The Devil is in the Decoder: Classification, Regression and GANs

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Abstract Many machine vision applications, such as semantic segmentation and depth prediction, require predictions for every pixel of the input image. Models for such problems usually consist of encoders which decrease spatial resolution while learning a high-dimensional representation, followed by decoders who recover the original input resolution and result in low-dimensional predictions. While encoders have been studied rigorously, relatively few studies address the decoder side. This paper presents an extensive comparison of a variety of decoders for a variety of pixel-wise tasks ranging from classification, regression to synthesis. Our contributions are: (1) Decoders matter: we observe significant variance in results between different types of decoders on various problems. (2) We introduce new residuallike connections for decoders. (3) We introduce a novel decoder: bilinear additive upsampling. (4) We explore prediction artifacts.

Keywords Machine Vision \cdot Computer Vision \cdot Neural Network Architectures \cdot Decoders \cdot 2D imagery \cdot perpixel prediction \cdot semantic segmentation \cdot depth prediction \cdot GANs

1 Introduction

Many important machine vision applications require predictions for every pixel of the input image. Examples include but are not limited to: semantic segmentation (e.g. Long et al. (2015); Tighe and Lazebnik (2010)), boundary detection (e.g. Arbeláez et al. (2011); Uijlings and Ferrari (2015)), human keypoints estimation (e.g. Newell et al. (2016)), super-resolution (e.g. Ledig et al. (2017)), colorization (e.g. Iizuka et al. (2016)), depth estimation (e.g. Silberman et al. (2012)), normal surface estimation (e.g. Eigen and Fergus (2015)), saliency prediction (e.g. Pan et al. (2016)), image generation with Generative Adversarial Networks (GANs) (e.g. Goodfellow et al. (2014); Nguyen et al. (2017)), and optical flow (e.g. Ilg et al. (2017)). Modern CNNbased models for such applications are usually composed of a feature extractor that decreases spatial resolution while learning high-dimensional representation and a decoder that recovers the original input resolution.

Feature extractors have been rigorously studied in the context of image classification, where individual network improvements directly affect classification results. This makes it relatively easy to understand their added value. Important improvements are convolutions (LeCun et al., 1989), Rectified Linear Units (Nair and Hinton, 2010), Local Response Normalization (Krizhevsky et al., 2012), Batch Normalization (Ioffe and Szegedy, 2015), and the use of Skip Layers (Bishop, 1995; Ripley, 1996) for Inception modules (Szegedy et al. 2016), ResNet (He et al., 2016), and DenseNet (Huang et al., 2017).

In contrast, decoders have been studied in relatively few works. Furthermore, these works focus on different problems and results are also influenced by choices and

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modifications of the feature extractor. This makes it difficult to compare existing decoder types. Our work, therefore, presents an extensive analysis of a variety of decoding methods on a broad range of machine vision tasks. For each of these tasks, we fix the feature extractor which allows a direct comparison of the decoders. In particular, we address seven machine vision tasks spanning a classification, regression, and synthesis:

- \star Classification
 - Semantic segmentation
 - Instance edge detection
- ★ Regression
 - Human keypoints estimation
 - Depth prediction
 - Colorization
 - Super-resolution
- ★ Synthesis
 - Image generation with generative adversarial networks

We make the following contributions: (1) Decoders matter: we observe significant variance in results between different types of decoders on various problems. (2) We introduce residual-like connections for decoders which yield improvements across decoder types. (3) We propose a new bilinear additive upsampling layer, which, unlike other bilinear upsampling variants, results in consistently good performance across various problem types. (4) We investigate prediction artifacts.

2 Decoder Architecture

Dense problems which require per pixel predictions are typically addressed with an encoder-decoder architecture (see Fig. 1). First, a feature extractor downsamples the spatial resolution (usually by a factor 8-32) while increasing the number of channels. Afterwards, a decoder upsamples the representation back to the original input size. Conceptually, such decoder can be seen as a reversed operation to what encoders do. One decoder module consists of at least one layer that increases spatial resolution, which is called an upsampling layer, and possibly layers that preserve spatial resolution (e.g., standard convolution, a residual block, an inception block).

The layers in the decoder which preserve spatial resolution are well studied in the literature in the context of neural architectures for image classification (e.g., Szegedy et al. (2016); Chollet (2017); Alvarez and Petersson (2016); He et al. (2016)). Therefore, we only analyze those layers which increase spatial resolution: the upsampling layers.



Fig. 1 General schematic architecture used for dense prediction problems. Typically, the resolution is reduced by a factor 2 in each of several encoder steps, after which they are upsampled by a factor 2. In this illustration, the lowest resolution is $0.125 \times$ the input size.

2.1 Related work

Many decoder architectures were previously studied in the context of a single machine vision task. The most common decoder is a transposed convolution, discussed in detail in (Dumoulin and Visin, 2016), used for feature learning (Zeiler et al., 2010; Matthew D. Zeiler and Fergus, 2011) and pixel-wise prediction tasks such as image segmentation (Ronneberger et al., 2015), semantic segmentation (Long et al., 2015), optical flow (Dosovitskiy et al., 2015a), depth prediction (Laina et al., 2016), image reconstruction from its feature representation (Dosovitskiy and Brox, 2016), and image synthesis (Dosovitskiy et al., 2015b).

Several convolution variants were proposed which trade model capacity for speed. In particular, in (Szegedy et al., 2016) and (Romera et al., 2018) a 2D convolution is decomposed into two 1D convolutions in the context of image classification and semantic segmentation. In (Chollet, 2017) separable convolutions are applied per channel to image classification. We test both variants in the context of transposed convolutions.

Bilinear and/or bi-cubic upsampling were studied in the context of super-resolution by (Dong et al., 2016). The depth-to-space upsampling method was proposed by (Shi et al., 2016) to improve the task of superresolution. Our paper compares all these decoders, adds one extra, and explores modifications, which are detailed in Sec. 2.2.

A previous study by (Odena et al., 2016) examined transposed convolution and bilinear upsampling qualitatively in the context of GANs (Goodfellow et al., 2014). We provide quantitative results comparing many more decoders on seven different machine vision tasks.

Recently, stacked hourglass networks were proposed (Newell et al., 2016), which are multiple encoder-decoder networks stacked in sequence. We include stacked hourglass networks in our analysis.

2.2 Upsampling layers

Below we present and compare several ways of upsampling the spatial resolution in convolution neural networks, a crucial part of any decoder. We limit our study to upsampling the spatial resolution by a factor of two, which is the most common setup in the literature (see e.g. Ronneberger et al. (2015) and Yu and Porikli (2016)).

2.2.1 Existing upsampling layers

Transposed Convolution. Transposed convolutions are the most commonly used upsampling layers and are also sometimes referred to as 'deconvolution' or 'upconvolution' (Long et al., 2015; Dumoulin and Visin, 2016; Zeiler et al., 2010; Dosovitskiy and Brox, 2016; Dosovitskiy et al., 2015a; Ronneberger et al., 2015; Laina et al., 2016). A transposed convolution can be seen as a reversed convolution in the sense of how the input and output are related to each other. However, it is not an inverse operation, since calculating the exact inverse is an under-constrained problem and therefore ill-posed. Transposed convolution is equivalent to interleaving the input features with 0's and applying a standard convolutional operation. The transposed convolution is illustrated in Fig. 2 and 4.

Decomposed Transposed Convolution. Whereas decomposed transposed convolution is similar to the transposed convolution, conceptually it splits the main convolution operation into multiple low-rank convolutions. For images, it simulates a 2D transposed convolution using two 1D convolutions (Fig. 5). Regarding possible feature transformations, decomposed transposed convolution is strictly a subset of regular transposed convolution. As an advantage, the number of trainable parameters is reduced (Tab. 1).

Decomposed transposed convolution was successfully applied in the inception architecture (Szegedy et al., 2016), which achieved state of the art results on ILSVRC 2012 classification (Russakovsky et al., 2015). It was also used to reduce the number of parameters of the network in (Alvarez and Petersson, 2016).

Conv + Depth-To-Space. Depth-to-space (Shi et al., 2016) (also called subpixel convolution) shifts the feature channels into the spatial domain as illustrated in Fig. 3. Depth-to-space preserves perfectly all floats inside the high dimensional representation of the image, as it only changes their placement. The drawback of this approach is that it introduces alignment artifacts. To be comparable with other upsampling layers which have

learnable parameters, before performing the depth-tospace transformation we apply a convolution with four times more output channels than for other upsampling layers.

Bilinear Upsampling + Convolution. Bilinear Interpolation is another conventional approach for upsampling the spatial resolution. To be comparable to other methods, we assume there is additional convolutional operation applied after the upsampling. The drawback of this strategy is that it is both memory and computationally intensive: bilinear interpolation increases the feature size quadratically while keeping the same amount of "information" as measured in the number of floats. Because bilinear upsampling is followed by a convolution, the resulting upsampling method is four times more expensive than a transposed convolution.

Bilinear Upsampling + Separable Convolution. Separable convolution was used to build a simple and homogeneous network architecture (Chollet, 2017) which achieved superior results to inception-v3 (Szegedy et al., 2016). A separable convolution consists of two operations, a per channel convolution and a pointwise convolution with 1×1 kernel which mixes the channels. To increase the spatial resolution, before separable convolution, we apply bilinear upsampling in our experiments.

2.2.2 Bilinear Additive Upsampling

To overcome the memory and computational problems of bilinear upsampling, we introduce a new upsampling layer: bilinear additive upsampling. In this layer, we propose to do bilinear upsampling as before, but we also add every N consecutive channels together, effectively reducing the output by a factor N. This process is illustrated in Fig. 6. Please note that this process is deterministic and has zero tunable parameters (similarly to depth-to-space upsampling, but does not cause alignment artifacts). Therefore, to be comparable with other upsampling methods we apply a convolution after this upsampling method. In this paper, we choose N in such a way that the final number of floats before and after bilinear additive upsampling is equivalent (we upsample by a factor 2 and choose N = 4), which makes the cost of this upsampling method similar to a transposed convolution.



Fig. 2 Visual illustration of (Decomposed) Transposed Convolution. The resolution is increased by interleaving the features with zeros. Afterwards either a normal or decomposed convolution is applied which reduces the number of channels by a factor 4.

Fig. 3 Visual illustration of Depth-to-Space. Fist a normal convolution is applied, keeping both the resolution and the number of channels. Then features of each four consecutive channels are re-arranged spatially into a single channel, effectively reducing the number of channels by a factor 4.



Output Dependencies on Weights

Fig. 4 Spatial dependency for the outputs of a Transposed Convolution with kernel size 3 and stride 2.



Fig. 5 Spatial dependency for the outputs of a Decomposed Transposed Convolution with kernel size 3 and stride 2.



Fig. 6 Visual illustration of our bilinear additive upsampling. We upsample all features and take the average over each four consecutive channels. For our upsampling layer this operation is followed by a convolution (not visualized).

2.3 Skip Connections and Residual Connections

2.3.1 Skip Connections

Skip connections have been successfully used in many decoder architectures (Lin et al., 2017a; Ronneberger et al., 2015; Pinheiro et al., 2016; Lin et al., 2017b; Kendall et al., 2015). This method uses features from the encoder in the decoder part of the same spatial resolution, as illustrated in Fig. 1. For our implementation of skip connections, we apply the convolution on the last layer of encoded features for a given spatial resolution and concatenate them with the first layer of decoded features (Fig. 1).

2.3.2 Residual Connections for decoders

Residual connections (He et al., 2016) have been shown to be beneficial for a variety of tasks. However, residual connections cannot be directly applied to upsampling methods since the output layer has a higher spatial resolution than the input layer and a lower number of feature channels. In this paper, we introduce a transformation which solves both problems.

In particular, the bilinear additive upsampling method which we introduced above (Fig. 6) transforms the input layer into the desired spatial resolution and number of channels without using any parameters. The resulting features contain much of the information of the original features. Therefore, we can apply this transformation (this time without doing any convolution) and add its result to the output of any upsampling layer, resulting in a residual-like connection. We present the graphical representation of the upsampling layer in Fig. 7. We demonstrate the effectiveness of our residual connection in Section 4.

Fig. 7 Visual illustration of our residual connection for upsampling layers. The "identity" layer is the sum of each four consecutive upsampled layers in order to get the desired number of channels and resolution.

3 Tasks and Experimental Setup

3.1 Classification

3.1.1 Semantic segmentation

We evaluate our approach on the standard PASCAL VOC-2012 dataset (Everingham et al., 2012). We use both the training dataset and augmented dataset (Hariharan et al., 2011) which together consist of 10,582images. We evaluate on the VOC Pascal 2012 validation dataset of 1,449 images. We follow a similar setup to Deeplab-v2 (Chen et al., 2018). As the encoder network, we use ResNet-101 with stride 16. We replace the first 7x7 convolutional layer with three 3x3 convolutional layers. We use [1, 2, 4] atrous rates in the last three convolutional layers of the block5 in Resnet-101 as in (Chen et al., 2017). We use batch normalization, have single image pyramid by using [6, 12, 18] atrous rates together with global feature pooling. We initialize our model with the pre-trained weights on ImageNet dataset with an input size of 513×513 . Our decoder upsamples four times by a factor 2 with a single 3x3convolutional layer in-between. We train the network with stochastic gradient descent on a single GPU with a momentum of 0.9 and batch size 16. We start from learning rate 0.007 and use a polynomial decay with the power 0.9 for 30,000 iterations. We apply L2 regularization with weight decay of 0.0001. We augment the dataset by rescaling the images by a random factor between 0.5 and 2.0. During evaluation, we combine the prediction from multiple scales [0.5, 0.75, 1.0, 1.25, 1.5, 1.75] and the flipped version of the input image. We train the model using maximum likelihood estimation per pixel (softmax cross entropy) and use mIOU (mean Intersection Over Union) to benchmark the models.

Upsampling method	# of parameters	# of operations	Comments
Transposed	whIO	whWHIO	
Decomposed Transposed	(w+h)IO	(w+h)WHIO	Subset of the Transposed method
Conv + Depth-To-Space	whI(4O)	whWHI(4O)	
Bilinear Upsampling $+$ Conv	whIO	wh(2W)(2H)IO	
Bilinear Upsampling $+$ Separable	whI + IO	(2W)(2H)I(wh+O)	
Bilinear Additive Upsampling + Conv	whIO	wh(2W)(2H)(I/4)O	

Table 1 Comparison of different upsampling methods. W, H - feature width and height, w, h - kernel width and height, I, O- number of channels for input and output features.

3.1.2 Instance boundaries detection

For instance-wise boundaries, we use PASCAL VOC 2012 segmentation (Everingham et al., 2012). This dataset contains 1,464 training and 1,449 validation images, annotated with contours for 20 object classes for all instances. The dataset was originally designed for semantic segmentation. Therefore, only interior object pixels are marked, and the boundary location is recovered from the segmentation mask. Similar to (Uijlings and Ferrari, 2015) and (Khoreva et al., 2016), we consider only object boundaries without distinguishing semantics, treating all 20 classes as one.

As an encoder or feature extractor we use ResNet-50 with stride 8 and atrous convolution. We initialize from pre-trained ImageNet weights. The input to the network is of size 321×321 . The spatial resolution is reduced to 41×41 , after which we use 3 upsampling layers, with an additional convolutional layer in-between, to make predictions in the original resolution.

During training, we augment the dataset by rescaling the images by a random factor between 0.5 and 2.0 and random cropping. We train the network with asynchronous stochastic gradient descent for 40,000 iterations with a momentum of 0.9. We use a learning rate of 0.0003 with a polynomial decay of power 0.99. We apply L2 regularization with a weight decay of 0.0002. We use a batch size of 5. We use sigmoid cross entropy loss per pixel (averaged across all pixels), where 1 represents an edge pixel, and 0 represents a non-edge pixel.

We use two measures to evaluate edge detection: fmeasure for the best-fixed contour threshold across the entire dataset and average precision (AP). During the evaluation, predicted contour pixels within three pixels from ground truth pixels are assumed to be correct (Martin et al., 2001).

3.2 Regression

3.2.1 Human keypoints estimation

We perform experiments on the MPII Human Pose dataset (Andriluka et al., 2014) which consists of around 25k images with over 40k annotated human poses in terms of keypoints of the locations of seven human body parts. The images cover a wide variety of human activities. Since test annotations are not provided and since there is no official train and validation split, we make such split ourselves (all experiments use the same split). In particular, we divide the training set randomly into 80% training images and 20% validation images. Following (Andriluka et al., 2014), we measure the Percentage of Correct Points with a matching threshold of 50% of the head segment length (PCKh), and report the average over all body part on our validation set.

For the network, we re-implement the stacked hourglass network (Newell et al., 2016) with a few modifications. In particular, we use as input a cropped the area around the center of human of size 353x272 pixels. We resize predictions from every hourglass subnetwork to their input resolution using bilinear interpolation and then we apply a Mean Squared Error (MSE) loss. The target values around each keypoint are based on a Gaussian distribution with a standard deviation of 6 pixels in the original input image, which we rescale such that its highest value is 10 (which we found to work better with MSE). We do not perform any test-time data augmentation such as horizontal flipping.

3.2.2 Depth prediction

We apply our method to the NYUDepth v2 dataset by (Silberman et al., 2012). We train on the entire NYUDepth v2 raw data distribution, using the official split. There are 209, 822 train and 187, 825 test images. We ignore the pixels that have a depth below a threshold of 0.3 meters in both training and test, as these reads are not reliable in the Kinect sensor.

As for the encoder network, we use ResNet-50 with stride 32, initialized with pre-trained ImageNet weights.

We use an input size of 304×228 pixels. We apply 1x1convolution with 1024 filters. Following (Laina et al., 2016), we upsample four times with a convolutional layer in between to get back to half of the original resolution. Afterwards, we upsample with bilinear interpolation to the original resolution (without any convolutions).

We train the network with asynchronous stochastic gradient descent on 3 machines with a momentum of 0.9, batch size 16, starting from a learning rate of 0.001, decaying by 0.92 every 72926 steps and train for 640,000 iterations. We apply L2 regularization with weight decay 0.0005. We augment the dataset through random changes in brightness, hue, and saturation, through put greyscale image. The output pixel value targets are random color removal and through mirroring.

For depth prediction, we use the reverse Huber loss following (Laina et al., 2016).

$$Loss(y, \hat{y}) = \begin{cases} |y - \hat{y}| & for \quad |y - \hat{y}| <= c\\ |y - \hat{y}|^2 & for \quad |y - \hat{y}| > c \end{cases}$$
(1)

$$c = \frac{1}{5} \max_{(b,h,w) \in [1...Batch][1...Height][1...Width]} |y_{b,h,w} - y_{\hat{b},\hat{h},w}|$$
(2)

The reverse Huber loss is equal to the L1 norm for $x \in$ [-c, c] and equal to L2 norm outside this range. In every gradient descent step, c is set to 20% of the maximal pixel error in the batch.

For evaluation, we use the metrics from (Eigen and Fergus, 2015), i.e., mean relative error, root mean squared error, root mean squared log error, the percentage of correct prediction within three relative thresholds: 1.25, 1.25^2 and 1.25^3 .

3.2.3 Colorization

We train and test our models on the ImageNet dataset (Russakovsky et al., 2015), which consists of 1,000,000 training images and 50,000 validation images.

For the network architecture, we follow (Iizuka et al., 2016), where we swap their original bilinear upsampling method with the methods described in Section 2. In particular, these are three upsampling steps of factor 2.

This model combines joint training of image classification and colorization, where we are mainly interested in the colorization part. We resize the input image to 224×224 pixels. We train the network for 30,000 iterations using the Adam optimizer with a batch size of 32 and we fix the learning rate to 1.0. We apply L2 regularization with a weight decay of 0.0001. During training, we randomly crop and randomly flip the input image. For the skip connections, we concatenate the decoded

features with the feature extractor applied to the original input image (as there are two encoder networks employed on two different resolutions).

As loss function we use the averaged L1 loss for pixel-wise color differences for the colorization part, and a softmax cross entropy loss for the classification part.

$$Loss(y, \hat{y}, y_{cl}, \hat{y_{cl}}) = 10|y - \hat{y}| - y_{cl} \log \hat{y_{cl}}$$
(3)

Color predictions are made in the YPbPr color space (luminance, blue - luminance, red - luminance). The luminance is ignored in the loss function during both training and evaluation as is it provided by the inscaled to the range [0,1]. $y_c l$ is the one hot encoding of the predicted class label and $\hat{y_{cl}}$ are the predicted classification logits.

To evaluate colorization we follow (Zhang et al., 2016). We compute the average root mean squared error between the color channels in the predicted and ground truth pixels. Then, for different thresholds for root mean squared errors, we calculate the accuracy of correctly predicted colored pixels within given range. From these we compute the Area Under the Curve (AUC) (Zhang et al., 2016). Additionally, we calculate the top-1 and top-5 classification accuracy for the colorized images on the ImageNet Russakovsky et al. (2015) dataset, motivated by the assumption that better recognition corresponds to more realistic images.

3.2.4 Super resolution

For super-resolution, we test our approach on the CelebA dataset, which consists of 167, 483 training images and 29,249 validation images (Liu et al., 2015). We follow the setup from (Yu and Porikli, 2016): the input images of the network are 16×16 images, which are created by resizing the original images. The goal is to reconstruct the original images which have a resolution of 128×128 .

The network architecture used for super-resolution is similar to the one from (Kim et al., 2016). We use six ResNet-v1 blocks with 32 channels after which we upsample by a factor of 2. We repeat this three times to get to a target upsampling factor of 8. On top of this, we add 2 pointwise convolutional layers with 682 channels with batch normalization in the last layer. Note that in this problem there are only operations which keep the current spatial resolution or which upsample the representation. We train the network on a single machine with 1 GPU, batch size 32, using the RM-SProp optimizer with a momentum of 0.9, a decay of 0.95 and a batch size of 16. We fix the learning rate at 0.001 for 30000 iterations. We apply L2 regularization

with weight decay 0.0005. The network is trained from scratch.

As loss we use the averaged L2 loss between the predicted residuals \hat{y} and actual residuals y. The ground truth residual y in the loss function is the difference between original 128×128 target image and the predicted upsampled image. All target values are scaled to [-1,1]. We evaluate performance using standard metrics for super-resolution: PSNR and SSIM.

3.3 Synthesis

3.3.1 Generative Adversarial Networks

To test our decoders in the generator network for Generative Adversarial Networks (Goodfellow et al., 2014) we follow the setup from (Lucic et al., 2017). We benchmark different decoders (generators) on the Cifar-10 dataset which consists of 50000 training and 10000 testing 32x32 color images. Additionally, we also test our approach on the CelebA dataset described in Sec. 3.2.4.

For Cifar-10 we use the GAN architecture with spectral normalization of Miyato et al. (2018). For CelebA, we use the InfoGan architecture (Chen et al., 2016). For both networks, we replace the transposed convolutions in the generator with our upsampling layers and add 3x3 convolutional layers between them. In the discriminator network, we do not use batch normalization. We train the model with batch size 64 through 200k iterations on a single GPU using the Adam optimizer. We perform extensive hyperparameter search for each of the studied decoder architectures: We vary the learning rate from a logarithmic scale between [0.00001, 0.01], we vary the beta1 parameter for the Adam optimizer from range [0, 1], and the λ used in the gradient penalty term (Eq. (4)) from a logarithmic scale in range [-1, 2]. We try both 1 and 5 discriminator updates per generator update.

As the discriminator and generator loss functions we are using the Wasserstein GAN Loss with Gradient Penalty (Gulrajani et al., 2017):

$$\mathcal{L}_D = -\mathbb{E}_{x \sim p_d}[D(x)] + \mathbb{E}_{\hat{x} \sim p_g}[D(\hat{x})] + \lambda \mathbb{E}_{\hat{x} \sim p_g}[(||\nabla D(\alpha x + (1 - \alpha \hat{x}))||_2 - 1)^2]$$
(4)

$$\mathcal{L}_G = -\mathbb{E}_{\hat{x} \sim p_g}[D(\hat{x})] \tag{5}$$

where p_d is data distribution, p_g is the generator output distribution, and D is the output of the discriminator. α is uniformly sampled in every iteration from range [0, 1]. We evaluate the performance of the generator architectures using the Frechet Inception Distance (FID) (Heusel et al., 2017).

4 Results

We first compare the upsampling types described in Section 2.2 without residual connections (Tab. 1(a)). Next, we discuss the benefits of adding our residual connections (Tab. 1(b)). Since all evaluation metrics are highly correlated, this table only reports a single metric per problem. A table with all metrics can be found in the supplementary material.

4.1 Results without residual-like connections.

For semantic segmentation, the depth-to-space transformation is the best upsampling method. For instance edge detection and human keypoints estimation, the skip-layers are necessary to get good results. For instance edge detection, the best performance is obtained by transposed, depth-to-space, and bilinear additive upsampling. For human keypoints estimation, the hourglass network uses skip-layers by default and all types of upsampling layers are about as good. For both depth prediction and colorization, all upsampling methods perform similarly, and the specific choice of upsampling matters little. For super-resolution, networks with skiplayers are not possible because there are no encoder modules which high-resolution (and relatively low-semantic) features. Therefore, this problem has no skip-layer entries. Regarding performance, only all transposed variants perform well on this task; other layers do not. Similarly to super-resolution, GANs have no encoder and therefore it is not possible to have skip connections. Separable convolutions perform very poorly on this task. There is no single decoder which performs well on both Cifar 10 and CelebA datasets: bilinear additive upsampling performs well on Cifar 10, but poorly in CelebA. In contrast, decomposed transposed performs well on CelebA, but poorly on Cifar 10.

Generalizing over problems, we see that no single upsampling scheme provides consistently good results across all the tasks.

4.2 Results with residual-like connections

Next, we add our residual-like connections to all upsampling methods. Results are presented in Tab. 1(b). For

(a) F	les	ult	s w	rith	lou	t r	esio	lua	ıl c	on	nec	tio	ns.	
Ns	CelebA	FID		27.3				32.0		34.9		42.4		37.7	
GA	Cifar 10	FID		34.1		35.9		36.7		36.5		39.3		30.0	
Super-resolution	CelebA	SSIM		0.675		0.681		0.596		0.593		0.677		0.594	
Colorization	ImageNet	AUC		0.952	0.954	0.951	0.954	0.950	0.954	0.950	0.950	0.952	0.949	0.950	0.953
Depth prediction	NYUv2 Depth	MRE		0.146	0.145	0.148	0.144	0.145	0.143	0.147	0.145	0.148	0.149	0.147	0.144
Human Keypoints	IIdw	PCK/0.50		0.780	0.853	0.782	0.853	0.772	0.853	0.782	0.853	0.784	0.854	0.779	0.853
Instance Edge detection	VOC Pascal 2012	f-measure		0.248	0.640	0.458	0.522	0.501	0.648	0.451	0.566	0.570	0.570	0.620	0.654
Semantic Segmentation	VOC Pascal 2012	mloU		0.773	0.770	0.766	0.769	0.782	0.777	0.771	0.773	0.775	0.244	0.776	0.780
			Skip	z	≻	z	≻	z	≻	z	≻	z	≻	z	≻
			Residual	z	z	z	z	z	z	z	z	z	z	z	z
Problem	Dataset	Metric		Transposed	Transposed	Decomposed Transposed	Decomposed Transposed	Conv + Depth To Space	Conv + Depth To Space	Bilinear Upsampling + Conv	Bilinear Upsampling + Conv	Bilinear Upsampling + Separable	Bilinear Upsampling + Separable	Bilinear Additive Upsampling + Conv	Bilinear Additive Upsampling + Conv

Problem			Semantic Segmentation	Instance Edge detection	Human Keypoints	Depth prediction	Colorization	Super-resolution	GAI	sk
Dataset			VOC Pascal 2012	VOC Pascal 2012	MPII	NYUv2 Depth	ImageNet	CelebA	Cifar 10	CelebA
Metric			Uolm	f-measure	PCK/0.50	MRE	AUC	SSIM	FID	FID
	Residual	Skip								
Transposed	۲	z	0.780	0.622	0.782	0.147	0.952	0.687	27.4	32.0
Transposed	۲	≻	0.780	0.296	0.853	0.147	0.953			
Decomposed Transposed	۲	z	0.781	0.244	0.784	0.149	0.954	0.683		26.9
Decomposed Transposed	۲	≻	0.781	0.531	0.853	0.146	0.952			
Conv + Depth To Space	۲	z	0.784	0.625	0.769	0.143	0.953	0.686	29.2	25.3
Conv + Depth To Space	۲	≻	0.782	0.629	0.851	0.142	0.945			
Bilinear Upsampling + Conv	۲	z	0.777	0.531	0.779	0.147	0.950	0.684	29.8	28.9
Bilinear Upsampling + Conv	۲	≻	0.780	0.537	0.854	0.143	0.955			
Bilinear Upsampling + Separable	۲	z	0.783	0.610	0.783	0.151	0.945	0.684	34.3	32.8
Bilinear Upsampling + Separable	۲	≻	0.778	0.517	0.853	0.149	0.947			
Bilinear Additive Upsampling + Conv	۲	z	0.774	0.626	0.777	0.149	0.952	0.684	30.1	38.4
Bilinear Additive Upsampling + Conv	۲	≻	0.775	0.644	0.852	0.145	0.952			

(b) Results with residual connections.

 Table 2
 Our main results comparing a variety of decoders on seven machine vision problems. The colors represent relative performance: red means top performance, yellow means reasonable performance, blue and green means poor performance.

Method	Measure	Our method	Recent work	Recent work
Semantic segmentation	mIoU	0.77	0.76	Chen et al. (2018)
Instance boundaries detection	f-measure	0.63	0.62	Khoreva et al. (2016)
Human keypoints estimation	PCKh	0.85^{*}	0.88*	Newell et al. (2016)
Depth prediction	MRE	0.15	0.15	Laina et al. (2016)
Colorization	AUC	0.95	0.90	Zhang et al. (2016)
Super-resolution	SSIM	0.68	0.70	Yu and Porikli (2016)
GANs on Cifar 10	FID	30	53	Lucic et al. (2017)

Table 3 Comparison of our bilinear additive upsampling + conv + res results with other methods from the literature. Comparing semantic segmentation results with (Chen et al., 2018), we did not use MS COCO detection dataset for pretraining and multiscale training. The performance of the depth prediction task is compared on all the scene frames from the test dataset and not on small subset as in (Laina et al., 2016), therefore we report the numbers of our reimplementation of the method. *Numbers on human keypoints estimation are not directly comparable since we use a slightly smaller training set and evaluate on a different random validation set. We see that for all tasks we achieve good results.

the majority of combinations, we see that adding residual connections is beneficial. In fact, for semantic segmentation, depth prediction, colorization, and superresolution, adding residual connections results in consistently high performance across decoder types. For instance edge detection, transposed convolutions, depthto-space, and bilinear additive upsampling work well when no skip connections are used. The only task which is unaffected by adding residual connections is human keypoints estimation. This is because there are already residual connections over each hourglass in the stacked hourglass network; each hourglass can be seen as refining the previous predictions by estimating residuals. Adding even more residual connections to the network does not help.

To conclude: (1) Residual-like connections almost always improve performance. (2) With residual connections, we can identify three consistently good decoder types: transposed convolutions, depth-to-space, and bilinear additive upsampling.

4.3 Decoder Artifacts

The work of (Odena et al., 2016) identified checkerboard artifacts in GANs by giving qualitative examples followed by an analysis of the decoder as to why these artifacts appear. In particular, they showed that transposed convolutions have "uneven overlap", which means that convolutions at different places in the image operate on a different number of features, as is also demonstrated in Fig. 4 (output dependencies on weights). Of course, this observation extends to all transposed convolution variants. We also note that a similar analysis holds for the depth-to-space decoder, where our explanation in Fig.3 visually shows how the artifacts arise.

We can indeed observe artifacts by looking at a few qualitative results: Fig. 8 and Fig. 9 show several out-

put results of respectively GANs and the depth prediction task. We show outputs for transposed convolutions, depth-to-space, and our new bilinear additive upsampling method. For GANs, when not having residual connections, depth-to-space upsampling frequently results in artifacts, as shown in the two left-most images. For transposed convolutions there are fewer and generally more subtle artifacts (see the right-most image), while we could not find these for our bilinear additive upsampling method. When adding residual connections, the depth-to-space and transposed methods stop producing visible artifacts. Similarly, for the depth-prediction task, without residual connections, depth-to-space and transposed convolutions result in clearly visible artifacts, while our method yields much better looking results.

For the depth-prediction problem, we can also directly measure a certain type of artifact. To do this, we observe that object surfaces are mostly smooth. This means that if we move over the pixels of an object in x- or y-direction, we expect the depth on these objects to either monotonically increase or decrease. In particular, we consider triplets of pixels in either horizontal or vertical direction on surfaces which face the camera (ground-truth depth difference between adjacent pixels is smaller than 1 cm) and for which the ground truth is monotonically increasing/decreasing. We then measure the percentage of triplets in which the predictions are non-monotonic. Results are presented in Tab. 4, where a higher percentage means more artifacts.

Results confirm our qualitative observations and what we expect from the decoder methods: the bilinear upsampling variants have significantly fewer artifacts than the transposed variants and than depth-to-space. We also observe that for transposed variants and depthto-space, residual connections help to reduce artifacts. We conclude that the bilinear upsampling methods are preferred for minimizing prediction artifacts.



Fig. 8 Visualizations of GANs

Upsampling layer	Skip	Res	artifacts	artifacts
			x-axis	y-axis
Transposed	Ν	Ν	27.6%	24.8%
Transposed	Y	N	30.6%	23.5%
Transposed	N	Y	23.1%	21.9%
Transposed	Y	Y	22.8%	22.8%
Dec. Transposed	Ν	N	27.0%	24.1%
Dec. Transposed	Y	N	33.4%	24.0%
Dec. Transposed	N	Y	25.5%	23.9%
Dec. Transposed	Y	Y	24.5%	24.1%
Depth-To-Space	Ν	Ν	28.4%	26.9%
Depth-To-Space	Y	N	26.3%	23.2%
Depth-To-Space	N	Y	25.6%	23.7%
Depth-To-Space	Y	Y	24.2%	22.9%
Bilinear Ups. $+$ Conv.	Ν	N	14.3%	14.6%
Bilinear Ups. $+$ Conv.	Y	N	19.5%	19.5%
Bilinear Ups. $+$ Conv.	N	Y	14.7%	14.8%
Bilinear Ups. $+$ Conv.	Y	Y	17.4%	18.3%
Bilinear Ups. $+$ Sep.	Ν	Ν	13.8%	14.6%
Bilinear Ups. $+$ Sep.	Y	Ν	20.5%	20.3%
Bilinear Ups. $+$ Sep.	N	Y	17.3%	16.0%
Bilinear Ups. $+$ Sep.	Y	Y	19.7%	18.3%
Bilinear Add. Ups.	Ν	Ν	14.8%	15.2%
Bilinear Add. Ups.	Y	N	19.0%	17.7%
Bilinear Add. Ups.	N	Y	14.7%	15.0%
Bilinear Add. Ups.	Y	Y	17.5%	18.4%

Table 4 Analysis of the artifacts for depth regression for1000 test images, totalling approximately 27 million tripletsper experiment.

4.4 Nearest Neighbour and Bicubic Upsampling

The upsampling layers in our study used bilinear upsampling if applicable. But one can also use nearest neighbour upsampling (e.g. Berthelot et al. (2017); Jia et al. (2017); Odena et al. (2016)) or bicubic upsampling (e.g. Dong et al. (2016); Hui et al. (2016); Kim et al. (2016); Odena et al. (2016)). We performed two small experiments to examine these alternatives.

In particular, on the Human keypoints estimation task (Sec. 3.2.1) we started from the bilinear upsampling + conv method with residual connections and skip layers. Then we changed the bilinear upsampling to nearest neighbour and bicubic. Results are in Tab. 5. As can be seen, the nearest neighbour sampling is slightly worse than bilinear and bicubic, which perform about the same.



Fig. 9 Input image and errors for three different models. Top row: input image. Second row: Transposed. Third row: Depth-to-Space. Fourth row: Bilinear additive upsampling. We observe that our proposed upsampling method produces smoother results with less arifacts.

	Res	Skip	PCK/0.50
NN Ups. + Conv	Y	Y	0.846
Bilinear Ups. $+$ Conv	Y	Y	0.854
Bicubic Ups. Conv	Υ	Y	0.855

Table 5 Comparison of Nearest Neighbour Upsamping, Bi-linear Upsampling, and Bicubic Usampling on Human Key-point Estimation.

	Res	Skip	FID
NN Ups. + Conv	Y	Ν	28.0
Bilinear Ups. + Conv	Y	Ν	29.8
Bicubic Ups. Conv	Y	Ν	28.1

Table 6Comparison of Nearest Neighbour Upsamping, Bi-linear Upsampling, and Bicubic Usampling on generating synthetic images using GANs for Cifar 10.

We also compared the same three decoder types to synthesize data using GANs (Sec. 3.3.1) on Cifar 10 (without skip connections since these are not applicable). Results are in Tab. 6. As can be seen, nearest neighbour upsampling and bicubic have similar performance. This is in contrast to Odena et al. (2016) who found that nearest neighbour upsampling worked better than bicubic upsampling. In Odena et al. (2016) they suggested that their results could have been caused by the hyperparameters being optimized for nearest neighbour upsampling. Our result seems to confirm this. Bilinear upsampling performs slightly worse than the other two methods.

To conclude, in both experiments the differences between upsampling methods are rather small. Bicubic upsampling slightly outperforms the two others. Intuitively, this is the more accurate non-parametric upsampling method but one that has a higher computational cost.

5 Conclusions

This paper provided an extensive evaluation for different decoder types on a broad range of machine vision applications. Our results demonstrate: (1) Decoders matter: there are significant performance differences among different decoders depending on the problem at hand. (2) We introduced residual-like connections which, in the majority of cases, yield good improvements when added to any upsampling layer. (3) We introduced the bilinear additive upsampling layer, which strikes a right balance between the number of parameters and accuracy. Unlike the other bilinear variants, it gives consistently good performance across all applications. (4) Transposed convolution variants and depth-to-space decoders have considerable prediction artifacts, while bilinear upsampling variants suffer from this much less. (5) Finally, when using residual connections, transposed convolutions, depth-to-space, and our bilinear additive upsampling give consistently strong quantitative results across all problems. However, since our bilinear additive upsampling suffers much less from prediction artifacts, it should be the upsampling method of choice.

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6 Appendix



Conv + Conv + Bilinear

ilinear

Silinear



Table 7 Our main results comparing a variety of decoders on seven machine vision problems. The colors represent relativeperformance: red means top performance, yellow means reasonable performance, blue and green means poor performance.