

REMEDI: Corrective Transformations for Improved Neural Entropy Estimation

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Motivation

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- Often, the quantity of concern $X \sim \mathbb{P}$ has a continuous distribution (probability measure) in \mathbb{R}^d . In other words $\mathbb{P} \ll \lambda$ (the Lebesgue measure) and \mathbb{P} has a density function p_X .
- For such a quantity we seek to estimate the **differential entropy**

$$H(\mathbb{P}) := \mathbb{E}[-\log p_X(X)].$$

Motivation

- Classical methods for estimating $H(\mathbb{P})$ based on samples $\{x_i\}_{i=1}^n$ from \mathbb{P} include: kernel density estimation (KDE), k -nearest neighbors estimates, methods based on sample spacings.

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- These possess nice asymptotic properties, like consistency, but fail in moderately high dimensions¹.

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- KDE may be improved by increasing the model class, e.g. going to Gaussian mixture models (GMMs).

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- These possess nice asymptotic properties, like consistency, but fail in moderately high dimensions¹.
- KDE may be improved by increasing the model class, e.g. going to Gaussian mixture models (GMMs).
- A recent development is entropy estimation with GMMs using gradient-based optimization of a cross-entropy target (KNIFE)².

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This work:

- Shows empirically that the problems of KDE estimates persist in for such GMM estimates.
- Introduces a deep learning-based correction to the estimates called REMEDI, demonstrating good performance in moderate dimensions.
- Proves theoretically that it satisfies a desirable consistency property.
- Investigates its performance in the Information Bottleneck context.

GMMs fail in moderate dimension

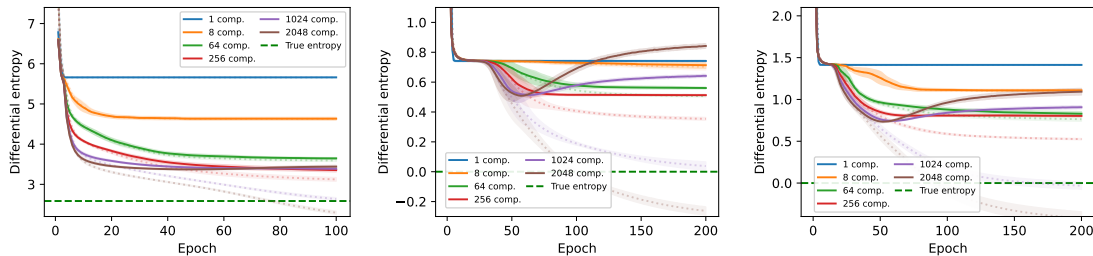


Figure: KNIFE training curves with error bars on 8-dimensional triangle and uniform ball/cube datasets. It is observed that increasing the number of components M for KNIFE leads to overfitting in all datasets.

Method

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- The difference between $H(\mathbb{P})$ and the cross-entropy $C(\mathbb{P}||\mathbb{Q})$ is then given by the relative entropy (KL-divergence) $R(\mathbb{P}||\mathbb{Q})$.

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- The difference between $H(\mathbb{P})$ and the cross-entropy $C(\mathbb{P}||\mathbb{Q})$ is then given by the relative entropy (KL-divergence) $R(\mathbb{P}||\mathbb{Q})$.
- Estimate $R(\mathbb{P}||\mathbb{Q})$ using Donsker-Varadhan's formula.

Method

Donsker-Varadhan's formula: For \mathbb{P}, \mathbb{Q} probability measures on \mathbb{R}^d such that $\mathbb{P} \ll \mathbb{Q}$ we have

$$R(\mathbb{P}||\mathbb{Q}) = \sup_{T \in \mathcal{C}_b} \mathbb{E}^{\mathbb{P}}[T] - \log \mathbb{E}^{\mathbb{Q}}[e^T]. \quad (1)$$

Such estimation has been used for mutual information³.

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- Questions:
 - Can we use empirical estimates in Eq. (1)?
 - How to optimize over \mathcal{C}_b ?
 - Can we obtain theoretical guarantees?

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Loss function

REMED I loss function: n samples from the data (\mathbb{P}) and m independent samples from the base (\mathbb{Q}) distribution,

$$\hat{\mathcal{L}}_{\text{REMED I}} = \underbrace{\frac{1}{n} \sum_{i=1}^n -\log q(x_i)}_{\hat{\mathcal{L}}_{\text{KNIFE}}} - \underbrace{\left(\frac{1}{n} \sum_{i=1}^n T(x_i) - \log \left(\frac{1}{m} \sum_{i=1}^m e^{T(\tilde{x}_i)} \right) \right)}_{\hat{\mathcal{L}}_{\text{DV}}} \quad (2)$$

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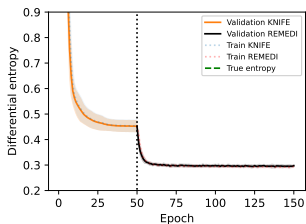
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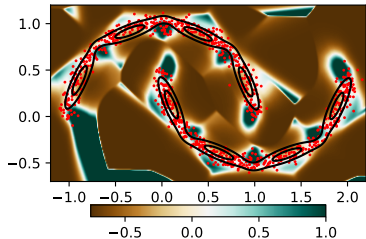
We select KNIFE⁴ as the base distribution.

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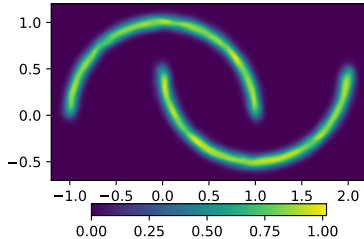
Entropy estimation



(a) Training curve



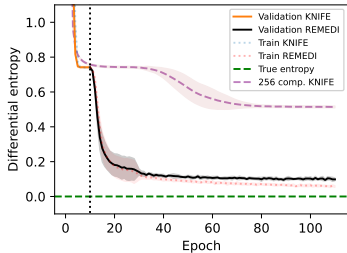
(b) T (image) vs. q (contours).



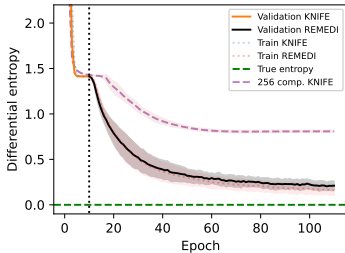
(c) Unnormalized density $qe^T(x)$

Figure: Results on two moons dataset. In the middle we see what direction (positive or negative) REMEDI affects the base distribution. To the right is the unnormalized distribution implied by $q(x)e^T(x)$.

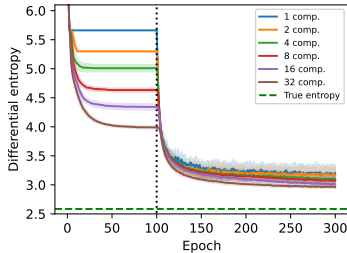
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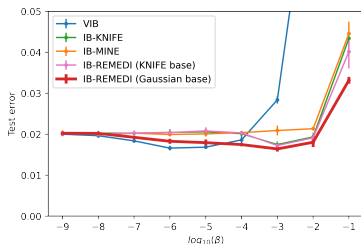
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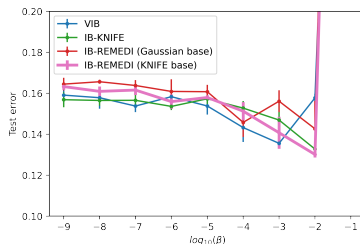
(c)

Figure: REMEDI training curves with error bars on 8-dimension uniform ball (a) and cube (b) datasets with 256-comp. KNIFE for reference. (c) The experiment on an 8-dimensional triangle dataset shows the effect of varying the number of components. REMEDI significantly improves the entropy estimation compared to KNIFE.

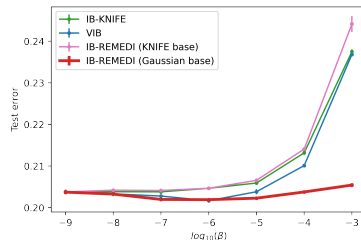
Information Bottleneck



(a) MNIST



(b) CIFAR10



(c) ImageNet

Figure: Plot of test error of the Information Bottleneck methods vs β on benchmark image classification datasets with error bars. For most β values, REMEDI outperforms the other methods on MNIST and ImageNet. On CIFAR10, the classification errors are similar for all the methods. However, REMEDI exhibits the lowest classification error across the β values.

Information Bottleneck

Information bottleneck latent space:

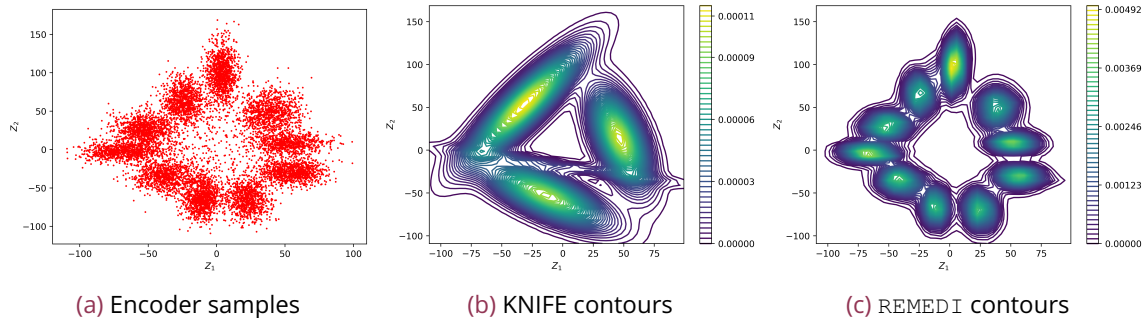


Figure: REMEDI marginal distribution of 2-d latent space on MNIST.

Generative capabilities

Rejection sampling: A sample X from \mathbb{Q} is accepted with probability $\phi(X)$, where $\phi(x) = \frac{e^{T(x)}}{\hat{C}}$.

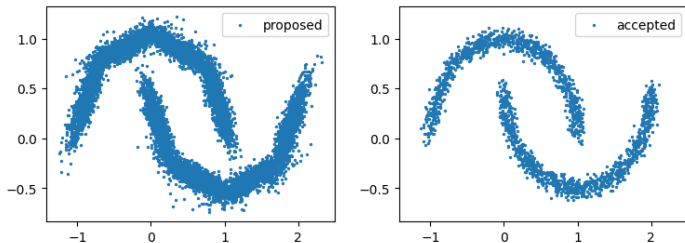


Figure: Left: 10000 proposals from \mathbb{Q} . Right: 1989 accepted samples.

Generative capabilities

Langevin diffusion:

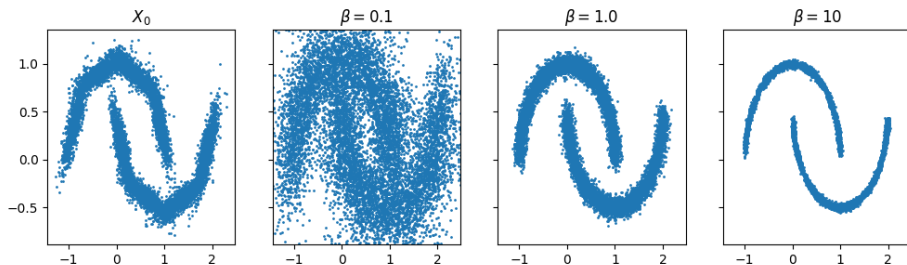





Figure: \mathbb{Q} -samples X_0 (leftmost) and X_{t_H} after simulating (3) with different β .

$$dX_t = -\nabla V(X_t)dt + \sqrt{2\beta^{-1}}dW_t, \quad X_0 = x_0, \quad V(x) = -(\log q(x) + T(x)) \quad (3)$$

Thank you!

-  Belghazi, Mohamed Ishmael et al. (2018). “Mutual information neural estimation”. In: *International conference on machine learning*. PMLR, pp. 531–540.
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-  Wasserman, Larry (2004). *All of statistics: a concise course in statistical inference*. Vol. 26. Springer.