

Generative Diffusion Models on Graphs: Methods and Applications

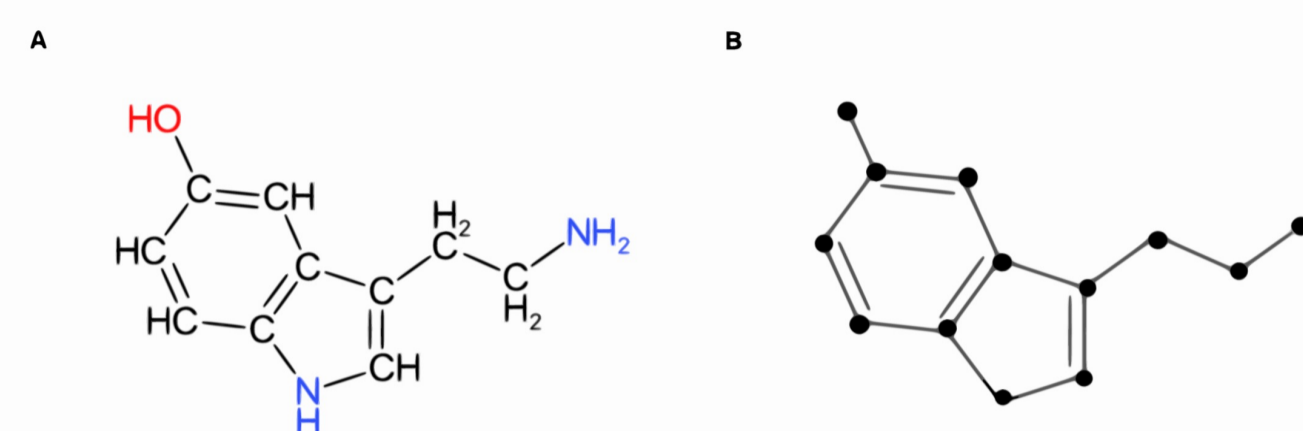
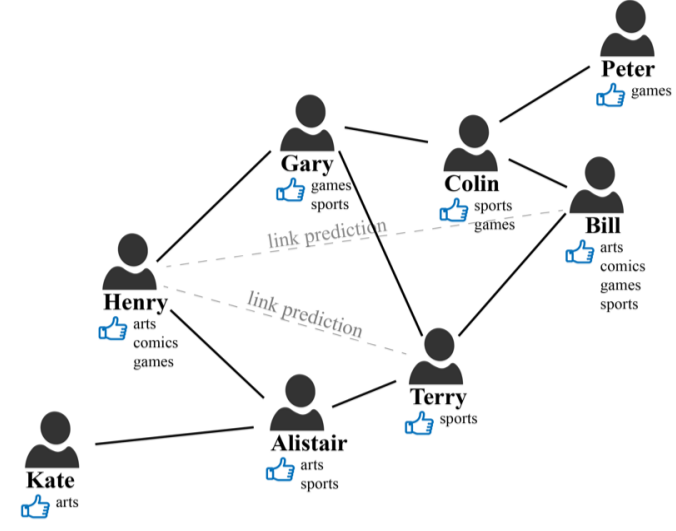
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Introduction

Graphs

Graphs can represent the rich variety of relationships between real-world entities. They have been widely used in a diversity of domains, aiming to model association information and structural patterns among various real-world objects.



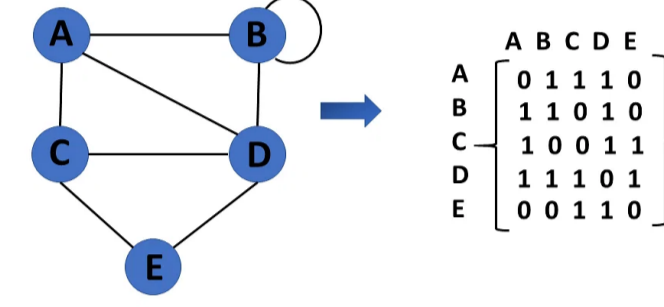
Graph Generation

Graph generative models, with the goal of learning the given graph distributions and generating novel graphs, can be categorized into two generation patterns:

- autoregressive generation
- one-shot generation

In general, graph generation faces three fundamental challenges:

- *Discreteness*
- *Complex Intrinsic Dependencies*
- *Permutation Invariant*



Diffusion Models

Three representative diffusion frameworks:

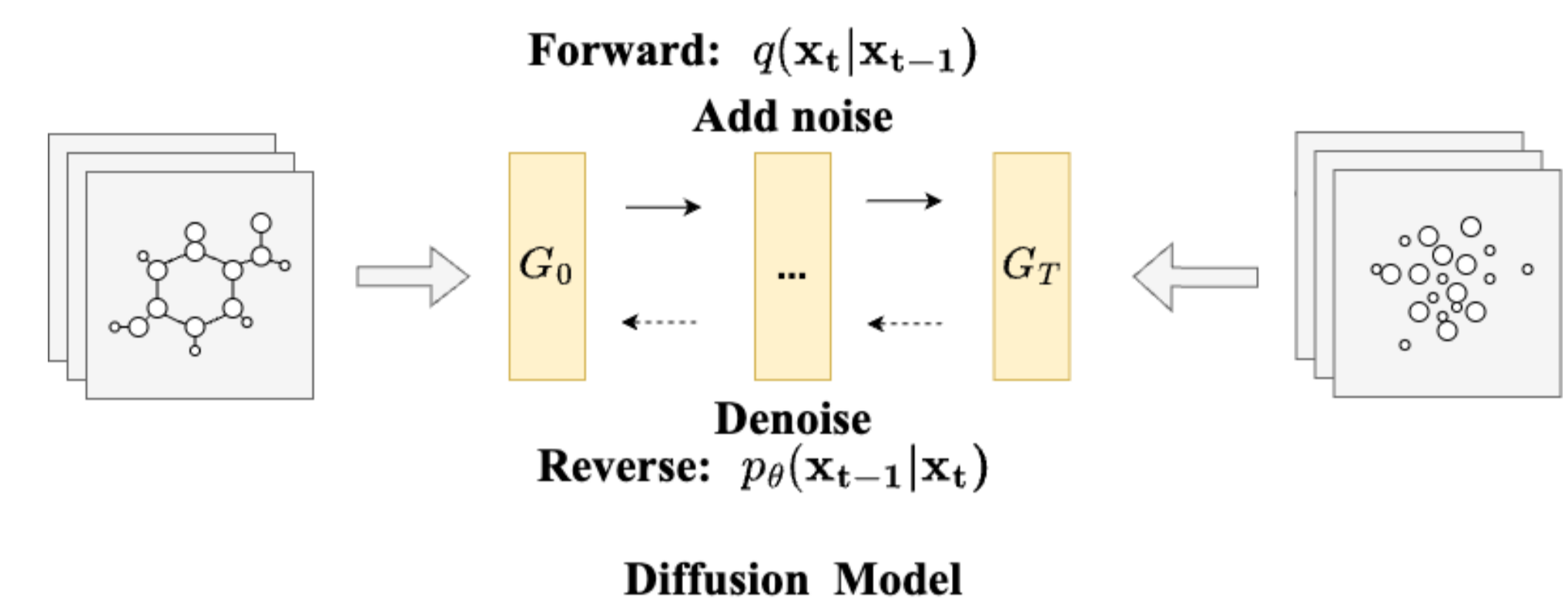
- Score Matching with Langevin Dynamics (SMLD)
- Denoising Diffusion Probabilistic Model (DDPM) on Graphs
- Diffusion Models

Two main stages:

- ◆ Forward diffusion: perturb the original data by adding random noise (generally Gaussian noise)
- ◆ Reverse diffusion: recover the original input data from the random noise.

Advancement:

- ✓ Solid theoretical foundation
- ✓ Easy-to-tractable probabilistic parameters



Generative Diffusion Models on Graphs

Score Matching with Langevin Dynamics (SMLD) on Graphs

- ◆ Forward: a sequence of incremental noise: $q_\sigma(\tilde{x}|x) := \mathcal{N}(\tilde{x}|x, \sigma^2 I)$
- ◆ Reverse: learning the gradient of the data distribution $\nabla_x \log p(x)$

EDP-GNN:

- The very first score-matching diffusion method, which is for undirected graph.

Denoising Diffusion Probabilistic Model (DDPM) on Graphs

Constructs two parameterized Markov chains:

- ◆ Forward: $q(x_t|x_{t-1}) = \mathcal{N}(x_t; \sqrt{1 - \beta_t}x_{t-1}, \beta_t I)$
- ◆ Reverse: $p_\theta(x_t|x_{t-1}) = \mathcal{N}(x_t; \mu_\theta(x_t, t), \Sigma_\theta(x_t, t))$

DiGress

- Simplify the task to a sequence of classification by incorporating the cross entropy.

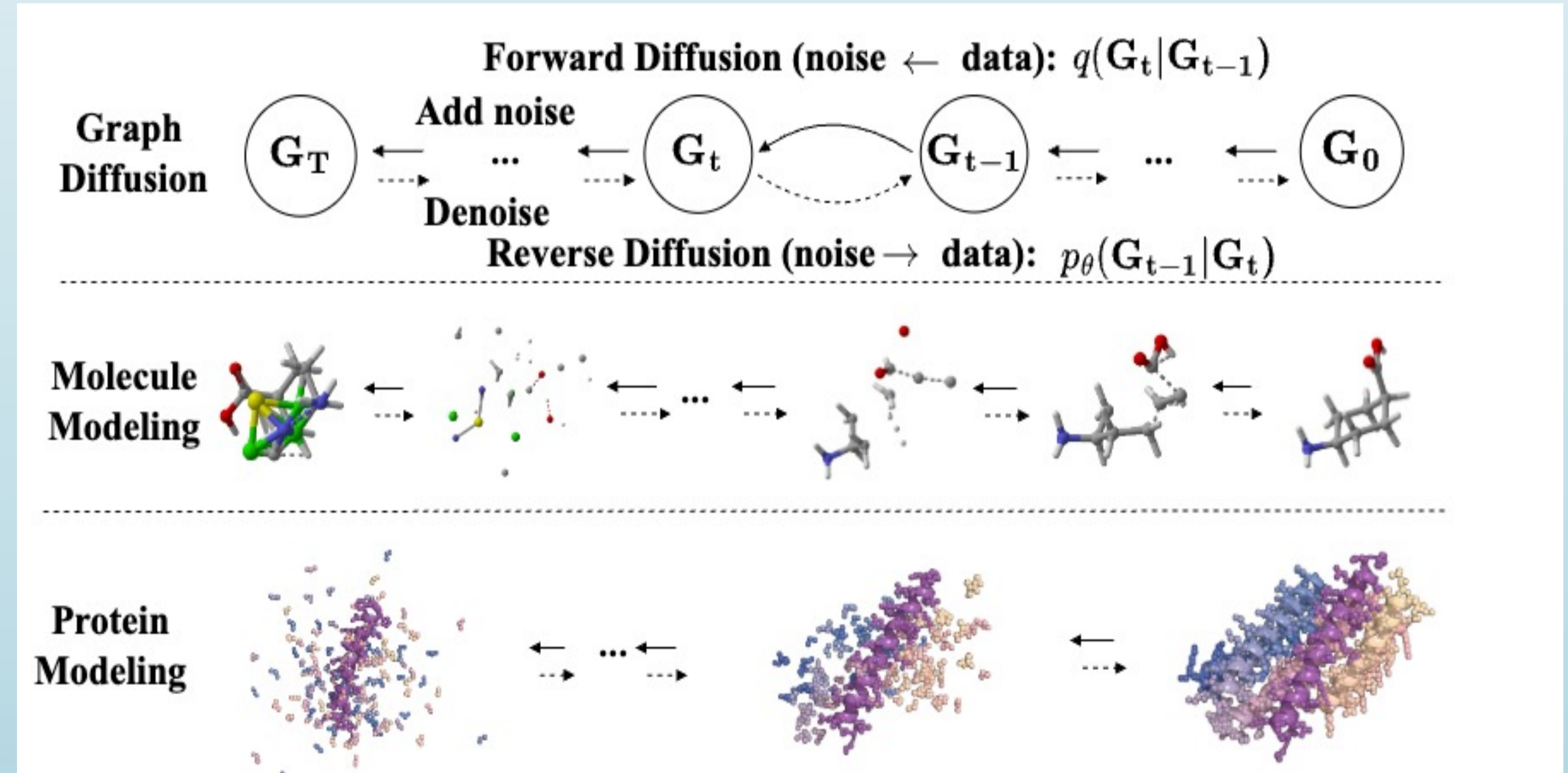
Score-based Generative Models (SGM) on Graphs

The score SDE formula describes the diffusion process in continuous time steps.

- ◆ Forward: $dx = f(x, t)dt + g(t)$
- ◆ Reverse: $dx = [f(x, t) - g(t)^2 \nabla_x \log p_t(x)] dt + g(t)d\bar{w}$

GDSS:

- Model the nodes and edges simultaneously
- The very first diffusion framework that enables the generation of a whole graph



Application

Molecule Modelling

To employ graph learning techniques for the purpose of representing to better perform downstream tasks.

Molecule conformation generation

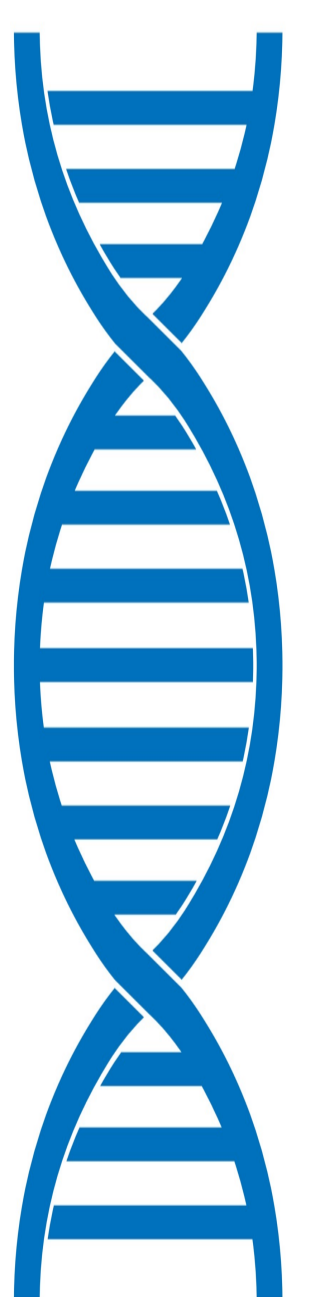
- The biological and physical characteristics of the molecule are significantly influenced by its 3-D structure.

Molecule docking

- A computational method for predicting the preferred orientation of one molecule to a second molecule (typically a protein)
- Drug discovery

Protein Modeling

- Generate and predict the structure of proteins with specific structural and functional properties.
- Predict the protein-ligand complex structure.



Tasks	Applications	Frame	Representative Methods
Molecule Modeling	Molecule Conformation Generation	SMLD	MDM [Huang et al., 2022b], GeoDiff [Xu et al., 2022], EDMs [Hoogeboom et al., 2022], EEGSDE [Bao et al., 2023], DiGress [Vignac et al., 2023]
		DDPM	Torsional Diffusion [Jing et al., 2022], MOOD [Lee et al., 2022], GDSS [Jo et al., 2022], DGSM [Luo et al., 2021a], DiffBridges [Wu et al., 2022b], FragDiff [Peng et al., 2023], DiffLink [Igashov et al., 2022], TargetDiff [Guan et al., 2023], DiffBP [Lin et al., 2022]
	Molecular Docking	DDPM	DiffDock [Corso et al., 2022]
		SGM	SMCDiff [Trippe et al., 2023], SiamDiff [Zhang et al., 2022], DiffFold [Wu et al., 2022a], ProSSDG [Anand and Achim, 2022], DiffAntigen [Luo et al., 2022a], RFdiffusion [Watson et al., 2022], ProteinSGM [Luo et al., 2022a]
Protein Modeling	Protein Generation	DDPM	DiffEE [Nakata et al., 2022]
		SGM	NeuralPLexer [Qiao et al., 2022]
	Protein-ligand Complex Structure Prediction	DDPM	

4 Future Challenges and Opportunities

Conditional Generation for Graph Diffusion Models.

- Incorporating conditions into generative models

Trustworthiness for Graph Diffusion Models.

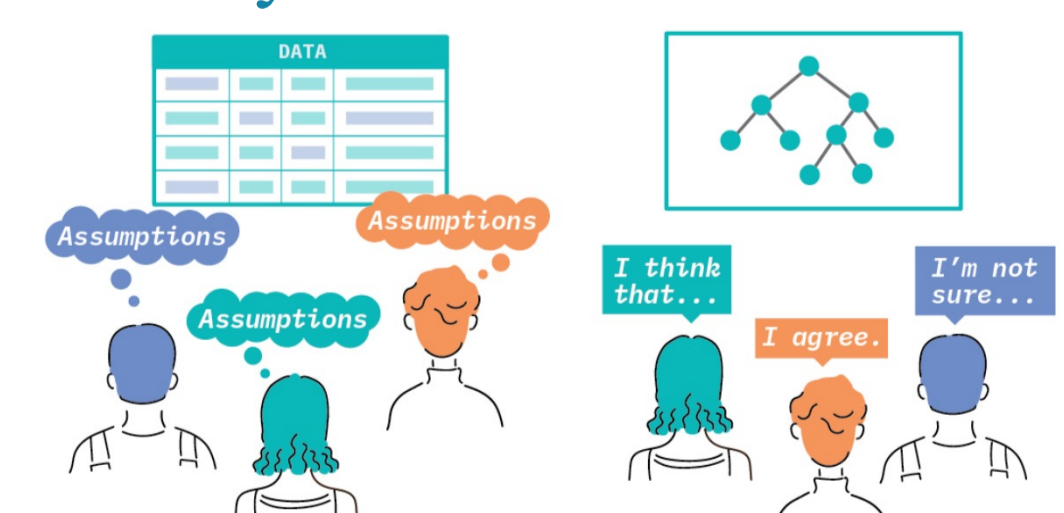
- Unintentional harm to users and society in various real-world tasks
- Safety-critical fields such as drug discovery.

Evaluation Metrics.

- Metrics based on graph statistics and properties are not fully trustworthy.
- Validity and diversity for graph generation

Graph Diffusion Applications.

- *Recommender Systems*
- *Graph Anomaly Detection*
- *Causal Graph Generation*



For arXiv version and reference

