

GUIDING DIFFUSION WITH LOGICAL CONSTRAINTS: MOLECULAR GRAPH GENERATION UNDER LIPINSKI'S RULES

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CONTEXT & MOTIVATION

De novo drug discovery

Generative diffusion models

Generating new drug-like molecules.

SLOW and **COSTLY!**

- Strong theoretical foundations,
 - Efficiency,
 - SOTA on molecular datasets,
 - Conditional variants.

RESEARCH GAP



CURRENT GUIDANCE CAPABILITIES

Individual properties,

Conjunctions of properties.

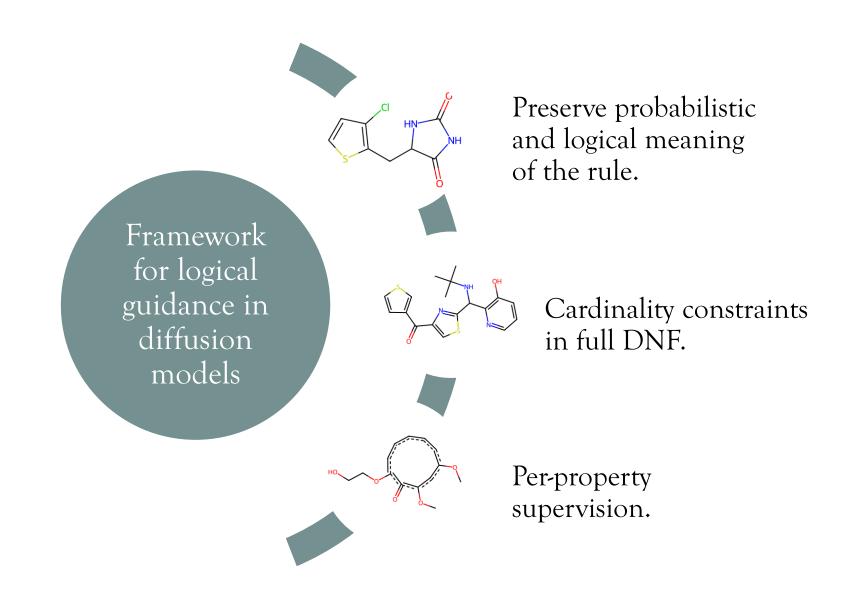


OPEN PROBLEMS

Complex constraints involving multiple properties are treated as black-boxes;

Guided diffusion models have not been systematically used to generate molecules at scale.

THESIS CONTRIBUTIONS



THESIS METHODS

Diffusion Model

• DiGress [Vignac, 2023].

Dataset

• GuacaMol [Brown, 2019].

Constraint

• Lipinski's Rule of Five [Lipinski, 2000].

Evaluation Metrics

- Rule compliance,
- Validity,
- Distributional similarity to the training set,
- Diversity (within the generated set and relative to the training set).

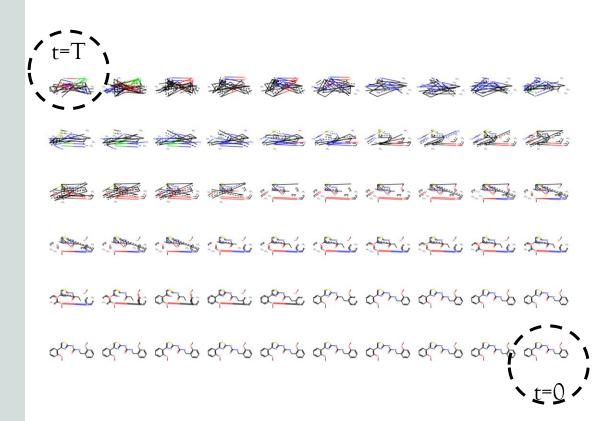
Baselines

- Simple conjunction (all K properties),
- Black-box guidance.

DIGRESS: DIFFUSION FOR GRAPHS

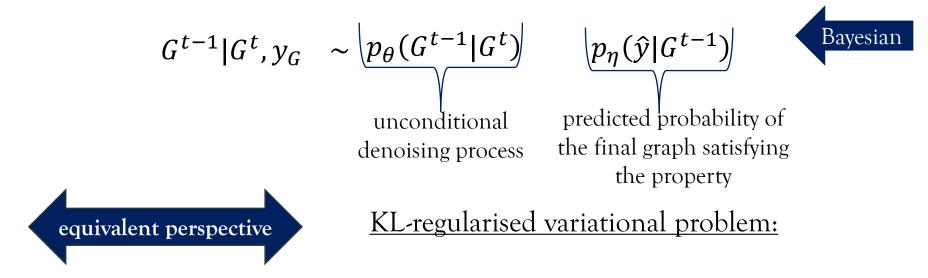
Main idea of every diffusion model:

- 1. Forward process: gradually corrupt data with noise;
- 2. Reverse process: train a neural network (θ) to denoise step by step;
- 3. Sampling: generate a clean realistic sample from pure noise $p_{\theta}(G^{t-1}|G^t)$.



CLASSIFIER GUIDANCE WITH DIGRESS

Conditional denoising process:



- Maximise the similarity with the unconditional denoising distribution,
- Maximise the utility function i.e., linear approximation of the log-probability.

$$p_{\eta}(\hat{y}|G^{t-1}) \propto \exp\left(\lambda \left(\nabla_{G^{t}} \log \dot{q}_{\eta}(y_{G}|G^{t}), G^{t-1}\right)\right)$$
guidance strength
knob
utility

LOGICAL GUIDANCE

Consider a logical formula defined on a set of Boolean variables $\mathcal{X} = \{X_1, ..., X_K\}$. Treat each X_i as a Bernoulli random variable.

The probability of satisfying the rule can then be modelled as a Bernoulli random variable, obtained by aggregating the Bernoulli's of the individual X_i 's.

If:

- the satisfying assignments of the formula are known (i.e., its full DNF is known), and
 - the relationship between the properties is known, then the satisfaction probability can be computed.



CARDINALITY CONSTRAINTS

Consider <u>cardinality constraints</u> of the form $\sum_{i=1}^{K} X_i \ge r$.

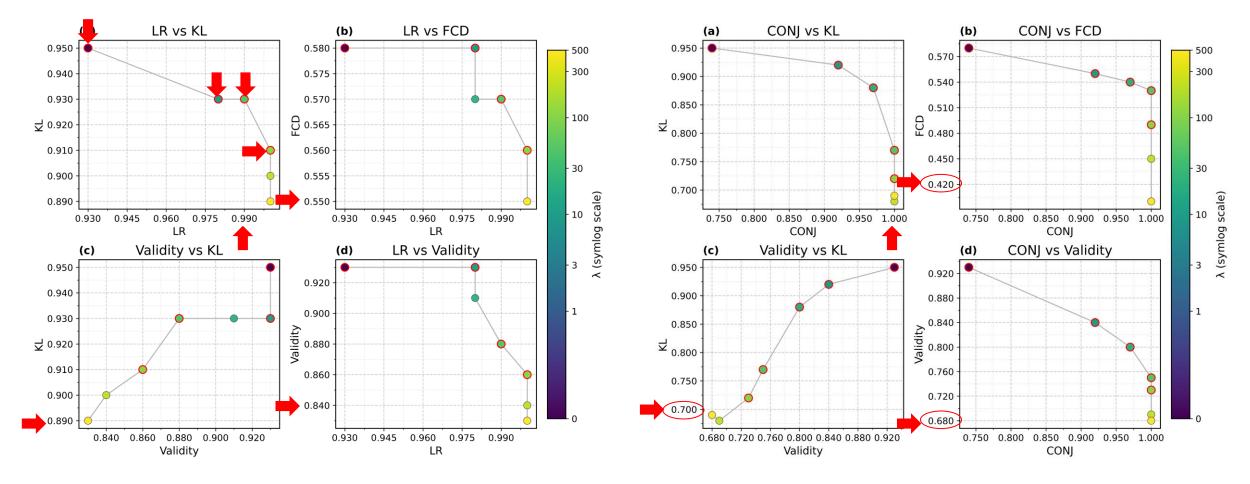
Assume that the variables (properties) are conditionally independent given G^t , and that their probabilities can differ.

$$q(y_G|G^t) = Bernoulli(\pi(G^t)_r)$$

$$\pi(G^t)_r = \sum_{\substack{S \subseteq [K] \\ |S| \ge r}} \left(\prod_{i \in S} p_i \right) \left(\prod_{j \notin S} (1 - p_j) \right)$$

Lipinski's rule: at least 3 of $\log P \le 5$, $molWt \le 500 (Da)$, $HBD \le 5$, $HBA \le 10$.

LIPINSKI'S VS. CONJUNCTIVE RULES

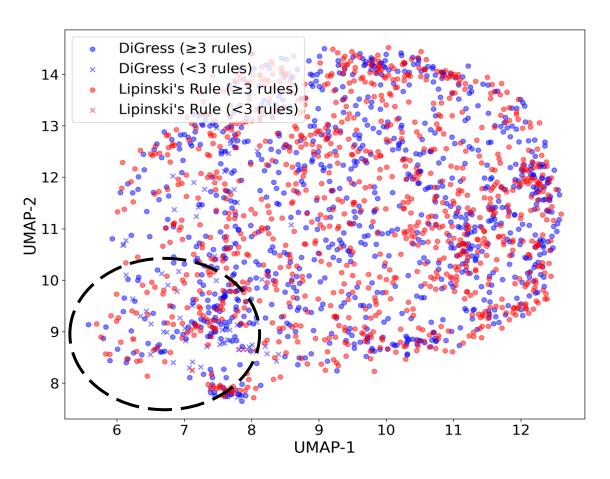


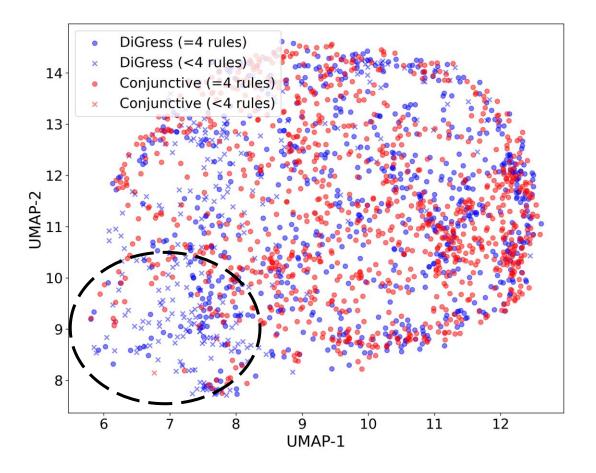
(a) Lipinski's rule

(b) Conjunctive rule

4 independent classifiers, one for each property

LIPINSKI'S VS. CONJUNCTIVE RULES



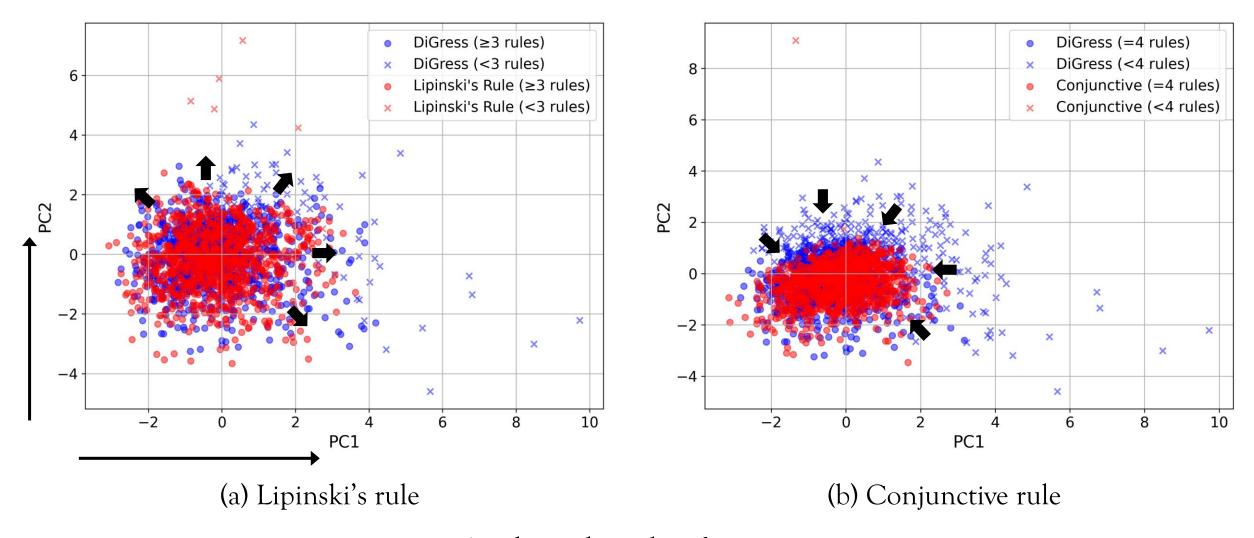


(a) Lipinski's rule

(b) Conjunctive rule

4 independent classifiers, one for each property; $\lambda = 47.82$

LIPINSKI'S VS. CONJUNCTIVE RULES



4 independent classifiers, one for each property; $\lambda = 47.82$

THEORETICAL PROPERTIES

Assume that

- i. the predictors are not negatively correlated under q_{λ} , i.e. $Cov_{q_{\lambda}}(p_j, p_k) \ge 0$ for all j, k;
 - ii. the rule probability is non-decreasing in each p_i (true for cardinality rules).

Then, for any fixed reverse step t and any $\lambda \ge 0$ we have

$$\frac{d}{d\lambda}\mathbb{E}_{q_{\lambda}}[p_k] \geq 0.$$

Furthermore, let $S = \sum_{i=1}^K X_i$ with mean $\mu_S(p) = \sum_{i=1}^K p_i$. It follows that $\frac{d}{d\lambda} \mathbb{E}_{q_\lambda}[\mu_S] \ge 0.$

THEORETICAL PROPERTIES

Define $p_r(p) := \mathbb{P}(S = r | p)$ and assume that $\sigma_{\mu_S}^2 = \mathbb{V}_{q_{\lambda}}[\mu_S] < +\infty$.

Using the assumptions in the previous slide it follows that, for every r and c > 0,

- $\frac{d}{d\lambda}\mathbb{E}_{q_{\lambda}}[p_r] \geq 0$ whenever $\mathbb{E}_{q_{\lambda}}[\mu_S] \leq r-1-c\sigma_{\mu_S}$, and
 - $\frac{d}{d\lambda}\mathbb{E}_{q_{\lambda}}[p_r] \leq 0$ whenever $\mathbb{E}_{q_{\lambda}}[\mu_S] \geq r + 1 + c\sigma_{\mu_S}$

up to a tail probability of order $^{1}/_{c^{2}}$ by Chebyshev's inequality.

Let $p_K(p) = \prod_{j=1}^K p_j$. Using the assumptions in the previous slide it follows that $\frac{d}{d\lambda} \mathbb{E}_{q_\lambda}[p_K] \ge 0$.

COMPOSITION OF THE PROPERTY DISTRIBUTIONS

• Increase in 4-of-4 proportion.

- 3-of-4 proportion not guaranteed to vanish;
- 3-of-4 proportion decreases as $\mu_S \ge 3$.



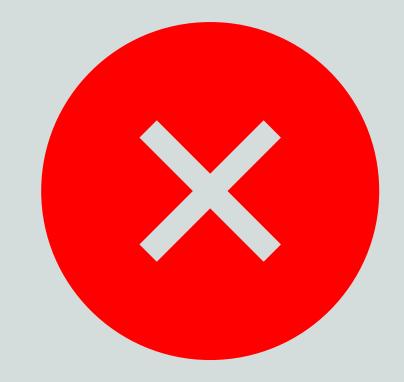
- Independent-2L: [77.9, 82.9]
- Shared-1L: [79.2, 90.0]
- Shared-2L: [77.7, 88.0]

- Independent-2L [17.1%, 19.9%]
- Shared-1L: [9.9%, 20%]
- Shared-2L: [11.9%, 21%]

Better preserved when $\lambda \in [10, 104]$.

LOGICAL GUIDANCE VS. BLACK-BOX GUIDANCE

- For Lipinski's rule, it achieves the <u>same</u> compliance as our logical method.
- The proportion of molecules that satisfies exactly 3 properties increases with λ: [19.4%, 26.1%].
- The proportion of molecules that satisfies exactly 4 properties decreases with λ: [73.9%, 79.6%].



Shortcut learning!

Learns to satisfy molWt, HBD and HBA, which are the most frequent in the training samples that satisfy the rule.

This neglects part of the satisfying set!

CONCLUSIONS & FUTURE WORK

Key Takeaways

• First framework for embedding logical rules (full DNF cardinality constraints) into graph diffusion models for conditional generation at scale.

Future Directions

- Test and build scalability for larger rules;
- Investigate and mitigate constraint overfitting;
- Relax the conditional independence assumption to reduce bias;
- Extend beyond DiGress (e.g. score-based SDEs).
- Experimental:
 - latent variable control of k-of-K satisfaction patterns.

<u>Vision</u>

• Probabilistic-symbolic integration in graph generative modelling beyond molecular design.

THANK YOU!

QUESTIONS?

Clement Vignac et al. "DiGress: Discrete Denoising Diffusion for Graph Generation". In: The 11th International Conference on Learning Representations. 2023.

Nathan Brown et al. "GuacaMol: Benchmarking Models for de novo Molecular Design". In: *Journal of Chemical Information and Modelling* 59.3 (2019). DOI: 10.1021/acs.jcim.8b00839.

Christopher A. Lipinski. "Drug-like Properties and the Causes of Poor Solubility and Poor Permeability". In: *Journal of Pharmacological and Toxicological Methods* 44.1 (2000). DOI: 10.1016/S1056-8719(00)00107-6.

