1 **Deep neural networks**

Deep learning learns multiple (hierarchical) layer of data representation (feature). Neural networks scale with compute, data/model size (vs. other ML approaches). Artificial neural networks is a simplified version of biological NN.



- Training dataset $\{(\mathbf{x}_1, \mathbf{y}_1), \cdots, (\mathbf{x}_n, \mathbf{y}_n)\}$.
- NN $f(\mathbf{x}; \Theta) \in \mathbb{R}$ parameterized by Θ .
- Forward propagation $\hat{y} = \sigma(\Theta_k^T \sigma(\cdots \sigma(\Theta_1^T x)))$

NNs with ≥ 2 layers i.e. 1 hidden layer can model complex functions.

1.1 Universal approximation theorem

One hidden layer (with enough width) is enough to approximate all continuous functions.

1.1.1 Arbitrary width case

Let $C(X, \mathbb{R}^m)$ denote the set of continuous functions from a subset *X* of a Euclidean \mathbb{R}^n space to \mathbb{R}^m space. Let $\sigma \in C(\mathbb{R}, \mathbb{R})$. Note that $(\sigma \circ x)_i = \sigma(x_i)$, so $\sigma \circ x$ denotes σ applied to each component of x. Then σ is polynomial if and only if $\forall n \in \mathbb{N}, m \in \mathbb{N}$, compact $K \subseteq$ $\mathbb{R}^{n}, f \in C(K, \mathbb{R}^{m}), \epsilon > 0$ there exist $k \in \mathbb{N}, A \in \mathbb{R}^{k \times n}, b \in \mathbb{R}^{k}, C \in \mathbb{R}^{k}$ $\mathbb{R}^{m \times k} \text{ such that } \sup \|f(x) - g(x)\| < \epsilon \text{ where } g(x) = C \circ (\sigma \odot (A \cdot x + b)).$



(a) We can approximate continuous function with piece-wise linear functions.



(b) Can construct by subtracting 2 step functions.

Proof sketch

1.1.2 Bounded depth and bounded width case

There exists an activation function σ which is analytic, strictly increasing and sigmoidal and has the following property: For any $f \in C[0,1]^d$ and $\epsilon > 0$ there exists constant $d_i, c_{ii}, \theta_{ii}, \gamma_i$ and vectors $\boldsymbol{w}^{ij} \in \mathbb{R}^d$ for which

$$\left|f(\boldsymbol{x}) - \sum_{i=1}^{6d+3} d_i \sigma \left(\sum_{j=1}^{3d} c_{ij} \sigma(\boldsymbol{w}^{ij} \cdot \boldsymbol{x} - \theta_{ij}) - \gamma_i\right)\right| < \epsilon$$

for all $\mathbf{x} = (x_1, \cdots, x_d) \in [0, 1]^d$.

Training DNNs 1.2

Objective: find a parameter that minimizes error (or empirical risk)

$$\min_{\Theta} \frac{1}{n} \sum_{i=1}^{n} \ell(f(\boldsymbol{x}_i; \Theta), y_i) := L(\Theta) \qquad \Theta^{(t+1)} = \Theta^{(t)} - \gamma \nabla L(\Theta^{(t)})$$

where $\ell(\cdot, \cdot)$ is a loss function. Gradient descent (GD) updates parameters iteratively to the gradient direction.

1.2.1 Backpropagation

TL;DR Adjust the last layer weights Θ_k . Propagate error back to each previous layers. Repeat for $\Theta_{k-1}, \dots, \Theta_1$.

Consider the input (\mathbf{x}_i, y_i) . Forward propagation to compute $\hat{y}_i =$ $f(\mathbf{x}_i; \Theta)$. *i*-th layer intermediate output $s_i = \Theta_i^T \mathbf{h}_{i-1}$. Compute MSE loss $\ell(\hat{y}_i, y_i) = 1/2 (y_i - \hat{y}_i)^2 = E_i$.

$$\frac{\partial E_i}{\partial \hat{y}_i} = \frac{\partial}{\partial \hat{y}_i} \frac{1}{2} (y_i - \hat{y}_i)^2 = -(y_i - \hat{y}_i)$$
$$\frac{\partial E_i}{\partial s_k} = \frac{\partial E_i}{\partial \hat{y}_i} \frac{\partial \hat{y}_i}{\partial s_k} = \frac{\partial E_i}{\partial \hat{y}_i} \frac{\partial}{\partial s_k} \sigma(s_k) = (y_i - \hat{y}_i) \sigma'(s_k)$$

1.2.2 Activation functions



 $tanh(x) = 2\sigma(2x) - 1$ and $dtanh(x)/dx = 1 - tanh^2(x)$

1.2.3 Subdifferential

Set of all subgradients of f at x is called the **subdifferential** of f at x, written $\partial f(x)$ if

- $\partial f(x)$ is a closed set
- $\partial f(x)$ nonempty (if f convex, and finite near x)
- $\partial f(x) = \{\nabla f(x)\}$ if f is differentiable at x
- if $\partial f(x) = \{g\}$, then *f* is differentiable at *x* and $g = \nabla f(x)$



The absolute value function (left), and its subdifferential (right)

In many cases, don't need complete $\partial f(x)$; sufficient to find one $g \in \partial f(x)$.

1.2.4 GD in practice

	Batch	Stochastic
Calculate gradients with	All data	Subset of data
Computation	Heavy	Less
Convergence	Quick	Long
Avoid local optimum	Hard	Sometimes

• Standardization can be helpful; Increase convergence speed.

- e.g. Max-min or z-score normalization
- Mini-batch training + standardization can be a good option.

1.2.5 Optimizers

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Momentum accelerates GD when we have surface that curves more steeply in one direction than in another.

Momentum dampens the oscillation.

Adam calculates individual adaptive learning rate for each parameter from estimates of first and second moments of gradients:

$$m_{t} = \beta_{1}m_{t-1} + (1 - \beta_{1})g_{t}, \quad v_{t} = \beta_{2}v_{t-1} + (1 - \beta_{2})g_{t}^{2}$$
$$\hat{m}_{t} = \frac{m_{t}}{1 - \beta_{1}^{t}}, \quad \hat{v}_{t} = \frac{v_{t}}{1 - \beta_{2}^{t}} \quad \text{(Biased corrected estimates)}$$
$$\theta_{t+1} = \theta_{t} - \frac{\eta\hat{m}_{t}}{\sqrt{\hat{v}_{t} + \epsilon}} \quad \text{(Parameter update)}$$



1.2.6 Parameter initialization

 $W \sim N(0, \operatorname{Var}(W))$ where

$$\operatorname{Var}(W) = \begin{cases} \sqrt{1/n_{\text{in}}} & \text{(LeCun normal init.)} \\ \sqrt{2/(n_{\text{in}} + n_{\text{out}})} & \text{(Xavier normal init.)} \\ \sqrt{2/n_{\text{out}}} & \text{(He normal init.)} \end{cases}$$

Batch normalization 1.2.7

- Allow higher LR and reduce strong • dependence on initialization.
- Activations have different distributions. BN makes them similar.
 - After FC/Conv layer and before non-linearity layer.

Batch 2

Input: Values of x over a mini-batch: $\mathcal{B} = \{x_{1...m}\};$ Parameters to be learned: γ , β **Output:** $\{y_i = BN_{\gamma,\beta}(x_i)\}$



1.3 **Convolutional neural networks**

- Convolution: Weight sharing and local connectivity
 - + Translation invariance
 - + Reduce the number of parameters (less overfitting)
 - + Learn local features
- Pooling (subsampling): operates on each activation map independently
 - + Translation invariance ↑ (to small transformations), Regularization
 - + Reduce the number of parameters and computation
- ConvNet is sequence of conv layers followed by non-linearity.



1.4 **Recurrent neural networks**

Markov chain $Pr(w_{i+1}|w_i)$. Language model $Pr(w_{i+1}|w_1, \dots, w_i)$ becomes a large-scale classification task at every time *i* since its vocabulary size is large.

RNN models temporal information. Hidden states as a function of inputs and previous time step information.



Process sequence of vectors by applying recurrence formula at every t In simple RNN (or vanilla RNN), the state consists of single hidden vector. Recurrence formula becomes $h_t = \tanh(\Theta_{hh}h_{t-1} + \Theta_{xh}x_t)$. Then compute $y_t = \Theta_{hv} h_t$.

1.4.1 Gradient vanishing

RNN with very long sequence suffer from gradient vanishing, where gradients become zeros during backpropagation. ReLU is sometimes problematic.

1.4.2 Backpropagation through time (BPTT)

Most common method used to train RNNs.

- The unfolded network (used during forward pass) is treated as one big FFN that accepts the whole time series as input
- The weight updates are computed for each copy in the unfolded network, then summed (or averaged) and applied to RNN weights
- In practice, truncated BPTT is used: run the RNN forward k_1 time steps, propagate backward for k_2 time steps



BPTT and computation graph

1.4.3 Long short-term memory (LSTM)

Add a memory cell that is not subject to matrix multiplication or squishing (e.g., sigmoid), thereby avoiding gradient decay.



1.4.4 Examples

- Many-to-many: machine translation
- Many-to-one: sentence classification
- One-to-many: image captioning

Residual network (ResNet) 1.5



(a) ResNet

(b) This direct path helps maintain the gradient's magnitude and prevents it from vanishing.





2 Ordinary differential equations

2.1 Differential equations

Let h(t) a state vector. dh(t)/dt is a differential equation describing how h(t) change over time. We are interested in solving the following initial value problem (IVP) to know the state in future.

$$\boldsymbol{h}(T) = \boldsymbol{h}(0) + \int_0^T \frac{d\boldsymbol{h}}{dt} dt$$

IVPs are sometimes analytically solved. Otherwise we rely on a solver to approximate the solution.

- T_1 has 100 liters of water, and T_2 has 100 liters of fertilizer.
- $z(t)=(z_1(t), z_2(t))$ means the amount of fertilizer at time t.

 z_1' = inflow per minute – outflow per minute = $-0.03 z_1 + 0.03 z_2 z_2'$ = inflow per minute – outflow per minute = $0.03 z_1 - 0.03 z_2$

:. z' = Az or z' - Az = 0, where $A = \begin{bmatrix} -0.03 & 0.03 \\ 0.03 - 0.03 \end{bmatrix}$

 T_1

 T_2

When we have an initial value of z(0)=(0, 100), what is z(2)? This kind of problem is called initial value problem (IVP) or forward problem.
Given data, what is A? This kind of problem is called backward problem.

Example of ODE

2.2 ODE solvers



Euler method. Look similar to residual connection

$$h(t + h) = h(0) + hf(h(t)),$$

$$h(t + 2h) = h(t + h) + hf(h(t + h)), \quad \cdots$$
Now pick a step-size $h > 0$ and define
$$y_{n+1} = y_n + \frac{1}{6}h(k_1 + 2k_2 + 2k_3 + k_4),$$

$$t_{n+1} = t_n + h$$
for $n = 0, 1, 2, 3, ..., using^{[3]}$

$$k_1 = f(t_n, y_n),$$

$$k_2 = f\left(t_n + \frac{h}{2}, y_n + h\frac{k_1}{2}\right),$$

$$k_3 = f\left(t_n + \frac{h}{2}, y_n + h\frac{k_2}{2}\right),$$

$$k_4 = f(t_n + h, y_n + hk_3).$$

Runge-Kutta (RK) method

Dormand–Prince (DOPRI) method. After comparing the RK4 and RK5 results, use a large step-size h if the difference is small, and small h if the difference is large. In other words, the (adaptive) size-size is inversely proportional to the estimated difference.

2.3 Neural ODE

Parameterize the hidden units using ODE specified by neural network:

$$\frac{d\boldsymbol{h}(t)}{dt} = f(\boldsymbol{h}(t), t, \theta)$$

Starting from the input layer h(0), we can define the output layer h(T)to be the solution to this ODE initial value problem at some time *T*. This can be computed by black-box differential equation solver.



Left: ResNet defines a discrete sequence of finite transformations. **Right**: A ODE network defines a vector field, which continuously transforms state. **Circles**: evaluation locations.

Defining/evaluating models using ODE solvers has several benefits:

- Memory efficiency: not store intermediate quantities of forward $\rightarrow O(1)$ memory learning
- Modern ODE solvers provide **error estimate** and evaluation (step size) **adaptive** to given resource

2.3.1 NODE-based image classifier

- A typical construction: feature extraction \rightarrow NODE \rightarrow output
- NODE layer is analogous to (continuous) residual layers
- Can use standard backpropagation algorithm to train.



2.3.2 Adjoint sensitivity method

Differentiating through the operations of forward pass is straightforward, but incurs a high memory cost and introduces numerical error. For example, depth of DOPRI frequently becomes large.

We treat the ODE solver as a black box, and compute gradients using the adjoint sensitivity method. Consider optimizing L(), whose input is the result of an ODE solver.

$$L(z(t_1)) = L\left(z(t_0) + \int_{t_0}^{t_1} f(z(t), t, \theta) dt\right) = L(\text{ODESolve}(z(t_0), t_0, T, \theta))$$

We first determine the *adjoint* $\mathbf{a}(t) = \partial L/\partial z(t)$. Its dynamics are given by another ODE, which can be thought of as the instantaneous analog of the chain rule:

$$\frac{d\boldsymbol{a}(t)}{dt} = -\boldsymbol{a}(t)^T \frac{\partial f(\boldsymbol{z}(t), t, \theta)}{\partial \boldsymbol{z}}$$

We can compute $\partial L/\partial z(t_0)$ by another call to an ODE solver. This solver must run backwards, starting from initial value of $\partial L/\partial z(t_1)$. This will require knowing value of z(t) along its entire trajectory, but we can simply recompute z(t) backwards in time together with the adjoint, starting from its final value $z(t_1)$. We can calculate the gradients with a **reverse-mode integral**,

$$\frac{dL}{d\theta} = -\int_{t_1}^{t_0} \boldsymbol{a}(t)^T \frac{\partial f(\boldsymbol{z}(t), t, \theta)}{\partial \theta} dt$$

No need to maintain computation graph of NODEs $\rightarrow O(1)$ space.

2.3.3 Analogy to ResNet



3 Transformers

3.1 Background

Context vector c is often just h_T . Input sequence is bottlenecked through fixed-sized vector. What if sequence is very long?

3.1.1 Image captioning with CNN and RNN

- Transfer learning: take last layer of CNN trained to ImageNet
- Final representation v of CNN is provided to RNN. Now $h = \tanh(W_{xh}x + W_{hh}h + W_{ih}v)$
- Sample word and copy to input. Stop after sampling <END> token

3.2 Watson-Nadaraya estimator

Data $\{x_1, \dots, x_m\}$ and ground-truths $\{y_1, \dots, y_m\}$. Estimate *y* at a new location *x*. Naive way is just average. Watson-Nadaraya estimator weigh the ground truths: $y = \sum_i a(x, x_i)y_i$ where *x* is query, x_i is key, and y_i is value.

- Consistency: given enough data, converges to optimal solution
- Simplicity: no free parameters: information is in data not weights
- Deep learning variant: learn the weighting function

3.2.1 Seq2Seq with RNN and attention



RNN need attention for parallelization and deal with long-range dependencies. Here, decoder doesn't use the fact that h_i form an ordered sequence–it just treats them as an unordered set $\{h_i\}$.

3.3 Attention

- Basic attention layer. Given query q ∈ ℝ^d, input X ∈ ℝ^{n×d}. Similarities e_i = f_{att}(q, x_i), e ∈ ℝⁿ. Attention weights a = softmax(e) ∈ ℝⁿ. Output vector y = ∑ a_ix_i ∈ ℝ^d.
- Commonly separate key and value: $K = W_k X, V = W_v X$. Similarities $E = f_{\text{att}}(K, Q) \in \mathbb{R}^{n \times n}$, attention weights $A = \text{softmax}(E) \in \mathbb{R}^{n \times n}$. Output vector $y_j = \sum a_{ij} v_i, y \in \mathbb{R}^d$
- Similarity functions f_{att} : $q^T x_i$ (dot product), $q^T x_i / \sqrt{d}$ (scaled dot product), $f(Q, X) = Q^T X$ (multiple queries product).
- Self-attention layer has one query per input. Q = X ∈ ℝ^{n×d}. Uses scaled dot product. Permutation equivariant i.e. f(s(x)) = s(f(x)) and works on sets of vectors.

3.3.1 Advantages of attention

- Allows decoder to focus on certain parts of source → Significantly improved NMT and time series performance
- Shortcut between faraway states \rightarrow Mitigates vanishing gradient
- Provides some interpretability

Weighted sum is a selective summary of the information contained in values. Way to **obtain a fixed-size representation** of an arbitrary set of representations (values) dependent on other (query).

Scaled dot-product attention attends to one or few entries in the input key-value pairs. (\leftrightarrow human) **Multi-head SA** split inputs, use *H* independent heads in parallel, and concatenate ouptuts.

3.3.2 Positional encoding

Unlike RNN, attention encoder outputs do not depend on the order of inputs, which is important. Concatenate positional information of input token to input embedding.

$$PE_{pos,2i} = \sin(pos/10000^{2i/d}),$$

$$PE_{pos,2i+1} = \cos(pos/10000^{2i/d})$$

3.4 Transformers

SA is the only interaction between vectors. LayerNorm and MLP work independently per vector. \rightarrow Highly scalable and parallelizable.

4 Generative models

A generative task aim to learn p(x) and generate fake samples from learned distribution $p_{\theta}(x)$.

- Quality vs. diversity dilemma:
 - Likelihood-based GMs directly learns the *pdf* of training data e.g. normalizing flows, VAE.
 - Implicit GMs do not directly maximize the likelihood of training data but implicitly and internally learn it.
- Score-based models (SGMs) propose a novel paradigm i.e., learning the gradient of the log *pdf*, a quantity often known as the (Stein) score function.

4.1 Flow-based models

4.1.1 Mathematical background

Consider a 2-d coordinate (x, y) and invertible transformation *T*. (u, v) = T(x, y) and $(x, y) = T^{-1}(u, v)$. Jacobian matrix **J** is all first-order partial derivatives of this transformation

$$\mathbf{J} = \begin{bmatrix} \frac{\partial x}{\partial u} & \frac{\partial x}{\partial v} \\ \frac{\partial y}{\partial u} & \frac{\partial y}{\partial v} \end{bmatrix}$$

Change of Variable theorem (CVT) states

$$\int \int_{S} f(x, y) \cdot dx dy = \int \int_{T(S)} f(T^{-1}(u, v)) \cdot \left[\frac{\partial x}{\partial u} \frac{\partial y}{\partial v} - \frac{\partial x}{\partial v} \frac{\partial y}{\partial u} \right] du dv$$

CVT for probability density estimation

1

$$\log p(u, v) = \log p(x, y) + \log \left| \det \frac{\partial T}{\partial(x, y)} \right|$$



Interpretation of $dxdy = |\det \partial T/\partial(x, y)| dudv$. *f* sends a small square to a distorted parallelogram. The Jacobian at a point gives the best linear approximation of the distorted parallelogram near that point, and the det **J** gives the ratio of the area of the approximating parallelogram to that of the original square.

4.1.2 Density estimation in NODEs

Suppose we design a generator using NODEs. z(0) typically follows a unit Gaussian. So we know log p(z(0)). We can estimate p(z(1)) as log $p(z(1)) = \log p(z(0)) + \log |\det of \operatorname{Jacobian} \operatorname{atz}(0)|$, then p(z(2)), and so on. Suppose z(2) is specific image. We know the probability that this specific image generated by the generator.



Transformer block

4.2 Generative adversarial networks (GAN)

Zero-sum minimax game between two players

$$\underset{G}{\operatorname{minmax}} V(G, D) = \mathbb{E}[\log D(x)]_{x \sim p_{\text{data}}(x)} + \mathbb{E}[\log(1 - D(G(x)))]_{z \sim p(z)}]_{z \sim p(z)}$$

To maximize, D(x) = 1 and D(G(z)) = 0. To minimize, D(G(z)) = 1.

4.2.1 Equilibrium state proof of GANs

First consider the optimal discriminator D for any given generator G.

Proposition 1. For G fixed, the optimal discriminator D is

$$D_G^*(\boldsymbol{x}) = \frac{p_{data}(\boldsymbol{x})}{p_{data}(\boldsymbol{x}) + p_g(\boldsymbol{x})}$$
(1)

Proof. The training criterion for the discriminator D, given any generator G, is to maximize the quantity V(G, D)

$$V(G, D) = \int_{\mathbf{x}} p_{\text{data}}(\mathbf{x}) \log(D(\mathbf{x})) dx + \int_{z} p_{z}(z) \log(1 - D(g(z))) dz$$
$$= \int_{\mathbf{x}} p_{\text{data}}(\mathbf{x}) \log(D(\mathbf{x})) + p_{g}(\mathbf{x}) \log(1 - D(\mathbf{x})) dx \qquad (2)$$

For any $(a, b) \in \mathbb{R}^2 \setminus \{0, 0\}$, the function $y \to a \log(y) + b \log(1 - y)$ achieves its maximum in [0, 1] at a/(a + b). The discriminator does not need to be defined outside of $\text{Supp}(p_{\text{data}}) \cup \text{Supp}(p_g)$. \Box

Note that the training objective for *D* can be interpreted as maximizing the log-likelihood for estimating the conditional probability $P(Y = y | \mathbf{x})$, where *Y* indicates whether \mathbf{x} comes from p_{data} (with y = 1) or from p_g (with y = 0). The minimax game is now:

$$C(G) = \max_{D} V(G, D)$$

= $\mathbb{E}_{\mathbf{x} \sim p_{\text{data}}} [\log D_{G}^{*}(\mathbf{x})] + \mathbb{E}_{z \sim p_{z}} [\log(1 - D_{G}^{*}(G(z)))]$ (3)

$$=\mathbb{E}_{\mathbf{x}\sim p_{\text{data}}}\left[\log D_{G}^{*}(\mathbf{x})\right] + \mathbb{E}_{\mathbf{x}\sim p_{g}}\left[\log(1 - D_{G}^{*}(\mathbf{x}))\right]$$
$$=\mathbb{E}_{\mathbf{x}\sim p_{\text{data}}}\left[\log \frac{p_{\text{data}}(\mathbf{x})}{P_{\text{data}}(\mathbf{x}) + p_{g}(\mathbf{x})}\right] + \mathbb{E}_{\mathbf{x}\sim p_{g}}\left[\log \frac{p_{g}(\mathbf{x})}{p_{\text{data}}(\mathbf{x}) + p_{g}(\mathbf{x})}\right]$$

Theorem 1. The global minimum of the virtual training criterion C(G) is achieved if and only if $p_g = p_{data}$. At that point, C(G) achieves the value $-\log 4$.

Proof. For $p_g = p_{data}$, $D_G^*(\mathbf{x}) = 1/2$, (consider Eq. 1). Hence, by inspecting Eq. 3 at $D_G^*(\mathbf{x}) = 1/2$, we find $C(G) = \log(1/2) + \log(1/2) = -\log 4$. To see that **this is the best possible value** of C(G), reached only for $p_g = p_{data}$, observe that

$$\mathbb{E}_{\boldsymbol{x} \sim p_{\text{data}}} \left[-\log 2 \right] + \mathbb{E}_{\boldsymbol{x} \sim p_g} \left[-\log 2 \right] = -\log 4$$

and that by subtracting this expression from $C(G) = V(D_G^*, G)$, we obtain:

$$C(G) = -\log(4) + KL\left(p_{\text{data}} \left\| \frac{p_{\text{data}} + p_g}{2} \right) + KL\left(p_g \left\| \frac{p_{\text{data}} + p_g}{2} \right)\right)$$
(4)

where KL is the Kullback–Leibler divergence. We recognize in the previous expression the Jensen–Shannon divergence between the model's distribution and the data generating process:

$$C(G) = -\log(4) + 2 \cdot JSD\left(p_{\text{data}} \| p_g\right)$$
(5)

Since the Jensen–Shannon divergence between two distributions is always non-negative and zero only when they are equal, we have shown that $C^* = -\log(4)$ is the global minimum of C(G) and that the only solution is $p_g = p_{\text{data}}$, i.e., the generative model perfectly replicating the data generating process.



4.2.2 Why Gaussian prior?

Gaussian distributions have the following favorable characteristics:

- The mean of many independent random variables will converge to a Gaussian distribution (*cf.* the central limit theorem)
- Encodes the least amount of prior knowledge i.e., the max entropy) into a model

4.3 Score-based generative models

The pdf is defined as $p_{\theta}(\mathbf{x}) = e^{-f_{\theta}(\mathbf{x})}/Z_{\theta}$. For training, we maximize the log-likelihood max $\sum \log p_{\theta}(\mathbf{x})i$) which is undesirable. We can bypass the intractable Z_{θ} by only considering the *score function*, which is the gradient of the log-density *w.r.t*. the random variable.

$$s_{\theta}(\mathbf{x}) = \nabla_{\mathbf{x}} \log p_{\theta}(\mathbf{x}) = -\nabla_{\mathbf{x}} f_{\theta}(\mathbf{x}) - \underbrace{\nabla_{\mathbf{x}} \log Z_{\theta}}_{0} = -\nabla_{\mathbf{x}} f_{\theta}(\mathbf{x})$$

Score function provides numerical benefits: range $[-\infty, \infty]$ and no need to normalize.

4.3.1 Score matching

Given the ground-truth score function, we can minimize the following Fisher divergence.

$$\mathbb{E}_{p(\boldsymbol{x})}[\|\nabla_{\boldsymbol{x}} \log p(\boldsymbol{x}) - s_{\theta}(\boldsymbol{x})\|_{2}^{2}]$$

After training, rely on Langevin dynamics to generate samples.

$$\mathbf{x}_{i+1} \leftarrow \mathbf{x}_i + \epsilon \nabla_{\mathbf{x}} \log p(\mathbf{x}) + \sqrt{2\epsilon z_i} \quad z_i \sim \mathcal{N}(0, I)$$

Pitfalls of score matching: Hard to learn correct score function for low-density regions since the L2 distance is weighted by p(x).

4.4 Noise conditional score-based models (NCSMs)

TL;DR: Diffuse data with Brownian motion i.e. perturb with Gaussian noise, and reverse this process.

NCSM minimize the following training objective where *i* is perturbation index:

$$\sum_{i=1}^{L} \lambda(i) \mathbb{E}_{p_{\sigma_i}(\boldsymbol{x})} [\|\nabla_{\boldsymbol{x}} \log p_{\sigma_i}(\boldsymbol{x}) - s_{\theta}(\boldsymbol{x}, i)\|_2^2]$$

$$\log p_{\sigma_i} = \int \underbrace{p(\boldsymbol{y})}_{\text{GT; hidden}} \underbrace{\mathcal{N}(\boldsymbol{x}; \boldsymbol{y}, \sigma_i^2 I) d\boldsymbol{y}}_{\text{Gaussian around } \boldsymbol{y}; \text{ transition probability}}$$

This is equivalent as learning score function for each σ_i . NCSM mathematically decompose original loss to constant plus loss of only transition probability, named **denoising score matching loss**.

4.5 Score-based generative models

SGMs continuously generalized diffusion models using SDEs.

$$d\mathbf{x} = \underbrace{f(\mathbf{x}, t)dt}_{\text{drift}} + \underbrace{g(t)d\mathbf{w}}_{\text{diffusion; }d\mathbf{w} \text{ is Gaussian nois}}$$



The authors propose 3 types of drift/diffusion: one of them is DDPM, one of them is NCSM.

At t = T, original distribution forgotten, noise term dominates; it is single Gaussian. If we know the score function at time t of forward process, we can stochastically reverse that forward process.

- Train score network. GT score function replaced to s_θ(x, t).
 Also rely on denoising score matching loss
- 2. Stochastically reverse single Gaussian to target distribution using Euler-Maruyama method
- \bullet Stochastically reverse \rightarrow high diversity
- If High quality (↔ GAN). This minimizes the upper bound of KL. JSD is more stable than KL, but global optimum of GAN is impossible to achieve in practice.
- \P Query score function T times \rightarrow high complexity

May use probability flow ODE for sampling, which deterministically reverse. Both have the same marginal probability $p_t(\mathbf{x}) \forall t$.



4.6 Poisson flow generative models

4.6.1 GNN basics

Degree matrix has number of neighbors at diagonal. **Laplacian** is degree matrix - adjacency matrix. Consider a graph *G* with Laplacian *L* and a graph signal (feature) $x \in \mathbb{R}^{N \times D}$ on *G*. Signal y = Lx i.e. $y_i = \sum_{j \in \mathcal{N}_i} (x_i - x_j)$. y_i measures difference between *x* at a node and its neighborhood, i.e. difference operator.

4.6.2 Heat diffusion over graph

Via heat equation $x_t = -Lx$, we say the signal diffuses through graph. Lx is positive if my temperature > neighbor \rightarrow this equation means 'cooling down'. Temperature at each location is averaged with its neighbors. Converges to average temperature.

The Euler method for temperature update is x(t+h) = x(t) - hLx(t). *h* can be interpreted as learning rate in DL but step size of Euler method in physics.

4.6.3 Graph convolutional network (GCN)

Let x(t) the hidden vector for each node at layer t. Consider a row-wise normalized Laplacian \tilde{L} . Euler discretized heat equation was $x(t + 1) = x(t) - \tilde{L}x(t) = (1 - \tilde{L})x(t)$

GCN uses diffusion process $x(t+1) = \sigma((1-\tilde{L})x(t)W)$. A trainable parameter (or diffusivity/conductivity) $W \in \mathbb{R}^{N \times N}$ represents how close two nodes are. This is just W multiplied to Euler discretized heat equation. The inductive bias of GCN is heat diffusion equation and is influenced by physics (*physics bias*). At the same time, one can consider that this is a first-order graph filtering approach. \tilde{A} is normalized adjacency matrix.

Given
$$H = \sum_{\ell=0}^{L} h_{\ell} S^{\ell}$$
, $\ell = 1$ and $S = I - \tilde{L} = \tilde{A}$

4.6.4 Oversmoothing problem

One major problem is oversmoothing problem: all nodes' last **hidden vectors become similar to each other** when the number of GCN layers is large. \rightarrow mean avg distance (MAD) decreases and node classification accuracy decreases as layer go deeper than some threshold.

Self attention is also GCN, since $1 - \tilde{L} = \tilde{A}$. So transformers also subject to oversmoothing problem. But in case of transformers \tilde{A} is learned not given.

4.6.5 Poisson flow models

Idea: at the beginning of sampling, move straight.

- Diffusion model is inspired from **thermodynamics**: any localized distribution of a gas will eventually spread out to fill an entire space evenly simply through random motion.
- Poisson flow generative models are inspired from electrostatics: any distribution of electrons in a hyperplane generates a uniform hemisphere.



A charge distribution (purple) and the electric field lines (black) it generates. If we let the charge distribution evolve along the field lines, it will transform into a uniform hemispherical distribution. z-axis corresponds to time of diffusion model.

4.6.6 Training Poisson flow models

- 1. Augment data with z = 0
- 2. Calculate the empirical field in random x, y, z-O(N) where N is number of training samples
- Calculate loss and update function approximator of empirical field, dx = -E(x)dt

4.6.7 Sampling Poisson flow models

- 1. Uniformly sample data on a large hemisphere
- 2. Use an ODE solver to evolve the points backwards along the Poisson field
- 3. Evolve backwards until we reach z = 0, at which point we have generated novel data from the training distribution.