GENHOP: An Image Generation Method Based on Successive Subspace Learning

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Abstract—Being different from deep-learning-based (DL-based) image generation methods, a new image generative model built upon successive subspace learning principle is proposed and named GenHop (an acronym of Generative PixelHop) in this work. GenHop consists of three modules: 1) high-to-low dimension reduction, 2) seed image generation, and 3) low-to-high dimension expansion. In the first module, it builds a sequence of high-to-low dimensional subspaces through a sequence of whitening processes, each of which contains samples of joint-spatial-spectral representation. In the second module, it generates samples in the lowest dimensional subspace. In the third module, it finds a proper high-dimensional sample for a seed image by adding details back via locally linear embedding (LLE) and a sequence of coloring processes. Experiments show that GenHop can generate visually pleasant images whose FID scores are comparable or even better than those of DL-based generative models for MNIST, Fashion-MNIST and CelebA datasets.

I. INTRODUCTION

Unconditional image generation has received increasing attention recently due to impressive results offered by deep-learning (DL) based methods such as generative adversarial networks (GANs), variational auto-encoders (VAEs), and flow-based methods. Yet, DL-based methods are blackbox tools. The end-to-end optimization of networks is a non-convex optimization problem, which is mathematically intractable. Being motivated by the design of other generative models that allow mathematical interpretation, a new image generative model is proposed in this work. Our method is developed based on the successive subspace learning (SSL) principle [1], [2], [3], [4] and built upon the foundation of the PixelHop++ architecture [5]. Thus, it is called Generative PixelHop (or GenHop in short). Its high-level idea is sketched below.

Since high-dimensional input images have complicated statistical correlations among pixel values, it is difficult to generate images directly in the pixel domain. To address this problem, GenHop contains three modules: 1) high-to-low dimension reduction, 2) seed image generation, and 3) low-to-high dimension expansion. In the first module, it builds a sequence of high-to-low dimensional subspaces through a sequence of whitening processes called the channel-wise Saab transform, where high frequency components are discarded to lower the dimension. In the second module, the sample distribution in the lowest dimensional subspace can be analyzed and generated by white Gaussian noise, which is called seed image generation. In the third module, GenHop attempts to find the corresponding source image of a seed image through dimension expansion and a coloring mechanism. For dimension expansion, discarded high frequency components are recovered via locally linear embedding (LLE). The coloring process is the inverse of the whitening process, which is achieved by the inverse Saab transform. Experiments are conducted on MNIST, Fashion-MNIST and CelebA three datasets to demonstrate that GenHop can generate visually pleasant images whose FID scores are comparable with (or even better than) those of DL-based generative models.

II. REVIEW OF RELATED WORK

DL-based Generative Models. An image generative model learns the distribution of image samples from a certain domain and then generates new images that follow the learned distribution. Generally speaking, the design of image generative models involves analysis and generation two pipelines. The former analyzes properties of training image samples while the latter generates new images after the training is completed. So far, the best performing image generative models are all DL-based. DL-based generative methods can be categorized into two categories: adversarial and non-adversarial models. For the adversarial category, generative adversarial networks (GANs) [6] demand that distributions of training and generated images are indistinguishable by training a generator/discriminator pair through end-to-end optimization. GANs exhibit good generalization capability and yield visually impressive images. For the non-adversarial category, examples include Variational Auto-Encoders (VAEs) [7], flow-based methods [8], [9] and GLANN [10]. VAEs learns an approximation of the density function with an encoder/decoder structure. Flow-based methods transform the Gaussian distribution into a complex distribution by applying a sequence of invertible transformation functions. GLANN [10] maps images to a feature space obtained by GLO [11] and maps the feature space to the noise space via IMLE [12]. It achieves the state-of-the-art performance among non-adversarial methods.

SSL. Traditional spectral analysis such as the Fourier transform and the principle component analysis (PCA) attempts to capture the global structure but sacrifices local detail (e.g. object boundaries) of images. In contrast, local detail can be well described in the spatial domain, yet the pure spatial representation cannot capture the global information well. To overcome these shortcomings, Kuo et al. [1], [2], [3], [4] proposed two affine transforms that determine a sequence of joint spatial-
An overview of the GenHop method is shown in Fig. 1, which contains three modules as elaborated below.

A. Module 1: High-to-Low Dimension Reduction

A sequence of high-to-low dimensional subspaces is constructed from source image space through PixelHop++ [5] as shown in the figure. Each PixelHop++ unit behaves like a whitening operation. It decouples a local neighborhood (i.e., a block) into DC and AC parts and conducts the principal component analysis (PCA) on the AC part. This is named the Saab transform. The reason to remove the DC first is that the ensemble mean of AC part can be well approximated by zero so that the PCA can be applied without the need to estimate the ensemble mean. The PCA is essentially a whitening process. It removes the correlation between AC components among pixels in the same block.

To give an example, for an input gray-scale image of size 28x28, we apply the Saab transform to 2x2 non-overlapping blocks, which offers one DC and three AC channels per block, in the first PixelHop unit. The output is a joint-spatial-spectral representation of dimension 14x14x4, which forms the first subspace. By setting an energy threshold, we can partition spectral channels into low- and high-frequency channels whose numbers are denoted by $K_{1,l}$ and $K_{1,h}$, respectively. Low-frequency channels have larger energy representing the main structure of an image while high-frequency channels has lower energy representing image details. Only low-frequency channels proceed to the next stage. In other words, high-frequency channels are discarded to lower the dimension and will be estimated via LLE as discussed in Sec. III-C. High-frequency channels with sufficiently small energy will not be estimated, for example, on MNIST dataset. This lead to the sum of $K_{1,l}$ and $K_{1,h}$ being less than 4. The cascade of several PixelHop++ units yields several subspaces. For images of small spatial resolutions, we adopt two PixelHop++ units as shown in Fig. 1 to ensure a proper spatial resolution in the subspace which has the lowest joint spatial-spectral dimension to capture the global structure of an image.

B. Module 2: Seed Image Generation

In the training phase, we conduct the following four steps to learn the sample distributions in the lowest dimension furthermore, as illustrated in Fig. 2.

1) Spatial PCA. There exist correlations between spatial pixels in the lowest dimension. They can be removed by applying PCA to the spatial dimension of each channel, called spatial PCA. Components with eigenvalues less than a threshold, $\gamma$, are discarded. After a sequence of whitening operations, elements of these vector samples are uncorrelated. However, they may still be dependent. Furthermore, they are not Gaussian distributed.

2) Sample Clustering. We perform k-means clustering on them to generate multiple clusters so that the sample
Independent Component Analysis (ICA). We would like to Cumulative Histogram Matching. The probability of selecting a cluster It rebuilds dependency among elements especially essential for multi-modal sample distributions. ICA in each cluster to ensure elements of vector samples are independent.

Cumulative Histogram Matching. We would like to match the cumulative histogram of each independent component in a cluster with that of a Gaussian random variable of zero mean and unit variance [26], [27]. In the generation phase, we conduct the following steps, which are the inverse of the operations as described above.

Cluster Selection. The probability of selecting a cluster is defined by the ratio of the number of samples in that cluster and the total number of samples. We randomly select a cluster based on its probability.

Sample Generation. We generate a random variable using the Gaussian density of zero mean and unit variance and map it to the corresponding value of the sample distribution in the cluster via inverse cumulative histogram matching.

Inverse ICA. It rebuilds dependency among elements of random vectors.

Inverse Spatial PCA. It rebuilds spatial correlations among pixels of each channel.

C. Module 3: Low-to-High Dimension Expansion

Recovering Discarded Details via LLE. The generated sample in the lowest dimension contains only low-frequency (LF) components since high-frequency (HF) components are discarded to simplify the seed generation procedure. HF responses should be generated along the reverse direction to enhance details. We adopt LLE [28] to achieve this task. LLE is a commonly used technique to build the correspondence between the manifolds of low- and high-resolution images in super-resolution [29] or restoration [30], assuming two manifolds have similar local geometries. Here, LLE is used to ensure two things. First, generated LF samples are located on the manifold of training LF samples. Second, we determine the correspondence between samples of LF components and samples of HF components. LLE is implemented in small regions of spatial resolutions 2x2 or 3x3 to reduce complexity.

Neighborhood Coloring via inverse Saab Transform. Finally, we build the correlations among spatial pixels via the inverse Saab transform, which can be interpreted as a coloring process. The Saab transform parameters are determined by PCA of AC components of a local neighborhood. The parameters of the inverse transform can be derived accordingly.

IV. EXPERIMENTS

Experimental Setup. We conduct experiments on three datasets: MNIST, Fashion-MNIST and CelebA. They are often used for unconditional image generation. MNIST and Fashion-MNIST contain gray-scale images (i.e. $K_0 = 1$) while CelebA contains RGB color images. To remove the correlation between R, G, B three color channels, we perform pixel-wise PCA to decouple them, yielding three uncorrelated channels denoted by P, Q and R. We discard the R channel that has the smallest eigenvalue to reduce the dimension. To recover the RGB channels, we apply LLE conditioned on generated P and Q channels. As a result, $K_0 = 2$ for CelebA. Hyperparameters $(K_{1,l}, K_{1,h}, K_{2,l}, K_{2,h})$ are set to $(2, 1, 4, 3), (2, 2, 4, 4)$ and $(3, 1, 4, 4)$ for MNIST, Fashion-MNIST and CelebA, respectively. They are chosen to ensure the gradual dimension transition between two successive subspaces. The eigenvalue threshold, $\gamma$, is set to 0.01, 0.01 and 0.03 for MNIST, Fashion-MNIST and CelebA, respectively. The number of nearest neighbors in LLE is adaptively chosen and upper bounded by 3. For CelebA, since the number of training samples of LLE is high, we perform LLE at one location at a time.

Performance Comparison. We compare the performance of GenHop with several representative DL-based generative models in Table I. The performance metric is the Fréchet Inception Distance (FID) score. It is commonly used since both diversity and fidelity of generated images are considered. By following the procedure described in [36], we extract the embedding of 10K generated and 10K real images from the test set obtained by the Inception network and fit them into two multivariate Gaussians, respectively. The difference between the two Gaussians are measured by the Fréchet TABLE I: Comparison of FID scores of the GenHop model and representative adversarial and non-adversarial models. The lowest FID scores are shown in bold while the second lowest FID scores are underlined.

|                      | MNIST | Fashion | CelebA |
|----------------------|-------|---------|--------|
| MM GAN [6]           | 9.8   | 29.6    | 65.6   |
| NS GAN [6]           | 6.8   | 26.5    | 55.0   |
| LSGAN [31]           | 7.8   | 30.7    | 53.9   |
| WGAN [32]            | 6.7   | 21.5    | 41.3   |
| WGAN-GP [33]         | 20.3  | 24.5    | 80.0   |
| DRAGAN [34]          | 7.6   | 27.7    | 42.3   |
| BEGAN [35]           | 13.1  | 22.9    | 38.9   |
| VAE [7]              | 23.8  | 58.7    | 85.7   |
| GLO [11]             | 49.6  | 57.7    | 52.4   |
| GLANN [10]           | 8.6   | 13.0    | 46.3   |
| Ours (GenHop)        | 6.1   | 18.1    | 40.3   |
distance with their mean vectors and covariance matrices. A smaller FID score means better performance. The FID scores of representative GAN-based models (listed in the first section of Table I) are collected from [36] while those of non-adversarial models (the second section) are taken from [10]. We see from the table that Genhop has the best FID score 5.1 for MNIST and the second best 18.1 for Fashion-MNIST falling behind GLANN and 40.3 for CelebA falling behind WGAN-GP.

V. CONCLUSION AND FUTURE WORK

A non-DL-based image generation method, called GenHop, was proposed in this work. To summarize, GenHop conducted the following tasks: 1) removing correlations among pixels in a local neighborhood via the Saab transform, 2) discarding high frequency components for dimension reduction, 3) generating seed images in the lowest dimensional space using white Gaussian noise, 4) adding back discarded high frequency components for dimension expansion based on LLE and 5) recovering correlations of pixels via the inverse Saab transform. All tasks except seed generation are performed in multiple stages to control dimension change. GenHop achieved state-of-the-art performance for MNIST, Fashion-MNIST and CelebA the datasets in FID scores. As future extension, it is desired to use GenHop to generate images of higher resolution and more complicated content. It is also interesting to apply GenHop to the context of transfer learning (e.g., transfer between horses and zebras) and image inpainting.
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