FLOW++: IMPROVING FLOW-BASED GENERATIVE MODELS WITH VARIATIONAL DEQUANTIZATION AND ARCHITECTURE DESIGN

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ABSTRACT

Flow-based generative models are powerful exact likelihood models with the benefit of efficient sampling and inference. Despite their computational efficiency, flow-based models generally have much worse density modeling performance compared to state-of-the-art autoregressive models. In this paper, we carefully investigate 3 design choices employed by prior flow-based models that turn out to be limiting: (1) uniform noise is a sub-optimal dequantization choice that hurts both training and generalization loss; (2) commonly used affine coupling flows are not expressive enough; (3) conv-net based conditioning architecture of flows fails to capture the global image context. Based on our findings, we propose Flow++, a set of alternative design choices that significantly improve the density modelling capacity of flow-based models. Our validation metrics - 3.13 bits per dim (b.p.d) on CIFAR10, 3.91 b.p.d on 32x32 Imagenet, have started to close the significant performance gap that has so far existed between autoregressive models and flow-based models.

1 Introduction

Advances in deep generative models – such as latent variable models like variational autoencoders (Kingma & Welling, 2013), implicit generative models like GANs (Goodfellow et al., 2014), and exact likelihood models like PixelRNN/CNN (van den Oord et al., 2016a;c), Image Transformer (Parmar et al., 2018), PixelSNAIL (Chen et al., 2017), NICE, RealNVP, and Glow (Dinh et al., 2014; 2016; Kingma & Dhariwal, 2018) – have enabled us to model high dimensional raw observations from complex real-world datasets, from natural images and videos, to audio signals and natural language (Karras et al., 2017; Kalchbrenner et al., 2016b; Oord et al., 2016a; Kalchbrenner et al., 2016a; Vaswani et al., 2017).

Autoregressive models, a certain subclass of exact likelihood models, have recently achieved state-of-the-art density estimation performance on many challenging real-world datasets, but suffer from slow sampling time due to their autoregressive structure (Oord et al., 2016b; Salimans et al., 2017; Chen et al., 2017; Parmar et al., 2018). Non-autoregressive flow-based models (which we will refer to as "flow models") form another subclass of exact likelihood models. These models, such as NICE, RealNVP, and Glow, are efficient for sampling, but have so far lagged behind autoregressive models in density estimation benchmarks (Dinh et al., 2014; 2016; Kingma & Dhariwal, 2018).

Within the family of exact likelihood models, it is currently impossible to have all 3 of the following attractive characteristics: (a) Fast enough generation for real-world use cases; (b) Efficient inference, likelihood evaluation at any given data point x, for maximum-likelihood (M-projection) training; (c) Expressive modelling capacity as measured by standard density modelling benchmarks like CIFAR10 and 32x32 ImageNet. Autoregressive models achieve (b) and (c) but are very slow at generation due to the sequential dependency between dimensions; Inverse Autoregressive models (Kingma et al., 2016b) have (a) and possibly (c) (Oord et al., 2017) but can't be trained efficiently under maximum-likelihood framework; flow models enjoy efficient generation (a) and inference (b) but are lacking behind in terms of capacity.

In the hope of creating an ideal likelihood-based generative model that achieves (a,b,c), we seek to close the density modelling performance gap between flow models and autoregressive models by

first identifying the inefficiency in common flow model designs and then presenting better alternatives. In subsequent sections, we will first formally define the generic flow model family and then describe our new flow model, Flow++, which is powered by an improved training procedure for continuous likelihood models and a number of architectural extensions of the coupling layer defined by Dinh et al. (2014; 2016); Kingma & Dhariwal (2018).

2 FLOW MODELS

A flow model f is constructed as an invertible transformation that maps observed data \mathbf{x} to a standard Gaussian latent variable $\mathbf{z}=f(\mathbf{x})$, as in nonlinear independent component analysis (Bell & Sejnowski, 1995; Hyvärinen et al., 2004; Hyvärinen & Pajunen, 1999). The key idea in the design of a flow model is to form f by stacking individual simple invertible transformations (Dinh et al., 2014; 2016; Kingma & Dhariwal, 2018; Rezende & Mohamed, 2015; Kingma et al., 2016b; Louizos & Welling, 2017). Explicitly, f is constructed by composing a series of invertible flows as $f(\mathbf{x})=f_1\circ\cdots\circ f_L(\mathbf{x})$, with each f_i having a tractable inverse and a tractable Jacobian determinant. This way, sampling is efficient, as it can be performed by computing $f^{-1}(\mathbf{z})=f_L^{-1}\circ\cdots\circ f_1^{-1}(\mathbf{z})$ for $\mathbf{z}\sim\mathcal{N}(\mathbf{0},\mathbf{I})$, and training by maximum likelihood is computationally efficient too, since

$$\log p(\mathbf{x}) = \log \mathcal{N}(f_1 \circ \cdots f_L(\mathbf{x}); \mathbf{0}, \mathbf{I}) + \sum_{i=1}^{L} \log \left| \frac{\partial f_i}{\partial f_{i-1}} \right|$$
(1)

is easy to compute and differentiate with respect to the parameters of the flows f_i .

3 FLOW++

In this section, we describe three main axes of modelling inefficiencies in prior flow models: (1) uniform noise is a sub-optimal dequantization choice that hurts both training loss and generalization; (2) commonly used affine coupling flows are not expressive enough; (3) conv-net based parametrization of flows fails to capture the global image context. Our proposed model, Flow++, consists of a set of improved design choices: (1) variational flow-based dequantization as opposed to uniform dequantization; (2) logistic mixture CDF coupling flows; (3) self-attention in the coupling flows.

3.1 DEQUANTIZATION VIA VARIATIONAL INFERENCE

Many real-world datasets, such as CIFAR10 and ImageNet, are recordings of continuous signals quantized into discrete representations. Fitting a continuous density model to discrete data, however, will produce a degenerate solution that places all probability mass on discrete datapoints (Uria et al., 2013). A common solution to this problem is to first convert the discrete data distribution into a continuous distribution via a process called "dequantization," and then model the resulting continuous distribution using the continuous density model (Uria et al., 2013; Dinh et al., 2016; Salimans et al., 2017).

3.1.1 Uniform dequantization

Dequantization is usually performed in prior work by adding uniform noise to the discrete data over the width of each discrete bin: if each of the D components of the discrete data ${\bf x}$ takes on values in $\{0,1,2,\ldots,255\}$, then the dequantized data is given by ${\bf y}={\bf x}+{\bf u}$, where ${\bf u}$ is drawn uniformly from $[0,1)^D$. Theis et al. (2015) note that training a continuous density model $p_{\rm model}$ on uniformly dequantized data ${\bf y}$ can be interpreted as maximizing a lower bound on the log-likelihood for a certain discrete model $P_{\rm model}$ on the original discrete data ${\bf x}$:

$$P_{\text{model}}(\mathbf{x}) := \int_{[0,1)^D} p_{\text{model}}(\mathbf{x} + \mathbf{u}) d\mathbf{u}$$
 (2)

The argument of Theis et al. (2015) proceeds as follows. Letting $P_{\rm data}$ denote the original distribution of discrete data and $p_{\rm data}$ denote the distribution of uniformly dequantized data, Jensen's

inequality implies that

$$\mathbb{E}_{\mathbf{y} \sim p_{\text{data}}} \left[\log p_{\text{model}}(\mathbf{y}) \right] = \sum_{\mathbf{x}} P_{\text{data}}(\mathbf{x}) \int_{[0,1)^D} \log p_{\text{model}}(\mathbf{x} + \mathbf{u}) d\mathbf{u}$$
 (3)

$$\leq \sum_{\mathbf{x}} P_{\text{data}}(\mathbf{x}) \log \int_{[0,1)^D} p_{\text{model}}(\mathbf{x} + \mathbf{u}) d\mathbf{u}$$
 (4)

$$= \mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \left[\log P_{\text{model}}(\mathbf{x}) \right] \tag{5}$$

Consequently, maximizing the log-likelihood of the continuous model on uniformly dequantized data cannot lead to the continuous model degenerately collapsing onto the discrete data, because its objective is bounded above by the log-likelihood of a discrete model.

3.1.2 Variational dequantization

While uniform dequantization successfully prevents the continuous density model $p_{\rm model}$ from collapsing to a degenerate mixture of point masses on discrete data, it asks $p_{\rm model}$ to assign uniform density to unit hypercubes ${\bf x} + [0,1)^D$ around the data ${\bf x}$. It is difficult and unnatural for smooth function approximators, such as neural network density models, to excel at such a task. To sidestep this issue, we now introduce a new dequantization technique based on variational inference.

Again, we are interested in modeling D-dimensional discrete data $\mathbf{x} \sim P_{\text{data}}$ using a continuous density model p_{model} , and we will do so by maximizing the log-likelihood of its associated discrete model $P_{\text{model}}(\mathbf{x}) \coloneqq \int_{[0,1)^D} p_{\text{model}}(\mathbf{x}+\mathbf{u}) \, d\mathbf{u}$. Now, however, we introduce a dequantization noise distribution $q(\mathbf{u}|\mathbf{x})$, with support over $\mathbf{u} \in [0,1)^D$. Treating q as an approximate posterior, we have the following variational lower bound, which holds for all q:

$$\mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \left[\log P_{\text{model}}(\mathbf{x}) \right] = \mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \left[\log \int_{[0,1)^D} q(\mathbf{u}|\mathbf{x}) \frac{p_{\text{model}}(\mathbf{x} + \mathbf{u})}{q(\mathbf{u}|\mathbf{x})} d\mathbf{u} \right]$$
(6)

$$\geq \mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \left[\int_{[0,1)^D} q(\mathbf{u}|\mathbf{x}) \log \frac{p_{\text{model}}(\mathbf{x} + \mathbf{u})}{q(\mathbf{u}|\mathbf{x})} d\mathbf{u} \right]$$
(7)

$$= \mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \mathbb{E}_{\mathbf{u} \sim q(\cdot | \mathbf{x})} \left[\log \frac{p_{\text{model}}(\mathbf{x} + \mathbf{u})}{q(\mathbf{u} | \mathbf{x})} \right]$$
(8)

We will choose q itself to be a conditional flow-based generative model of the form $\mathbf{u}=q_{\mathbf{x}}(\boldsymbol{\epsilon})$, where $\boldsymbol{\epsilon}\sim p(\boldsymbol{\epsilon})=\mathcal{N}(\boldsymbol{\epsilon};\mathbf{0},\mathbf{I})$ is Gaussian noise. In this case, $q(\mathbf{u}|\mathbf{x})=p(q_{\mathbf{x}}^{-1}(\mathbf{u}))\cdot\left|\partial q_{\mathbf{x}}^{-1}/\partial \mathbf{u}\right|$, and thus we obtain the objective

$$\mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \left[\log P_{\text{model}}(\mathbf{x}) \right] \ge \mathbb{E}_{\mathbf{x} \sim P_{\text{data}}, \boldsymbol{\epsilon} \sim p} \left[\log \frac{p_{\text{model}}(\mathbf{x} + q(\mathbf{x}, \boldsymbol{\epsilon}))}{p(\boldsymbol{\epsilon}) \left| \partial q_{\mathbf{x}} / \partial \boldsymbol{\epsilon} \right|^{-1}} \right]$$
(9)

which we maximize jointly over p_{model} and q. When p_{model} is also a flow model $\mathbf{x} = f^{-1}(\mathbf{z})$ (as it is throughout this paper), it is straightforward to calculate a stochastic gradient of this objective through pathwise derivative, as $f(\mathbf{x} + q(\mathbf{x}, \boldsymbol{\epsilon}))$ is differentiable with respect to the parameters of f and q.

Notice that the lower bound for uniform dequantization – eqs. (3) to (5) – is a special case our variational lower bound – eqs. (6) to (8) – when the dequantization distribution q is a uniform distribution that ignores dependence on \mathbf{x} . Because the gap between our objective (8) and the true expected log-likelihood $\mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \left[\log P_{\text{model}}(\mathbf{x}) \right]$ is exactly $\mathbb{E}_{\mathbf{x} \sim P_{\text{data}}} \left[D_{\text{KL}} \left(q(\mathbf{u}|\mathbf{x}) \parallel p_{\text{model}}(\mathbf{u}|\mathbf{x}) \right) \right]$, using a uniform q both forces the variational lower bound to stay loose due to inexpressiveness of q and forces p_{model} to unnaturally place uniform density over each hypercube $\mathbf{x} + [0, 1)^D$. Using an expressive flow-based q, on the other hand, allows p_{model} to place density in each hypercube $\mathbf{x} + [0, 1)^D$ according to a much more flexible distribution $q(\mathbf{u}|\mathbf{x})$. This is a more natural task for p_{model} to perform, improving both training and generalization loss.

3.2 IMPROVED COUPLING LAYERS

Recent progress in the design of flow models has involved carefully constructing flows to increase their expressiveness while preserving tractability of the inverse and Jacobian determinant computations. One example is the invertible 1×1 convolution flow, whose inverse and Jacobian determinant can be calculated and differentiated with standard automatic differentiation libraries (Kingma & Dhariwal, 2018). Another example, which we build upon in our work here, is the affine coupling layer (Dinh et al., 2016). It is a parameterized flow $\mathbf{y} = f_{\theta}(\mathbf{x})$ that first splits the components of \mathbf{x} into two parts $\mathbf{x}_1, \mathbf{x}_2$, and then computes $\mathbf{y} = (\mathbf{y}_1, \mathbf{y}_2)$, given by

$$\mathbf{y}_1 = \mathbf{x}_1, \qquad \mathbf{y}_2 = \mathbf{x}_2 \cdot \exp(\mathbf{a}_{\theta}(\mathbf{x}_1)) + \mathbf{b}_{\theta}(\mathbf{x}_1)$$
 (10)

Here, \mathbf{a}_{θ} and \mathbf{b}_{θ} are outputs of a neural network that acts on \mathbf{x}_1 in a complex, expressive manner, but the resulting behavior on \mathbf{x}_2 always remains an elementwise affine transformation – effectively, \mathbf{a}_{θ} and \mathbf{b}_{θ} together form a data-parameterized family of invertible affine transformations. This allows the affine coupling layer to express complex dependencies on the data while keeping inversion and log-likelihood computation tractable. Using \cdot and \exp to respectively denote elementwise multiplication and exponentiation,

$$\mathbf{x}_1 = \mathbf{y}_1, \quad \mathbf{x}_2 = (\mathbf{y}_2 - \mathbf{b}_{\theta}(\mathbf{y}_1)) \cdot \exp(-\mathbf{a}_{\theta}(\mathbf{y}_1)), \quad \log \left| \frac{\partial \mathbf{y}}{\partial \mathbf{x}} \right| = \mathbf{1}^{\top} \mathbf{a}_{\theta}(\mathbf{x}_1)$$
 (11)

3.2.1 Expressive coupling transformations with continuous mixture CDFs

We found in our experiments that density modeling performance of these coupling layers could be improved by augmenting the data-parameterized elementwise affine transformations by more general nonlinear elementwise transformations. For a given scalar component x of \mathbf{x}_2 , we apply the cumulative distribution function (CDF) for a mixture of K logistics – parameterized by mixture probabilities, means, and log scales π , μ , \mathbf{s} – followed by an inverse sigmoid and an affine transformation parameterized by a and b:

$$x \longmapsto \sigma^{-1}\left(\text{MixLogCDF}(x; \boldsymbol{\pi}, \boldsymbol{\mu}, \mathbf{s})\right) \cdot \exp(a) + b$$
 (12)

where
$$\operatorname{MixLogCDF}(x; \boldsymbol{\pi}, \boldsymbol{\mu}, \mathbf{s}) \coloneqq \sum_{i=1}^{K} \pi_i \sigma\left((x - \mu_i) \cdot \exp(-s_i)\right)$$
 (13)

The transformation parameters π , μ , \mathbf{s} , a, b for each component of \mathbf{x}_2 are produced by a neural network acting on \mathbf{x}_1 . This neural network must produce these transformation parameters for each component of \mathbf{x}_2 , hence it produces vectors $\mathbf{a}_{\theta}(\mathbf{x}_1)$ and $\mathbf{b}_{\theta}(\mathbf{x}_1)$ and tensors $\pi_{\theta}(\mathbf{x}_1)$, $\mu_{\theta}(\mathbf{x}_1)$, $\mathbf{s}_{\theta}(\mathbf{x}_1)$ (with last axis dimension K). The coupling transformation is then given by:

$$\mathbf{y}_1 = \mathbf{x}_1, \qquad \mathbf{y}_2 = \sigma^{-1}\left(\text{MixLogCDF}(\mathbf{x}_2; \boldsymbol{\pi}_{\theta}(\mathbf{x}_1), \boldsymbol{\mu}_{\theta}(\mathbf{x}_1), \mathbf{s}_{\theta}(\mathbf{x}_1))\right) \cdot \exp(\mathbf{a}_{\theta}(\mathbf{x}_1)) + \mathbf{b}_{\theta}(\mathbf{x}_1)$$
(14)

where the formula for computing y_2 operates elementwise.

The inverse sigmoid ensures that the inverse of this coupling transformation always exists: the range of the logistic mixture CDF is (0,1), so the domain of its inverse must stay within this interval. The CDF itself can be inverted efficiently with bisection, because it is a monotonically increasing function. Moreover, the Jacobian determinant of this transformation involves calculating the probability density function of the logistic mixtures, which poses no computational difficulty.

3.2.2 Expressive conditioning architectures by introducing self-attention

In addition to improving the expressiveness of the elementwise transformations on \mathbf{x}_2 , we found it crucial to improve the expressiveness of the conditioning on \mathbf{x}_1 – that is, the expressiveness of the neural network responsible for producing the elementwise transformation parameters π , μ , \mathbf{s} , \mathbf{a} , \mathbf{b} . Our best results were obtained by stacking convolutions and multi-head self attention into a gated residual network (Mishra et al., 2018; Chen et al., 2017), in a manner resembling the Transformer (Vaswani et al., 2017) with pointwise feedforward layers replaced by 3×3 convolutional layers. Our architecture is defined as a stack of blocks. Each block consists of the following two layers connected in a residual fashion, with layer normalization (Ba et al., 2016) after each residual connection:

Conv = input
$$\rightarrow$$
 nonlinearity \rightarrow conv_{3×3} \rightarrow nonlinearity \rightarrow gate
Attn = input \rightarrow conv_{1×1} \rightarrow MultiHeadSelfAttention \rightarrow gate

where gate refers to a 1×1 convolution that doubles the number of channels, followed by a gated linear unit (Dauphin et al., 2016). The convolutional layer is identical to the one used by PixelCNN++

(Salimans et al., 2017), and the multi-head self attention mechanism we use is identical to the one in the Transformer (Vaswani et al., 2017). (We always use 4 heads in our experiments, since we found it to be effective early on in our experimentation process.)

With these blocks in hand, the network that outputs the elementwise transformation parameters is simply given by stacking blocks on top of each other, and finishing with a final convolution that increases the number of channels to the amount needed to specify the elementwise transformation parameters.

4 EXPERIMENTS

So far we have identified 3 key areas that impact a flow model's performance: dequantization, coupling layer and conditioning architecture. In this section, we will first perform ablation experiments that confirm our improved design choice in each area independently contributes to improved performance. And then we show that Flow++, with all 3 improvements, achieve new flow model state-of-the-art density modelling performance on both CIFAR10 and 32x32 ImageNet. Finally we present example generative samples from Flow++ and compare them with autoregressive models' samples.

All experiments were run using weight normalization with data-dependent initialization (Salimans & Kingma, 2016). We also used the invertible 1 convolution flows of Kingma & Dhariwal (2018), as well as a variant of their "actnorm" flow that normalizes all activations independently (instead of normalizing per channel).

4.1 ABLATIONS

We ran the following ablations of our model on unconditional CIFAR10 density estimation: variational dequantization vs. uniform dequantization; logistic mixture coupling vs. affine coupling; and stacked self-attention vs. convolutions only.

Due to computation resource limit, we are not able to compare the performance of fully converged models of all variants. In fig. 5 and table 1, we compare the performance of these ablations relative to Flow++ at approximately 160 epochs of training, which was not enough for these models to converge, but far enough to see their relative performance differences. Switching from our variational dequantization to the more standard uniform dequantization costs the most: approximately 0.125 bits/dim. The remaining two ablations both cost approximately 0.05 bits/dim: switching from our logistic mixture coupling layers to affine coupling layers, and switching from our hybrid convolution-and-self-attention architecture to a pure convolutional residual architecture.

These experiments confirm that improvement in each axis independently contributes to better performance. The most surprising of them is probably the effect that different dequantization can have on both training and generalization loss. In fact, if we run Importance Sampling to further tighten the variational dequantization bound on our final CIFAR10 model, a modest number of importance samples (1024, compared to the 10000 used by Kingma et al. (2016b)) can reveal a gap of approximately 0.025 bits/dim, indicating that further gain might be possible given an even better dequantization proposal distribution.

Table 1: CIFAR10 ablation results after 164 epochs of training. Models not converged for the purposes of ablation study.

Ablation	bits/dim
uniform dequantization	3.305
no self-attention	3.233
affine coupling	3.226
Flow++ (not converged for ablation)	3.180

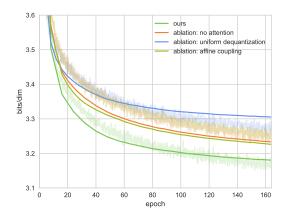


Figure 1: Ablation training (light) and validation (dark) curves on unconditional CIFAR10 density estimation. These runs are not fully converged, but the gap in density estimation performance is already visible.

Table 2: Unconditional image modeling results

Model family	Model	CIFAR10 bits/dim	ImageNet 32x32 bits/dim
Flows and VAEs	RealNVP (Dinh et al., 2016)	3.49	4.28
	Glow (Kingma & Dhariwal, 2018)	3.35	4.09
	Flow++ (ours)	3.13	3.91
	IAF-VAE (Kingma et al., 2016b)	3.11	-
Autoregressive	Multiscale PixelCNN (Reed et al., 2017)	_	3.95
C	PixelCNN/RNN (van den Oord et al., 2016a)	3.14	3.86
	Gated PixelCNN (van den Oord et al., 2016c)	3.03	3.83
	PixelCNN++ (Salimans et al., 2017)	2.92	_
	Image Transformer (Parmar et al., 2018)	2.90	3.77
	PixelSNAIL (Chen et al., 2017)	2.85	3.80

4.2 Density modeling

Here we show in table 2 that Flow++ is able to achieve new state-of-the-art density modelling results among Flow-based models and has performance that's on-par with the first generation of PixelCNN models (van den Oord et al., 2016a). As of submission, our models have not fully converged due to computational constraint and we expect further performance gain in future revision of this manuscript.

4.3 VISUALIZATIONS

We present the samples from our trained density models of Flow++ on CIFAR 10 and 32 x 32 Imagenet in Figures 2 and 3 respectively. We also import the generated samples of a PixelRNN (CNN) trained on these datasets from van den Oord et al. (2016b) (specifically, Figure 7 on page 7 in https://arxiv.org/pdf/1601.06759.pdf. We see that the samples generated by Flow++ on both the datasets match the perceptual quality of PixelCNN samples. This shows that the Flow++ model can capture both local and global dependencies at the level of a PixelCNN and is capable of generating diverse samples on large datasets.

To understand why this result is significant, one should take into account the sampling time comparison between PixelCNN and Flow++. See table 3. The numbers for O(N) and O(logN) PixelCNN



Figure 2: CIFAR 10 Samples. Left: samples from a PixelCNN taken from van den Oord et al. (2016a). Right: samples from Flow++. We see that Flow++ is able to capture local dependencies and generate good samples at the quality level of PixelCNN, but with the advantage of efficient sampling: 122x more efficient than a Fast PixelCNN++ - refer to table 3 for more details.

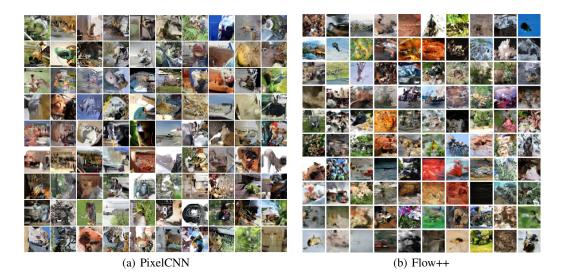


Figure 3: 32x32 Imagenet Samples. Left: samples generated from a PixelRNN taken from van den Oord et al. (2016a). Right: Flow++ samples. 32 x 32 Imagenet is a much larger dataset compared to CIFAR 10. Note that the samples from a flow-based generative model Flow++ can match the sample diversity of an autoregressive model like PixelRNN on such a large dataset.

are taken from the Multiscale PixelCNN paper (Reed et al., 2017) while the sampling time of Fast PixelCNN++ is taken from Ramachandran et al. (2017), which already uses advanced activations caching to improve on the original PixelCNN++'s sampling speed. Note that our density estimation metrics are better than that of Multiscale PixelCNN as shown in Table 2. As a caveat, these comparisons are not entirely accurate since the experiments corresponding to each model were carried out on different GPU configurations. However, it is well known in deep generative modeling that autoregressive models are much slower and inefficient for sampling when compared to flow-based generative models Kingma et al. (2016a); Kingma & Dhariwal (2018). With this context, it becomes

easy to appreciate the importance of why it makes sense to keep improving the sample quality and density estimation benchmarks of flow-based generative models and that Flow++ is a step in that direction.

Table 3: Sampling Time Comparisons for generating 32x32 images in parallel

Model	Wall Clock Time (secs)		
Flow++	0.324		
Fast PixelCNN++	39.58		
O(N) PixelCNN	120.0		
O(logN) PixelCNN	1.17		
O(logN) PixelCNN in-graph	1.14		

5 RELATED WORK

Likelihood-based models constitute a large family of deep generative models. There are methods based on variational inference (Kingma & Welling, 2013; Rezende et al., 2014; Kingma et al., 2016b) that allow for efficient approximate inference and sampling, but do not admit exact inference of log likelihood computation. There are also a large class of models that admit exact log likelihood evaluation, which we call exact likelihood models in this work. These exact likelihood models are typically specified as invertible transformations that are parametrized by neural networks (Deco & Brauer, 1995; Larochelle & Murray, 2011; Uria et al., 2013; Dinh et al., 2014; Germain et al., 2015; Oord et al., 2016b; Salimans et al., 2017; Chen et al., 2017).

There are prior works that aim to improve the sampling speed of deep autoregressive models. In particular, the Multiscale PixelCNN (Reed et al., 2017) modifies the PixelCNN to be non-fully-expressive by introducing conditional independence assumptions among pixels in a way that permits sampling in a logarithmic number of steps, rather than linear. Such a change in the autoregressive structure allows faster sampling but also makes some statistical patterns impossible to capture by definition, hence reducing the capacity of the density modeling.

There are also recent works that aim to improve the the expressiveness of the coupling layers used in flow models. Glow (Kingma & Dhariwal, 2018) shows improved density estimation performance on standard benchmarks by introducing an invertible 1x1 convolution coupling layer. Kingma & Dhariwal (2018) also shows that very large flow models can be trained to produce photorealistic faces. Müller et al. (2018) introduces piecewise polynomial coupling layers that are similar in spirit to our mixture of logistics coupling layer. They found piecewise polynomial couplings to be more expressive than affine couplings, but reported little performance gains in density estimation. We leave a detailed comparison between our coupling layer and the piecewise polynomial CDFs for future work.

6 Conclusion

We presented Flow++, a set of techniques that start to close the performance gap between flow models and autoregressive models. Aside from the specific design choices proposed here, we hope the key axes of considerations identified here will help guide future research of flow models: (a) dequantization choice; (b) coupling layer design; (c) conditioning architecture. We have shown through ablation studies that improvement along any axis will help improve a flow model's final performance and we believe more performance gains are possible as the research community continues to make strides in these directions.

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7 APPENDIX A: SAMPLES

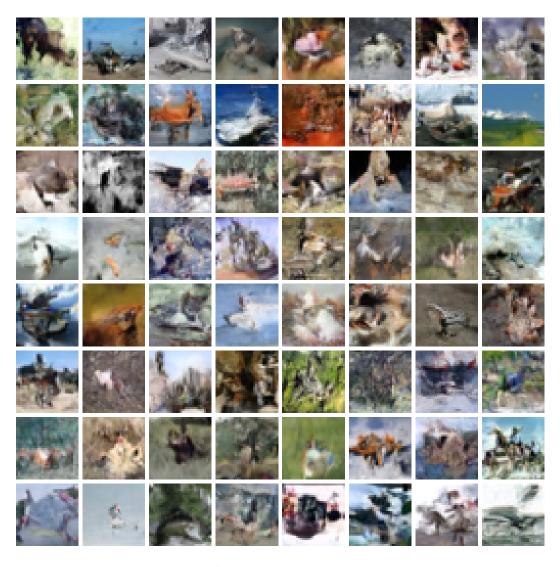


Figure 4: Samples from a Flow++ trained on CIFAR10 dataset

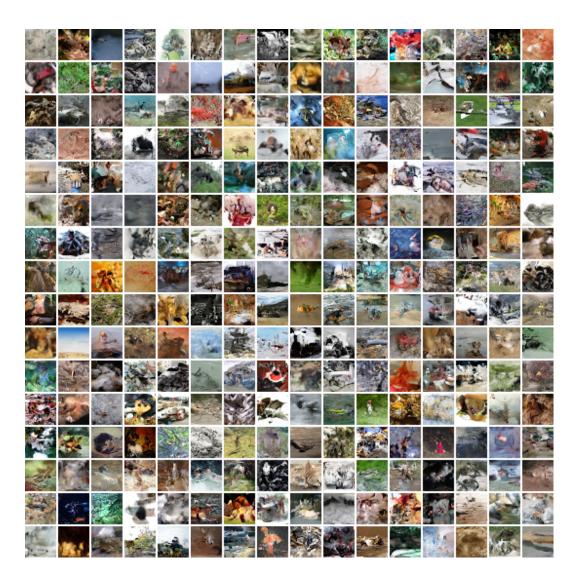


Figure 5: More Samples from a Flow++ trained on 32x32 Imagenet dataset

8 APPENDIX B: ARCHITECTURE

8.1 FLOWS

All our flows are described below (as self-contained as possible):

```
import math
import numpy as np
import tensorflow as tf
from tensorflow.contrib.framework.python.ops import arg_scope
from tqdm import tqdm
# nn module and its imports used is here are based on
   https://github.com/openai/pixel-cnn/blob/master/pixel_cnn_pp/nn.py
def safe_log(x):
   return tf.log(tf.maximum(x, 1e-22))
class tf_go:
   def __init__(self, x, debug=False):
      self.value = x
      self.debug = debug
   def __getattr__(self, name):
      def go(*args, **kwargs):
         try:
            if self.debug:
               print("Calling %s on %s" % (name, self.value))
            method = tf.__dict__[name]
            tf_val = method(*([self.value] + (list(args) if args else
                [])), **kwarqs)
            return tf_go(tf_val)
            print("Exception while calling %s on %s" % (name,
                self.value))
            raise
      return go
class Flow:
   def forward(self, x, **kwargs):
      raise NotImplementedError
   def inverse(self, y, **kwargs):
      raise NotImplementedError
class Inverse(Flow):
   def __init__(self, base_flow):
      self.base_flow = base_flow
   def forward(self, x, **kwargs):
      return self.base_flow.inverse(x, **kwargs)
   def inverse(self, y, **kwargs):
      return self.base_flow.forward(y, **kwargs)
class ImgProc(Flow):
   def forward(self, x, **kwargs):
      \# assert ((x >= 0) & (x <= 256)).all()
      x = x * (.9 / 256) + .05 # [0, 256] -> [.05, .95]
      x = -tf.log(1. / x - 1.) # inverse sigmoid
      logd = np.log(.9 / 256) + tf.nn.softplus(x) + tf.nn.softplus(-x)
```

```
logd = tf.reduce_sum(tf.reshape(logd, [int_shape(logd)[0], -1]), 1)
      return x, logd
  def inverse(self, y, **kwargs):
      y = tf.sigmoid(y)
      logd = tf.log(y) + tf.log(1. - y)
     y = (y - .05) / (.9 / 256) # [.05, .95] -> [0, 256]
      logd -= np.log(.9 / 256)
     logd = tf.reduce_sum(tf.reshape(logd, [int_shape(logd)[0], -1]), 1)
      return y, logd
class Normalize(Flow):
   def __init__(self, init_scale=1.):
      def f(input_, forward, init, ema):
         assert not isinstance(input_, list)
         if isinstance(input_, tuple):
            is_tuple = True
         else:
            assert isinstance(input_, tf.Tensor)
            input_ = [input_]
            is_tuple = False
         mus, invstds = [], []
         with arg_scope([nn.init_normalization], counters={}, init=init,
            ema=ema):
            for x in input_:
               mu, invstd = nn.init_normalization(x)
               assert mu.shape == invstd.shape == x.shape[1:]
               mus.append(mu)
               invstds.append(invstd * init_scale)
         logd = tf.fill([int_shape(input_[0])[0]],
            tf.add_n([tf.reduce_sum(tf.log(invstd)) for invstd in
            invstds]))
         if forward:
            out = [(x - mu[None]) * invstd[None] for (x, mu, invstd) in
                zip(input_, mus, invstds)]
         else:
            out = [x / invstd[None] + mu[None] for (x, mu, invstd) in
               zip(input_, mus, invstds)]
            logd = -logd
         if not is_tuple:
            assert len(out) == 1
            return out[0], logd
         return tuple(out), logd
      self.template = tf.make_template(self.__class__.__name__, f)
  def forward(self, x, init=False, ema=None, **kwargs):
      return self.template(x, forward=True, init=init, ema=ema)
   def inverse(self, y, init=False, ema=None, **kwarqs):
      return self.template(y, forward=False, init=init, ema=ema)
class TupleFlip(Flow):
  def forward(self, x, **kwargs):
     assert isinstance(x, tuple)
     a, b = x
     return (b, a), None
   def inverse(self, y, **kwargs):
     assert isinstance(y, tuple)
      a, b = y
```

```
return (b, a), None
class SpaceToDepth(Flow):
   def __init__(self, block_size=2):
      self.block_size = block_size
   def forward(self, x, **kwargs):
      return tf.space_to_depth(x, self.block_size), None
   def inverse(self, y, **kwargs):
      return tf.depth_to_space(y, self.block_size), None
class CheckerboardSplit(Flow):
   def __init__(self, bin=0, h_collapse=True):
      self.cf, self.ef, self.merge =
          self.checkerboard_condition_fn_gen(bin, h_collapse)
   def forward(self, x, **kwargs):
      return (self.cf(x), self.ef(x)), None
   def inverse(self, y, **kwargs):
      assert isinstance(y, tuple)
      cf_val, ef_val = y
      return self.merge(cf_val, ef_val), None
   @staticmethod
   def checkerboard_condition_fn_gen(bin, h_collapse):
      id = bin % 2
      def split_gen(bit):
         def go(x):
            shp = int\_shape(x)
            if len(shp) != 4:
               import IPython;
               IPython.embed()
            assert len(shp) == 4
            half = (
               tf_go(x).
                  transpose([0, 3, 1, 2]).
                  reshape([shp[0], shp[3], shp[1] * shp[2] // 2, 2]).
                  transpose([0, 2, 1, 3]).
                  value[:, :, :, bit]
            if h_collapse:
               return tf.reshape(
                  half,
                  [shp[0], shp[1], shp[2] // 2, shp[3]]
            else:
               # collapse vertically
               return (
                  tf_go(half).
                     reshape([shp[0], shp[1] // 2, 2, shp[2] // 2,
                         shp[3]]).
                     transpose([0, 1, 3, 2, 4]).
                     reshape([shp[0], shp[1] // 2, shp[2], shp[3]]).
                     value
               )
         return go
      def merge(condition, effect):
         shp = int_shape(condition)
         assert len(shp) == 4
```

```
xs = [condition, effect] if id == 0 else [effect, condition]
         if not h_collapse:
            xs = [
               tf_go(x).
                  reshape([shp[0], shp[1], shp[2] // 2, 2, shp[3]]).
                  transpose([0, 1, 3, 2, 4]).
                  reshape([shp[0], shp[1] * 2, shp[2] // 2, shp[3]]).
                  value
               for x in xs
            shp = int\_shape(xs[0])
         vs = [
            tf_go(x).
               transpose([0, 3, 1, 2]).
               reshape([shp[0], shp[3], shp[1], shp[2], 1]).
               value
            for x in xs
         return (
            tf_go(tf.concat(vs, axis=4)).
               reshape([shp[0], shp[3], shp[1], shp[2] * 2]).
               transpose([0, 2, 3, 1]).
               value
         )
      return split_gen(id), split_gen((id + 1) % 2), merge
class Pointwise(Flow):
   def __init__(self, noisy_identity_init=None):
      def f(input_, forward, ema):
         assert not isinstance(input_, list)
         if isinstance(input_, tuple):
            is_tuple = True
         else:
            assert isinstance(input_, tf.Tensor)
            input_ = [input_]
            is_tuple = False
         out, logds = [], []
         for i, x in enumerate(input_):
            _, img_h, img_w, img_c = x.shape.as_list()
            if noisy_identity_init:
               # identity + gaussian noise
               initializer = (
                     np.eye(img_c) + noisy_identity_init *
                         np.random.randn(img_c, img_c)
               ).astype(np.float32)
            else:
               # random orthogonal
               initializer = np.linalg.qr(np.random.randn(img_c,
                   img_c))[0].astype(np.float32)
            W = nn.get_var_maybe_avg(
               'W{}'.format(i), ema, dtype=tf.float32,
               initializer=initializer,
               trainable=True
            out.append(_nin(x, W if forward else tf.matrix_inverse(W)))
            logds.append(
               (1 \text{ if forward else } -1) * img_h * img_w *
               tf.to_float(tf.log(tf.abs(tf.matrix_determinant(tf.to_double(W)))))
         logd = tf.fill([input_[0].shape[0]], tf.add_n(logds))
         if not is_tuple:
```

```
assert len(out) == 1
            return out[0], logd
         return tuple(out), logd
      self.template = tf.make_template(self.__class__.__name___, f)
   def forward(self, x, init=False, ema=None, **kwargs):
      return self.template(x, forward=True, ema=ema)
   def inverse(self, y, init=False, ema=None, **kwargs):
      return self.template(y, forward=False, ema=ema)
class Sigmoid(Flow):
   def forward(self, x, **kwargs):
      y = tf.sigmoid(x)
      logd = -tf.nn.softplus(x) - tf.nn.softplus(-x)
      return y, tf.reduce_sum(tf.layers.flatten(logd), axis=1)
   def inverse(self, y, **kwargs):
      x = -safe_{log}(tf.reciprocal(y) - 1.) # inverse sigmoid
      logd = -safe_log(y) - safe_log(1. - y)
      return x, tf.reduce_sum(tf.layers.flatten(logd), axis=1)
class Compose(Flow):
   def __init__(self, flows):
      self.flows = flows
   def _maybe_tqdm(self, iterable, desc, verbose):
      return tqdm(iterable, desc=desc) if verbose else iterable
   def forward(self, x, **kwargs):
      bs = int((x[0] if isinstance(x, tuple) else x).shape[0])
      logd_terms = []
      for i, f in enumerate(self._maybe_tqdm(self.flows, desc='forward
          {}'.format(kwargs),
                                    verbose=kwarqs.get('verbose'))):
         assert isinstance(f, Flow)
         x, l = f.forward(x, **kwargs)
         if 1 is not None:
            assert l.shape == [bs]
            logd_terms.append(1)
      return x, tf.add_n(logd_terms) if logd_terms else tf.constant(0.)
   def inverse(self, y, **kwargs):
   bs = int((y[0] if isinstance(y, tuple) else y).shape[0])
      logd_terms = []
      for i, f in enumerate(
            self._maybe_tqdm(self.flows[::-1], desc='inverse
                {}'.format(kwargs), verbose=kwargs.get('verbose'))):
         assert isinstance(f, Flow)
         y, l = f.inverse(y, **kwargs)
         if 1 is not None:
            assert l.shape == [bs]
            logd_terms.append(1)
      return y, tf.add_n(logd_terms) if logd_terms else tf.constant(0.)
```

8.2 Utils: 1 - Mol CDF and its inversion

```
# logistic.py
import numpy as np
import tensorflow as tf
```

```
def logistic_logpdf(*, x, mean, logscale):
   log density of logistic distribution
  this operates elementwise
  z = (x - mean) * tf.exp(-logscale)
  return z - logscale - 2 * tf.nn.softplus(z)
def logistic_logcdf(*, x, mean, logscale):
   log cdf of logistic distribution
  this operates elementwise
   z = (x - mean) * tf.exp(-logscale)
  return tf.log_sigmoid(z)
def mixlogistic_logpdf(*, x, prior_logits, means, logscales):
   """logpdf of a mixture of logistics"""
  assert len(x.get_shape()) + 1 == len(prior_logits.get_shape()) ==
      len (means.get_shape()) == len(
      logscales.get_shape())
   return tf.reduce_logsumexp(
     tf.nn.log_softmax(prior_logits, axis=-1) + logistic_logpdf(
        x=tf.expand_dims(x, -1), mean=means, logscale=logscales),
     axis=-1
   )
def mixlogistic_logcdf(*, x, prior_logits, means, logscales):
   """log cumulative distribution function of a mixture of logistics"""
  assert (len(x.get_shape()) + 1 == len(prior_logits.get_shape()) ==
         len(means.get_shape()) == len(logscales.get_shape()))
   return tf.reduce_logsumexp(
      tf.nn.log_softmax(prior_logits, axis=-1) + logistic_logcdf(
         x=tf.expand_dims(x, -1), mean=means, logscale=logscales),
      axis=-1
   )
def mixlogistic_sample(*, prior_logits, means, logscales):
   # Sample mixture component
   sampled_inds = tf.argmax(
     prior_logits -
         tf.log(-tf.log(tf.random_uniform(tf.shape(prior_logits),
         minval=1e-5, maxval=1. - 1e-5))),
      axis=-1
  )
  sampled_onehot = tf.one_hot(sampled_inds, tf.shape(prior_logits)[-1])
   # Pull out the sampled mixture component
  means = tf.reduce_sum(means * sampled_onehot, axis=-1)
  logscales = tf.reduce_sum(logscales * sampled_onehot, axis=-1)
  # Sample from the component
  u = tf.random_uniform(tf.shape(means), minval=1e-5, maxval=1. - 1e-5)
  x = means + tf.exp(logscales) * (tf.log(u) - tf.log(1. - u))
  return x
def mixlogistic_invcdf(*, y, prior_logits, means, logscales, tol=1e-10,
   max_bisection_iters=100):
   """inverse cumulative distribution function of a mixture of
      logistics"""
  assert len(y.shape) + 1 == len(prior_logits.shape) ==
      len(means.shape) == len(logscales.shape)
   dtype = y.dtype
  with tf.control_dependencies([assert_in_range(y, min=0., max=1.)]):
     y = tf.identity(y)
```

```
def body(x, lb, ub, _last_diff):
   cur_y = tf.exp(mixlogistic_logcdf(x=x, prior_logits=prior_logits,
      means=means, logscales=logscales))
   gt = tf.cast(tf.greater(cur_y, y), dtype=dtype)
  lt = 1 - qt
  new_x = gt * (x + lb) / 2. + lt * (x + ub) / 2.
  new_lb = gt * lb + lt * x
  new\_ub = gt * x + lt * ub
  diff = tf.reduce_max(tf.abs(new_x - x))
  return new_x, new_lb, new_ub, diff
init_x = tf.zeros_like(y)
maxscales = tf.reduce_sum(tf.exp(logscales), axis=-1, keepdims=True)
   # sum of scales across mixture components
init_lb = tf.reduce_min(means - 20 * maxscales, axis=-1)
init_ub = tf.reduce_max(means + 20 * maxscales, axis=-1)
init_diff = tf.constant(np.inf, dtype=dtype)
out_x, _, _, = tf.while_loop(
   cond=lambda _x, _lb, _ub, last_diff: last_diff > tol,
  body=body,
  loop_vars=(init_x, init_lb, init_ub, init_diff),
  back_prop=False,
  maximum_iterations=max_bisection_iters
assert out_x.shape == y.shape
return out_x
```

8.3 Utils: 2 - Neural Network Architectural blocks

```
DEFAULT_FLOATX = tf.float32
STORAGE_FLOATX = tf.float32
def to_default_floatx(x):
  return tf.cast(x, DEFAULT_FLOATX)
def at_least_float32(x):
  assert x.dtype in [tf.float16, tf.float32, tf.float64]
   if x.dtype == tf.float16:
      return tf.cast(x, tf.float32)
  return x
def get_var(var_name, *, ema, initializer, trainable=True, **kwargs):
   """forced storage dtype"""
  assert 'dtype' not in kwargs
  if isinstance(initializer, np.ndarray):
     initializer = initializer.astype(STORAGE_FLOATX.as_numpy_dtype)
   v = tf.get_variable(var_name, dtype=STORAGE_FLOATX,
      initializer=initializer, trainable=trainable, **kwargs)
   if ema is not None:
     assert isinstance (ema, tf.train.ExponentialMovingAverage)
     v = ema.average(v)
  return v
def dense(x, *, name, num_units, init_scale=1., init, ema):
   # use weight normalization (Salimans & Kingma, 2016)
  with tf.variable_scope(name):
      assert x.shape.ndims == 2
```

```
_V = get_var('V', shape=[int(x.shape[1]), num_units],
         initializer=tf.random_normal_initializer(0, 0.05),
               ema=ema)
     _g = get_var('g', shape=[num_units],
         initializer=tf.constant_initializer(1.), ema=ema)
     _b = get_var('b', shape=[num_units],
         initializer=tf.constant_initializer(0.), ema=ema)
      _vinvnorm = tf.rsqrt(tf.reduce_sum(tf.square(_V), [0]))
     V, g, b, vinvnorm = map(to_default_floatx, [_V, _g, _b, _vinvnorm])
      # V, g, b = map(to_default_floatx, [_V, _g, _b])
      # vinvnorm = tf.rsqrt(tf.reduce_sum(tf.square(V), [0]))
     x0 = x = tf.matmul(x, V)
     x = (g * vinvnorm)[None, :] * x + b[None, :]
     if init: # normalize x
        m_init, v_init = tf.nn.moments(x, [0])
         scale_init = init_scale / tf.sqrt(v_init + 1e-8)
        with tf.control_dependencies([
            _g.assign(tf.cast(g * scale_init, dtype=_g.dtype)),
            _b.assign_add(tf.cast(-m_init * scale_init, dtype=_b.dtype))
        1):
           g, b = map(to_default_floatx, [_g, _b])
           x = (g * vinvnorm)[None, :] * x0 + b[None, :]
      return x
def conv2d(x, *, name, num_units, filter_size=(3, 3), stride=(1, 1),
   pad='SAME', init_scale=1., init, ema):
   # use weight normalization (Salimans & Kingma, 2016)
  with tf.variable_scope(name):
     assert x.shape.ndims == 4
     _V = get_var('V', shape=[*filter_size, int(x.shape[-1]),
         num_units],
               initializer=tf.random_normal_initializer(0, 0.05),
                   ema=ema)
     _g = get_var('g', shape=[num_units],
         initializer=tf.constant_initializer(1.), ema=ema)
     _b = get_var('b', shape=[num_units],
         initializer=tf.constant_initializer(0.), ema=ema)
      _vnorm = tf.nn.12_normalize(_V, [0, 1, 2])
     V, g, b, vnorm = map(to_default_floatx, [_V, _g, _b, _vnorm])
     W = g[None, None, None, :] * vnorm
      # calculate convolutional layer output
     input x = x
     x = tf.nn.bias_add(tf.nn.conv2d(x, W, [1, *stride, 1], pad), b)
     if init: # normalize x
        m_{init}, v_{init} = tf.nn.moments(x, [0, 1, 2])
        scale_init = init_scale * tf.rsqrt(v_init + 1e-8)
        with tf.control_dependencies([
           _g.assign(tf.cast(g * scale_init, dtype=_g.dtype)),
            _b.assign_add(tf.cast(-m_init * scale_init, dtype=_b.dtype))
        1):
           g, b = map(to_default_floatx, [_g, _b])
           W = g[None, None, None, :] * vnorm
           x = tf.nn.bias_add(tf.nn.conv2d(input_x, W, [1, *stride, 1],
               pad), b)
      return x
```

```
def nin(x, *, num_units, **kwargs):
  assert 'num_units' not in kwargs
  s = x.shape.as_list()
  x = tf.reshape(x, [np.prod(s[:-1]), s[-1]])
  x = dense(x, num_units=num_units, **kwargs)
  return tf.reshape(x, s[:-1] + [num_units])
def concat_elu(x):
   axis = len(x.get\_shape()) - 1
   return tf.nn.elu(tf.concat([x, -x], axis))
def gate(x, *, axis):
  a, b = tf.split(x, 2, axis=axis)
  return a * tf.sigmoid(b)
def gated_resnet(x, *, name, a, nonlinearity=concat_elu, conv=conv2d,
   use_nin, init, ema, dropout_p):
  with tf.variable_scope(name):
     num_filters = int(x.shape[-1])
     c1 = conv(nonlinearity(x), name='c1', num_units=num_filters,
         init=init, ema=ema)
      if a is not None: # add short-cut connection if auxiliary input
          'a' is given
         c1 += nin(nonlinearity(a), name='a_proj',
            num_units=num_filters, init=init, ema=ema)
      c1 = nonlinearity(c1)
      if dropout_p > 0:
         c1 = tf.nn.dropout(c1, keep_prob=1. - dropout_p)
      c2 = (nin if use_nin else conv)(c1, name='c2',
         num_units=num_filters * 2, init_scale=0.1, init=init, ema=ema)
      return x + gate(c2, axis=3)
def attn(x, *, name, pos_emb, heads, init, ema, dropout_p):
   with tf.variable_scope(name):
     bs, height, width, ch = x.shape.as_list()
     assert pos_emb.shape == [height, width, ch]
     assert ch % heads == 0
     timesteps = height * width
     dim = ch // heads
      # Position embeddings
     c = x + pos\_emb[None, :, :, :]
      # b, h, t, d == batch, num heads, num timesteps, per-head dim (C
         // heads)
      c = nin(c, name='proj1', num_units=3 * ch, init=init, ema=ema)
     assert c.shape == [bs, height, width, 3 * ch]
      \# Split into heads / Q / K / V
     c = tf.reshape(c, [bs, timesteps, 3, heads, dim]) # b, t, 3, h, d
     c = tf.transpose(c, [2, 0, 3, 1, 4]) # 3, b, h, t, d
     q_bhtd, k_bhtd, v_bhtd = tf.unstack(c, axis=0)
      assert q_bhtd.shape == k_bhtd.shape == v_bhtd.shape == [bs, heads,
         timesteps, dim]
      # Attention
     w_bhtt = tf.matmul(q_bhtd, k_bhtd, transpose_b=True) /
         np.sqrt(float(dim))
      w_bhtt = tf.cast(tf.nn.softmax(at_least_float32(w_bhtt)),
         dtype=x.dtype)
      assert w_bhtt.shape == [bs, heads, timesteps, timesteps]
      a_bhtd = tf.matmul(w_bhtt, v_bhtd)
      # Merge heads
```

```
a_bthd = tf.transpose(a_bhtd, [0, 2, 1, 3])
      assert a_bthd.shape == [bs, timesteps, heads, dim]
      a_btc = tf.reshape(a_bthd, [bs, timesteps, ch])
      # Project
      c1 = tf.reshape(a_btc, [bs, height, width, ch])
      if dropout_p > 0:
         c1 = tf.nn.dropout(c1, keep_prob=1. - dropout_p)
      c2 = nin(c1, name='proj2', num_units=ch * 2, init_scale=0.1,
         init=init, ema=ema)
      return x + gate(c2, axis=3)
def _norm(x, \star, axis, g=None, b=None, e=1e-5):
   u = tf.reduce_mean(x, axis=axis, keepdims=True)
   s = tf.reduce_mean(tf.square(x - u), axis=axis, keepdims=True)
  x = (x - u) * tf.rsqrt(s + e)
   if g is not None and b is not None:
      x = x * q + b
   return x
def norm(x, *, scope, ema):
   """Layer norm over last axis"""
   with tf.variable_scope(scope):
      shape = [1] * (x.shape.ndims - 1) + [int(x.shape[-1])]
      g = get_var("g", ema=ema, shape=shape,
         initializer=tf.constant_initializer(1))
      b = get_var("b", ema=ema, shape=shape,
          initializer=tf.constant_initializer(0))
      return _norm(x, g=g, b=b, axis=-1)
```

8.4 Mol Coupling Layer with Self-Attention

```
class MixLogisticAttnCoupling(Flow):
  CDF of mixture of logistics, followed by affine
  def __init__(self, filters, blocks, use_nin, components, with_affine,
      use_final_nin=False, init_scale=0.1):
      self.components = components
     self.with_affine = with_affine
     self.scale_flow = Inverse(Sigmoid())
     def f(x, init, ema, dropout_p, verbose, context):
         if init and verbose:
            # debug stuff
           xmean, xvar = tf.nn.moments(x,
               axes=list(range(len(x.get_shape()))))
           x = tf.Print(
               x, [tf.shape(x), xmean, tf.sqrt(xvar), tf.reduce_min(x),
                  tf.reduce_max(x)],
               message='{} (shape/mean/std/min/max)
                  '.format(self.template.variable_scope.name),
               summarize=10,
        B, H, W, C = x.shape.as_list()
        pos_emb = get_var('pos_emb', ema=ema, shape=[H, W, filters],
                      initializer=tf.random_normal_initializer(stddev=0.01),
                      trainable=True)
        x = conv2d(x, name='proj_in', num_units=filters, init=init,
            ema=ema)
         for i_block in range(blocks):
```

```
with tf.variable_scope(f'block{i_block}'):
            x = norm(gated_resnet(x, name='conv', a=context,
               use_nin=use_nin, init=init, ema=ema,
                            dropout_p=dropout_p), scope=f'ln1',
                                ema=ema)
            x = norm(attn(x, name='attn', pos_emb=pos_emb, heads=4,
               init=init, ema=ema, dropout_p=dropout_p),
                  scope=f'ln2', ema=ema)
     x = (nin if use_final_nin else conv2d)(x, name='proj_out',
         num\_units=C * (2 + 3 * components),
                                   init_scale=init_scale, init=init,
                                       ema=ema)
     assert x.shape == [B, H, W, C * (2 + 3 * components)]
     x = tf.reshape(x, [B, H, W, C, 2 + 3 * components])
     s, t = tf.tanh(x[:, :, :, :, 0]), x[:, :, :, 1]
     ml_logits, ml_means, ml_logscales = tf.split(x[:, :, :, :, 2:],
         3, axis=4)
     ml_logscales = tf.maximum(ml_logscales, -7.)
     assert s.shape == t.shape == [B, H, W, C]
     assert ml_logits.shape == ml_means.shape == ml_logscales.shape
         == [B, H, W, C, components]
     return s, t, ml_logits, ml_means, ml_logscales
   self.template = tf.make_template(self.__class__.__name___, f)
def forward(self, x, init=False, ema=None, dropout_p=0.,
   verbose=True, context=None, **kwargs):
   assert isinstance(x, tuple)
  cf, ef = x
   s, t, ml_logits, ml_means, ml_logscales = self.template(
     cf, init=init, ema=ema, dropout_p=dropout_p, verbose=verbose,
         context=context
  )
  out = tf.exp(mixlogistic_logcdf(x=ef, prior_logits=ml_logits,
      means=ml_means, logscales=ml_logscales))
  out, scale_logd = self.scale_flow.forward(out)
   if self.with_affine:
     assert out.shape == s.shape == t.shape
     out = tf.exp(s) * out + t
  logd = mixlogistic_logpdf(x=ef, prior_logits=ml_logits,
      means=ml_means, logscales=ml_logscales)
   if self.with_affine:
     assert s.shape == logd.shape
     logd += s
  logd = tf.reduce_sum(tf.layers.flatten(logd), axis=1)
   assert scale_logd.shape == logd.shape
  logd += scale_logd
  assert out.shape == ef.shape == cf.shape
  return (cf, out), logd
def inverse(self, y, init=False, ema=None, dropout_p=0.,
   verbose=True, context=None, **kwargs):
  assert isinstance(y, tuple)
  cf, ef = y
  s, t, ml_logits, ml_means, ml_logscales = self.template(
     cf, init=init, ema=ema, dropout_p=dropout_p, verbose=verbose,
         context=context
   )
```

```
out = ef
if self.with_affine:
   out = tf.exp(-s) * (ef - t)
out, invscale_logd = self.scale_flow.inverse(out)
out = tf.clip_by_value(out, 1e-5, 1. - 1e-5)
out = mixlogistic_invcdf(y=out, prior_logits=ml_logits,
   means=ml_means, logscales=ml_logscales)
logd = mixlogistic_logpdf(x=out, prior_logits=ml_logits,
   means=ml_means, logscales=ml_logscales)
if self.with_affine:
   assert s.shape == logd.shape
   logd += s
logd = -tf.reduce_sum(tf.layers.flatten(logd), axis=1)
assert invscale_logd.shape == logd.shape
logd += invscale_logd
assert out.shape == ef.shape == cf.shape
return (cf, out), logd
```

8.4.1 FLOW-BASED DEQUANTIZATION

```
class FlowbasedDequantWithPointwise(Flow):
  def __init__(self, *, pointwise_kwargs, coupling_kwargs):
      super(FlowbasedDequantWithPointwise, self).__init__()
      self.dequant_flow = Compose([
         CheckerboardSplit(),
         Normalize(), Pointwise(**pointwise_kwargs),
            MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
         Normalize(), Pointwise(**pointwise_kwarqs),
            MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
         Normalize(), Pointwise(**pointwise_kwargs),
            MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
         Normalize(), Pointwise(**pointwise_kwargs),
            MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
         Inverse(CheckerboardSplit()),
     ])
      def shallow_processor(x, init, ema, dropout_p):
         this = CheckerboardSplit.checkerboard_condition_fn_gen(0,
            True) [0] (x)
         that = CheckerboardSplit.checkerboard_condition_fn_gen(0,
            True) [1] (x)
         processed_context = conv2d(tf.concat([this, that], 3),
            name='proj', num_units=32, init=init, ema=ema)
         for i in range(3):
            processed_context = gated_resnet(
               processed_context, name=f'c{i}', init=init, ema=ema,
                   dropout_p=dropout_p, use_nin=False, a=None
            ) # , filter_size=[5, 5])
         return processed_context
      self.context_proc = tf.make_template("context_proc",
         shallow_processor)
   # x: [0, 256)
   def forward(self, x, init=False, ema=None, dropout_p=0.,
      verbose=True, **kwargs):
      from fun.sandboxes.hoj.tf_gm.flows import Gaussian, safe_log
     norm_x = x / 256
     noise_dist = Gaussian(dim=int(np.prod(x.shape[1:])))
      eps, eps_logli =
         noise_dist.sample_logli(noise_dist.prior_dist_info(x.shape[0]))
```

```
context = self.context_proc(norm_x, init, ema, dropout_p)
unbound_xd, logd = self.dequant_flow.forward(
    tf.reshape(eps, x.shape),
    context=context,
    init=init,
    ema=ema,
    dropout_p=dropout_p,
    verbose=verbose
)
xd = tf.nn.sigmoid(unbound_xd)
sigmoid_logd = safe_log(xd) + safe_log(1. - xd)
sigmoid_logd = tf.reduce_sum(tf.reshape(sigmoid_logd,
    [sigmoid_logd.shape[0], -1]), 1)
return x + xd, logd + sigmoid_logd - eps_logli
```

8.5 CIFAR 10 FLOW++ ARCHITECTURE:

```
def construct(*, components, with_affine, blocks):
   # see MixLogisticAttnCoupling constructor
  pointwise_kwargs = dict(noisy_identity_init=0.001)
   dequant_coupling_kwargs = dict(filters=96, blocks=2, use_nin=True,
      components=components, with_affine=with_affine)
  coupling_kwargs = dict(filters=96, blocks=blocks, use_nin=True,
      components=components, with_affine=with_affine)
  dequant_flow = FlowbasedDequantWithPointwise(
     pointwise_kwargs=pointwise_kwargs,
         coupling_kwargs=dequant_coupling_kwargs
   flow = Compose([
     ImgProc(),
      CheckerboardSplit(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
      Inverse(CheckerboardSplit()),
     SpaceToDepth(),
     ChannelSplit(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Inverse(ChannelSplit()),
     CheckerboardSplit(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(**pointwise_kwargs),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Inverse(CheckerboardSplit()),
   1)
   return dequant_flow, flow
```

```
max_lr = 1e-3
warmup_steps = 200

def lr_schedule(step):
    if step < warmup_steps:
        return max_lr * step / warmup_steps
    return max_lr

grad_clip = 1. # grad norm clipped at 1
dropout_p = 0.2
components = 32
with_affine = True
blocks = 10

init_batchsize=128 # for weight norm
total_batchsize=64 # across 8 GPUs
ema_decay = .999 # EMA for validation params (as in PixelCNN++)</pre>
```

8.6 32x32 Imagenet Flow++ Architecture:

```
def gaussian_sample_logp(shape, dtype):
  eps = tf.random_normal(shape)
  logp = Normal(0., 1.).log_prob(eps)
  assert logp.shape == eps.shape
  logp = tf.reduce_sum(tf.layers.flatten(logp), axis=1)
  return tf.cast(eps, dtype=dtype), tf.cast(logp, dtype=dtype)
class Dequantizer(Flow):
  def __init__(self, dequant_flow):
     super().__init__()
     assert isinstance (dequant_flow, Flow)
     self.dequant_flow = dequant_flow
     def shallow_processor(x, *, init, ema, dropout_p):
        this = CheckerboardSplit.checkerboard_condition_fn_gen(0,
            True) [0] (x)
        that = CheckerboardSplit.checkerboard_condition_fn_gen(0,
            True) [1] (x)
        processed_context = conv2d(tf.concat([this, that], 3),
            name='proj', num_units=32, init=init, ema=ema)
         for i in range(3):
            processed_context = gated_resnet(
               processed_context, name='c{}'.format(i),
               a=None, dropout_p=dropout_p, ema=ema, init=init,
              use_nin=False
            # TODO filter_size=[5, 5])
        return processed_context
      self.context_proc = tf.make_template("context_proc",
         shallow_processor)
  def forward(self, x, init=False, ema=None, dropout_p=0.,
      verbose=True, **kwarqs):
     eps, eps_logli = gaussian_sample_logp(x.shape,
         dtype=DEFAULT_FLOATX)
     unbound_xd, logd = self.dequant_flow.forward(
        eps,
         context=self.context_proc(x / 256.0 - 0.5, init=init, ema=ema,
            dropout_p=dropout_p),
         init=init, ema=ema, dropout_p=dropout_p, verbose=verbose
```

```
xd, sigmoid_logd = Sigmoid().forward(unbound_xd)
      assert x.shape == xd.shape and logd.shape == sigmoid_logd.shape ==
         eps_logli.shape
      return x + xd, logd + sigmoid_logd - eps_logli
def construct(*, filters, blocks, components, attn_heads, use_nin,
   use_ln):
   # see MixLogisticAttnCoupling constructor
   dequant_coupling_kwargs = dict(
      filters=filters, blocks=4, use_nin=use_nin, components=components,
         attn_heads=attn_heads, use_ln=use_ln
   dequant_flow = Dequantizer(Compose([
     CheckerboardSplit(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**dequant_coupling_kwargs),
         TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**dequant_coupling_kwargs),
         TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**dequant_coupling_kwargs),
         TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**deguant_coupling_kwargs),
         TupleFlip(),
      Inverse(CheckerboardSplit()),
   1))
   coupling_kwargs = dict(
      filters=filters, blocks=blocks, use_nin=use_nin,
         components=components, attn_heads=attn_heads, use_ln=use_ln
   flow = Compose([
      ImgProc(),
      CheckerboardSplit(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
      Inverse(CheckerboardSplit()),
      SpaceToDepth(),
     ChannelSplit(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
      Inverse(ChannelSplit()),
      CheckerboardSplit(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
     Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
      Normalize(), Pointwise(),
         MixLogisticAttnCoupling(**coupling_kwargs), TupleFlip(),
```

```
Inverse(CheckerboardSplit()),
   ])
   return dequant_flow, flow
max_lr = 1e-3
min_lr = 3e-4
warmup\_steps = 10000
bs = 32
def lr_schedule(step, *, decay=0.99999995, min_lr=3e-4):
   global curr_lr
   if step < warmup_steps:</pre>
      curr_lr = max_lr * step / warmup_steps
      return max_lr * step / warmup_steps
   #return max_lr
   elif step > (warmup_steps * 4) and curr_lr > min_lr:
      curr_lr *= decay
      return curr_lr
   return curr_lr
global curr_lr
curr_lr = max_lr
use_ln = True
dropout_p = 0. # Image Transformer (SOTA on Imagenet 32x32) uses 0.1
   dropout.
filters = 128
blocks = 20
components = 32 # logistic mixture components
attn_heads = 4
init_bs=128 # for weight norm
max_grad_norm=1
ema_decay=.999222
```