Generative Models and VAEs



CPSC/AMTH 663



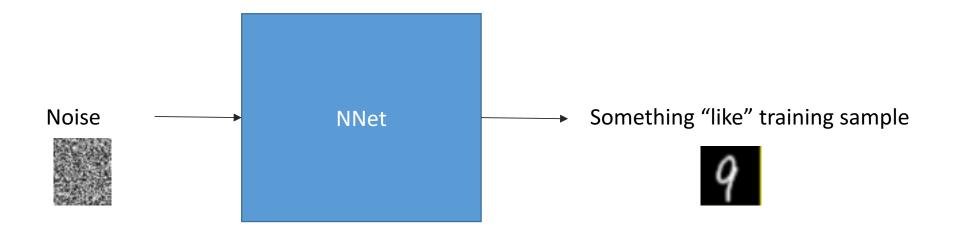
Outline



- 1. Generating with Neural Networks
- 2. MMDnet: Generating Neural Networks with Noise
- 3. Generalized Stochastic Autoencoders
- 4. VAEs

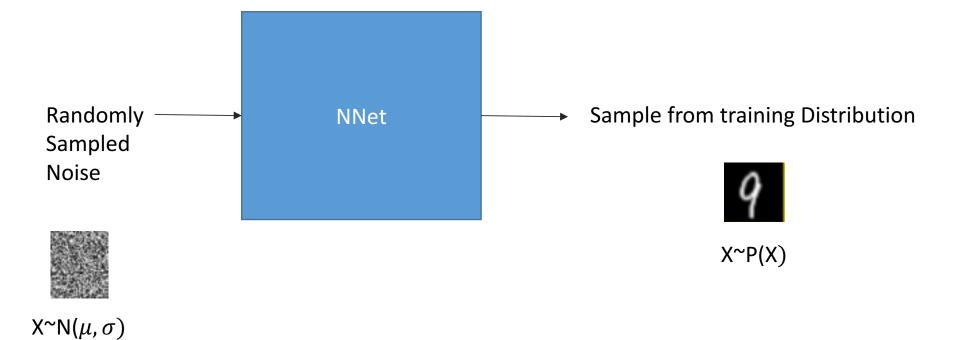
Generative Models





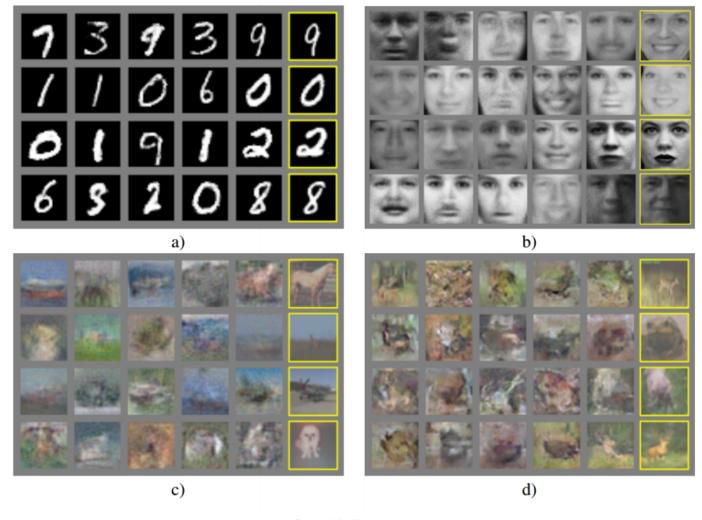
Probabilistic Interpretation





Images Generated from GANs

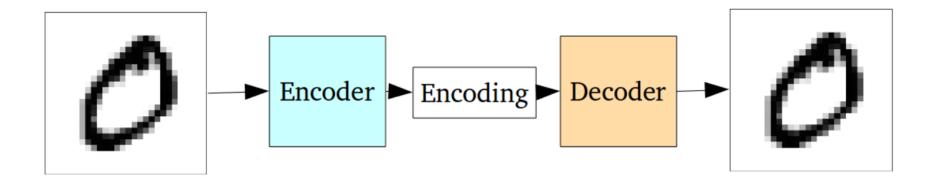




Goodfellow

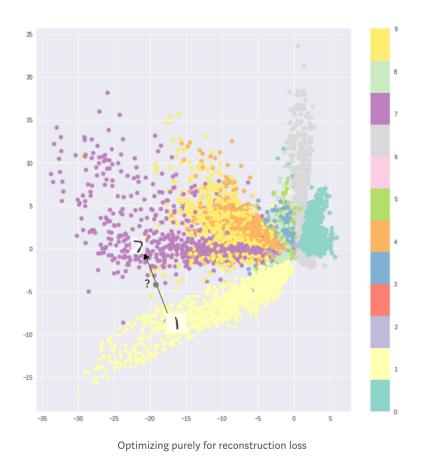
Why not autoencoders?





- Optimized with Reconstruction Loss
- Not explicitly penalized for Generative Purposes





Formation of clusters helps decoding

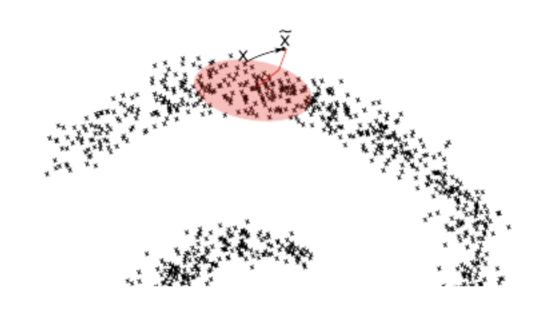
Does not help generation

Latent space cannot be interpolated

Can we modify autoencoder for generation?



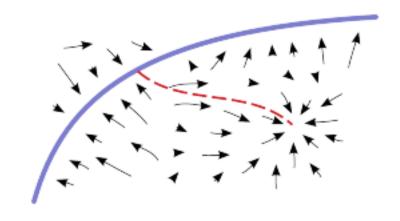
Generalized. Denoising Autoencoder [Bengio et al. 2013]



- Basic idea: add noise to samples to push them away from the data manifold and then have them pull it back.
- For each training sample X define a corruption process $C(\tilde{X},X)$ that creates a corrupted sample \tilde{X} .
- Train a denoising autoencoder to reverse this by using (X, \tilde{X}) as a training example

Lost away from the manifold?



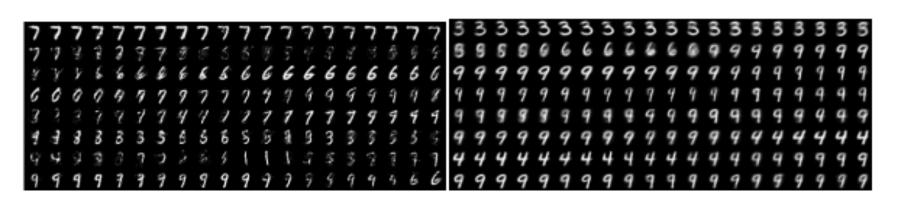


- Problem: spurious modes
- The DAE may not be able to walk back well enough if you get too far away from the sample space

Walkback training



- Train the neural network to walk back from several steps away
- Create longer range corruption processes using the original corruption process
- Sample a second step $C(\tilde{X}, \tilde{X})$
- Add the training sample (\tilde{X}, \tilde{X}) to the training



Trains the DAE to estimate conditional



- This way of training actually trains the DAE to estimate a conditional probability distribution $P(X|\tilde{X})$
- [Bengio et al. 2013] show that a consistent estimator of P(X), i.e. distribution of the training example can be recovered by alternating sampling from the corruption process. $C(\tilde{X},X)$ and the denoising process $P(X|\tilde{X})$
- Turns out that learning a conditional distribution is a lot simpler than learning the joint distribution!
 - This idea is used in style transfer and other places
- A conditional distribution can just be a Gaussian or something simple, but this alternation allows convergence towards the joint distribution

Probabilities



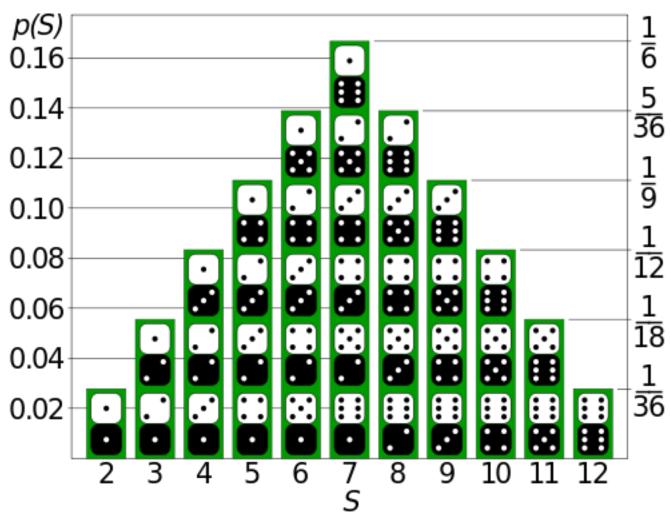
- Probability is the likelihood P(E) that an event E will occur
- Random experiment, with set of possible outcomes in S
- Probability is a function assigned to subsets E of S, in the range [0,1]

Axioms:

- Non-negativity $0 \le P(E) \le 1$
- Measure P(S) = 1
- Additivity of disjoint events $P(E_1 \cup E_1) = P(E_1) + P(E_2)$

Probability Mass Function

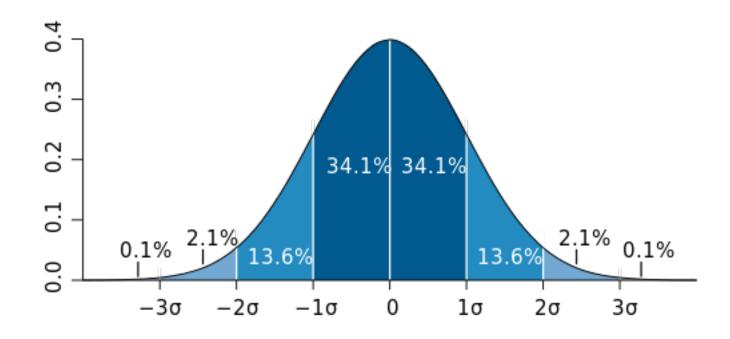




Assigns probabilities to each event

Probability Density Function





Describes the infinitesimal probability of any given value, and the probability that the outcome lies in a given interval can be computed by integrating the probability density function over that interval

Real Data



- Do not have PDF
- Need to estimate PDF
- Why?
 - Generative/predictive model
 - Estimate mutual information
 - Capture inherent stochasticity
 - Reason about outliers and noise

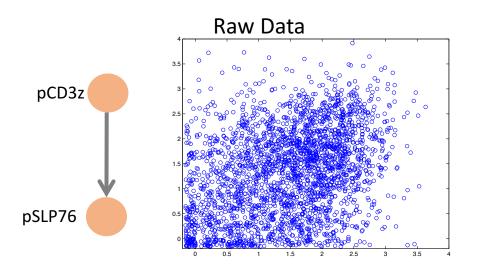
Estimating Probability Distributions

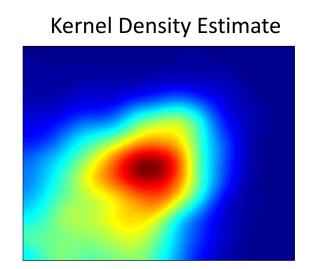


- Parametric distributions:
 - Maximum likelihood:
 - MAP
- Example $\underset{\theta}{\operatorname{arg\,max}} p(data \mid \theta)$
- Gaussian: $\underset{\theta}{\operatorname{arg \, max} \, p(\theta) p(data \, | \, \theta)}$
 - Parameters: μ Mean, σ variance
- Poisson
 - Parameter: λ (mean and variance)

Non-parametric Estimates





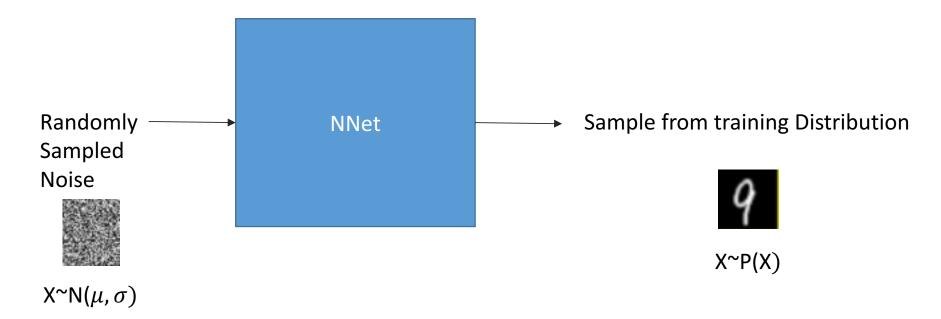


- A density estimate is a function $\widehat{f}(x)$ that is an approximation to the true probability density function
- Properties: real-valued, non-negative, integrates to 1
- Non-parametric estimate, no assumptions on shape

Neural Network Estimates



 Train a neural network with input stochasticity to mimic output distribution



 Penalize by a distribution distance or divergence: KL divergence, MMD distance, wasserstein distance

MMD nets



- Train the neural network to transmute noise into desired probability distribution using Maximum Mean Discrepancy optimization. [Dziugaite et al. 2015]
- MMD is a kernel-based 2-sample distribution test for distribution similarity
- This is a batch-level penalty, penalize a whole batch to look like the training distribution

Distances



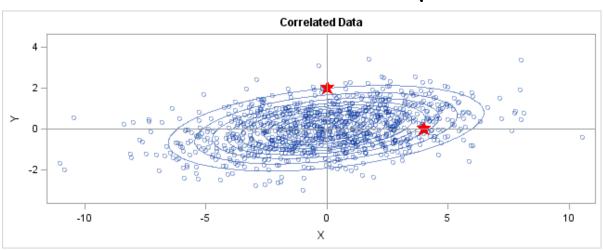
- Distances measure ways in which objects are different
- A distance metric is a real valued function d(x,y) such that
 - d(x,y) >= 0 non-negative
 - d(x,x) = 0 identity
 - d(x,y)=d(y,x) symmetric
 - d(x,z)<=d(x,y)+d(y,z) triangle inequality

Example Distances



- $A=[a_1, a_2, ... a_n], B=[b_{1,b_2}, ... b_n]$
- Euclidean Distance | |A-B||₂
- Minkowski Distance | | A-B | | P
- Mahalanobis Distance

$$\sqrt{(A-B)\Sigma^{-1}(A-B)^T}$$



Affinities



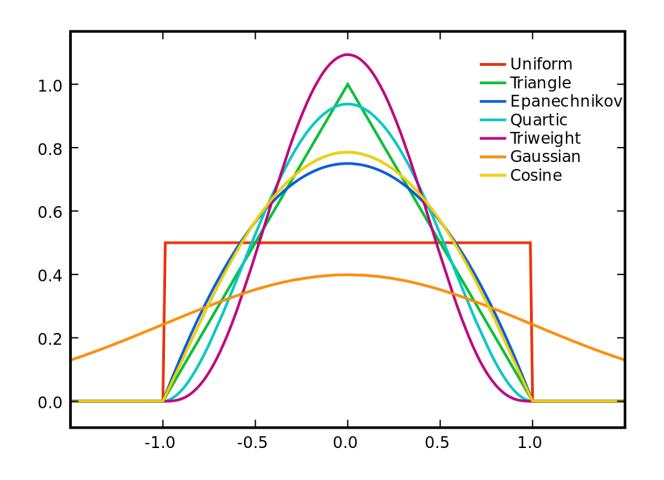
- Similarity is a function s(x,y) such that:
 - s(x,x) = 1
 - s(x,y) = s(y,x)
- Similarities between datapoints
- Correlation

$$\frac{\operatorname{cov}(x, y)}{\sigma^{x} \sigma^{y}}$$

$$\frac{\|x^{T} y\|}{\|x\| \|y\|}$$

Distance to Affinity via Kernels

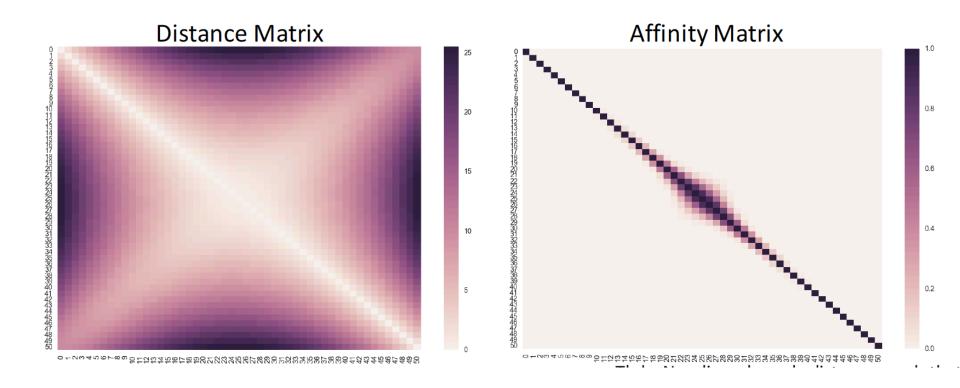




Real-valued Non-negative Symmetric

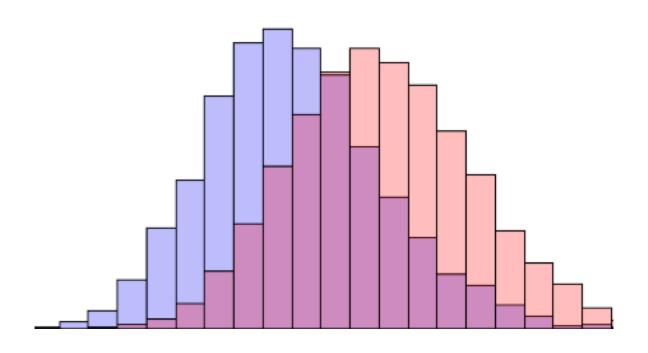
Affinities are correlations in this hidden hypothetical space

Distance->Affinity



$$sij := \exp\left(-\frac{d(x_i, x_j)^2}{2\sigma^2}\right)$$

Mean Maximal Discrepancy



$$MMD(p,q) = \frac{1}{m^2} \sum_{i,j \in m} K(p_i,p_j) - \frac{2}{mn} \cdot \sum_{i,j} K(p_i,q_j) + \frac{1}{n^2} \sum_{i,j \in n} K(q_i,q_j)$$

[Dziugaite, Roy, Ghahramani, UAI 2015]

MMD

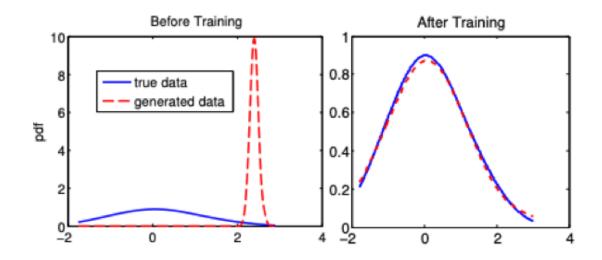
MMD quantifies, how similar are two sets of samples by

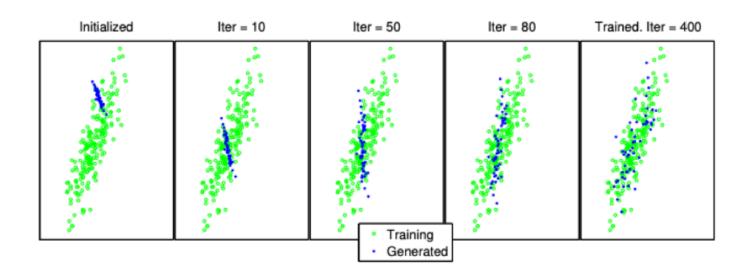
Picking a pair X, X' from distribution 1

Picking a pair Y, Y' from distribution 2

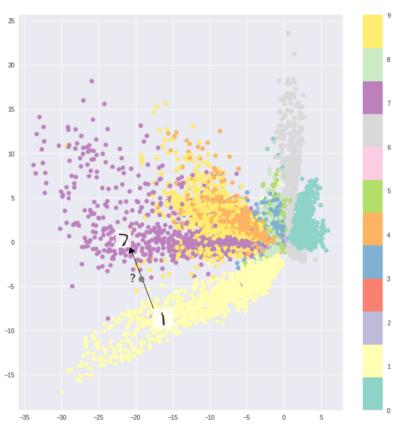
Testing how similar the with sample pairs are compared to the across-sample pairs X,Y and X',Y'

The average distance between the within sample similarities and across samples



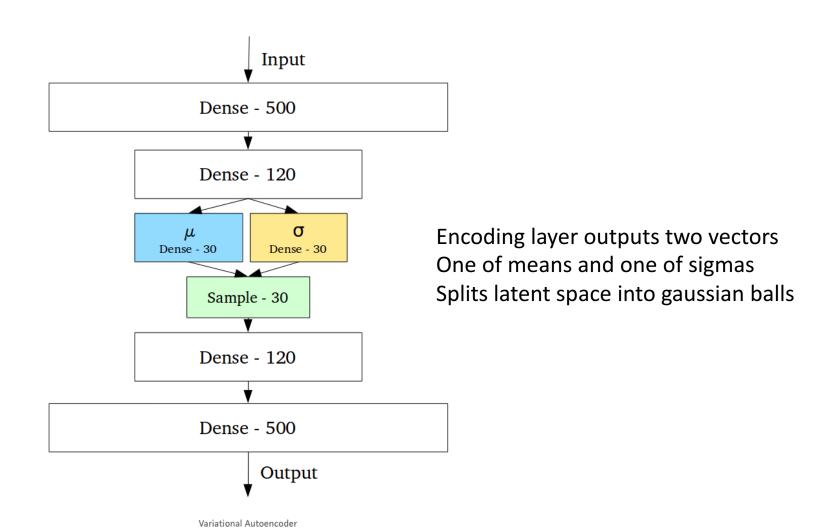


VAEs Revisited- Want Dense Middle Layer

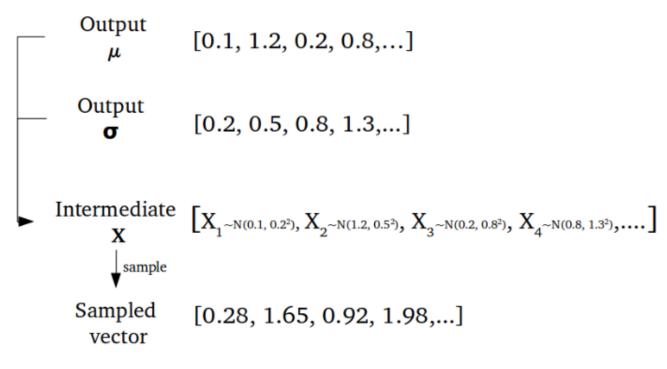


Optimizing purely for reconstruction loss

Designed for Sampling

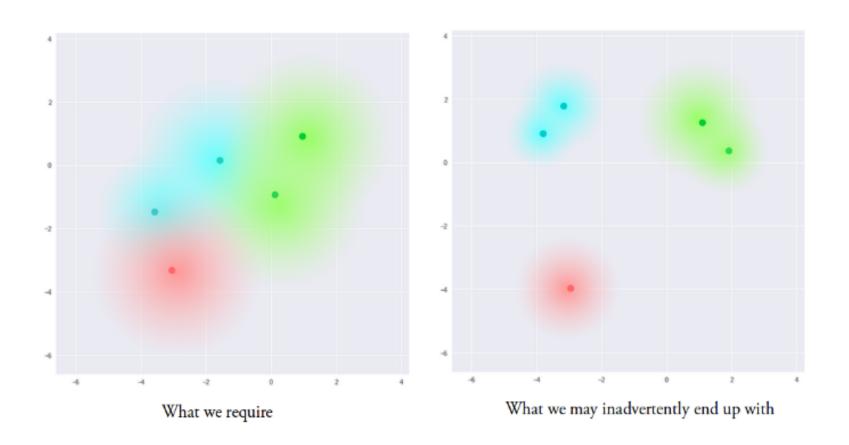


Sampling a vector in latent space



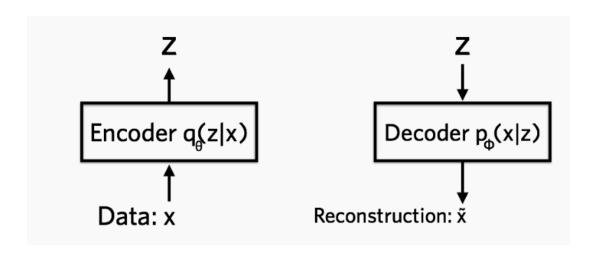
Stochastically generating encoding vectors

Discouraging Clusters



Loss Function of VAE

$$l_i(heta,\phi) = -\mathbb{E}_{z\sim q_{ heta}(z\mid x_i)}[\log p_{\phi}(x_i\mid z)] + \mathbb{KL}(q_{ heta}(z\mid x_i)\mid\mid p(z))$$



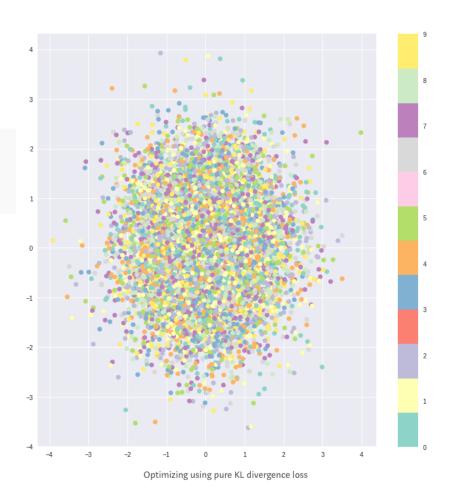
$$-\mathbb{E}_{z\sim q_{ heta}(z|x_i)}[\log p_{\phi}(x_i\mid z)]$$

KL Divergence Penalty

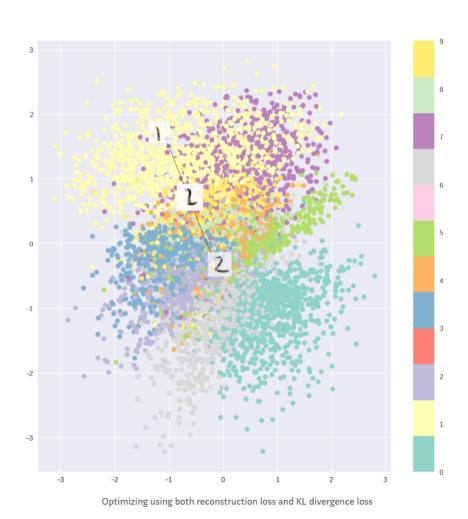
$$\mathbb{KL}(q_{ heta}(z\mid x_i)\mid\mid p(z))$$

Penalizing distributions
In the latent space from being
Too far from a standard normal

$$p(z) = Normal(0, 1).$$



KL+Reconstruction Loss



Suggested Reading Blogs

https://towardsdatascience.com/intuitivelyunderstanding-variational-autoencoders-1bfe67eb5daf

https://jaan.io/what-is-variational-autoencoder-vae-tutorial/