

# Generative Ratio Matching Networks

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# Introduction

Adversarial Generative Models(GANs, MMD-GANs)

✅ can generate high-dimensional data such as natural images.

❌ are very difficult to train due to the saddlepoint optimization problem

GRAM is a *stable* learning algorithm for *implicit* deep generative models that does **not** involve a saddlepoint optimization problem and therefore is easy to train 🎉

# Overview

1. Learn a low-dimensional manifold
  - that preserves the difference between the data ( $p_x$ ) and the model ( $q_x$ ) densities.
  - We use the ratio ( $r(x) = \frac{p_x}{q_x}$ ) of the two densities as the measure of this difference.
2. Train the model ( $G_\gamma$ ) in the low-dimensional manifold
  - using the *Maximum Mean Discrepancy* criterion as it work very well in low dimensional data.

## GRAM: Algorithm

1. Learn the manifold projection function  $f_\theta(x)$  by minimising the squared difference between the pair of density ratios:

$$\begin{aligned} D(\theta) &= \int q_x(x) \left( \frac{p_x(x)}{q_x(x)} - \frac{\bar{p}(f_\theta(x))}{\bar{q}(f_\theta(x))} \right)^2 dx \\ &= C - \text{PD}(\bar{q}, \bar{p}) \end{aligned}$$

2. Train the generator  $G_\gamma$  by minimizing the empirical estimator of MMD in the low-dimensional manifold,

$$\min_{\gamma} \left[ \frac{1}{N^2} \sum_{i=1}^N \sum_{i'=1}^N k(f_{\theta}(x_i), f_{\theta}(x_{i'})) - \frac{2}{NM} \sum_{i=1}^N \sum_{j=1}^M k(f_{\theta}(x_i), f_{\theta}(G_{\gamma}(z_j))) \right. \\ \left. + \frac{1}{M^2} \sum_{j=1}^M \sum_{j'=1}^M k(f_{\theta}(G_{\gamma}(z_j)), f_{\theta}(G_{\gamma}(z_{j'}))) \right]$$

# Pearson Divergence Maximisation and Density Ratio Estimation

Monte Carlo approximation of PD,

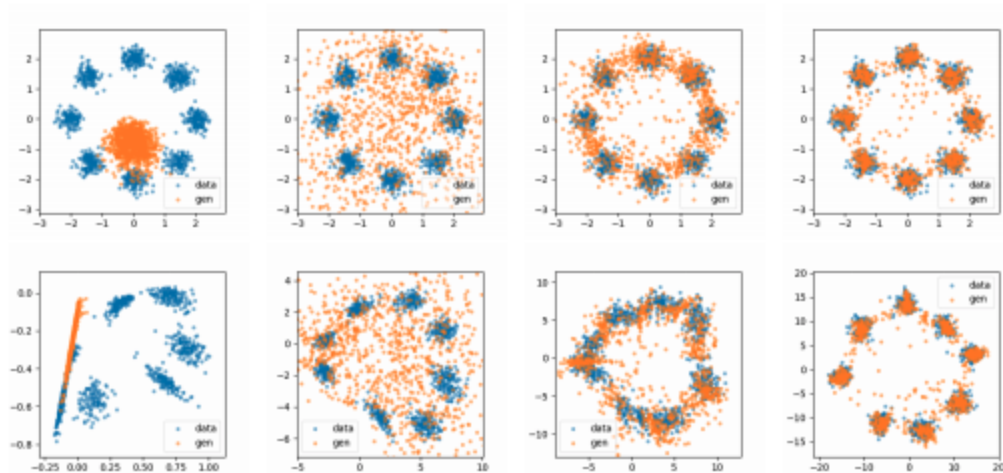
$$\text{PD}(\bar{q}, \bar{p}) \approx \frac{1}{N} \sum_{i=1}^N \left( \frac{\bar{p}(f_{\theta}(x_i))}{\bar{q}(f_{\theta}(x_i))} \right)^2 - 1$$

where  $x_i^q \sim q_x$ .

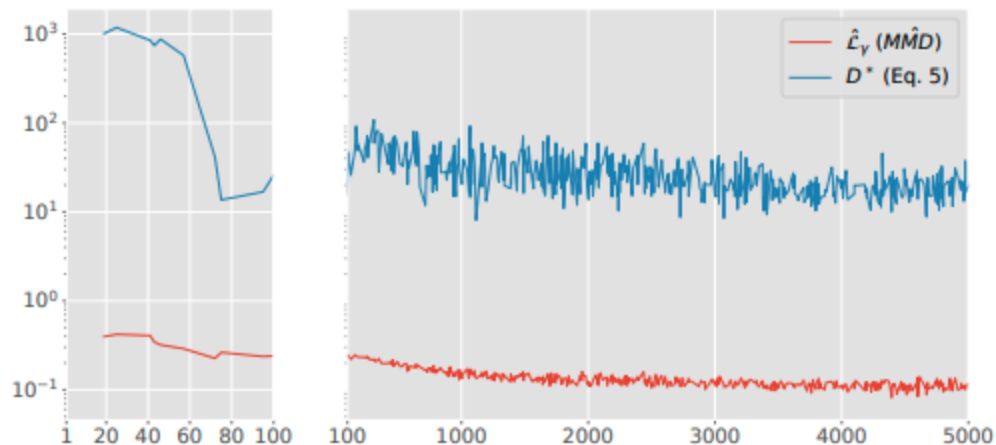
We use a MMD based density ratio estimator (Sugiyama et al., 2012) under the fixed-design setup:  $\hat{r}_q = \mathbf{K}_{q,q}^{-1} \mathbf{K}_{q,p} \mathbf{1}$ .

- $\mathbf{K}_{q,q}$  and  $\mathbf{K}_{q,p}$  are Gram matrices defined by  $[\mathbf{K}_{q,q}]_{i,j} = k(f_{\theta}(x_i^q), f_{\theta}(x_j^q))$  and  $[\mathbf{K}_{q,p}]_{i,j} = k(f_{\theta}(x_i^q), f_{\theta}(x_j^p))$ .

# The Ring dataset: Illustration of the Method and Stability

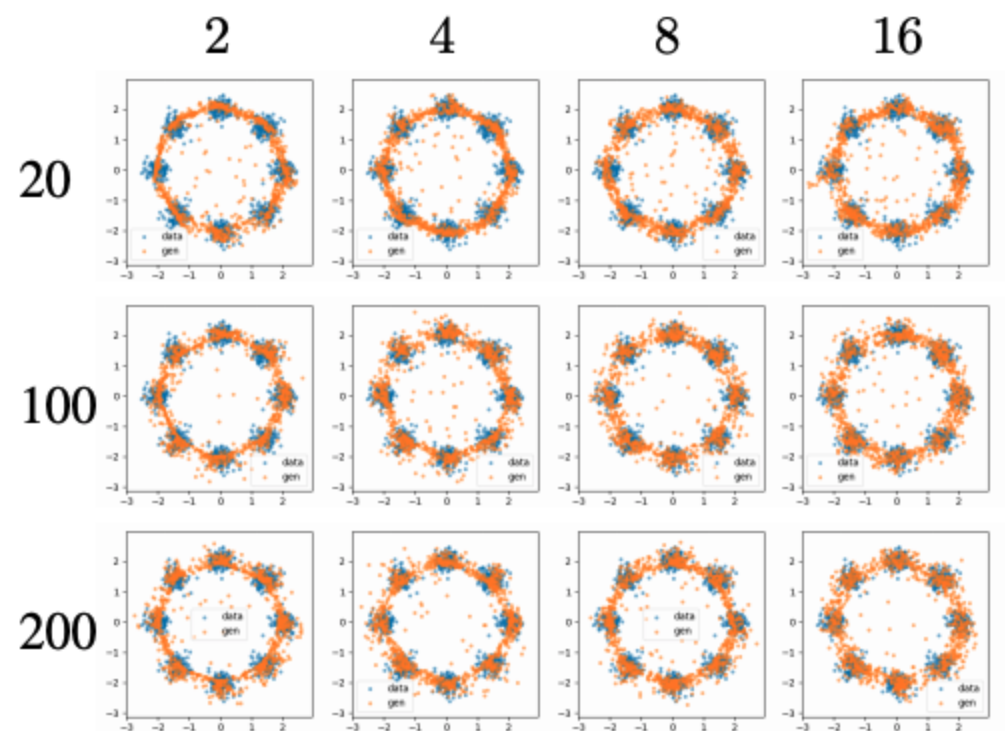


(a) Data and samples in the original (top) and projected space (bottom) during training; four plots are at iteration 10, 100, 1000 and 10,000 respectively. Notice how the projected space separates  $\bar{p}$  and  $\bar{q}$ .

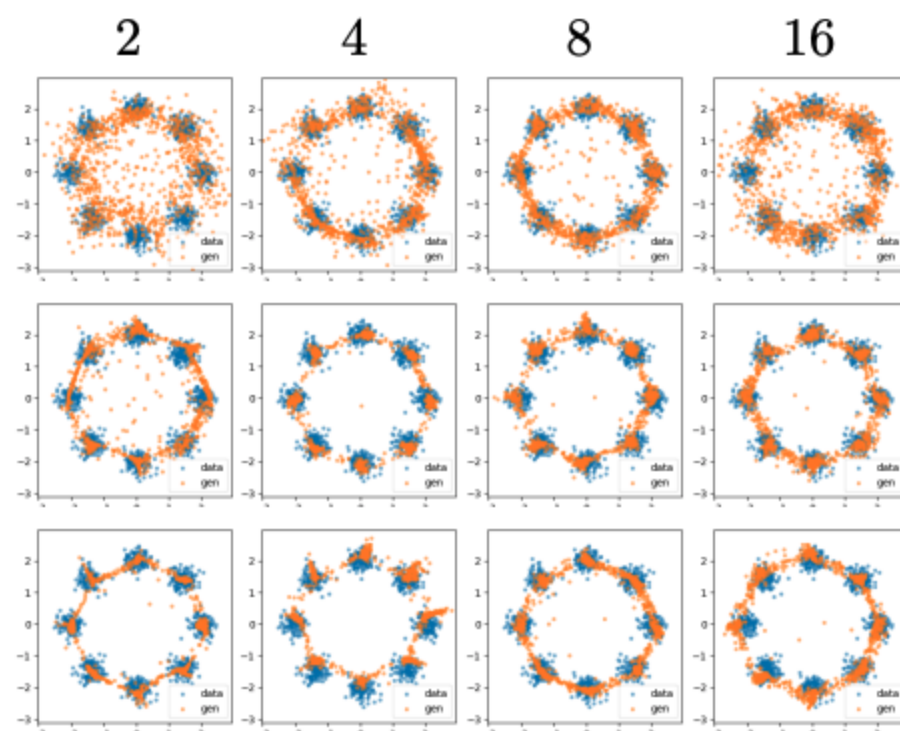


(b) Trace of  $\hat{\mathcal{L}}_\gamma$  and  $D^*$  (equation (5)) during training. The left plot is for iteration 1 to 100 and the right plot is for 100 to 5,000, with the same y-axes in the log scale.

Figure 1: Training results with projected dimension fixed to 2.

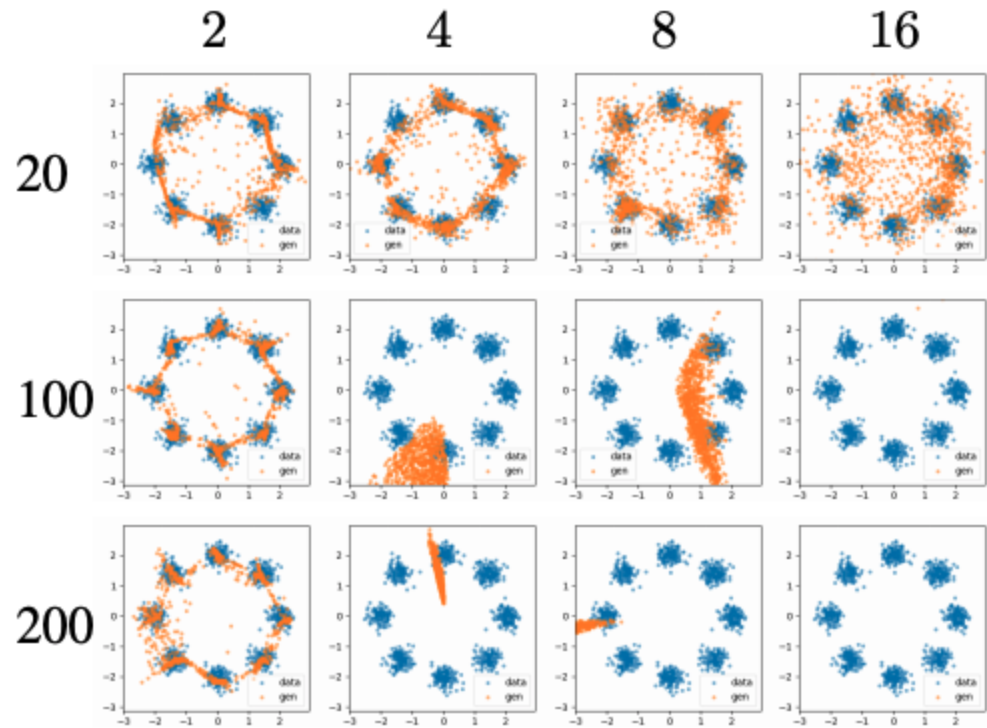


(a) MMD-nets

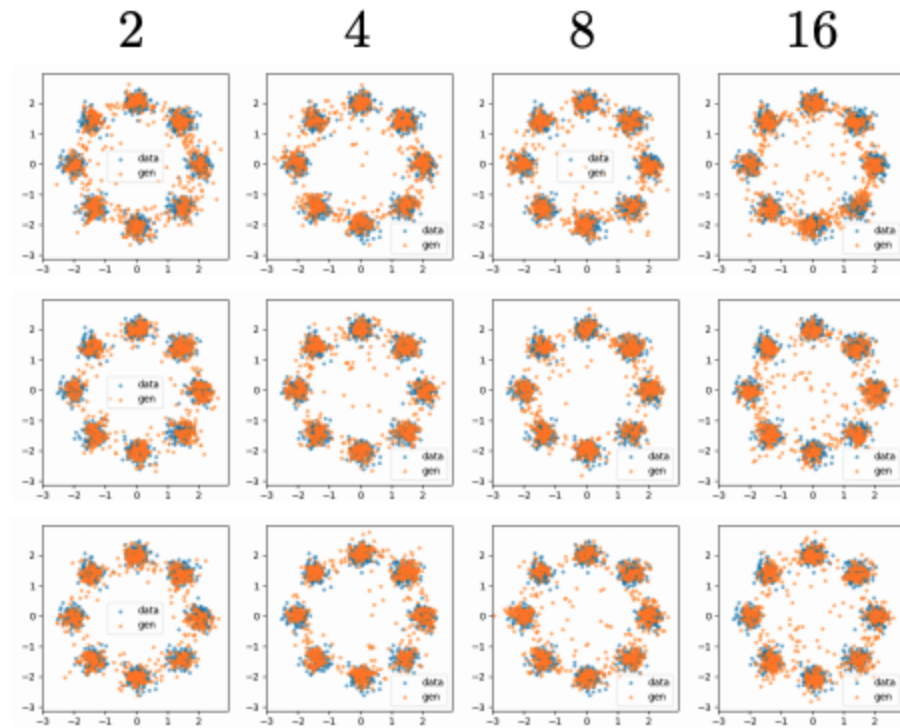


(b) MMD-GANs





(a) GAN



(b) GRAM-net

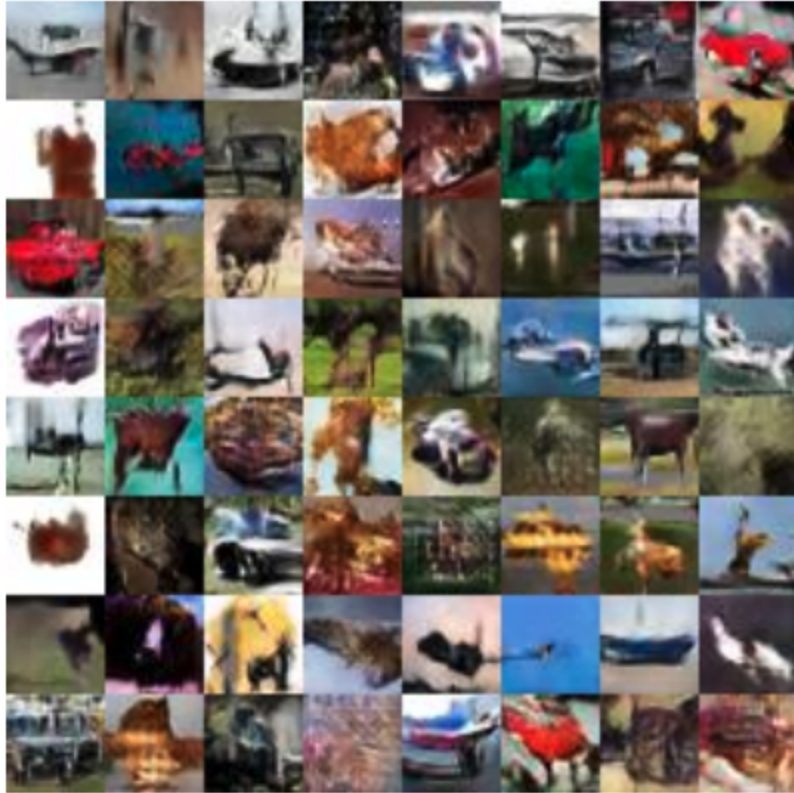
Figure 2: Training after 2,000 epochs by varying noise dimension  $h$  and the hidden layer size of critic model. For each model, each row is a different layer size in  $[20, 100, 200]$  and each column is a different  $h$  in  $[2, 4, 8, 16]$ . Half of the GAN training diverges while all GRAM training converges.

## Quantitative Results: Sample Quality

Table 1: Sample quality (measured by FID; lower is better) of GRAM-nets compared to GANs.

Arch.	Dataset	MMD-GAN	GAN	GRAM-net
DCGAN	Cifar10	$40.00 \pm 0.56$	$26.82 \pm 0.49$	<b><math>24.85 \pm 0.94</math></b>
Weaker	Cifar10	$210.85 \pm 8.92$	$31.64 \pm 2.10$	<b><math>24.82 \pm 0.62</math></b>
DCGAN	CelebA	$41.105 \pm 1.42$	$30.97 \pm 5.32$	<b><math>27.04 \pm 4.24</math></b>

# Qualitative Results: Random Samples



(a) CIFAR10



(b) CelebA

# The End!

Extra slides to follow...

# Density Ratio Estimation via (Infinite) Moment Matching

*Maximum mean discrepancy*

$$\text{MMD}_{\mathcal{F}}(p, q) = \sup_{f \in \mathcal{F}} (\mathbb{E}_p[f(x)] - \mathbb{E}_q[f(x)])$$

Gretton et al. (2012) show that it is sufficient to choose  $\mathcal{F}$  to be a unit ball in an reproducing kernel Hilbert space  $\mathcal{R}$  with a characteristic kernel  $k$ .

Using this definition of MMD, the density ratio estimator  $r(x)$  can be derived as the solution to

$$\min_{r \in \mathcal{R}} \left\| \int k(x; \cdot) p(x) dx - \int k(x; \cdot) r(x) q(x) dx \right\|_{\mathcal{R}}^2.$$

## Generator Training

- The generator  $G_\gamma$  is trained by minimizing the empirical estimator of MMD,

$$\min_{\gamma} \left[ \frac{1}{N^2} \sum_{i=1}^N \sum_{i'=1}^N k(f_{\theta}(x_i), f_{\theta}(x_{i'})) - \frac{2}{NM} \sum_{i=1}^N \sum_{j=1}^M k(f_{\theta}(x_i), f_{\theta}(G_{\gamma}(z_j))) \right. \\ \left. + \frac{1}{M^2} \sum_{j=1}^M \sum_{j'=1}^M k(f_{\theta}(G_{\gamma}(z_j)), f_{\theta}(G_{\gamma}(z_{j'}))) \right]$$

with respect to its parameters  $\gamma$ .