

A Diffusion Model Framework for Unsupervised Neural Combinatorial Optimization

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Abstract

Learning to sample from intractable distributions over discrete sets without relying on corresponding training data is a central problem in a wide range of fields, including Combinatorial Optimization. Currently, popular deep learning-based approaches rely primarily on generative models that yield exact sample likelihoods. This work introduces a method that lifts this restriction and opens the possibility to employ highly expressive latent variable models like diffusion models. Our approach is conceptually based on a loss that upper bounds the reverse Kullback-Leibler divergence and evades the requirement of exact sample likelihoods. We experimentally validate our approach in data-free Combinatorial Optimization and demonstrate that our method achieves a new state-of-the-art on a wide range of benchmark problems.*

1. Introduction

Sampling from a known but intractable, high-dimensional target distribution like the Boltzmann distribution is of high relevance in many scientific fields like the prediction of molecule configurations (Noé and Wu, 2018), lattice models in physics (Wu et al., 2019) and Monte Carlo integration (Müller et al., 2019). Recently, the works of Hibat-Allah et al. (2021) and Sanokowski et al. (2023) showed that Combinatorial Optimization (CO) can be concisely formulated as such a distribution learning problem where the resulting samples correspond to solutions of a CO problem. In all of these domains, the energy function associated to the distributions of interest is known but obtaining unbiased samples

represents a formidable challenge. Problem areas like CO or lattice models in physics are characterized by discrete target distributions. In these applications, the approximation of the target distribution is predominantly based on products of categorical distributions or autoregressive models. While product distributions are computationally convenient they lack expressivity due to their inability to represent statistical inter-dependencies. Autoregressive models rely on sequentially generating the components of the samples. For the high-dimensional distributions that are frequently encountered in the aforementioned application areas, this procedure becomes prohibitively expensive. In addition, there is typically no canonical ordering among the components of samples. Hence, an autoregressive approach appears as an unnatural approach. Intuitively, an issue with the sequential sample generation procedure of autoregressive models is that there is no possibility of correcting sub-optimal decisions once they are made. More formally, it can be shown that autoregressive models are less expressive than energy-based models or latent variable models from a computational complexity point of view (Lin et al., 2021). Approximate likelihood models that utilize latent variables, like Variational Autoencoders (VAE) (Kingma and Welling, 2014) and diffusion models (Sohl-Dickstein et al., 2015; Ho et al., 2020; Song et al., 2021), are trained using data samples and a loss that is based on the Evidence Lower Bound (ELBO). These models have the advantage that they are more expressive and applicable in the discrete setting. However, with these models, it is in general not feasible to evaluate exact sample likelihoods. This problem precludes their application in problems of data-free approximation of target distributions which typically rely on exact sample likelihoods. Recent works that proposed methods to apply diffusion models in data-free approximation of target distributions are limited to the continuous setting (Berner et al., 2022; Richter et al., 2023; Vargas et al., 2023; 2024). To the best of our knowledge, the discrete setting remains unexplored.

In this work, we propose in Sec. 3 an approach that allows for the application of latent variable models like diffusion models in the problem of data-free approximation of discrete distributions. We demonstrate our method on paradigmatic problems in this area from the field of CO and obtain state-

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*Code is available at <https://github.com/ml-jku/DiffUCO>.

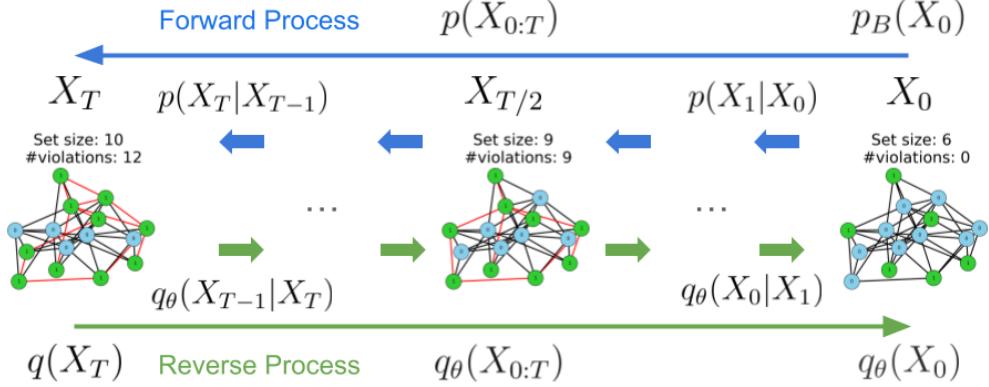


Figure 1. Illustration of DiffUCO’s solution generation process on the example of finding the Maximum Independent Set (see Sec. 6.1) of a graph. Solutions are generated in the reverse diffusion process that is modeled by iteratively sampling the time-conditioned model q_θ .

of-the-art performance over a wide range of benchmarks. To achieve this our method *Diffusion for Unsupervised Combinatorial Optimization* (DiffUCO) uses an upper bound on the reverse Kullback-Leibler divergence as a loss function. We show that the performance of the model consistently improves as we increase the number of diffusion steps used during training. Additionally, we find that during inference the solution quality can be further improved by applying more diffusion steps than during training (see Sec. 6.2). In Sec. 4.1 we propose a significantly more efficient version of a frequently used sampling strategy called *Conditional Expectation*. We show that combined with our diffusion model this method allows for time-efficient generation of high-quality solutions to CO problems. The proposed framework yields a highly efficient and general way of employing latent variable models like diffusion models in the ubiquitous challenge of data-free approximation of discrete distributions.

2. Problem Description

Following Lucas (2014) we will represent CO problems by a corresponding energy function $H : \{0, 1\}^N \rightarrow \mathbb{R}$ which assigns a scalar value called energy to a given solution $X \in \{0, 1\}^N$. The dimensionality of X is denoted as N and is called the problem size. The Boltzmann distribution associated to H is defined by:

$$p_B(X, \beta) = \frac{\exp(-\beta H(X))}{Z}, \quad Z = \sum_X \exp(-\beta H(X)),$$

where the parameter \mathcal{T} is typically referred to as the temperature and $\beta = 1/\mathcal{T}$ as the inverse temperature. In the context of CO the relevant feature of $p_B(X, \beta)$ is that as $\beta \rightarrow \infty$ it places all the probability mass on the solutions with the lowest energy, i.e. on the *optimal* solutions of H . It can be shown that the sample complexity of learning $p_B(X, \beta)$ is in $O(\beta^2)$ (Sanokowski et al., 2023). Thus, the more probability mass the Boltzmann distribution puts on optimal solutions the harder it becomes to learn. For large N

this distribution is in general intractable, i.e. sampling from this distribution is difficult for a wide range of frequently encountered energy functions in statistical physics and CO. Approximate sampling from such an intractable distribution can be realized with Markov Chain Monte Carlo methods but in complex, multi-modal distributions these methods often exhibit insufficient mixing (Nicoli et al., 2020). There is a growing interest in training neural networks $q_\theta(X)$ to approximate such a distribution. In these approaches, deep generative models are used to efficiently obtain unbiased samples from Boltzmann distributions (see Sec. 5). We call the problem of data-free approximation of a target distribution with a deep generative model *Neural Probabilistic Optimization* (NPO). A common approach to NPO is to minimize the reverse Kullback–Leibler divergence (KL) $D_{KL}(q_\theta(X) \parallel p_B(X, \beta))$ with respect to the model parameters θ . Multiplying this objective with \mathcal{T} yields an expression that is proportional to the Variational Free Energy $F_\theta(X, \mathcal{T})$, which is given by:

$$L(\theta) = \mathbb{E}_{X \sim q_\theta(X)} [H(X) + \mathcal{T} \log q_\theta(X)] \quad (1)$$

$$= \mathbb{E}_{X \sim q_\theta(X)} [\widehat{F}_\theta(X, \mathcal{T})] := F_\theta(X, \mathcal{T}), \quad (2)$$

where $\widehat{F}_\theta(X, \mathcal{T}) := H(X) + \mathcal{T} \log q_\theta(X)$ (Wu et al., 2019). Minimizing this objective requires exact evaluation of $\log q_\theta(X)$ not only to calculate the Variational Free Energy objective but also to backpropagate through the expectation of this objective. This can for example be done with REINFORCE (Williams, 1992) gradient estimator:

$$\nabla_\theta L(\theta) = \mathbb{E}_{X \sim q_\theta(X)} [\widehat{F}_\theta(X, \mathcal{T}) \nabla_\theta \log q_\theta(X)].$$

To minimize this objective it is necessary to choose a generative model for which the sample likelihood $q_\theta(X)$ can be efficiently evaluated. Thus, optimizing this objective with a latent variable model like VAEs or diffusion models is not possible since these models do not allow an exact evaluation

of $q_\theta(X)$. In Sec. 3, we propose an objective based on an upper bound of the reverse KL divergence that mitigates this issue. In principle, also other divergences could be used to approximate a given target distribution. For example, both the reverse and the forward KL divergence are special cases of Renyi divergences (van Erven and Harremoës, 2014) which all require exact evaluation of $\log q_\theta(X)$. By introducing an upper bound on the divergence as we do in Sec. 3, the evaluation of $\log q_\theta(X)$ is avoided. Which divergence is the most suitable depends on the application as Renyi divergences are either mass covering or mode seeking (Minka et al., 2005). We will focus on the reverse KL divergence and note that our framework can be applied to any other Renyi divergence.

2.1. Unsupervised Neural Combinatorial Optimization

In Combinatorial Optimization (CO) the task is to find the solution $X \in \{0, 1\}^N$ that minimizes an objective $O : \{0, 1\}^N \rightarrow \mathbb{R}$ under the constraint that X is in the set of feasible solutions f . In Lucas (2014) it is shown that many of these CO problems can be described by identifying the objective O with an energy function H_Q in the form of:

$$H_Q(X) = \sum_{i,j} Q_{ij} X_i X_j, \quad (3)$$

Here, Q_{ij} is defined by the problem at hand, and for each CO problem, it is easy to choose $Q \in \mathbb{R}^{N \times N}$ in a way so that global minima correspond to the optimal solutions of the problem (Lucas, 2014). In the context of optimization and quantum computing, this kind of energy function is well known as the Quadratic Unconstrained Binary Optimization (QUBO) formulation of CO problems (Glover et al., 2022). In these terms, the goal of UCO is to train a generative model to sample low energy configurations X when conditioned on a CO problem instance Q which is sampled from a distribution $\omega(Q)$. The goal in UCO is to train a model without the use of any example solutions. The corresponding optimization task is to find parameters θ that minimize the following loss:

$$L(\theta) = \mathbb{E}_{Q \sim \omega(Q)} [\mathbb{E}_{X \sim q_\theta(X|Q)} [H_Q(X)]] . \quad (4)$$

The square matrix Q can be regarded as the adjacency matrix of a graph with weighted edges, which is why the dependence on Q in $q_\theta(X|Q)$ is often parameterized by a Graph Neural Network (GNN) (Cappart et al., 2021). Instead of minimizing Eq. 3 directly, numerous approaches (Hibat-Allah et al., 2021; Sun et al., 2022; Sanokowski et al., 2022; 2023) instead frame this as an NPO problem by minimizing Eq. 2. Here, ideas from Simulated Annealing (Kirkpatrick et al., 1983) are borrowed, i.e. the objective is first optimized at a high temperature and then the temperature is gradually reduced according to a predefined temperature schedule. In the limit of $\mathcal{T} \rightarrow 0$, Eq. 2 reduces to Eq. 4 (up

to the expectation over $\omega(Q)$) which indicates that in this limit the energy of the sampled solutions is minimized without any additional entropy regularization. Consequently, at the end of this annealing procedure, the average energy of the model’s samples can be considered as a measure of the residual bias of the learned approximation of the Boltzmann distribution. Sanokowski et al. (2023) motivate this technique theoretically in the form of a curriculum learning approach and it has been shown empirically (Hibat-Allah et al., 2021; Sun et al., 2022; Sanokowski et al., 2023) that this method leads to models with solutions of higher quality.

3. Neural Probabilistic Optimization Objective for Approximate Likelihood Models

Training an approximate likelihood model to learn to generate samples from a target distribution* $p_B(X)$ in the data-free setting is a challenging problem because the evaluation of the sample likelihood within the reverse KL divergence is intractable (see Sec. 5). In the following, we will derive an objective that can be used with approximate likelihood models like VAEs or diffusion models. For that consider the task of approximating a target distribution $p_B(X)$ by using the marginal sample probability of a latent variable model $q_\theta(X) = \int q_\theta(X|Z) q(Z) dZ$, where $q(Z)$ is an easy to sample prior distribution over latent variables Z . Due to the intractability of this marginalization it is not possible to directly minimize $D_{KL}(q_\theta(X) || p_B(X))$. Instead we use a simple tractable upper bound (see App. A.1 for a derivation) which is given by:

$$D_{KL}(q_\theta(X) || p_B(X)) \leq D_{KL}(q_\theta(X, Z) || p(X, Z)), \quad (5)$$

where $q_\theta(X, Z) = q_\theta(X|Z) q(Z)$ and $p(X, Z) = p(Z|X) p_B(X)$. $p(Z|X)$ can either be learned by parameterizing it with a neural network like it is done in a VAE or it can be simply defined in a way so that $p(Z|X)$ maps samples $X \sim p_B(X)$ onto a pre-defined stationary distribution $q(Z)$ as it is done in diffusion models. The advantage of this upper bound is that it is tractable as $q_\theta(X, Z)$ and $p(X, Z)$ can be exactly evaluated and samples $X, Z \sim q_\theta(X, Z)$ are easy to obtain. Therefore, as we have shown that the reverse KL of approximate likelihood models $D_{KL}(q_\theta(X) || p_B(X))$ is bounded from above by the reverse joint KL $D_{KL}(q_\theta(X, Z) || p(X, Z))$, we can decrease the intractable reverse KL divergence by decreasing the tractable reverse joint-KL divergence instead. According to Gibbs’ inequality the KL divergence between two distributions is non-negative and equals zero iff the distributions are identical. Thus, if the right-hand side of the inequality is zero, the inequality becomes an equality. In this case, the target distribution is exactly approximated.

*In contexts, where it is not relevant we suppress the dependence of the Boltzmann distribution on β and denote it as $p_B(X)$.

Otherwise, the right-hand side of the inequality bounds the approximation error given by the KL divergence on the left-hand side. In the following we call this objective $D_{KL}(q_\theta(X, Z) \parallel p(X, Z))$ for NPO the *Joint Variational Upper Bound*. Equation 5 is a special case of the data processing inequality for Renyi divergences (van Erven and Harremoës, 2014) and can therefore be applied analogously to other Renyi divergences.

3.1. Training Diffusion Models with the Joint Variational Upper Bound

To minimize the right-hand side of Eq. 5, we train a diffusion model. As illustrated in Fig. 1 the forward diffusion process transforms the target distribution $p_B(X_0)$ into a *stationary distribution* $q(X_T)$ through iterative sampling of a *noise distribution* $p(X_t|X_{t-1})$ for a total of T iterations. The diffusion model is supposed to model the reverse process, i.e. to map samples $X_T \sim q(X_T)$ to $X_0 \sim p_B(X_0)$ by iteratively sampling $q_\theta(X_{t-1}|X_t)$. The joint distribution of the reverse process is given by $q_\theta(X_{0:T}) = q(X_T) \prod_{t=1}^T q_\theta(X_{t-1}|X_t)$ and we can easily obtain samples $X_{0:T} \sim q_\theta(X_{0:T})$. The joint distribution of the forward process $p(X_{0:T}) = p_B(X_0) \prod_{t=1}^T p(X_t|X_{t-1})$ is analytically known, but in the data-free setting samples $X_{0:T} \sim p(X_{0:T})$ are not available. We apply the upper bound in Eq. 5 by substituting Z with $X_{1:T}$ and get the inequality:

$$D_{KL}(q_\theta(X_0) \parallel p_B(X_0)) \leq D_{KL}(q_\theta(X_{0:T}) \parallel p(X_{0:T})).$$

We use this Joint Variational Upper Bound as a loss function and minimize it with respect to θ using RAdam (Liu et al., 2020). The resulting expression can be simplified (see App. A.2) to:

$$\begin{aligned} \mathcal{T} D_{KL}(q_\theta(X_{0:T}) \parallel p(X_{0:T})) &= \\ &- \mathcal{T} \cdot \sum_{t=1}^T \mathbb{E}_{X_{T:t} \sim q_\theta(X_{T:t})} [S(q_\theta(X_{t-1}|X_t))] \\ &- \mathcal{T} \cdot \sum_{t=1}^T \mathbb{E}_{X_{T:t-1} \sim q_\theta(X_{T:t-1})} [\log p(X_t|X_{t-1})] \\ &+ \mathbb{E}_{X_{T:0} \sim q_\theta(X_{T:0})} [H_Q(X_0)] + C, \end{aligned} \quad (6)$$

where $S(q_\theta(X_{t-1}|X_t))$ is the Shannon entropy of $q_\theta(X_{t-1}|X_t)$. The logarithm of the normalization constant Z of the Boltzmann distribution is absorbed into the constant C as it depends neither on X nor on θ .

The first term on the right-hand side of Eq. 6 represents an entropy regularization that encourages our model to explore the solution space in the initial phase of training, i.e. when annealing starts with high values of \mathcal{T} . The second term represents a coupling between the forward diffusion and the reverse diffusion. Essentially, it ensures that the model

yields reverse diffusion paths that are likely under the forward diffusion process. The third term represents the CO objective given in Eq. 4 and the fourth term is a collection of additive constants that are independent of θ .

As it is done in Ho et al. (2020) we use a time-conditioned diffusion model $q_\theta(X_{t-1}|X_t, t)$, where we provide the model with a one-hot encoding of the current time step t . In the following, we omit the conditional dependence on t for notational convenience.

3.2. Noise Distributions

Due to the discrete nature of UCO, noise distributions have to be discrete. In this work, we apply the following two noise distributions.

Categorical Noise Distribution. In Austin et al. (2021) the categorical distribution is proposed as a noise distribution in discrete diffusion models.

In the binary setting $X \in \{0, 1\}^N$, this distribution is given by $p(X_t|X_{t-1}) = \prod_{i=1}^N p(X_{t,i}|X_{t-1})$. Here, $X_{t,i}$ is the i -th component of $X_t = (X_{t,1}, \dots, X_{t,N})$ and $p(X_{t,i}|X_{t-1})$ is a Bernoulli distribution which is given by

$$p(X_{t,i}|X_{t-1}) = \begin{cases} (1 - \beta_t)^{1-X_{t-1,i}} \cdot \beta_t^{X_{t-1,i}} & \text{for } X_{t,i} = 0 \\ (1 - \beta_t)^{X_{t-1,i}} \cdot \beta_t^{1-X_{t-1,i}} & \text{for } X_{t,i} = 1 \end{cases}$$

where $\beta_t = \frac{1}{T-t+2}$. Effectively, this distribution flips the value of $X_{t-1,i}$ with a probability of β_t , independently of the energy function and the value of $X_{t-1,i}$. At $t = T$ this noise distribution yields the uniform distribution which is taken to be the stationary distribution.

Annealed Noise Distribution. When the Categorical Noise Distribution is used, the second term on the right-hand side of Eq. 6 becomes zero at $\mathcal{T} = 0$ and does not contribute to training. Therefore, we propose another noise distribution such that the resulting logarithm of the noise distribution scales inversely with the temperature. That way \mathcal{T} cancels out and this loss term does not vanish anymore at $\mathcal{T} = 0$. We use the target distribution at higher temperatures $1/(\beta \beta_t)$ to construct a noise distribution according to $p(X_t|X_{t-1}) = p_B(X_t, \beta \beta_t)$, where $\beta_t \in [0, 1[$. At $\beta_T = 0$ the Boltzmann distribution is equal to a uniform distribution, which is used as the stationary distribution $q(X_T)$. In the forward diffusion process, i.e. going from $t = 0$ to $t = T$, we sequentially decrease β_t with a linear schedule according to $\beta_t = 1 - \frac{t}{T}$. The resulting Annealed Noise Distribution reads:

$$p(X_t|X_{t-1}) \propto \exp(-\beta \beta_t X_t^T Q X_t),$$

where the normalization constant of $p_B(X, \beta \beta_t)$ is omitted since it is irrelevant to the optimization of the model for the reason that is given in Sec. 3.1. In contrast to the Categorical Noise Distribution, this noise distribution depends on the

energy function of the considered problem. In the forward direction, it initially favors to enter low-energy states and gradually turns into a uniform random selection of solutions. An empirical comparison between the Categorical Noise Distribution and the Annealed Noise Distribution is presented in Sec. 6.2.

4. Joint Variational Diffusion Models in Neural Combinatorial Optimization

In the following, we discuss the calculation of the individual terms in Eq. 6.

Each diffusion step is represented by a conditional mean-field distribution:

$$q_\theta(X_{t-1}|X_t) = \prod_i \hat{q}_\theta(X_t)_i^{X_{t-1,i}} (1 - \hat{q}_\theta(X_t)_i)^{1-X_{t-1,i}},$$

where $\hat{q}_\theta(X_t)_i := q_\theta(X_{t-1,i} = 1|X_t)_i$.

With these product distributions, the expectation of $H_Q(X)$ with respect to $q_\theta(X_{t-1}|X_t)$ can be conveniently calculated in closed form (see App. A.4):

$$\begin{aligned} \mathbb{E}_{X_{t-1} \sim q_\theta(X_{t-1}|X_t)}[H_Q(X_{t-1})] &= \sum_{i \neq j} Q_{ij} \hat{q}_\theta(X_t)_i \hat{q}_\theta(X_t)_j \\ &\quad + \sum_i Q_{ii} \hat{q}_\theta(X_t)_i. \end{aligned}$$

Likewise, we can compute the exact expectation of terms that include the noise distribution and the entropy (see App. A.4). The result for the entropy term reads:

$$\begin{aligned} S(q_\theta(X_{t-1}|X_t)) &= - \sum_i [\hat{q}_\theta(X_t)_i \log \hat{q}_\theta(X_t)_i \\ &\quad - (1 - \hat{q}_\theta(X_t)_i) \log (1 - \hat{q}_\theta(X_t)_i)]. \end{aligned}$$

The ultimate goal of UCO is to train a model on a dataset of CO problem instances $G_m \sim \omega(G_m)$, which can then be used at test time to generate solutions to i.i.d. CO problems via a time-efficient inference process. In our approach, this is realized by conditioning a diffusion model on each CO problem instance and by using the empirical mean over M_ω problem instances. We implement such a conditional generative model via the GNN-based architecture described in App. C.4. The corresponding overall loss function becomes:

$$L_\theta = \frac{\mathcal{T}}{M_\omega} \sum_{m=1}^{M_\omega} D_{KL}(q_\theta(X_{0:T}|G_m) || p(X_{0:T}|G_m)),$$

where the energy function is contained in the joint-target distribution $p(X_{0:T}|G_m)$ and depends on the specific CO problem instance at hand*. The expectations with respect to $q_\theta(X_{0:T}|G_m)$ in the loss above are approximated with a

*In the following we suppress the conditional dependencies on G_m to keep the notation simple.

Monte Carlo estimate based on M_{KL} samples drawn from the reverse process. We discuss the computation of the gradients of our loss in App. A.3 and details on the optimizer for the model parameters θ in App. C.5. In all of our experiments, we use annealing, i.e. we linearly decrease the temperature \mathcal{T} from $\mathcal{T}_{\text{start}}$ to zero. We show in Sec. 6.2 that this indeed improves the solution quality of our model.

4.1. Conditional Expectation

Conditional Expectation (CE) is an iterative strategy to obtain samples from a mean-field distribution that have a better-than-average solution quality (Raghavan, 1988; Karalias and Loukas, 2020). In the following, we consider a mean-field distribution $q_\theta(X) = \prod_i q_\theta(X_i)$ that is defined by the vector of Bernoulli probabilities v with components $v_i = q_\theta(X_i)$. In the first step of CE, we sort the components of v in descending order. Then starting from $i = 0$ the i -th component of the resulting vector is set to 0 to obtain a vector ω_0 and set to 1 to obtain another vector ω_1 . So in this case $\omega_0 = (0, p_1, \dots, p_N)$ and $\omega_1 = (1, p_1, \dots, p_N)$, where p_i is the i -th component of v after it is sorted. Then $H(\omega_0)$ and $H(\omega_1)$ are computed and v is updated to the configuration ω_j , where $j = \operatorname{argmin}_{l \in \{0,1\}} H(\omega_l)$ and retains the resulting value of v_i in the following steps. Next i is incremented to $i + 1$ and the process is repeated until all v_i are set to either 0 or 1.

Tokenized Conditional Expectation. In this work we propose to combine CE with Subgraph Tokenization (ST) (Sanokowski et al., 2023), where $k \in \mathbb{N}^+$ solution variables are grouped together to form a subgraph token. In the case of a token size k there are 2^k possible configurations of a given token. The CE procedure described above is carried out analogously by consecutively selecting token configurations ω_j with $j = \operatorname{argmin}_{l \in \{1, \dots, 2^k\}} H(\omega_l)$. Then i is incremented to $i + k$ and the next token gets its value assigned. This process above is repeated until all components of v are set. When k is not exceedingly large the values $H(\omega_l)$ can be computed in parallel for all of the 2^k configurations. This leads to a computational speed up with respect to CE since the number of iterations is reduced by a factor of k . The corresponding efficiency improvement is demonstrated experimentally in Sec. 6.1 and App. B.3.

4.2. EGN is secretly a one-step Joint Variational Diffusion Model

The UCO methods Erdős Goes Neural (EGN, Karalias and Loukas (2020)) and its annealed variant EGN-Anneal (Sun et al., 2022) use a mean-field distribution $q_\theta(X)$ which is parameterized by a GNN. They also use Eq. 2 as a loss function, which corresponds to minimizing $D_{KL}(q_\theta(X) || p_B(X))$. However, due to their reliance on the deterministic CE sampling procedure, their mod-

els would only generate one solution, which is why they use additional random node features $Z \sim q(Z)$ so that more than a single solution can be sampled from the models. Consequently, these methods are actually optimizing $\mathbb{E}_{Z \sim q(Z)}[D_{KL}(q_\theta(X|Z) || p_B(X))]$ which is equal to $D_{KL}(q_\theta(X, Z) || p_B(X)) - \hat{C}$, where $\hat{C} = \int q(Z) \log q(Z) dZ$ is constant. It can then be shown that:

$$\begin{aligned} D_{KL}(q_\theta(X, Z) || p_B(X)) + C \\ = D_{KL}(q_\theta(X, Z) || p(X, Z)). \end{aligned} \quad (7)$$

The right-hand side of this equation is equivalent to our loss in Eq. 6 when only one diffusion step is used, i.e. $T = 1$. Finally, since for diffusion models, C and \hat{C} are independent of model parameters (see App. A.5) the gradients of the left-hand side and right-hand side of the Eq. 7 are identical. Consequently, it can be concluded that DiffUCO generalizes EGN and EGN-Anneal.

5. Related Work

Variational Autoencoders. Variational Autoencoders (VAEs) are latent variable models, where samples X are drawn by first sampling latent variables Z from a prior distribution $q(Z)$ which are then processed by a stochastic decoder $q_\theta(X|Z)$ that is parameterized by a neural network (Kingma and Welling, 2014). The goal is to train the model on data X drawn from a data distribution $p_D(X)$ by minimizing the forward KL divergence $D_{KL}(p_D(X) || q_\theta(X)) = -\mathbb{E}_{X \sim p_D(X)}[\log q_\theta(X)] + C$. As $\log q_\theta(X) = \log(\int q_\theta(X|Z)q(Z) dZ)$ cannot be efficiently evaluated, the negative ELBO is estimated via an encoder network $p_\Phi(Z|X)$ and used as an upper bound on the forward KL divergence. Up to an additive constant, the ELBO objective is equivalent to the joint forward KL divergence $D_{KL}(p_{D,\Phi}(X, Z) || q_\theta(X, Z))$ (Kingma et al., 2019). In Ambrogioni et al. (2018) VAEs are used in variational inference by using KL divergences between joint-probability distributions.

Diffusion Models. Diffusion models (Sohl-Dickstein et al., 2015; Ho et al., 2020; Song et al., 2021) are latent variable models that are typically trained with samples X_0 from a data distribution $p_D(X_0)$. In the forward diffusion process noise is sequentially added at time step t on X_t by sampling a predefined noise distribution $p(X_t|X_{t-1})$. Starting from X_0 this process is repeated T times until the resulting data follows approximately a known stationary distribution $q(X_T)$. The diffusion model $q_\theta(X_{t-1}|X_t)$ learns the reverse diffusion process, i.e. to iteratively remove the noise from samples of the stationary distribution. Once trained, the repeated application of this model to samples of the stationary distribution yields a generative process that approximates sampling from $p_D(X_0)$. As in the case of VAEs, sample probabilities cannot be efficiently computed. There-

fore, these models are typically trained via a loss that is given by the negative ELBO. Discrete diffusion models are studied in Austin et al. (2021), where several different discrete noise distributions like the Categorical Noise Distribution used in this work are proposed.

Neural Optimization. Optimization using neural networks is a problem that is extensively researched. The objective is to find the optimal solution η of an objective, i.e. $\min_\eta O(\eta)$, where η can either be either a function or a variable. Usually, the goal is to find η without using training data. In UCO, for example, $\eta \in \{0, 1\}^N$ is a vector with binary variables. As described in Sec. 2 this is frequently framed as a Neural Probabilistic Optimization (NPO) problem, where the goal is to approximate a target distribution with a neural network. Wu et al. (2019) use autoregressive models to approximate discrete probability distributions related to Ising models. Another example of NPO are Boltzmann Generators (Noé and Wu, 2018), where Normalizing Flows are used to learn the equilibrium distribution of molecule configurations at a given temperature. However, their method cannot be used in CO as Normalizing Flows are not applicable in the discrete setting. In Müller et al. (2019) a Normalizing Flow is used for Monte Carlo Integration. Another example of NPO is Variational Monte Carlo, where neural networks are used to approximate the lowest energy state of a quantum system (Carleo and Troyer, 2017).

Approximate Likelihood Models in Neural Probabilistic Optimization In Wu et al. (2020) Normalizing Flows are trained to approximate an annealed target distribution in continuous NPO problems. Here, MCMC is used to introduce randomness in the iterative sampling process and the data processing inequality is used to obtain an upper bound of the reverse KL divergence to train the model. In Berner et al. (2022); Ghio et al. (2023); Richter et al. (2023); Vargas et al. (2023; 2024); Akhoud-Sadegh et al. (2024) diffusion models are used in the continuous setting of NPO.

Neural Combinatorial Optimization. Neural CO aims at the fast generation of high-quality solutions to CO problems. In supervised CO, a generative model is typically trained using solutions to CO problems that are obtained by classical solvers such as Gurobi (Gurobi Optimization, LLC, 2023) or KaMIS (Lamm et al., 2017). In Sun and Yang (2023), for example, discrete diffusion models are applied by training them on data from solvers. In Li et al. (2018) (INTEL) and Böther et al. (2022) (DGL) supervised learning is combined with tree search refinement. As highlighted in Yehuda et al. (2020), these supervised approaches suffer from the problem of expensive data generation, leading to a growing interest in Unsupervised CO (UCO). In UCO, the aim is to train a model that learns how to solve CO problems without relying on labeled training data. These methods commonly

rely on exact likelihood models, such as mean-field models (Karalias and Loukas, 2020; Sun et al., 2022; Wang and Li, 2023). These works highlight that the calculation of expectation values in the context of UCO is particularly convenient with mean-field models due to their assumption of statistical independence among the modeled random variables. As outlined in Sec. 3 our diffusion model approach also enjoys this feature. In Ahn et al. (2020) (LwtD) a mean-field model is combined with Reinforcement Learning (RL) to iteratively refine solutions. Sanokowski et al. (2023) demonstrate that the statistical independence assumption in mean-field models restricts the performance of these approaches on particularly hard CO problems. They show that more expressive exact likelihood models like autoregressive models exhibit performance benefits. However, these models suffer from high memory requirements and long sampling time which slows down the training process. Sanokowski et al. (2023) (VAG-CO) argue that autoregressive methods should be combined with RL to reduce the memory requirements and speed up training. Khalil et al. (2017) use autoregressive models combined with Q-learning to select the next solution variable together with its value. Sanokowski et al. (2023) address the problem of slow sampling in autoregressive models by introducing Subgraph Tokenization. Zhang et al. (2023) use GFlow networks (Bengio et al., 2021b), which realizes autoregressive solution generation in UCO via GFlowNets (Bengio et al., 2021a).

6. Experiments

In this work, we evaluate our method DiffUCO on five different CO problem types, namely Maximum Independent Set (MIS), Maximum Clique (MaxCl), Minimum Dominating Set (MDS), Maximum Cut (MaxCut), and Minimum Vertex Cover (MVC). The energy functions used by DiffUCO for these CO problems can be found in Tab. 2. To compare our method to recently published state-of-the-art autoregressive models, we compare on MIS, MaxCl, MDS, and MaxCut to published results reported in Let The Flows Tell (LTFT, Zhang et al. (2023)) (see Tab. 1 and Tab. 3) and on MVC we compare our method to published results from Sanokowski et al. (2023) (see. Fig. 2, Left). For the experiments on MIS and MaxCl, graphs are generated by the so-called RB-Model (Xu et al., 2005), which is known to yield hard instances of CO problems like MIS, MVC and MaxCl (Tönshoff et al., 2020). On MDS and MaxCut randomly generated Barabasi-Albert (BA) graphs (Barabási and Albert, 1999) are used. For both of these graph types, 4000 graphs are used for training and 500 graphs are used for the validation and testing, respectively. In either case, we follow Zhang et al. (2023) and generate a small and a large dataset with uniformly random sampled graph sizes between 200-300 and 800-1200 nodes. In accordance with Zhang et al. (2023) we label the methods in our tables either

as Operations Research (OR), Supervised Learning (SL), or Unsupervised Learning (UL). On BA datasets, two Heuristic (H) baselines namely Greedy and mean-field annealing (MFA) are additionally used (Bilbro et al., 1988). The OR methods have substantially longer running times than the other methods, as they are usually run until the optimal solution is found. Therefore, these results cannot be used to compare the OR methods to the other methods. The OR results are merely provided to showcase the quality of optimal solutions to the CO problem datasets. For each method we provide the inference time required to solve the test set problems (see App. C.7).

First, we report the performance of our method DiffUCO with and without CE and we investigate the runtime advantage of combining CE with ST (see Sec. 4.1). We denote the combination of CE and ST as DiffUCO: CE-ST_k, where k is the token size used for ST. As DiffUCO: CE-ST₁ equates to CE without using ST, we will just denote it as DiffUCO: CE. For the experiments in Tab. 1 and Tab. 3 we report our results as a test set average over eight samples per CO instance which is then averaged over three training seeds. For more details about the experiments on MVC we refer to the corresponding paragraphs. In Sec. 6.2 we show that DiffUCO’s solution quality can further be improved when more diffusion steps are applied during evaluation time than we use during training. Therefore, on benchmark datasets, we always evaluate our method with three times more diffusion steps than we have used during training (see App. C.2).

6.1. Benchmarks

Maximum Independent Set. The MIS problem is the problem of finding the largest set within a graph under the constraint that neighboring nodes are not both within the set. We evaluate our method on RB-small and RB-large and compare to published results from Zhang et al. (2023). Results are shown in Tab. 1 (Left). The CE and CE-ST variants of DiffUCO achieve the best results on RB-large and marginally outperform LTFT on RB-small.

Minimum Dominating Set. On the MDS problem, the goal is to find the set with the lowest number of vertices in a graph so that at least one neighbor of each node is within the set. We evaluate our method on the BA-small and BA-large datasets and compare in Tab. 1 (Right) our results to published results of Zhang et al. (2023). Here, DiffUCO and its variants outperform all other methods on both datasets.

Maximum Clique. In MaxCl the goal is to find the largest set of nodes, in which every node is connected to all other nodes within the set. The MaxCl problem can be solved by solving the MIS problem on the complementary graph (see Tab. 2). We evaluate our method on the RB-small datasets. Results are shown in Tab. 3 (Left). DiffUCO CE-ST achieves the best results and outperforms LTFT in-

MIS		RB-small		RB-large		MDS		BA-small		BA-large	
Method	Type	Size ↑	time ↓	Size ↑	time ↓	Method	Type	Size ↓	time ↓	Size ↓	time ↓
Gurobi (r)	OR	19.98	47:34	40.90	2:10:26	Gurobi (r)	OR	27.89	1:47	103.80	13:48
KaMIS (r)	OR	20.10	1:24:12	43.15	2:03:36	Greedy (r)	H	37.39	2:13	140.52	35:01
LwtD (r)	UL	19.01	1:17	32.32	7:33	MFA (r)	H	36.36	2:56	126.56	36:31
INTEL (r)	SL	18.47	13:04	34.47	20:17	EGN: CE (r)	UL	30.68	1:00	116.76	3:56
DGL (r)	SL	17.36	12:47	34.50	23:54	EGN-Anneal: CE (r)	UL	1:01	3:55	111.50	3:55
LTFT (r)	UL	19.18	0:32	37.48	4:22	LTFT (r)	UL	28.61	2:20	110.28	32:12
DiffUCO	UL	18.88 ± 0.06	0:07	38.10 ± 0.13	0:10	DiffUCO	UL	28.30 ± 0.10	0:05	107.01 ± 0.33	0:05
DiffUCO: CE	UL	19.24 ± 0.05	0:54	38.87 ± 0.13	4:57	DiffUCO: CE	UL	28.20 ± 0.09	0:54	106.61 ± 0.30	3:28
DiffUCO: CE-ST ₈	UL	19.24 ± 0.05	00:13	38.87 ± 0.13	1:05	DiffUCO: CE-ST ₈	UL	28.20 ± 0.09	0:13	106.61 ± 0.31	0:40

Table 1. Left: Average independent set size on the whole test dataset on the RB-small and RB-large dataset. The higher the better. Right: Average dominating set size on the whole test dataset on the BA-small and BA-large datasets. The lower the set size the better. Left and Right: Total evaluation time is shown in h:m:s. (r) indicates that results are reported as in Zhang et al. (2023). (CE) indicates that results are reported after applying Conditional Expectation and (ST_k) indicates that results are reported by combining CE and ST at a specific k for faster CE. The best neural method is marked as bold.

Problem Type	Objective: $\min_{X \in \{0,1\}^N} H(X)$
MIS	$H(X) = -A \sum_{i=1}^N X_i + B \sum_{(i,j) \in \mathcal{E}} X_i \cdot X_j$
MDS	$H(X) = A \sum_{i=1}^N X_i + B \sum_{i=1}^N (1 - X_i) \prod_{j \in \mathcal{N}(j)} (1 - X_j)$
MaxCl	$H(X) = -A \sum_{i=1}^N X_i + B \sum_{(i,j) \notin \mathcal{E}} X_i \cdot X_j$
MaxCut	$H(\sigma) = -\sum_{(i,j) \in \mathcal{E}} \frac{1 - \sigma_i \sigma_j}{2}$ where $\sigma_i = 2X_i - 1$
MVC	$H(X) = A \sum_{i=1}^N X_i + B \sum_{(i,j) \in \mathcal{E}} (1 - X_i) \cdot (1 - X_j)$

Table 2. Table with energy functions of the MIS, MDS, MaxCl, MaxCut and MVC problems (Lucas, 2014). Choosing $A < B$ ensures that all minima of the energy function are feasible solutions. In all of our Experiments, we chose $A = 1.0$ and $B = 1.01$.

significantly.

Maximum Cut. The goal of the MaxCut problem is to separate a graph into two different sets of nodes so that the number of edges between these two sets is as high as possible. On MaxCut we evaluate our method on BA-small and BA-large graphs, where we see in Tab. 3 (Right) that DiffUCO and its variants are the best-performing UCO methods. On BA-large our methods even outperform Gurobi with a long time limit of 300 seconds per graph.

Minimum Vertex Cover. The MVC problem aims at finding the smallest set of nodes so that every edge is connected to at least one node in the set. On MVC we evaluate our method on the RB-200 dataset (see App. C.3) as reported in Sanokowski et al. (2023), where the method VAG-CO is compared to EGN, EGN-Anneal, the greedy baseline DB-Greedy (Tönshoff et al., 2020) and Gurobi at different time limits. We compare to reported results from Sanokowski et al. (2023), where the test set average of the best approximation ratio AR^* (the lower the better, see App. C.1) is reported at different generation parameter values of p . The parameter p controls the hardness of the CO problem on RB-graphs, where lower values of p yield harder instances (Tönshoff et al., 2020). Results are shown in Fig. 2 (Left),

where we see that DiffUCO significantly outperforms all other UCO methods on all values for p . On hard instances, i.e. low values of p , DiffUCO achieves a significantly lower AR^* than Gurobi, while needing only a fraction of its runtime. However, we note that a fair comparison of our method to Gurobi is difficult as Gurobi runs on CPUs and DiffUCO mostly on GPUs.

Combination of CE and ST. Our experiments show that on MIS and MaxCl the use of CE improves the performance of our model significantly and that by using ST the computational cost of CE can be reduced by a large margin (see Tab. 1 and Tab. 3).

6.2. Ablations

Figure 2 (Middle) shows that increasing the number of diffusion steps during training consistently improves the performance of DiffUCO. For this, we train the model for different numbers of diffusion steps $T \in \{1, 2, 3, 4, 5\}$ on the RB-100 MIS dataset (see App. C.3) and observe in Fig. 2 that the relative error ϵ_{rel} (the lower the better, see App. C.1) decreases as T is increased, irrespectively of the used noise distribution. We also see that the Categorical Noise Distribution (CND) (blue) performs similarly to the Annealed Noise Distribution (AND) (orange) and that the use of annealing is highly beneficial. If we compare CND w/o annealing to AND w/o annealing we find that the results of the CND have a higher variance but similar average results. A possible reason for this is that the second term on the right-hand side of Eq. 6 is zero for the CND when no annealing is used. Therefore, this noise loss term provides no guidance at intermediate diffusion steps which may make the training process more prone to being stuck in different local minima depending on different weight initialization. We conclude from this that the AND might be the preferable noise distribution when no or only a little annealing is used.

More evaluation diffusion steps improve solution qual-

MaxCI		RB-small		MaxCut		BA-small		BA-large	
Method	Type	Size \uparrow	time \downarrow	Method	Type	Size \uparrow	time \downarrow	Size \uparrow	time \downarrow
Gurobi (r)	OR	19.05	1:55	Gurobi t_{lim}	OR	730.87 ± 2.35	8:30:00	2944.38 ± 0.86	1:17:35:00
Greedy (r)	H	13.53	0:25	Greedy (r)	H	688.31	0:13	2786.00	3:07
MFA (r)	H	14.82	0:27	MFA (r)	H	704.03	1:36	2833.86	7:16
EGN: CE (r)	UL	12.02	0:41	EGN: CE (r)	UL	693.45	0:46	2870.34	2:49
EGN-Anneal: CE (r)	UL	14.10	2:16	EGN-Anneal: CE (r)	UL	696.73	0:45	2863.23	2:48
LTFT (r)	UL	16.24	0:42	LTFT (r)	UL	704	2:57	2864	21:20
DiffUCO	UL	14.51 ± 0.39	0:04	DiffUCO	UL	727.11 ± 2.31	0:04	2947.27 ± 1.50	0:04
DiffUCO: CE	UL	16.22 ± 0.09	1:00	DiffUCO: CE	UL	727.32 ± 2.33	1:00	2947.53 ± 1.48	3:47
DiffUCO: CE-ST ₈	UL	16.30 ± 0.08	0:13	DiffUCO: CE-ST ₈	UL	727.33 ± 2.31	0:13	2947.53 ± 1.49	0:53

Table 3. Left: Testset average clique size on the whole RB-small dataset. The larger the set size the better. Right: Average test set cut size on the BA-small and BA-large datasets. The larger the better. Left and Right: Total evaluation time is shown in d:h:m:s. See Tab. 1 for the meaning of (CE) and (CE-ST₈). Gurobi t_{lim} denotes that Gurobi was run with a time limit. On BA-small the time limit is set to 60 and on BA-large to 300 seconds per graph. The best neural method is marked as bold.

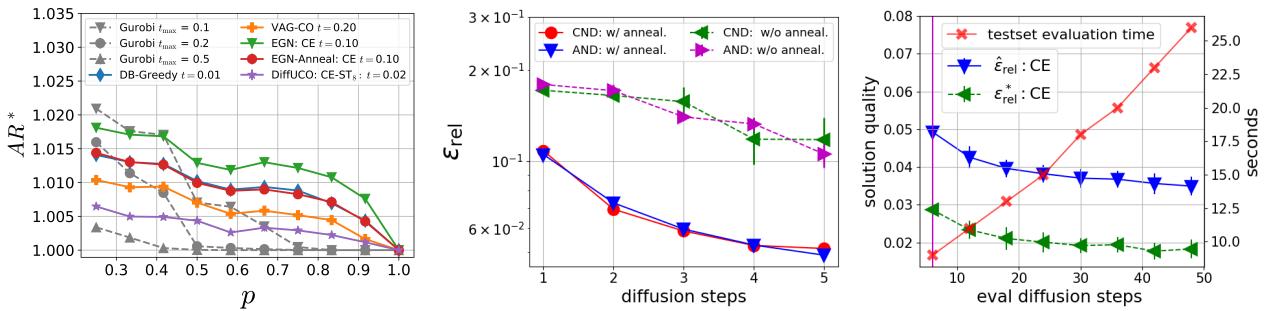


Figure 2. Left: Comparison of AR^* at different generation values p on the RB-small MVC dataset. t is the time that each method takes to generate the solutions. Middle: Relative error over the number of diffusion steps used during training for the Categorical Noise Distribution (CND) and the Annealed Noise Distributions (AND) in the setting with (w/ anneal) and without annealing (w/o anneal.). Right: Solution quality (left axis) on the RB-small MIS dataset over the number of diffusion steps during evaluation. $\hat{\epsilon}_{rel}$ is the average relative error on the test set and ϵ_{rel}^* is the best test set relative error out of 8 model samples. The model is trained with 6 diffusion steps (purple vertical line).

ity. In Fig. 2 (Right) we investigate whether the solution quality of DiffUCO can be improved by applying more diffusion steps during evaluation than during training. Here, the model is trained with 6 diffusion steps (purple vertical line) on the RB-small MIS dataset. Figure 2 (Right) shows on the left axis the test dataset average relative error $\hat{\epsilon}_{rel}$ and the best relative error ϵ_{rel}^* out of 8 solutions per problem instance over an increasing number of diffusion steps during evaluation (for details see App. C.2). We also plot on the additional y-axis the evaluation time on the whole dataset. The results show that both, $\hat{\epsilon}_{rel}$ and ϵ_{rel}^* consistently improve as the number of steps is increased. This improvement comes at the cost of a linearly increasing evaluation time cost.

7. Limitations and Future Work

While our method yields a model that can be used to generate high-quality solutions, the learned probability distribution will in practice always have a bias, i.e. the model samples do not exactly follow the target distribution. In several interesting applications, a learned probability function is only useful when it allows to obtain unbiased samples

from the target distribution (Wu et al., 2021; Nicoli et al., 2020; McNaughton et al., 2020). Training DiffUCO on large graph datasets with high connectivity is memory- and time-expensive (see App. C.6). Therefore, it would be interesting to consider approaches that improve upon these two aspects. This could be achieved by reducing the computational graph during backpropagation. The concept of latent diffusion models (Rombach et al., 2022) and U-Net inspired GNN architectures (Ronneberger et al., 2015) represent promising research directions in this context.

8. Conclusion

In this work, we introduce the Joint Variational Upper Bound as an efficiently computable loss that allows the application of latent variable models like diffusion models in Neural Probabilistic Optimization. Specifically, we demonstrate the excellent performance of diffusion models in Unsupervised Combinatorial Optimization. Our method outperforms recently presented methods on a wide range of benchmarks. Finally, we show in experiments that the application of variational annealing and additional diffusion

steps during inference consistently improves the solution quality of the model.

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Impact Statement

In this work, we aim to improve data-free neural optimization algorithms. We hope our work will contribute to advances in various scientific fields such as Combinatorial Optimization.

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A. Derivations

A.1. Joint Variational Upper Bound

Consider the simple case of a one-step diffusion model. Here in the forward process the distribution $p_B(x)$ is directly mapped to the stationary distribution $q(z)$ by the application of a noise distribution $p(z|x)$. In the reverse process, the parameterized distribution $q_\theta(x|z)$ directly maps samples from the stationary distribution $q(z)$ to the target distribution $p_B(x)$.

In the following we will derive that $D_{KL}(q_\theta(x) \parallel p_B(x))$ is upper bounded by $D_{KL}(q_\theta(x, z) \parallel p_B(x, z))$, where $p_B(x, z) = p(z|x) p_B(x)$ and $q_\theta(x, z) = q_\theta(x|z) q(z)$.

Derivation via the chain-rule for KL divergences:

This inequality follows directly from the chain-rule for KL divergences which states that:

$$\begin{aligned} D_{KL}(q_\theta(x, z) \parallel p_B(x, z)) &= \int q_\theta(x, z) \log \frac{q_\theta(x, z)}{p_B(x, z)} dx dz \\ &= \int q_\theta(x, z) \left[\log \frac{q_\theta(x)}{p(x)} + \log \frac{q_\theta(z|x)}{p(z|x)} \right] dx dz \\ &= D_{KL}(q_\theta(x) \parallel p_B(x)) + \mathbb{E}_{q_\theta(x)} [D_{KL}(q_\theta(z|x) \parallel p(z|x))]. \end{aligned}$$

According to Gibbs' inequality, the second term on the right-hand side is non-negative which is sufficient to show the upper bound.

Derivation via the Evidence Upper Bound:

Alternatively, the upper bound can be derived using the Evidence Upper Bound (EUBO) (Ji and Shen, 2019), where we use $q_\theta(x) = \frac{q_\theta(x, z)}{q_\theta(z|x)}$ and $\int q_\theta(z|x) dz = 1$ and therefore

$$\log q_\theta(x) = \int q_\theta(z|x) \log q_\theta(x) dz = \int q_\theta(z|x) [\log q_\theta(x, z) - \log q_\theta(z|x)] dz$$

In the first step, we will derive an upper bound for $\log q_\theta(x)$ by using the EUBO that is based on the Gibbs' inequality.

$$\log q_\theta(x) = \int q_\theta(z|x) [\log q_\theta(x, z) - \log q_\theta(z|x)] dz \leq \int q_\theta(z|x) [\log q_\theta(x, z) - \log p(z|x)] dz$$

By using this inequality we can now show that $D_{KL}(q_\theta(x) \parallel p_B(x)) \leq D_{KL}(q_\theta(x, z) \parallel p_B(x, z))$.

$$D_{KL}(q_\theta(x) \parallel p_B(x)) = \int q_\theta(x) \log \frac{q_\theta(x)}{p_B(x)} dx \leq \int q_\theta(x) \left[\int q_\theta(z|x) (\log q_\theta(x, z) - \log p(z|x)) dz - \log p_B(x) \right] dx$$

By using

$$\int q_\theta(x) \log p_B(x) dx = \int \int q_\theta(x, z) \log p_B(x) dz dx,$$

we can then write with $-\log p(z|x) - \log p_B(x) = -\log p_B(x, z)$

$$\begin{aligned} D_{KL}(q_\theta(x) \parallel p_B(x)) &\leq \int \int q_\theta(x, z) [\log q_\theta(x, z) - \log p(z|x) - \log p_B(x)] dz dx \\ &= D_{KL}(q_\theta(x, z) \parallel p_B(x, z)), \end{aligned}$$

which proves the inequality.

A.2. Joint Variational Diffusion Model Objective

In this section, we aim to derive the objective shown in Eq.6.

Starting from

$$\begin{aligned} D_{KL}(q_\theta(X_{0:T}) \parallel p(X_{0:T})) &= \sum_{X_{0:T}} q_\theta(X_{0:T}) \left[\log q_\theta(X_{0:T}) - \log p(X_{0:T}) \right] \\ &= \sum_{X_{0:T}} q_\theta(X_{0:T}) \left[\sum_{t=1}^T \left[\log q_\theta(X_{t-1}|X_t) - \log p(X_t|X_{t-1}) \right] + \log q(X_T) - \log p_B(X_0) \right] \end{aligned}$$

we can first simplify the first term with

$$\sum_{X_{T:0}} q_\theta(X_{T:0}) \log q_\theta(X_{t-1}|X_t) = \sum_{X_{T:t-1}} q_\theta(X_{T:t-1}) \log q_\theta(X_{t-1}|X_t)$$

where we have used that the sum over all X_τ where $\tau < t-1$ sums up to one because $\log q_\theta(X_{t-1}|X_t)$ is independent of samples coming from these time steps. Remember that in the reverse process the sample process starts at X_T and ends at X_0 . So samples, where $\mathcal{T} < t-1$ are future events and therefore $\log q_\theta(X_{t-1}|X_t)$ does not depend on these future events. In the second step we pull out the sum over X_{t-1} , so that we arrive at

$$\begin{aligned} \sum_{X_{T:t-1}} q_\theta(X_{T:t-1}) \log q_\theta(X_{t-1}|X_t) &= \sum_{X_{T:t}} q_\theta(X_{T:t}) \sum_{X_{t-1}} q_\theta(X_{t-1}|X_t) \log q_\theta(X_{t-1}|X_t) \\ &= - \sum_{X_{T:t}} q_\theta(X_{T:t}) S(q_\theta(X_{t-1}|X_t)), \end{aligned}$$

where $S(q_\theta(X_{t-1}|X_t))$ is the entropy over $q_\theta(X_{t-1}|X_t)$. Analogously, this simplification can be done with $\tilde{H}(X_t, X_{t-1}) := -\log p(X_t|X_{t-1})$, so that we arrive at:

$$\sum_{X_{T:0}} q_\theta(X_{T:0}) \tilde{H}(X_t, X_{t-1}) = \sum_{X_{T:t-1}} q_\theta(X_{T:t-1}) \tilde{H}(X_t, X_{t-1})$$

Furthermore, we use that $\log p_B(X_0) = -\beta H(X_0) - \log \mathcal{Z}$ and absorb $\log(\mathcal{Z})$ and $\log q(X_T)$ into the constant C as these terms neither depend on θ nor on $X_{0:T-1}$. Finally, we arrive at:

$$\begin{aligned} D_{KL}(q_\theta(X_{0:T}) \parallel p(X_{0:T})) &= \sum_{t=1}^T \sum_{X_{T:t}} q_\theta(X_{T:t}) \left[\left[S(q_\theta(X_{t-1}|X_t)) - \log p(X_t|X_{t-1}) \right] + \log q(X_t) - \log p_B(X_0) \right] \\ &= - \sum_{t=1}^T \mathbb{E}_{X_{T:t} \sim q_\theta(X_{T:t})} [S(q_\theta(X_{t-1}|X_t))] \\ &\quad + \sum_{t=1}^T \mathbb{E}_{X_{T:t-1} \sim q_\theta(X_{T:t-1})} [\tilde{H}(X_{t-1}, X_t)] \\ &\quad + \beta \mathbb{E}_{X_{T:0} \sim q_\theta(X_{T:0})} [H(X_0)] + C, \end{aligned}$$

where we have exchanged the sums over $X_{T:t}$ with the expectation over $X_{T:t} \sim q_\theta(X_{T:t})$. Finally, we can arrive at Eq. 6 by multiplying the whole objective with the temperature \mathcal{T} .

A.3. Gradient of the Joint Variational Upper Bound

When $X_{0:T}$ is discrete, the gradient of the Joint Variational Upper Bound has to be decomposed into two parts via the application of the chain rule. The gradient through exact expectations (see Sec. 4) can be computed by backpropagating through $q_\theta(X_{t-1}|X_t)$ directly. To backpropagate through the expectations over $X_{T:t-1} \sim q_\theta(X_{T:t-1})$ we use the log derivative trick with a baseline for variance reduction.

As an example, we show the gradient through the entropy term which is given by:

$$\begin{aligned}\nabla_\theta \mathbb{E}_{X_{T:t} \sim q_\theta(X_{T:t})} [S(q_\theta(X_{t-1}|X_t))] &= \mathbb{E}_{X_{T:t} \sim q_\theta(X_{T:t})} \left[[S(q_\theta(X_{t-1}|X_t)) - b_S] \nabla_\theta \log q_\theta(X_{T:t}) \right] \\ &\quad + \mathbb{E}_{X_{T:t} \sim q_\theta(X_{T:t})} [\nabla_\theta S(q_\theta(X_{t-1}|X_t))],\end{aligned}$$

where the baseline $b_S = \mathbb{E}_{X_{T:t} \sim q_\theta(X_{T:t})} [S(q_\theta(X_{t-1}|X_t))]$ is used for variance reduction. Here, the first term is the REINFORCE gradient through the expectation and the second term is the gradient which flows through the exact expectation of the entropy.

A.4. Exact expectations of mean-field distributions

Exact Expectation of Energy Functions. In the following, we will derive the exact expectations from Sec. 4. Suppose we have a mean-field distribution $p(X) = \prod_{i=1}^N p(X_i)$ and want to calculate $\mathbb{E}_{X \sim p(X)} [Q_{ij} X_i X_j]$, where $X_i \in \{0, 1\}$, then when $i \neq j$

$$\begin{aligned}\mathbb{E}_{X \sim p(X)} [Q_{ij} X_i X_j] &= \sum_X p(X) Q_{ij} X_i X_j = \sum_X \prod_{i=1}^N p(X_i) Q_{ij} X_i X_j \\ &= \sum_{X_i} \sum_{X_j} p(X_i) p(X_j) Q_{ij} X_i X_j = Q_{ij} p(X_i = 1) p(X_j = 1) := Q_{ij} p_i p_j.\end{aligned}$$

In the third equality, we have integrated out the probabilities that do not depend on X_i and X_j and in the fourth equality we have used that only terms where $X_i = X_j = 1$ will remain. For $i = j$ it can be similarly shown that $\mathbb{E}_{X \sim p(X)} [Q_{ii} X_i X_i] = Q_{ii} p_i$, by using that $X_i^2 = X_i$.

Exact Expectation of Bernoulli Noise Distribution. As explained in Sec. 4 for the binary case $X_i \in \{0, 1\}$ the Categorical Noise Distribution can be reduced to a Bernoulli Noise Distribution.

The logarithm of this noise distribution can be written as:

$$\begin{aligned}\tilde{H}(X_t, X_{t-1}) &:= \sum_i (1 - X_{t,i}) \left[(1 - X_{t-1,i}) \log(1 - \beta_t) + X_{t-1,i} \log \beta_t \right] \\ &\quad + X_{t,i} \left[(1 - X_{t-1,i}) \log(1 - \beta_t) + X_{t-1,i} \log \beta_t \right]\end{aligned}\tag{8}$$

Similarly, as done before we can calculate the expectation with respect to a mean-field distribution $p(X_t)$ exactly, where we arrive at

$$\begin{aligned}p(X_t|X_{t+1}) p(X_{t-1}|X_t) \tilde{H}(X_t, X_{t-1}) &= p(X_t|X_{t+1}) \sum_i (1 - X_{t,i}) \left[p_{t-1,i} \log(1 - \beta_t) + (1 - p_{t-1,i}) \log \beta_t \right] \\ &\quad + X_{t,i} \left[(1 - p_{t-1,i}) \log(1 - \beta_t) + p_{t-1,i} \log \beta_t \right],\end{aligned}$$

where we define $p_{t-1,i} := p(X_{t-1,i} = 1|X_t)$. The expectation over $p(X_t|X_{t+1})$ cannot be computed in closed form because $p_{t-1,i}$ depends nontrivially on X_t .

Exact Expectation of the Entropy. For the exact expectation of the entropy we can similarly show that:

$$\begin{aligned}
S(p(X)) &= - \sum_X p(X) \log p(X) = \sum_X \prod_i p(X_i) \sum_j \log p(X_j) \\
&= - \sum_X \sum_j \prod_i p(X_i) \log p(X_j) \\
&= - \sum_j \sum_{X_j} p(X_j) \log p(X_j) = \sum_j \left[p_j \log p_j - (1 - p_j) \log(1 - p_j) \right]
\end{aligned}$$

A.5. Equivalence of one step DiffUCO and EGN

In diffusion models, the last step of the forward diffusion process typically corresponds to sampling independent random noise. For a one-step diffusion model as considered in Sec. 4.2 this implies that $p(Z|X)$ does not depend on X . With this insight it follows that:

$$C = \sum_{X,Z} q_\theta(X|Z) q(Z) \log p(Z|X) = \sum_Z q(Z) \log p(Z).$$

This shows that C is independent of θ .

B. Additional Experiments

B.1. Experiments on the Gset Dataset

We furthermore evaluate our method on the Gset Maxcut dataset, where models are trained and validated on Erdős–Rényi graphs and then evaluated out-of-distribution on Gset graphs (Ye, 2003). We train our method on Erdős–Rényi (ER) graphs between 200 to 500 nodes that are generated with a uniformly sampled edge probability $p \in \{0.05, 0.3\}$. The neural baseline methods ECORD (Barrett et al., 2022), ECO-DQN (Barrett et al., 2020), RUNCSO (Tönshoff et al., 2020), and ANYCSP (Tönshoff et al., 2023) use graphs with the same edge probability but train on graphs with vertex sizes of 500, 200, 100, and 100 respectively. All methods are validated on ER graphs with 500 vertices. Results are also compared to a greedy construction algorithm (Greedy) and semidefinite programming (SDP) (Goemans and Williamson, 1995) which is a well-known approximation algorithm on MaxCut. Results are shown in Tab. 5, where the difference to the best-known cut value is shown. We compare to results reported as in ANYCSP (Tönshoff et al., 2023). Here, Gset graphs are grouped according to their vertex size $|V|$ and then on each graph models run with 20 parallel processes for a time limit of 180 seconds. Results show that our model achieves the second-best results and ANYCSP achieves the best results in all groups. However, a comparison between DiffUCO and ANYCSP cannot be considered to be fair since DiffUCO is a solution generation method whereas ANYCSP is a search method. The former category of methods is trained to generate solutions without iteratively improving a solution, which differentiates them from search methods. Therefore, search-based methods such as ANYCSP have an advantage in the setting of sampling 20 solutions in parallel within the time limit of 180 seconds per graph. Furthermore, the comparison between neural methods in this setting may not be fair because the test dataset is out-of-distribution (OOD) and the training datasets of the neural methods also differ in some cases. For a more detailed discussion on OOD generalization experiments, we refer to App. B.2.

B.2. Out of distribution generalization

We present in Tab. 6 OOD results on Barabási-Albert (BA) graphs for both the MaxCut and the Minimum Dominating Set (MDS) problems. Our approach is trained on BA-small graphs and then evaluated on datasets comprising larger graphs, including BA-large, BA-huge, and BA-giant with 800-1200, 1200-1800, and 2000-3000 nodes, respectively. In this assessment, we benchmark our method against Gurobi with a generous time constraint of 300 seconds per graph. We observe remarkably good generalization capabilities on these datasets and even outperform Gurobi on BA-large, BA-huge, and BA-giant MaxCut at the given time limit. However, OOD results have to be treated with great caution as it is well known (Le and Jegelka, 2023; Zhang et al., 2021) that OOD generalization on graph problems does not only depend on the method but also the GNN architecture and the node-to-degree distribution of the graphs. Our method was by no means designed for robust OOD generalization, which would be an attractive avenue for future work.

B.3. Effect of different token sizes in CE-ST

In Fig. 3 (Left) we show how the solution quality and the inference speed of our model changes on the RB-100 MIS dataset when we vary the token size k within CE-ST. Here, the relative error ϵ_{rel} is plotted on the left axis and the evaluation time in seconds on the right axis. We see that k does not influence the solution quality of the model and the inference time has an optimum at $k = 8$. Because some operations within our implementation of CE-ST scale exponentially with k the evaluation time increases at $k > 8$. By further optimizing the CE-ST code in this respect, the negative time effects of these operations might be reduced.

B.4. DiffUCO learns to generate samples that are more likely under the forward diffusion process

In Fig. 3 (Right) we show that our model indeed learns a diffusion process. To show that samples are related to the diffusion process we plot the evolution of the unnormalized forward process likelihood $\mathbb{E}_{X_{0:T} \sim q_\theta(X_{0:T})} [\log p(X_{0:T})]$ averaged over model samples throughout training. As this measure increases during training, this indicates that the model indeed learns to generate samples that are related to the diffusion process, i.e. it shows that the generative process is increasingly reconstructing the forward diffusion process.

B.5. Factoring in training time

In Tab. 4 we quantify the number of n_{Query} , where $t_{\text{train}} + n_{\text{Query}} \cdot t_{\text{eval}} < t_{\text{Gurobi}}$. Under the assumption that the solution quality of Gurobi and the model are the same, this quantity is supposed to show at which amount of samples it is more useful to use our model instead of Gurobi. Our results show that the number of n_{Query} on some problems is quite large, while on others it is comparably low. On MaxCut, for example, Gurobi takes a lot of time to obtain a good solution quality, and due to the low connectivity of the BA graphs our method does not take a lot of time for training. In contrast to that n_{Query} is very large on MaxCl because, as we have mentioned in App B.6, the MaxCl experiments are the most expensive due to the high connectivity of the complementary RB graphs.

CO problem type	MaxCut		MIS		MDS		MaxCl
Dataset	BA-small	BA-large	RB-small	RB-large	BA-small	BA-large	RB-small
n_{Query}	482	197	8.596	48.216	348.830	31.525	1.752.059

Table 4. The number of n_{Query} , where $t_{\text{train}} + n_{\text{Query}} \cdot t_{\text{eval}} < t_{\text{Gurobi}}$. Here, t_{train} is the training time of DiffUCO, t_{eval} is DiffUCO’s evaluation time per graph and t_{Gurobi} is the evaluation time per graph of Gurobi.

Method	$ V = 800$	$ V = 1K$	$ V = 2K$	$ V \geq 3K$
Greedy (r)	411.44	359.11	737.00	774.25
SDP (r)	245.44	229.22	N/A	N/A
RUNCSP (r)	185.89	156.56	357.33	401.00
ECO-DQN (r)	65.11	54.67	157.00	428.25
ECORD (r)	8.67	8.78	39.22	187.75
ANYCSP (r)	1.22	2.44	13.11	51.63
DiffUCO: CE-ST ₈	4.11	6.33	31.67	116.75

Table 5. Comparison to MaxCut results from Tönshoff et al. (2023). Models are evaluated according to the average deviation from the best-known cut size on the Gset dataset. The graphs are grouped according to their node size $|V|$. (r) indicates that the results are taken from Tönshoff et al. (2023).

MDS Size ↓				
Method	BA-small	BA-large	BA-huge	BA-giant
Gurobi - $t_{lim} = 300s$	27.91	105.05	153.89	253.53
DiffUCO: CE-ST ₈	28.31	106.08	157.46	259.02
MaxCut Size ↑				
Gurobi - $t_{lim} = 300s$	735.67	2945.43	4414.29	7319.47
DiffUCO: CE-ST ₈	731.72	2949.02	4444.47	7390.99

Table 6. Out-of-distribution (OOD) results on MDS and MaxCut. DiffUCO is trained on BA-small and then the average set size is evaluated on three larger OOD datasets BA-large, BA-huge with an average of 2000 nodes, and BA-giant with an average of 3000 nodes. Gurobi is evaluated at a time limit of 300s per graph. The best method is marked as bold.

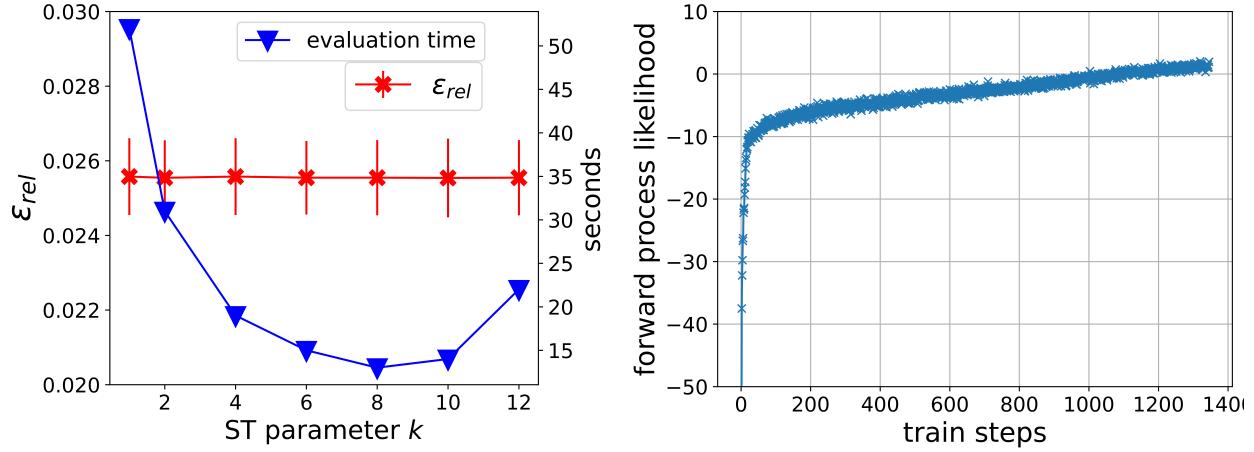


Figure 3. Left: Evaluation of the solution quality of CE-ST over k on the RB-small MIS dataset. The plot shows that the relative error does not change when k is increased, whereas the evaluation time decreases up to $k = 8$. As some operations scale exponentially with k we observe that the evaluation time increases at $k > 8$. Right: The unnormalized forward process likelihood $\mathbb{E}_{X_{0:T} \sim g_\theta(X_{0:T})} [\log p(X_{0:T})]$ over the number of training steps on the RB-100 MIS dataset. The model is trained at a constant temperature. The plot shows that the model learns to generate samples that become more likely under the forward diffusion process.

C. Experimental Details

C.1. Metrics

Relative Error The relative error is defined as

$$\hat{\epsilon}_{rel} = \frac{|E_{opt} - \hat{E}_{model}|}{|E_{opt}|},$$

where E_{opt} is the optimal energy and \hat{E}_{model} is the average energy of the model.

Similarly, ϵ_{rel}^* is the best relative error, where the best out of eight samples E_{model}^* instead of the average model energy \hat{E}_{model} is used.

Approximation Rate The best approximation rate is defined as

$$AR^* = \frac{E_{model}^*}{E_{opt}},$$

where E_{opt} is the optimal energy and E_{model}^* is the energy of the best out of eight samples from the model.

For approximation rate and relative error, lower values are better. Our experiments are always averaged over three independent seeds.

C.2. Evaluation diffusion steps

In some experiments (see Sec. 6.1 and see Sec. 6.2) we evaluate DiffUCO on the test dataset by applying more diffusion steps than we have used during training.

We do this in the following way:

When the diffusion model is trained with T diffusion steps, at diffusion step t the diffusion model is additionally conditioned on a one-hot encoding of this time step. Therefore, if we instead want to run DiffUCO with $n \times T$ diffusion steps, we repeat each time step n times. So during evaluation, at each time step $t \in \{1, \dots, n \times T\}$ a X_t is sampled from the diffusion model and reprocessed at the next time step. At each time step the model is time conditioned on the value $\lfloor \frac{t+1}{n} \rfloor$. Here, $\lfloor \cdot \rfloor$ is an integer operation.

C.3. Datasets

RB-100 MIS dataset The RB-100 MIS dataset is generated by specifying generation parameters n , k' , and p . With n the number of cliques, i.e. a set of fully connected nodes, and with k' the number of nodes within the clique are specified. p serves as a parameter that regulates the interconnectedness between cliques. The lower the value of p the more connections are randomly drawn between cliques. If $p = 1$ there are no connections between the cliques at all. To generate the RB-100 dataset with graphs of an average node size of 100, we generate graphs with $n \in \{9, 15\}$, $k' \in \{8, 11\}$, and $p \in \{0.25, 1\}$.

RB-200 MVC dataset On the RB-200 dataset $k \in \{9, 10\}$ and $n \in \{20, 25\}$, and $p \in \{0.25, 1\}$. We use 2000 graphs for training, 500 for validation, and for each p we use 100 graphs for testing.

RB-small and RB-large On the RB-small dataset $k \in \{5, 12\}$ and $n \in \{20, 25\}$ and graphs that are smaller than 200 nodes or larger than 300 nodes are resampled. On BA-large $k \in \{20, 25\}$ and $n \in \{40, 55\}$ and graphs that are smaller than 800 nodes or larger than 1200 nodes are resampled. For both of these datasets $p \in \{0.3, 1\}$.

C.4. Architecture

We use a simple GNN Architecture, where in the first step input each node feature is transformed by a linear layer with n_h neurons. These node embeddings are then multiplied with a weight matrix with n_h neurons followed by a sum aggregation over the neighborhood. Additionally, a skip connection on each node is added. Afterward, a NodeMLP processes the aggregated nodes together with the skip connection. After n message passing steps each node embedding is fed into a final three-layer MLP which computes the probabilities of each solution variable X_i . Layernorm (Ba et al., 2016) is applied after every MLP layer, except for the last layer within the final MLP. We always use $n_h = 64$ in all of our experiments.

C.5. Hyperparameters

A table with all hyperparameters on each dataset is given in Tab. 7. M_ω is the batch size of different CO problem instances and M_{KL} is the batch size with which the Joint Variational Upper Bound is estimated. RAdam (Liu et al., 2020) is used as an optimizer. We clip the gradients to a norm of 1.0.

In experiments on the RB dataset, except for MaxCl, we additionally use 5 random node features to increase the expressivity of the GNN in critical cases. This is necessary to improve the performance on the RB dataset at low p values.

C.6. Computational Requirements

Experiments on RB graphs are quite memory expensive, because of the high connectivity of these graphs. Especially experiments on MaxCl are expensive because the energy aggregation on the complementary graph leads to even higher connectivity and therefore to a large computational graph. On RB-large MIS experiments are for example conducted on one A100 GPU with 80 GB. Experiments on BA graphs have much fewer computational requirements because the connectivity of these graphs is quite low. Comparing the training time of our method to baselines taken from Zhang et al. (2023) is difficult as the authors do not report the training time nor the computational resources of their method or the other methods used in their paper. They also do not provide a list of hyperparameters such as the number of epochs for any method, which makes the reproducibility of their results difficult. We hypothesize that LTFT might need shorter training and fewer computational resources on problems like MIS and MaxCl because, for these CO problems, the authors design problem-specific Markov Decision Processes that are very efficient for these kinds of problems. However, this makes their

Dataset	lr	T_{start}	GNN layers	diffusion steps	N_{anneal}	noise distr.	M_ω	M_{KL}
RB-large MIS	0.002	0.3	7	4	3000	annealed noise distr.	18	5
RB-small MIS	0.002	0.4	8	6	4000	annealed noise distr.	30	10
RB-small MaxCl	0.002	0.5	8	5	3000	cat. noise distr.	20	8
BA-large MaxCut	0.002	0.2	4	6	1000	annealed noise distr.	20	10
BA-small MaxCut	0.002	0.2	8	4	2000	annealed noise distr.	20	10
BA-large MDS	0.003	0.3	8	3	2000	annealed noise distr.	20	10
BA-small MDS	0.003	0.3	8	5	2000	annealed noise distr.	20	10
RB-200 MVC	0.001	0.4	8	4	4500	annealed noise distr.	30	10

Table 7. Table with all hyperparameters on each dataset.

approach in contrast to our approach less general because these tricks for example cannot be applied to MDS and MaxCut. Therefore, we expect LTFT to have long training times on MaxCut and MDS. This assumption is supported by the long evaluation times of LTFT on these CO problems as reported in Tab. 1 and Tab. 3. On MaxCut their evaluation time is more than 21 times longer and on MDS the evaluation time is more than 32 times longer compared to our method. If we compare the training time of our method to VAG-CO on the RB-200 dataset, we can state that our method takes 2d 2h for training and VAG-CO takes 3d 7h for training. Other baselines like EGN and EGN-Anneal converge quite fast but do not achieve a competitive solution quality due to a lack of expressivity (Sanokowski et al., 2023). In EGN-Anneal the training duration highly depends as in all annealing-based methods on the annealing schedule. The more annealing steps are used the longer the training time. However, here the solution quality with longer annealing also saturates due to a lack of expressivity.

C.7. Time Measurement

All time measurements were conducted on an A100 NVIDIA GPU. We only include the time of the forward pass and the time needed to perform CE in the time measurement. Therefore, we do not include the time that is necessary to load the graphs from the dataloader. For the forward pass, we measure the time on jitted functions. Different states from the diffusion model can in principle be computed in a parallelized manner. Therefore, we measure the time sequentially on each state and then compute the average of this measured time. Self conducted Gurobi evaluations on MaxCut are run on a Intel Xeon Platinum 8168 @ 2.70GHz CPU with 24 cores.

C.8. Code

The code for this research project is based on jax (Bradbury et al., 2018).