

INTRODUCTION

In this work, we propose a regularization framework for VAEs to generate **semantically valid** graphs. Contributions:

1. A new deep generative framework for general graph generation with validity constraints.
2. Formulation of constraints for **I. molecular graphs** and **II. node-compatible graphs**.
3. An efficient inference algorithm for generating valid graphs.

VISUALIZATION

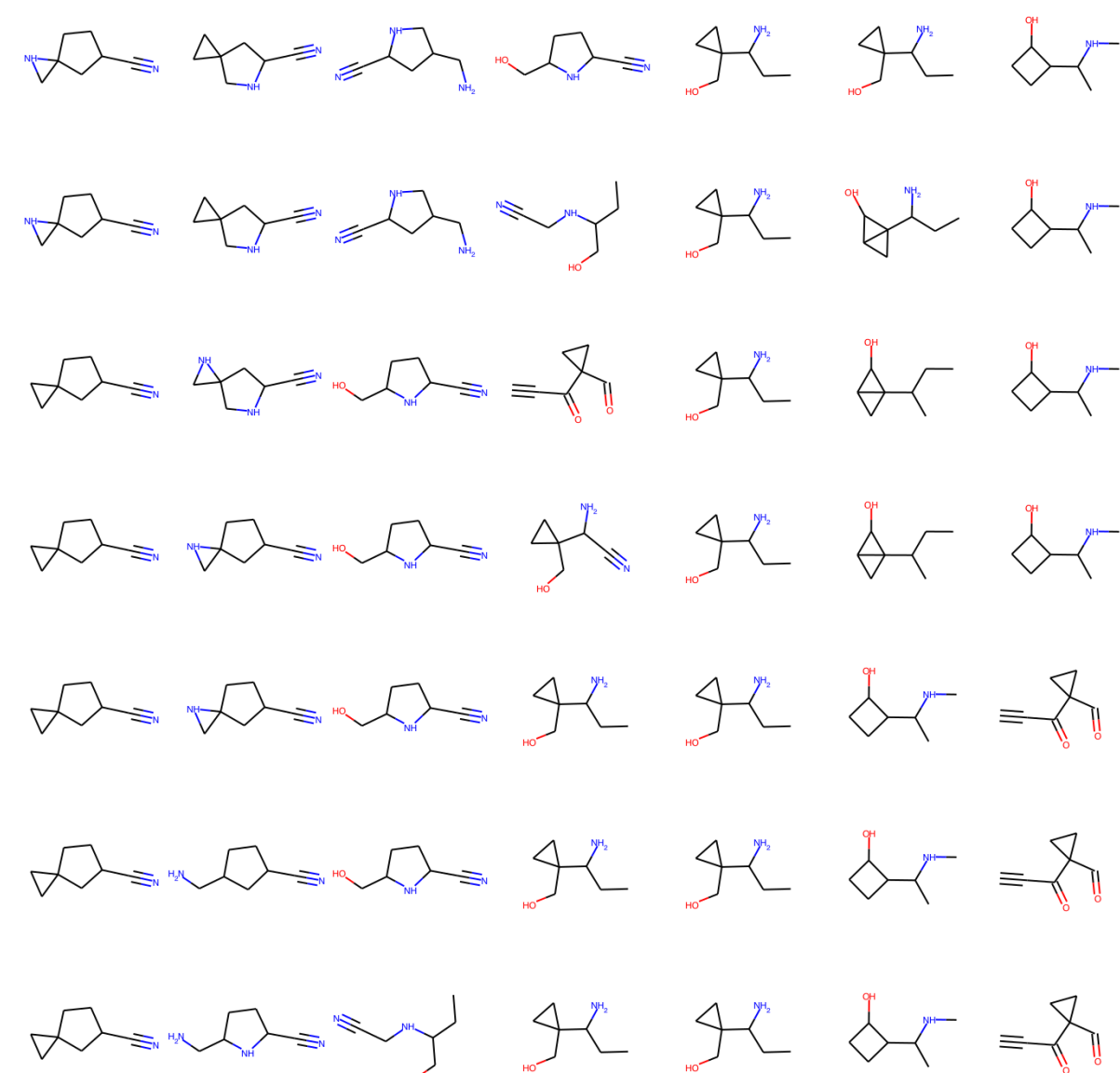


Figure 3: 2D interpolation of latent space.

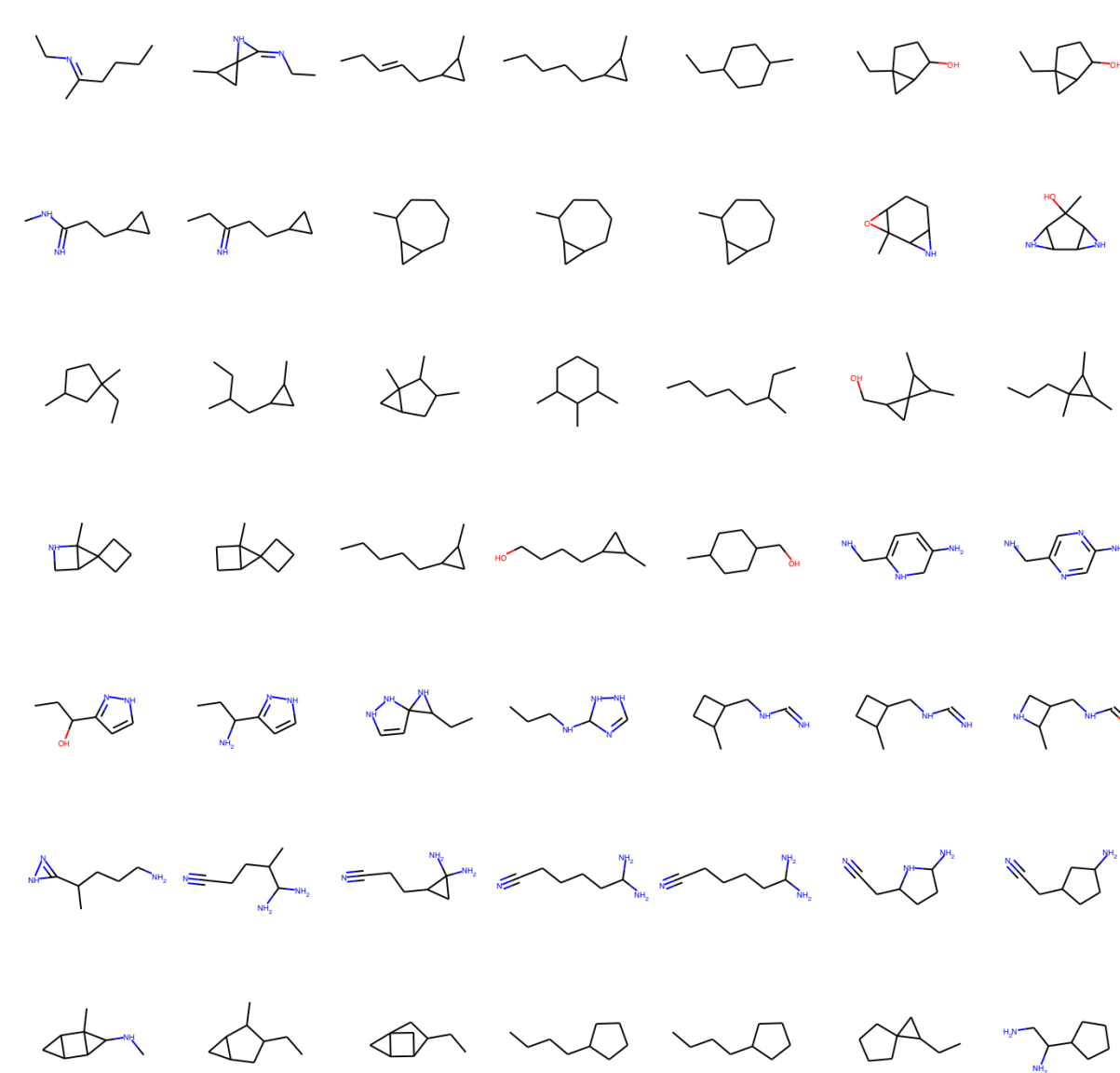


Figure 4: 1D interpolation of latent space (each row).

NETWORK ARCHITECTURE

For input representation, we unfold the edge-label tensor $E \in \mathbb{R}^{N \times N \times (1+t)}$ and concatenate it with the node-label matrix $F \in \mathbb{R}^{N \times (1+d)}$ to form a wide matrix. The encoder is a 4-layer convolutional neural net. The decoder is a 4-layer deconvolutional neural net similar to DCGAN.

EXAMPLES

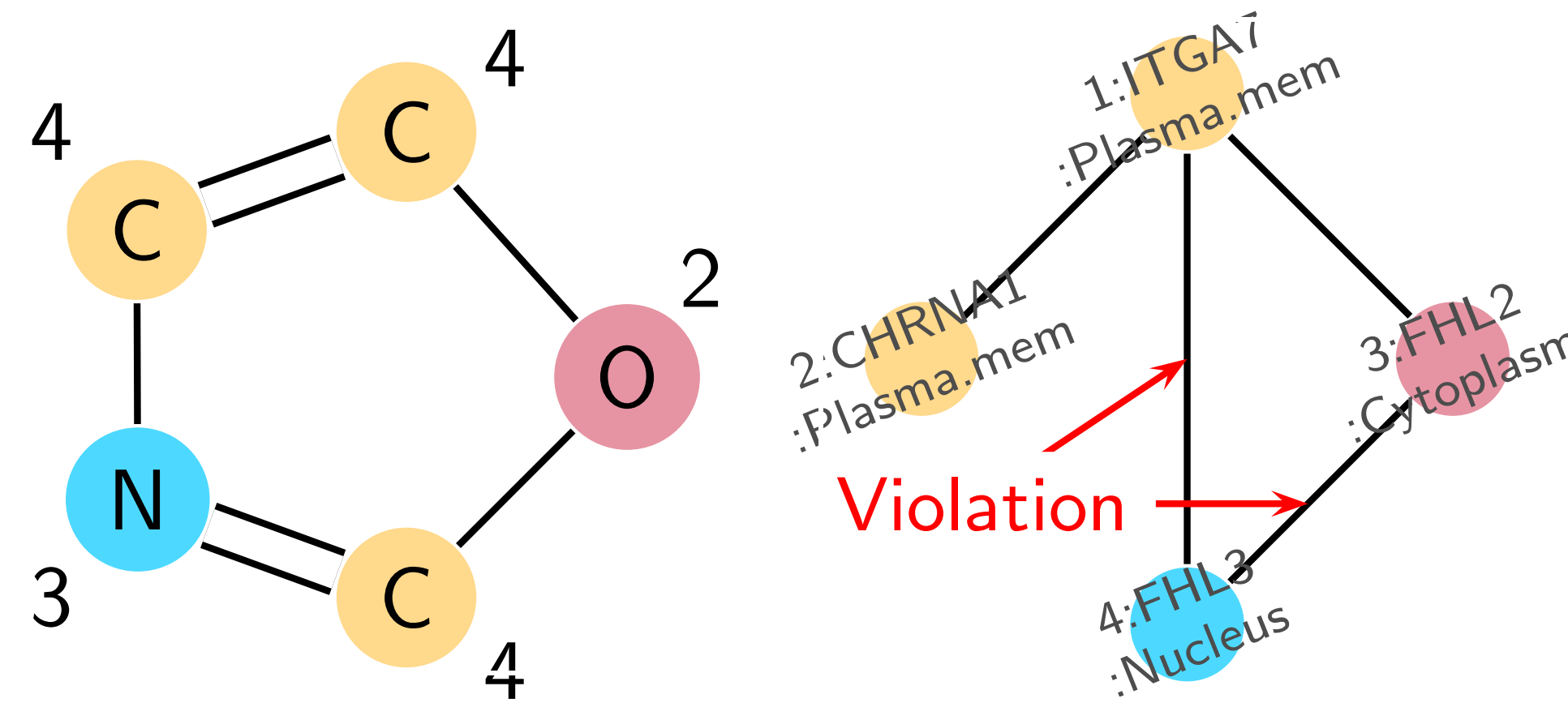


Figure 1: Left: Molecular graph. Right: Protein network.

RESULTS

Table 1: Standard VAE versus regularized VAE.

QM9		
Method	% Valid	ELBO
Standard	83.2	-17.3
Regul.	96.6	-18.5
Node-compatible		
Method	% Valid	ELBO
Standard	40.2	-42.5
Regul.	98.4	-51.2

Table 2: Comparison with other VAEs.

QM9			
Method	% Valid	% Novel	% Recon.
Proposed	96.6	97.5	61.8
GVAE	60.2	80.9	96.0
CVAE	10.3	90.0	3.61
ZINC			
Method	% Valid	% Novel	% Recon.
Proposed	34.9	100	54.7
GVAE	7.2	100	53.7
CVAE	0.7	100	44.6

Table 3: Denoising node-incompatible graphs.

Standard VAE	Regularized VAE
11.2	93.8

FRAMEWORK

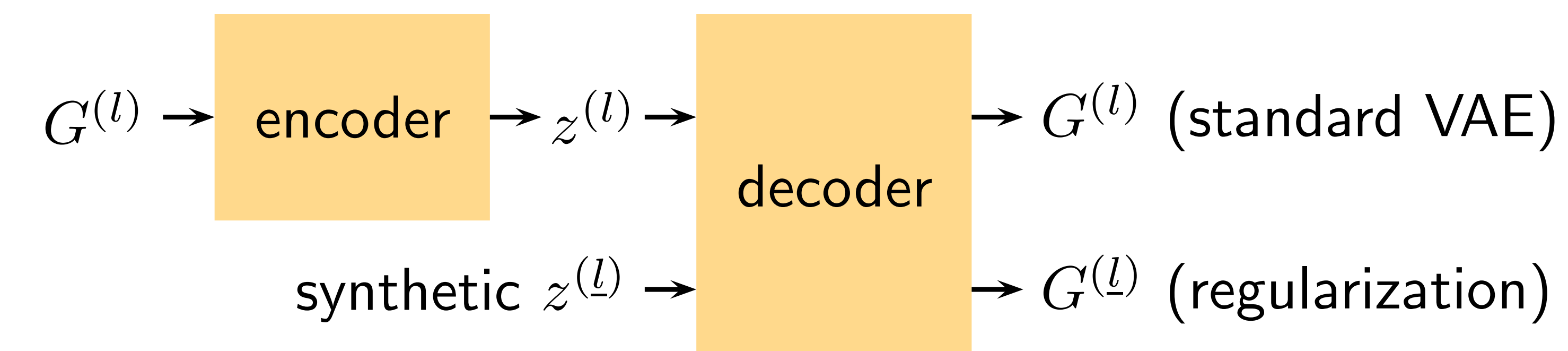


Figure 2: Overview of the Regularized VAE

The central contribution of this work is an approach to imposing validity constraints in the training of VAEs. Besides the usual loss function $f(x)$ of VAE, we want the graph samples produced by the generative network to be valid, regardless of what latent value z one starts with:

$$\begin{aligned} \min_x f(x) \\ \text{subject to} \quad & \text{for almost all } z \sim p_x(z), \\ & h_1(x, z) = 0, \dots, h_m(x, z) = 0, \\ & g_1(x, z) \leq 0, \dots, g_r(x, z) \leq 0. \end{aligned} \quad (1)$$

Lagrangian: To solve (1), we generalize the usual notion of Lagrangian function to

$$\mathcal{L}(x, \lambda, \mu) = f(x) + \sum_{i=1}^m \lambda_i \tilde{h}_i(x) + \sum_{j=1}^r \mu_j \tilde{g}_j(x), \quad (2)$$

where $\tilde{g}_j(x) = [\int g_j(x, z)^2 p_x(z) dz]^{\frac{1}{2}}$ and similarly for $\tilde{h}_i(x)$.

Training: Let us write the i -th validity constraint as $g_i(\theta, z) \leq 0$ for all z , then the *loss function* is

$$-L_{\text{ELBO}}(\theta, \phi) + \mu \sum_i \left[\int g_i(\theta, z)_+^2 p_\theta(z) dz \right]^{\frac{1}{2}}, \quad (3)$$

where $g_+ = \max(g, 0)$ is the ramp function. This regularization will not penalize the desirable case $g_i \leq 0$. **In practice**, the integral in the regularization may be intractable, and hence we appeal to *Monte Carlo approximation* for evaluating the loss in each parameter update:

$$-L_{\text{ELBO}}(\theta, \phi) + \mu \sum_i g_i(\theta, z)_+, \quad \text{where } z \sim p_\theta(z). \quad (4)$$

REGULARIZATION

Molecules:

- **Ghost Nodes and Valence.** Denote by $V(i)$ the capacity of a node i and by $U(i)$ the valence (upper bound of the capacity). The valence constraint is $g_i = V(i) - U(i)$.
- **Connectivity:** If A is the adjacency matrix, then the (i, j) element of $B = I + A + A^2 + \dots + A^{N-1}$ is nonzero iff i and j are connected by a path. Let $q(i) = 0$ indicate the ghost node, then the connectivity constraint is $q(i)q(j) \cdot \mathbf{1}\{B(i, j) = 0\} + [1 - q(i)q(j)] \cdot \mathbf{1}\{B(i, j) \neq 0\} \leq 0$. To be differentiable, in our framework $g_{ij} = q(i)q(j) \cdot [1 - C(i, j)] + [1 - q(i)q(j)] \cdot C(i, j)$ where $C = \sigma(B)$.

Node-compatible Graphs: Consider the matrix $P = \tilde{F}D\tilde{F}^T$. The (i, j) element of P is the probability that nodes i and j have compatible types. We want node pairs with low compatibility to be disconnected. Hence, the constraint is $g_{ij} = [1 - \tilde{E}(i, j, 0)][1 - P(i, j)] - \alpha$ where $\alpha \in (0, 1)$ is a tunable hyperparameter.

REFERENCES

- [1] Diederik P Kingma and Max Welling. Auto-encoding variational Bayes. In *ICLR*, 2014.
- [2] Matt J Kusner, Brooks Paige, and José Miguel Hernández-Lobato. Grammar variational autoencoder. In *ICML*, 2017.