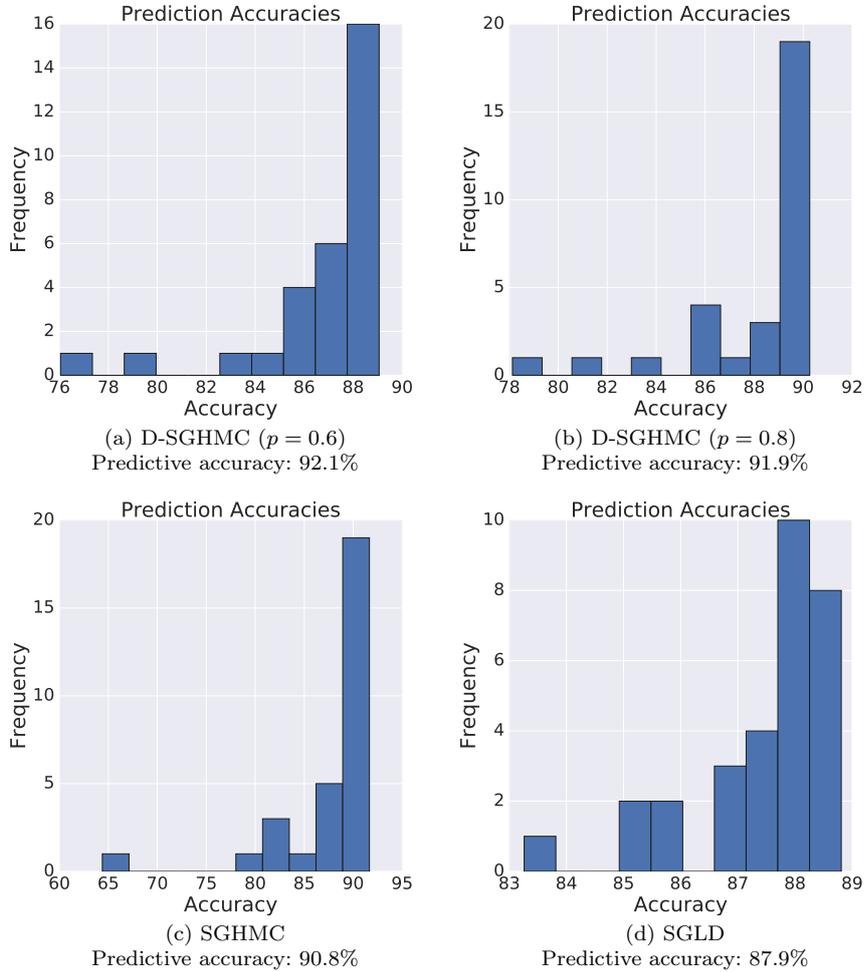


Fig. 4 Histogram of predictive uncertainty.

5.1.1 Confusing classes

One of the biggest challenges of the MNIST data set is to separate highly confusing digits such as '4' and '9'. This problem has been addressed earlier in feature selection challenges [15].

We analyze the behavior between total uncertainty and predictive accuracy for one of these digits in each method. We can observe that there is less over-confidence and at the same time the classification results improve. Figure 5 shows the mean expected probabilities for digit 9 in the test set.

As the Dropout rate decreases, the uncertainty around similar classes becomes more visible. However, low Dropout probabilities does not guarantee better classification results or improved uncertainty estimates. As shown in the Figure 5, models generated with low Dropout rates aggressively reduce the number of variables used for prediction.

5.1.2 Confusing Example (digit 9)

In the first example (Fig. 6.a), both state-of-the-art SGHMC (Fig. 6.b) and SGLD methods (Fig. 6.c) maintain a high confidence.

In the example we can see how the proposed method decreases the over-confidence generated by SGHMC and SGLD and allows to classify the example effectively, producing higher uncertainty (Fig. 6.d, 6.e and 6.f).

5.1.3 Confusing Example (digit 8)

In the second example (Fig. 7.a), we observe that it is similar to the digits '4', '5' and even '1', for thus SGHMC (Fig. 7.b) and SGLD (Fig. 7.c) maintain a high confidence in this digits.

In this example, D-SGHMC does not significantly improve the classification results (Fig. 7.d and 7.f). However, it does improve the uncertainty estimates, generating higher confidence in the true label (Fig. 7.e).

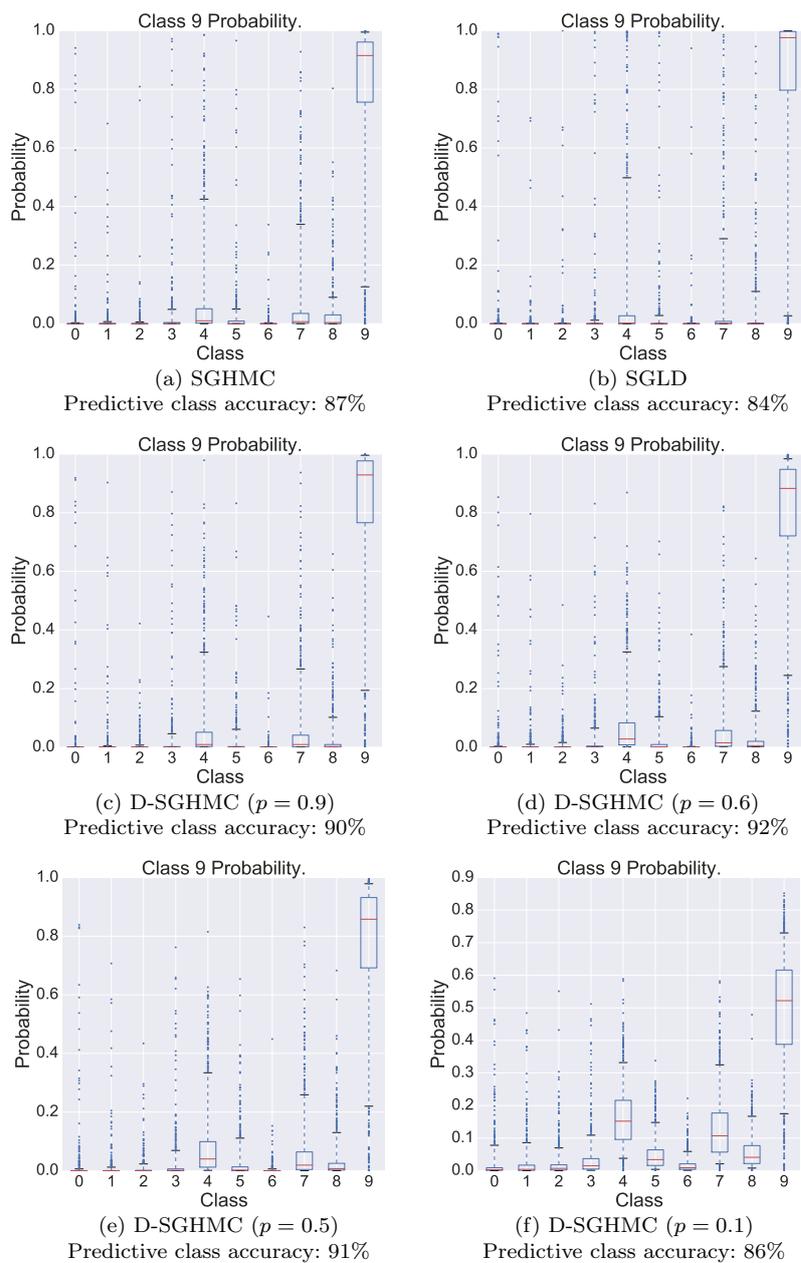


Fig. 5 Box plot of the error rates for class 9.

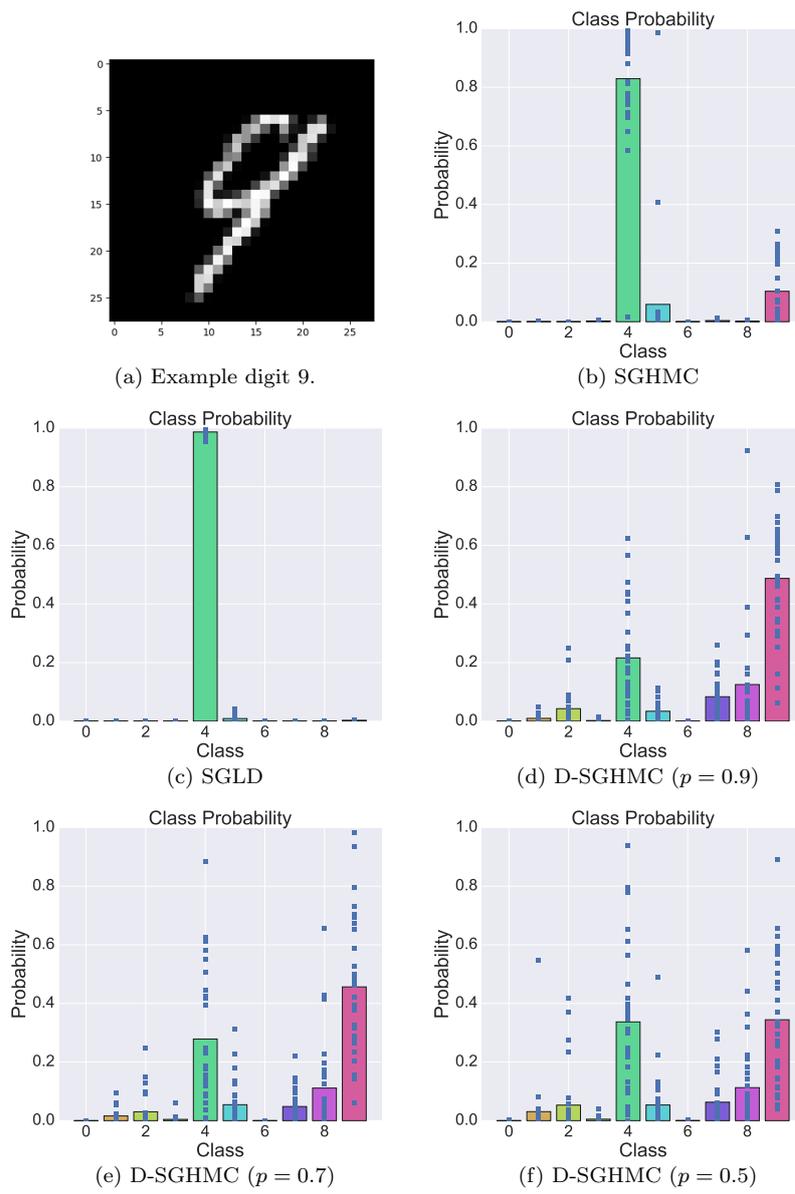


Fig. 6 Bar plot of error rates for confusing example 9.

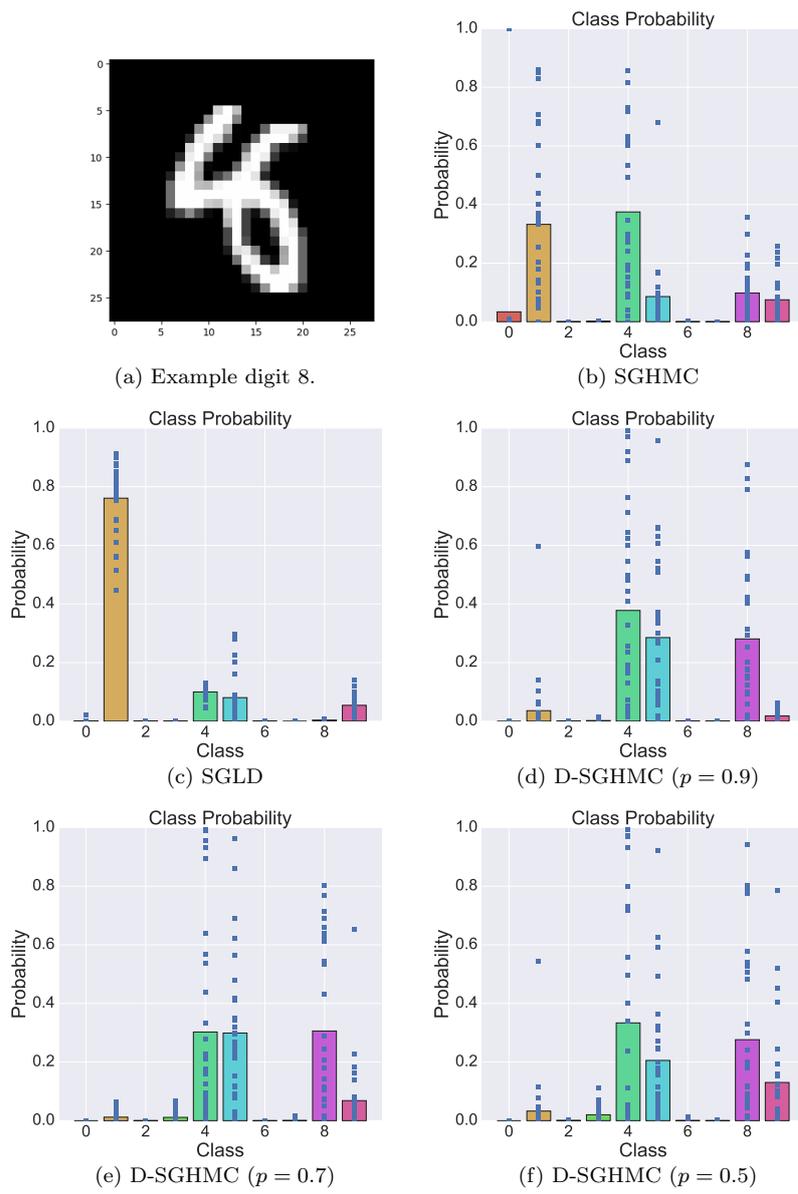


Fig. 7 Bar plot of error rates for confusing example 8.



Fig. 8 Examples from ADIANCE database.

5.2 Age Recognition on ADIANCE

As part of the many challenges of facial recognition systems, age recognition has been recognized as difficult problem [10]. The ADIANCE dataset (see Fig. 8) contains 26.580 images of 2.284 different subjects and has been used to study the performance of age and gender recognition systems. After pre-processing and cleaning, the resulting data set is partitioned into 13.000 training examples and 3.534 test examples. The final number of samples can be seen in Table 3.

Age	ID Class	Train	Test	Total
[0 – 2]	0	1.201	199	1.400
[4 – 6]	1	1.566	573	2.139
[8 – 13]	2	1.942	343	2.285
[15 – 20]	3	1.385	255	1.640
[25 – 32]	4	3.940	1.099	5.039
[38 – 43]	5	1.794	546	2.340
[48 – 53]	6	579	246	825
[60–]	7	593	273	866
Total	8	13.000	3.534	16.534

Table 3 Final examples distribution of the ADIANCE data set.

Transfer learning from VGG-Face CNN (Convolutional Neural Networks)[28] with AVG pooling is used as a convolutional descriptor. For each example of ADIANCE database, VGG-Face computes a descriptor of 512 features.

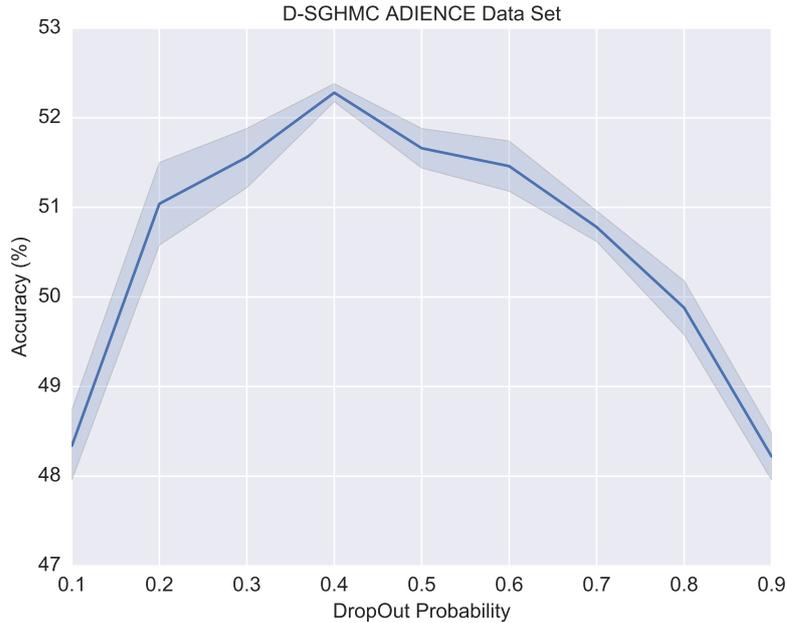


Fig. 9 Sensitivity analysis for D-SGHMC with Dropout rate p .

A total number of 5 independent runs is used for each one of the methods. Using the hyper-parameter setting set in table 1, where state-of-art results are achieved [21]. Comparison with baseline methods can be seen in Table 4.

Method	Total Accuracy (%)
SGHMC	45.6 ± 1.23
SGLD	44.0 ± 1.10
D-SGHMC($p = 0.9$)	48.2 ± 0.56
D-SGHMC($p = 0.5$)	51.6 ± 0.51
D-SGHMC($p = 0.1$)	48.3 ± 0.88

Table 4 Test accuracy for ADIENCE data Set.

The role of the Dropout rate p on the performance is analyzed. As shown in Figure 9, best results are obtained with values between 0.3 and 0.5.

In ADIENCE, uncertainty is increased for neighboring classes. This can be seen in Figure 10, which shows the true and the predicted labels.

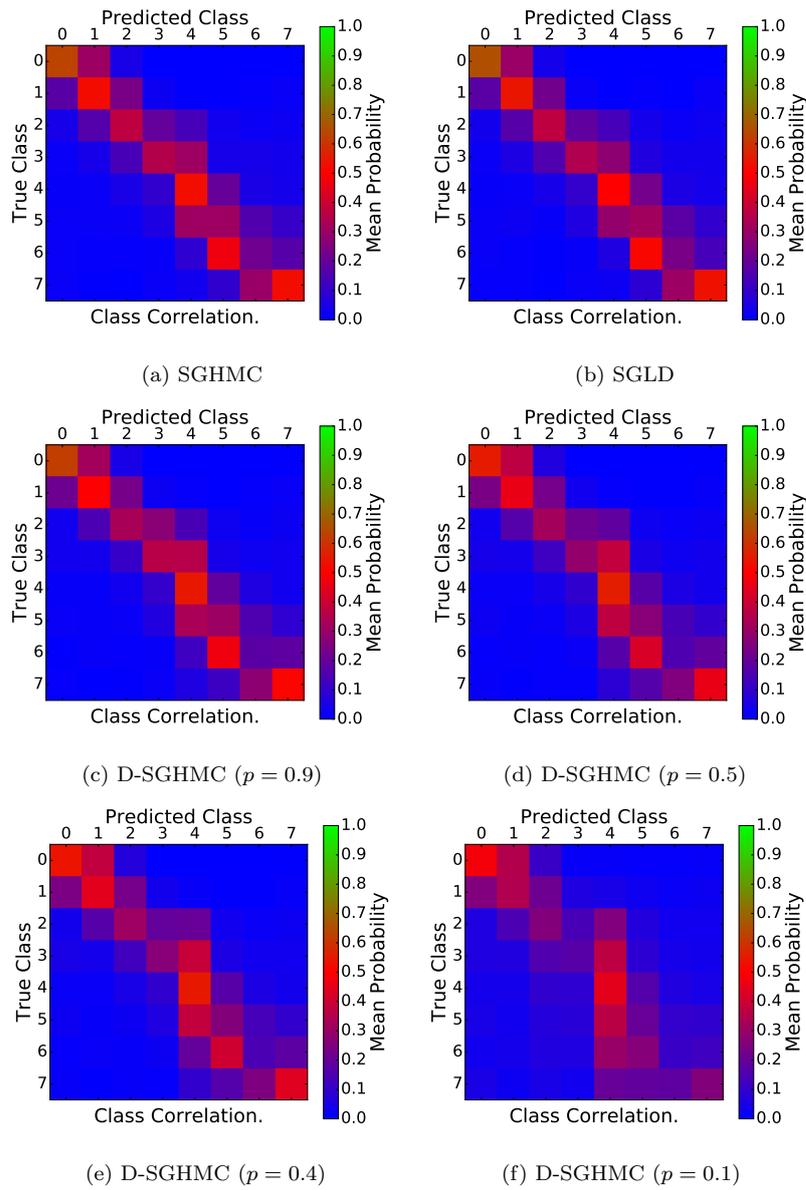


Fig. 10 Matrix Correlation plot of probabilities for all classes. 5 independent chains on ADIENCE.

5.2.1 Accuracy

Class imbalance plays an important role in the final classification results for the age recognition problem in ADIANCE. D-SGHMC (see Figure 11.d) improves results in most of classes, sacrificing to a lesser extent the accuracy for classes with less number of examples.

As the Dropout rate decreases, the imbalance problem becomes more evident and the accuracy results are also dropped (see Figure 9).

5.2.2 Predictive Uncertainty for the Adience Dataset

In the ADIANCE data set there are many confusing examples which are usually neighboring classes. One example for each class is randomly sampled and the predictive distributions for each model are evaluated. These results are shown in Figure 12. In general, SGHMC and SGLD produce over-confident probabilities, assigning high values to incorrect labels. In contrast, D-SGHMC reduces that confidence, increasing uncertainty in the class predictions.

In particular, high confidence estimates for both SGHMC and SGLD on one example of class 4 can be seen in Figure 12. On the other hand, the proposed method allows to improve the uncertainty estimates between the neighboring classes, which significantly improves the classification results.

This situation is also replicated on one example of class 7, where SGHMC and SGLD achieves over-confident misclassification (class 1). The proposed method allows to alleviate the misclassification error and return uncertainty estimates for neighboring classes. The predictive distribution improves the classification results and provides the improved uncertainty estimates.

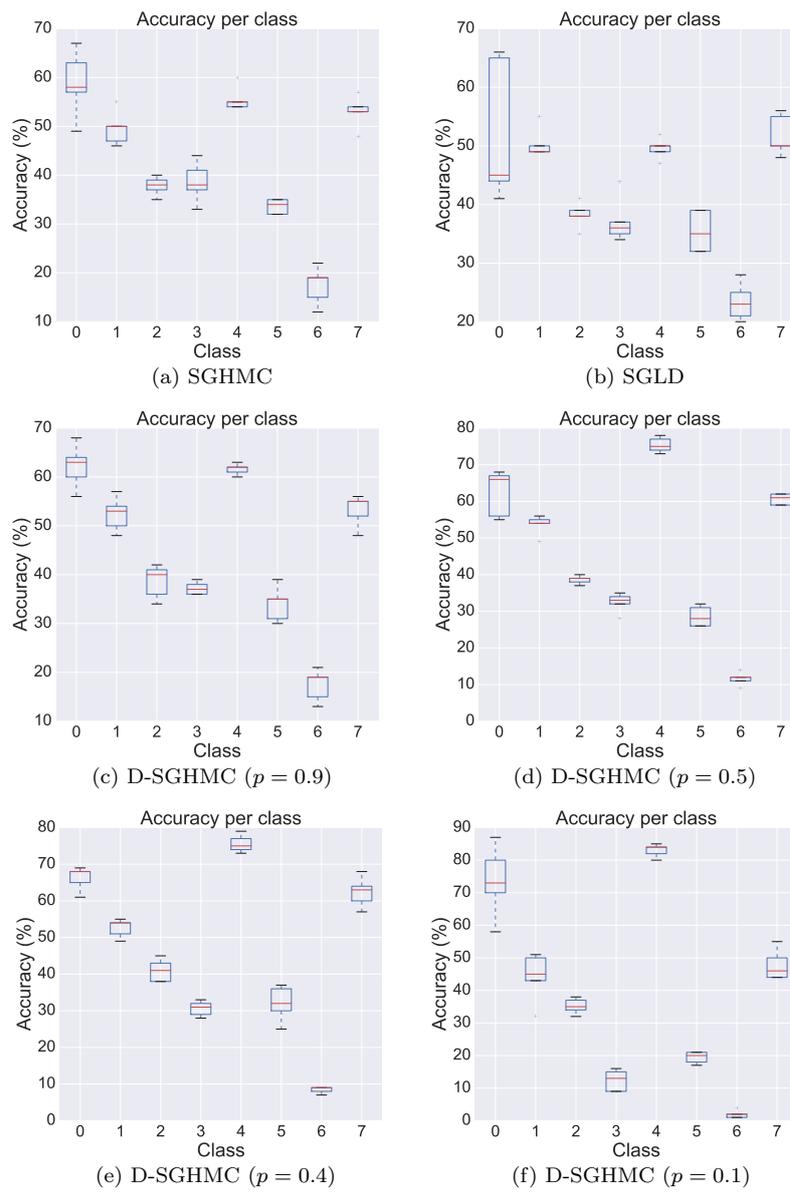


Fig. 11 Box plot of accuracies for all classes on ADIENCE.



Fig. 12 Predictive distribution for randomly selected samples from the ADIANCE dataset.

6 Conclusion

In many decision-making systems, the estimated uncertainty plays a crucial role in the final classification. Over-confident estimates can arise from noisy examples or when leading with classes that were not part of the training data set. The fundamental objective of achieving improved uncertainty estimates is to encourage decision-makers to question their decisions. In this way, compared to other state-of-the-art MCMC methods for large scale and high dimensional classification problems, the methodology presented in this paper improves the uncertainty estimated.

The proposed method is based on HMC and Dropout regularization. The experiments demonstrated that the method is capable of generating an approximation to the posterior distribution. In addition, the resulting predictive distributions also alleviate the misclassification error in difficult examples. However, it has not yet been proven that generated stochastic dynamics preserve the volume in its entirety. Future work will perform comparisons with other state-of-the-art variational methods. Moreover, the relationship between the proposed approach to approximate Bayesian model averaging can be also another line of research.

Compliance with Ethical Standards

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