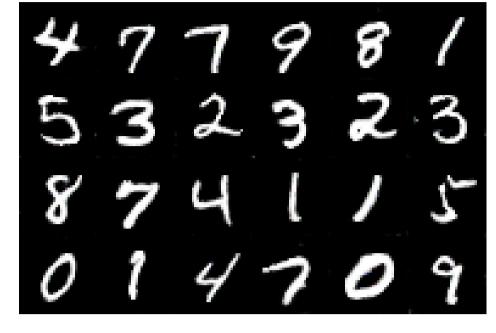
# Generative Models and Model Criticism via Optimized Maximum Mean Discrepancy

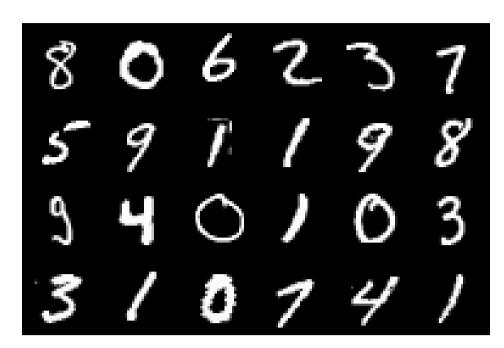


#### **Two-sample testing**

Say we observe two different datasets:



 $X \sim \mathbb{P} \pmod{6}$ Our question: is  $\mathbb{P} = \mathbb{Q}$ ?



 $Y \sim \mathbb{Q}$  (MNIST samples)

- Did my generative model actually learn the distribution I wanted it to?
- Do smokers and non-smokers have different distributions of cancers?
- Do these neurons fire differently when the subject is looking at image A instead of B?
- Are these different data sources the same? We want
- to be able to detect any possible difference,
- without making parametric assumptions,
- on high-dimensional data.

#### Maximum mean discrepancy

Distance between distributions [2] based on a kernel on sample points  $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ :

 $\mathrm{MMD}^{2}(\mathbb{P},\mathbb{Q}) = -2 \mathbb{E}_{X \sim \mathbb{P}, Y \sim \mathbb{Q}}[k(X,Y)]$  $+ \mathbb{E}_{X,X'\sim\mathbb{P}}[k(X,X')] + \mathbb{E}_{Y,Y'\sim\mathbb{O}}[k(Y,Y')].$ 

Estimate the MMD by taking sample means.

#### MMD tests

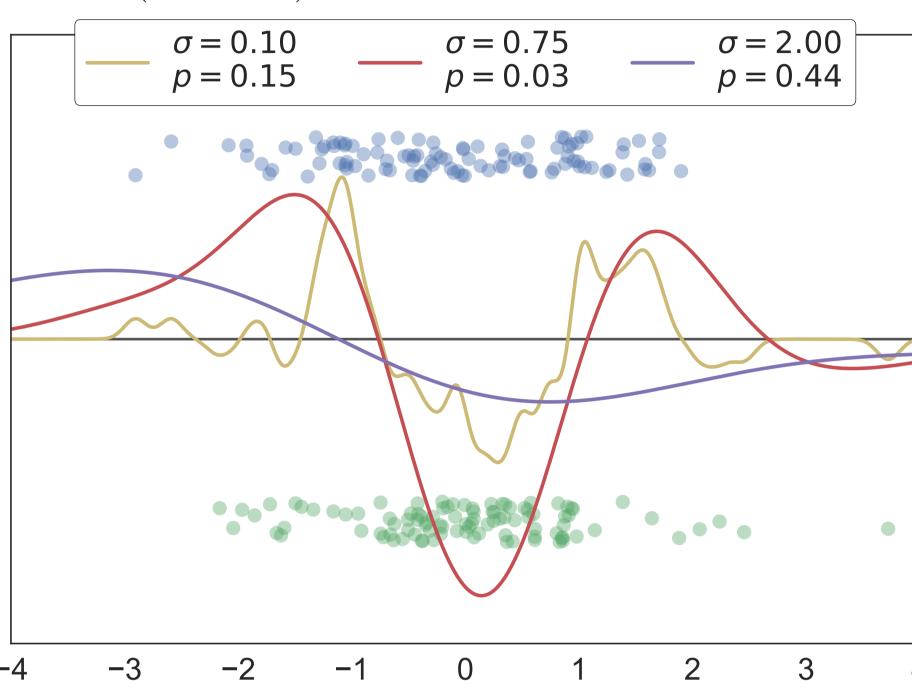
- Estimate  $MMD(\mathbb{P}, \mathbb{Q})$  with  $\widehat{MMD}(X, Y)$ .
- Estimate a threshold  $\hat{c}_{\alpha}$ :
- shuffle up  $X \cup Y$  into random halves many times;
- take  $\hat{c}_{\alpha}$  as the  $1 \alpha$ th quantile of the  $\widehat{\text{MMD}}^2$ s.
- Say  $\mathbb{P} \neq \mathbb{Q}$  if  $\widehat{m \text{MMD}}^2(X, Y) > \hat{c}_{\alpha}$ .

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### Kernel choice matters!

- Test blue  $\mathbb{P}$  versus green  $\mathbb{Q}$  with Gaussian kernels.
- Witness function  $\mathbb{E}_{X \sim \mathbb{P}}[k(X, \cdot)] \mathbb{E}_{Y \sim \mathbb{Q}}[k(Y, \cdot)]$ shows which locations more indicative of  $\mathbb{P}$  (top) or of  $\mathbb{Q}$  (bottom).

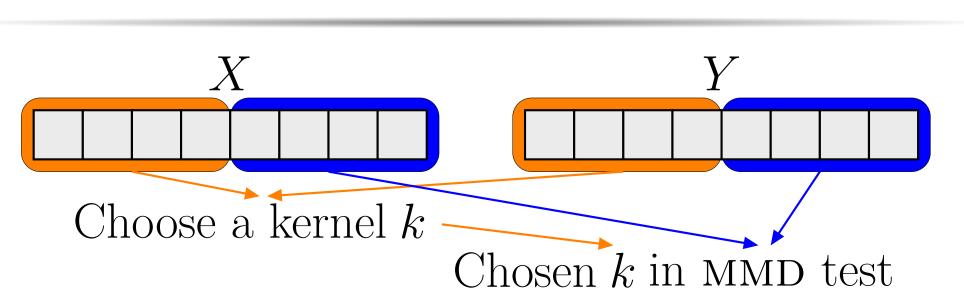


• Too small bandwidth: overfits to minor variation. • Too wide a bandwidth: not confident enough.

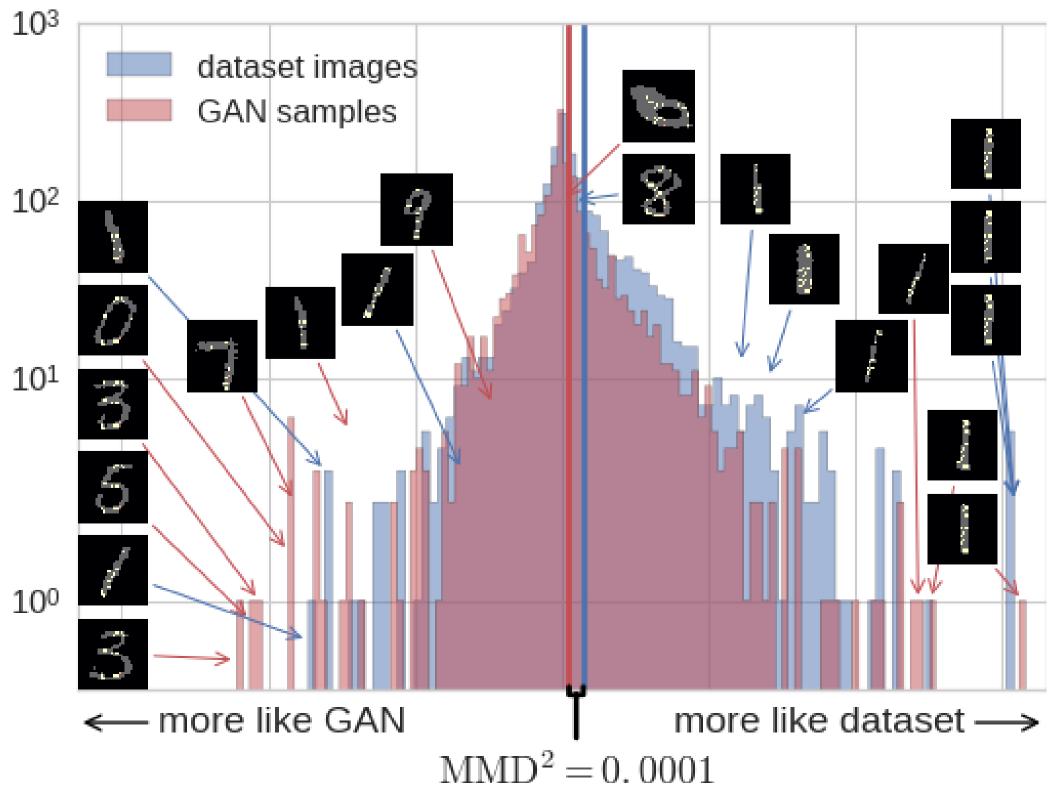
# Optimizing MMD test power

• When $\mathbb{P} \neq \mathbb{Q}$ , $\widehat{\text{MMD}}^2$ is asymptotically normal:
$\frac{\widehat{\mathrm{MMD}}^2(X,Y) - \mathrm{MMD}^2(\mathbb{P},\mathbb{Q})}{\sqrt{V_m(\mathbb{P},\mathbb{Q})}} \xrightarrow{D} \mathcal{N}(0,1).$
• Then test power $\Pr\left(m \widehat{MMD}^2(X, Y) > \hat{c}_\alpha\right)$ goes to
$\Phi\left(\frac{\mathrm{MMD}^2(\mathbb{P},\mathbb{Q})}{\sqrt{V_m(\mathbb{P},\mathbb{Q})}} - \frac{c_\alpha}{m\sqrt{V_m(\mathbb{P},\mathbb{Q})}}\right).$
• $V_m = O(m^{-1})$ ; MMD, $c_\alpha$ are constant in $m$ .
• So, maximize $\hat{t} = \widehat{\text{MMD}}^2(X, Y) / \sqrt{\hat{V}_m(X, Y)}$ .
• $\hat{V}_m$ : quadratic-time, unbiased estimator of $V_m$ .
• Maximize kernel parameters with backprop.

# **Train-test splits**



- Gaussian-ARD kernel: p-values almost exactly 0.
- Pixel weights (right) show where the model's distribution differs.
- Just optimizing bandwidth: 57% power at  $\alpha = .01$ .
- Median heuristic: 42% power.



 $10^{\circ}$ 

# Efficient permutation tests

• Current ways to compute permutations very slow. • Inefficient memory access pattern.

• Wrote a cache-aware implementation in Shogun.

— Our perm.

— MKL spectr

Problem size

• 15-30x the speed of existing implementations.

- Faster, more scalable than spectral approximations.

# Model criticism

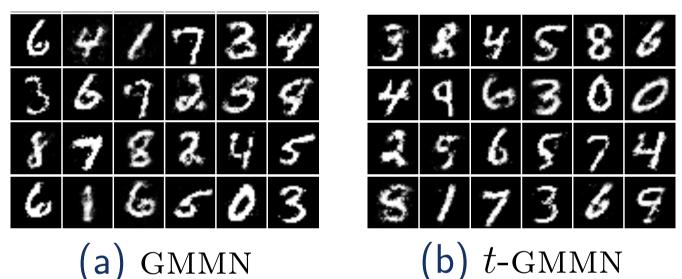
- [6]'s MNIST GAN is really good (top-left).
- Can we tell the distributions apart? Yes!

Looking at points with high/low witness function values from ARD kernel (like [5]) gives more insight: Model underproduces vertical 1s,

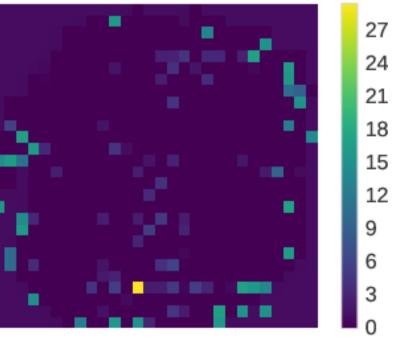
Overproduces right-slanted digits.

MMD value very small, but very consistent.

- sample at a time.



(a) GMMN



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# As a GAN objective

• Discriminators in standard GANs [3] look at one

• Problem: generator incentivized to produce just one sample that the discriminator likes, then gets stuck.

• Generator distribution should match true one.

• Use a two-sample test as the discriminator!

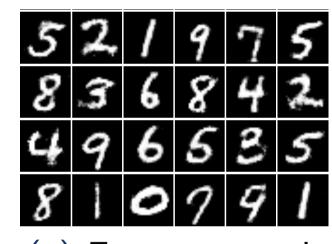
• [1, 4]: doing this by maximizing the MMD.

• Generative Moment Matching Network (GMMN)

• Instead, optimize  $\hat{t}$  criterion (*t*-GMMN).

• Or, do distributional feature matching (like [6]): • Train discriminator normally.

• Generator uses  $\hat{t}$  with kernel from discriminator.



(c) Feature match

• Used sum of Gaussian kernels:

• Not a great kernel on MNIST pixels.

• Nearly useless on natural image pixels.

• Gradients decay too fast.

• We're trying out better kernels.

#### References

[1] Dziugaite, Roy, and Ghahramani. Training generative neural networks via Maximum Mean Discrepancy optimization. UAI 2015.

[2] Gretton, Borgwardt, Rasch, Schölkopf, and Smola. A kernel two-sample test. JMLR 2012.

[3] Goodfellow, Pouget-Abadie, Mirza, Xu, Warde-Farley, Ozair, Bengio, Generative Adversarial Nets. NIPS 2014.

[4] Li, Swersky, and Zemel. Generative moment matching networks. UAI 2105.

[5] Lloyd and Ghahramani. Statistical model criticism using kernel two sample tests. NIPS 2015.

[6] Salimans, Goodfellow, Zaremba, Cheung, Radford, and Chen. Improved techniques for training GANs. NIPS 2016.