
Posterior Inference in Latent Space for Scalable Constrained Black-box Optimization

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Abstract

1 Optimizing high-dimensional black-box functions under black-box constraints is
2 a pervasive task in a wide range of scientific and engineering problems. These
3 problems are typically harder than unconstrained problems due to hard-to-find
4 feasible regions. While Bayesian optimization (BO) methods have been developed
5 to solve such problems, they often struggle with the curse of dimensionality. Re-
6 cently, generative model-based approaches have emerged as a promising alternative
7 for constrained optimization. However, they suffer from poor scalability and are
8 vulnerable to mode collapse, particularly when the target distribution is highly
9 multi-modal. In this paper, we propose a new framework to overcome these chal-
10 lenges. Our method iterates through two stages. First, we train flow-based models
11 to capture the data distribution and surrogate models that predict both function
12 values and constraint violations with uncertainty quantification. Second, we cast
13 the candidate selection problem as a posterior inference problem to effectively
14 search for promising candidates that have high objective values while not violating
15 the constraints. During posterior inference, we find that the posterior distribution
16 is highly multi-modal and has a large plateau due to constraints, especially when
17 constraint feedback is given as binary indicators of feasibility. To mitigate this
18 issue, we amortize the sampling from the posterior distribution in the latent space
19 of flow-based models, which is much smoother than that in the data space. We
20 empirically demonstrate that our method achieves superior performance on various
21 synthetic and real-world constrained black-box optimization tasks. Our code is
22 publicly available here.

23 1 Introduction

24 Optimizing high-dimensional black-box functions under black-box constraints is a fundamental task
25 across numerous scientific and engineering problems, including machine learning [1], drug discovery
26 [2, 3], control [4, 5], and industrial design [6, 7]. In most cases, these problems are much harder than
27 unconstrained problems due to analytically undefined and hard-to-find feasible regions [8].

28 Bayesian Optimization (BO) has been widely used to solve black-box optimization problems in
29 a sample-efficient manner [9, 10]. While most BO methods focus on unconstrained optimization
30 problems, some works address problems with black-box constraints by developing new acquisition
31 functions [1, 11] or relaxing the constraints [12, 13]. However, even without constraints, BO methods
32 scale poorly to high dimensionality [14]. Moreover, incorporating constraints makes the function
33 landscape highly complex, hindering accurate estimation of surrogate models.

34 Recently, generative models have emerged as an alternative solution for black-box optimization
35 problems with constraints [15, 16, 17]. For example, we can leverage generative models to sample
36 protein sequences that maximize the binding affinity while preserving the naturalness of the design.

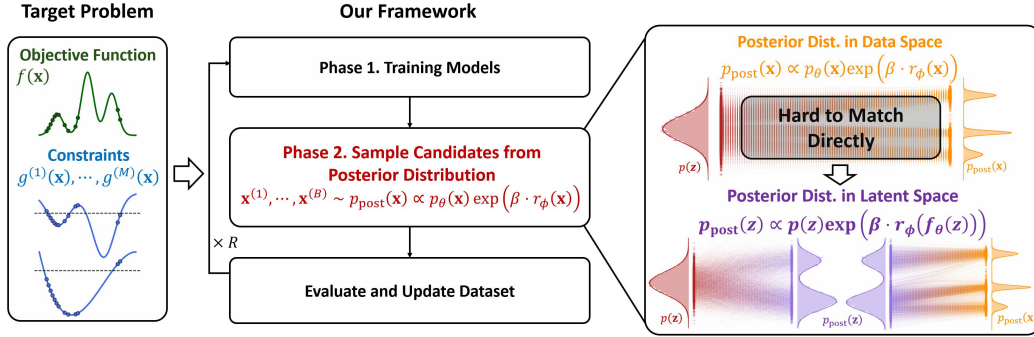


Figure 1: Motivating figure. In a high-dimensional setting, sampling from the posterior distribution is beneficial for selecting candidates. However, the posterior distribution is highly multi-modal and has a large plateau due to the constraints (orange one). We can mitigate this issue by sampling latents from the posterior distribution (purple one) of the latent space and projecting them into the data space.

37 However, existing methods rely on MCMC-based approaches [15], which limit scalability in high-
 38 dimensional spaces. While one can fine-tune pretrained generative models with reward functions
 39 [18, 19, 20], naive application of fine-tuning methods is vulnerable to mode collapse when the target
 40 distribution is highly multi-modal [21], which leads to a convergence on sub-optimal solutions.

41 In this paper, we propose a novel generative model-based framework for constrained black-box
 42 optimization to overcome the aforementioned limitations. To efficiently explore high-dimensional
 43 spaces, we first frame the candidate selection problem as sampling from the posterior distribution,
 44 which can be constructed by multiplying the prior distribution with a Lagrangian-relaxed objective.
 45 To sample candidates from the posterior distribution, our key idea is to amortize inference in the latent
 46 space of a flow-based model using an outsourced diffusion sampler [21], as illustrated in Figure 1.
 47 Since the posterior distribution in the latent space is much smoother than that in the data space, we
 48 can approximate the distribution more accurately and alleviate the mode collapse problem [22].

49 Our method iterates through two stages. First, we train a flow-based model to capture the current
 50 data distribution and surrogate models to predict the objective value and constraints, respectively.
 51 For the surrogate models, we use an ensemble to quantify the uncertainty of the prediction, as we
 52 have only a small amount of data that covers a tiny fraction of the whole search space. We treat a
 53 trained flow-based model as a prior, and Lagrangian relaxation of the objective as a reward function.
 54 Second, we sample candidates from the posterior distribution. As the posterior distribution is highly
 55 multi-modal and has a large plateau due to constraints, especially when constraint feedback is given
 56 as binary indicators of feasibility, we train a diffusion sampler that amortizes the posterior distribution
 57 in the latent space of flow models. Then, we sample latents from the diffusion sampler and project
 58 them into data space using a deterministic mapping derived from the trained flow model. By repeating
 59 these two stages, we can effectively capture high-scoring regions that satisfy the constraints.

60 We conduct extensive experiments on three synthetic and three real-world benchmarks to validate the
 61 superiority of our method on scalable constrained black-box optimization problems. We also consider
 62 a more challenging scenario where the feedback from the constraints is given as a binary value. We
 63 empirically show that our method outperforms several competitive baselines across different tasks.

64 2 Related Works

65 2.1 Constrained Black-box Optimization

66 Most scientific and engineering optimization problems involve black-box constraints, such as the
 67 synthesizability of molecules in chemical design [2] and safety constraints in robot control policies
 68 [4]. Existing BO methods solve this problem by either integrating the constraints directly into the
 69 acquisition function (cEI [23], LogcEI [24]) or by employing trust region approaches for scalability
 70 (SCBO [8], PCAGP-SCBO [7]). Another line of work utilizes evolutionary algorithms like CMA-
 71 ES [25, 26] with an augmented Lagrangian method to navigate constrained spaces. However, the
 72 performance of these methods often degrades as dimensionality and the number of evaluations
 73 increase, which motivates the need for a more scalable approach.

74 2.2 Generative Model-based Optimization

75 There are several attempts to utilize generative models for black-box optimization. In an offline
 76 setting, DDOM [27] trains a conditional diffusion model with classifier-free guidance and applies a
 77 loss-reweighting to emphasize samples with high objective values. DiffOPT [15] solves a constrained
 78 optimization problem. It applies diffusion to capture data distribution, followed by an iterative
 79 importance-sampling procedure. In an online setting, DiffBBO [28] and DiBO [29] both leverage
 80 diffusion models and incorporate uncertainty estimation during candidate selection. DiBO treats
 81 candidate selection as posterior inference to guide sampling toward regions of high reward and uncer-
 82 tainty, while DiffBBO selects conditioning targets by employing an uncertainty-based acquisition
 83 function. Unfortunately, constrained black-box optimization in the online setting remains unexplored.

84 2.3 Amortized Inference in Flow-based and Diffusion Models

85 Given a diffusion or flow prior $p_\theta(\mathbf{x})$ trained on a dataset and a reward function $r(\mathbf{x})$, sampling from
 86 the posterior $p_{\text{post}}(\mathbf{x}) \propto p_\theta(\mathbf{x})r(\mathbf{x})$ has numerous applications in downstream tasks [30, 18, 31, 32,
 87 19, 21]. However, direct sampling from the unnormalized posterior $p_\theta(\mathbf{x})r(\mathbf{x})$ is intractable [18, 33].

88 To address this problem, some approaches train classifiers directly within intermediate noised spaces
 89 [30, 34] while others approximate posterior sampling via Markov Chain Monte Carlo (MCMC)
 90 procedures [22, 32, 35, 36]. However, training classifiers in noisy data spaces and employing MCMC
 91 methods scale poorly to high dimensionality. On the other hand, several methods utilize reinforcement
 92 learning [37, 38] or stochastic optimal control [19] to fine-tune the pretrained model and amortize the
 93 posterior sampling. Meanwhile, naive implementations of fine-tuning methods can be prone to mode
 94 collapse when the target distribution is highly multi-modal and has a large plateau region [21].

95 To mitigate this issue, we adopt the outsourced diffusion sampler method proposed by Venkatraman
 96 et al. [21]. Matching the distribution within the latent space significantly simplifies the alignment
 97 task when the distribution is highly multi-modal and has a large flat region in the original data space.

98 3 Preliminaries

99 3.1 Constrained Black-box Optimization

100 In constrained black-box optimization, our problem is:

$$\begin{aligned} \text{find } \mathbf{x}^* = \arg \max_{\mathbf{x} \in \mathcal{X}} f(\mathbf{x}) \quad \text{s.t. } g^{(1)}(\mathbf{x}) \leq 0, \dots, g^{(M)}(\mathbf{x}) \leq 0 \\ \text{with } R \text{ rounds of } B \text{ batch of queries} \end{aligned} \quad (1)$$

101 The objective function $f : \mathcal{X} \rightarrow \mathbb{R}$ and constraints $g^{(1)}, \dots, g^{(M)} : \mathcal{X} \rightarrow \mathbb{R}$ are black-box functions.
 102 We also consider a more challenging scenario, only access to information on whether we violate
 103 constraints or not, i.e., $h^{(m)}(\mathbf{x}) = \mathbb{I}[g^{(m)}(\mathbf{x}) > 0]$. We refer to this as an indicator constraint.

104 3.2 Flow-based Models

105 Flow-based models [39, 40, 41] are a class of generative models for approximating a target distribution
 106 $q(\mathbf{x})$. Flow-based models are defined via the deterministic ordinary differential equation (ODE):

$$d\mathbf{x}_t = v_\theta(\mathbf{x}_t, t) dt \quad (2)$$

107 where $v_\theta(\mathbf{x}_t, t) : \mathbb{R}^d \times [0, 1] \rightarrow \mathbb{R}^d$ is a parametric velocity field.

108 For each given velocity field, the corresponding flow $\psi_\theta(\mathbf{x}_0, t) : \mathbb{R}^d \times [0, 1] \rightarrow \mathbb{R}^d$ satisfies:

$$\frac{d}{dt} \psi_\theta(\mathbf{x}_0, t) = v_\theta(\psi_\theta(\mathbf{x}_0, t), t), \quad \psi_\theta(\mathbf{x}_0, 0) = \mathbf{x}_0. \quad (3)$$

109 The velocity field $v_\theta(\mathbf{x}_t, t)$ defines a continuous probability path p_t induced by the flow:

$$\mathbf{x}_t = \psi_\theta(\mathbf{x}_0, t) \sim p_t, \quad \text{where } \mathbf{x}_0 \sim p_0. \quad (4)$$

110 **Training Flow-based Models.** We use Flow Matching [39] to learn the velocity field v_θ that generates
 111 a path interpolating smoothly between an initial distribution $p_0 = p$ and a target distribution $p_1 = q$.

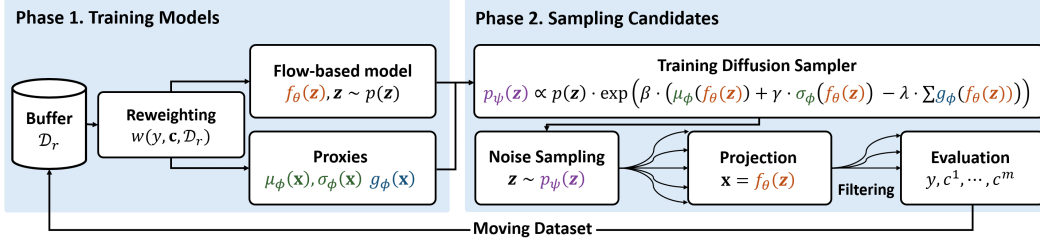


Figure 2: Overview of our method. **Phase 1:** Train flow-based models and proxies for the objective and constraints. **Phase 2:** Sample candidates from the posterior distribution using an outsourced diffusion sampler. After sampling, we utilize filtering to enhance sample efficiency. Then, we evaluate samples, update the dataset, and repeat the process until the evaluation budget is exhausted.

We employ the simplest linear interpolation path $\mathbf{x}_t = (1-t)\mathbf{x}_0 + t\mathbf{x}_1$, with derivative $\frac{d\mathbf{x}_t}{dt} = \mathbf{x}_1 - \mathbf{x}_0$, following [39]. The Flow Matching loss is expressed as:

$$\mathcal{L}_{\text{FM}}(\theta) = \mathbb{E}_{\mathbf{x}_0 \sim \mathcal{N}(0, I), \mathbf{x}_1 \sim q(\mathbf{x}), t \sim \text{Unif}(0, 1)} [\|v_\theta(\mathbf{x}_t, t) - (\mathbf{x}_1 - \mathbf{x}_0)\|^2]. \quad (5)$$

3.3 Posterior Inference in Flow-based and Diffusion Models

Given a pretrained flow-based prior $p_\theta(\mathbf{x})$, and a reward function $r(\mathbf{x})$, we consistently encounter a situation where we need to sample from the posterior distribution, $p_{\text{post}}(\mathbf{x}) \propto p_\theta(\mathbf{x})r(\mathbf{x})$. To sample from this intractable [33] distribution, we utilize the outsourced diffusion sampling [21].

We can interpret the sampling process of flow-based models into a noise generation $\mathbf{z} \sim p(\mathbf{z})$, followed by a deterministic transformation $\mathbf{x} = f_\theta(\mathbf{z})$, where $p(\mathbf{z})$ is standard normal and f_θ represents the learned mapping derived by prior. Under this formulation, by Proposition 3.1 of [21], we can sample from the posterior distribution by substituting noise generation as $\mathbf{z} \sim p_{\text{post}}(\mathbf{z}) \propto p(\mathbf{z})r(f_\theta(\mathbf{z}))$.

To approximate the target distribution $p_\psi(\mathbf{z}) \approx p_{\text{post}}(\mathbf{z})$, we can learn the parameters of diffusion sampler ψ with the trajectory balance (TB) objective [42, 43]:

$$\mathcal{L}_{\text{TB}}(\mathbf{z}_{0:1}; \psi) = \left(\log \frac{Z_\psi p(\mathbf{z}_0) \prod_{i=0}^{T-1} p_F(\mathbf{z}_{(i+1)\Delta t} | \mathbf{z}_{i\Delta t}; \psi)}{p(\mathbf{z}_1)r(f_\theta(\mathbf{z}_1)) \prod_{i=1}^T p_B(\mathbf{z}_{(i-1)\Delta t} | \mathbf{z}_{i\Delta t})} \right)^2, \quad (6)$$

where Z_ψ is the parameterized partition estimator, $(\mathbf{z}_0 \rightarrow \mathbf{z}_{\Delta t} \rightarrow \dots \mathbf{z}_1 = \mathbf{z})$ is the discrete time Markov chain of reverse-time stochastic differential equation (SDE) [44] with time increment $\Delta t = \frac{1}{T}$. p_F and p_B are transition kernels of the discretized reverse and forward SDE.

4 Method

In this section, we introduce **CiBO**, a new framework for scalable constrained black-box optimization by leveraging generative models. Our method consists of two iterative stages. First, we train a flow-based model to capture the data distribution and surrogate models to predict objective values and constraints with uncertainty quantification. Next, we sample candidates from the posterior distribution. To accomplish this, we train a diffusion sampler that draws samples from the posterior distribution in the latent space. After sampling, we evaluate candidates, update the dataset, and repeat the process until the evaluation budget is exhausted. Figure 2 illustrates the overview of our method.

4.1 Phase 1. Training Models

In each round r , we have a pre-collected dataset $\mathcal{D}_r = \{\mathbf{x}_i, y_i, \mathbf{c}_i\}_{i=1}^I$, where $y_i = f(\mathbf{x}_i)$, $\mathbf{c}_i = \{c_i^m | c_i^m = g^{(m)}(\mathbf{x}_i), \forall m = 1, \dots, M\}$, and I is the number of data points collected so far.

Training Prior. We first train a prior model p_θ to capture the current data distribution. As the search space is too high-dimensional, it is better to implicitly constrain the search space close to the current data distribution. We use flow-based models to learn this distribution using Equation (5).

Training Surrogates. We also train surrogate models to predict both objective values and constraints. As we are only able to access a small number of data points in the vast search space, we need to

properly quantify the uncertainty of the prediction. To this end, we train an ensemble of proxies to estimate objective values with uncertainty quantification [45]. Specifically, we train an ensemble of K proxies $f_{\phi_1}, \dots, f_{\phi_K}$ for objective values, and individual proxy $g_{\phi}^{(1)}, \dots, g_{\phi}^{(M)}$ for each constraint.

Rewighted Training. During training, we introduce a reweighted training scheme [27, 46, 47] to focus on promising data points with high objective values while not violating constraints. Specifically, the weight for each data point is computed as follows:

$$l(y, \mathbf{c}) = y - \lambda \sum_{m=1}^M \max(0, c^m), \quad w(y, \mathbf{c}, \mathcal{D}_r) = \frac{\exp(l(y, \mathbf{c}))}{\sum_{(y', \mathbf{c}') \in \mathcal{D}_r} \exp(l(y', \mathbf{c}'))}. \quad (7)$$

Then, our training objective for flow-based models and proxies can be described as follows:

$$\mathcal{L}(\theta) = \mathbb{E}_{\mathbf{x}_0 \sim \mathcal{N}(0, I), (\mathbf{x}, y, \mathbf{c}) \in \mathcal{D}_r, t \sim \text{Unif}(0, 1)} [w(y, \mathbf{c}, \mathcal{D}_r) \|v_{\theta}(\mathbf{x}_t, t) - (\mathbf{x} - \mathbf{x}_0)\|_2^2], \quad (8)$$

$$\mathcal{L}(\phi) = \sum_{(\mathbf{x}, y, \mathbf{c}) \in \mathcal{D}_r} w(y, \mathbf{c}, \mathcal{D}_r) \left[\sum_{k=1}^K (y - f_{\phi_k}(\mathbf{x}))^2 + \sum_{m=1}^M \left(c^m - g_{\phi}^{(m)}(\mathbf{x}) \right)^2 \right]. \quad (9)$$

4.2 Phase 2. Sampling Candidates

After training models, we proceed to select candidates for evaluation in the current round. As the search space is high-dimensional, the prediction of surrogate models is likely to be inaccurate in regions that are too far away from the dataset collected so far. Therefore, it is advantageous to sample candidates from the distribution that satisfies the two desiderata: (1) promote exploration towards high-scoring and feasible regions, and (2) prevent sampling candidates that deviate too far from the current data distribution. To accomplish these objectives, we cast the candidate selection problem as sampling from the target distribution p_{post} defined as follows:

$$p_{\text{post}}(\mathbf{x}) = \arg \max_{p \in \mathcal{P}} \mathbb{E}_{\mathbf{x} \sim p} [r_{\phi}(\mathbf{x})] - \frac{1}{\beta} \cdot D_{\text{KL}}(p \| p_{\theta}), \quad (10)$$

where \mathcal{P} is the space of all probability distributions over the domain \mathcal{X} , and

$$r_{\phi}(\mathbf{x}) = \mu_{\phi}(\mathbf{x}) + \gamma \cdot \sigma_{\phi}(\mathbf{x}) - \lambda \sum_{m=1}^M \max(0, g_{\phi}^{(m)}(\mathbf{x})). \quad (11)$$

$\mu_{\phi}(\mathbf{x})$ and $\sigma_{\phi}(\mathbf{x})$ represent the mean and standard deviation from the ensemble of surrogate models for the objective. γ controls exploration-exploitation trade-off, β is an inverse temperature, and λ is a Lagrange multiplier. Based on derivation from [48], our target distribution analytically derived as:

$$p_{\text{post}}(\mathbf{x}) \propto p_{\theta}(\mathbf{x}) \exp(\beta \cdot [r_{\phi}(\mathbf{x})]). \quad (12)$$

If we treat the flow-based model $p_{\theta}(\mathbf{x})$ as a prior and the exponential term $\exp(\beta \cdot [r_{\phi}(\mathbf{x})])$ as a reward $r(\mathbf{x})$, then our objective is to sample from the posterior distribution $p_{\text{post}}(\mathbf{x}) \propto p_{\theta}(\mathbf{x})r(\mathbf{x})$.

Amortized Inference in Latent Space. However, directly sampling from this posterior is intractable [33]. Also, the target posterior is highly multi-modal and has a large plateau due to the constraint penalties in (11), making finetuning-based methods [18, 37] susceptible to mode collapse. To this end, we utilize an amortized sampler in the latent space suggested by Venkatraman et al [21].

As introduced in Section 3.3, we can view the sampling procedure of flow-based models as drawing samples from the standard normal distribution $\mathbf{z} \sim p(\mathbf{z})$, followed by the deterministic transformation $\mathbf{x} = f_{\theta}(\mathbf{z})$. Within this framework, we can generate samples from the posterior distribution $p_{\text{post}}(\mathbf{x})$ by modifying the noise generation distribution as follows:

$$\mathbf{z} \sim p_{\text{post}}(\mathbf{z}) \propto p(\mathbf{z})r(f_{\theta}(\mathbf{z})). \quad (13)$$

To sample latents \mathbf{z} from the posterior distribution in the latent space $p_{\text{post}}(\mathbf{z})$, we train a diffusion model $p_{\psi}(\mathbf{z})$ to amortize $p_{\text{post}}(\mathbf{z})$ with the following Trajectory Balance (TB) objective:

$$\mathcal{L}_{\text{TB}}(\mathbf{z}_{0:1}; \psi) = \left(\log \frac{Z_{\psi} p(\mathbf{z}_0) \prod_{i=0}^{T-1} p_F(\mathbf{z}_{(i+1)\Delta t} | \mathbf{z}_{i\Delta t}; \psi)}{p(\mathbf{z}_1) r(f_{\theta}(\mathbf{z}_1)) \prod_{i=1}^T p_B(\mathbf{z}_{(i-1)\Delta t} | \mathbf{z}_{i\Delta t})} \right)^2. \quad (14)$$

By training an amortized sampler in the latent space of flow-based models, we can more accurately sample candidates from the target distribution as the posterior distribution in the latent space is smoother than that in the data space. We also adopt off-policy training, detailed in Appendix D.2.1

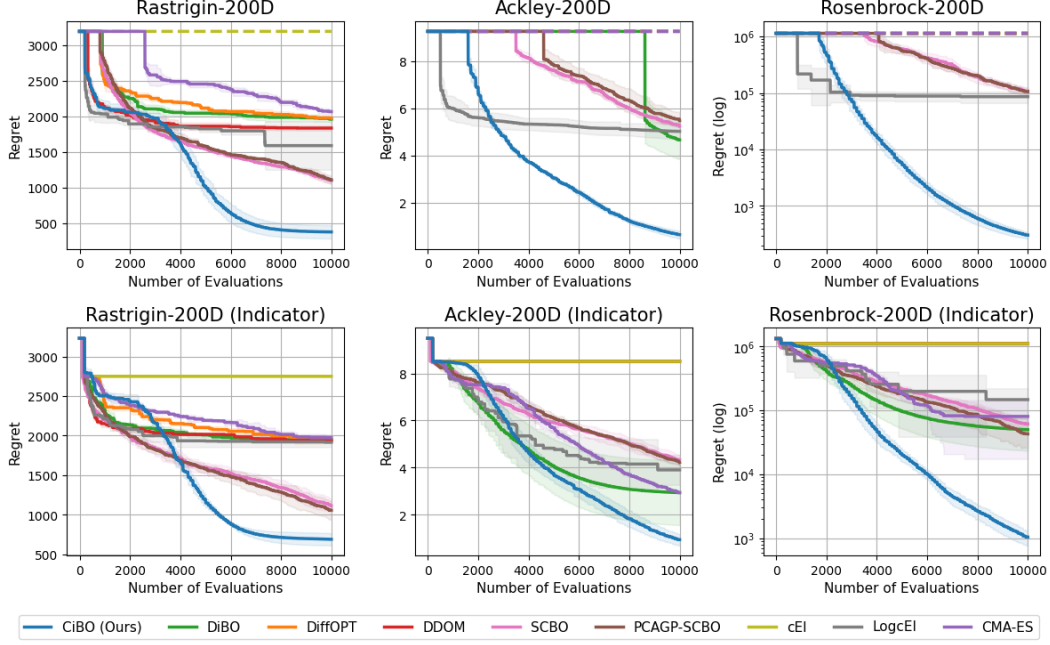


Figure 3: Comparison between our method and baselines in synthetic tasks. Experiments are conducted with four random seeds, and the mean and one standard deviation are reported. A dashed line means that no feasible solutions were found.

4.3 Filtering, Evaluation and Moving Dataset

Filtering. After sampling from the posterior distribution, we need to carefully select candidates for the sample efficiency of the algorithm. To do so, we generate $N \cdot B$ samples from the amortized sampler and select the top- B samples in terms of Lagrangian relaxation of objectives as candidates.

Evaluation and Moving Dataset. We evaluate the values of the objective function and constraint functions for each selected candidate, then update the dataset with new observations. During the update, we empirically find that taking only a subset of total observations is beneficial in terms of computational complexity. We remove the samples with the lowest Lagrangian-relaxed objective if the dataset size is larger than the buffer size L . The pseudocode of our method is in Algorithm 1.

5 Experiments

In this section, we report experimental results for scalable constrained black-box optimization tasks. First, we perform experiments on three 200-dimensional synthetic functions, which are the standard benchmarks in Bayesian Optimization (BO) studies [14]. Furthermore, we assess the performance of our method on a more challenging scenario, where the feedback from constraints is given as binary indicators of feasibility. We refer to this setting as the indicator constraint setting. Finally, we conduct experiments on three real-world optimization tasks: Rover Planning 60D [8, 49], Mopta 124D [6], and Lasso DNA 180D [50]. The detailed description of each task can be found in Appendix A.

For evaluation, we report the minimum regret of feasible solutions over the course of the training, and assign the largest regret found in all algorithms to the infeasible solutions, following [8, 51].

5.1 Baselines

We compare our method with several constrained BO baselines, including cEI [23], LogcEI [24], SCBO [8], PCAGP-SCBO [7], and the evolutionary search algorithm CMA-ES [25]. We also evaluate generative model-based approaches specifically designed for constrained optimization: DiffOPT [15], as well as methods that can be extended to constrained optimization via the Lagrangian relaxation: DDOM [27] and DiBO [29]. Detailed implementations of all baselines are provided in Appendix B.

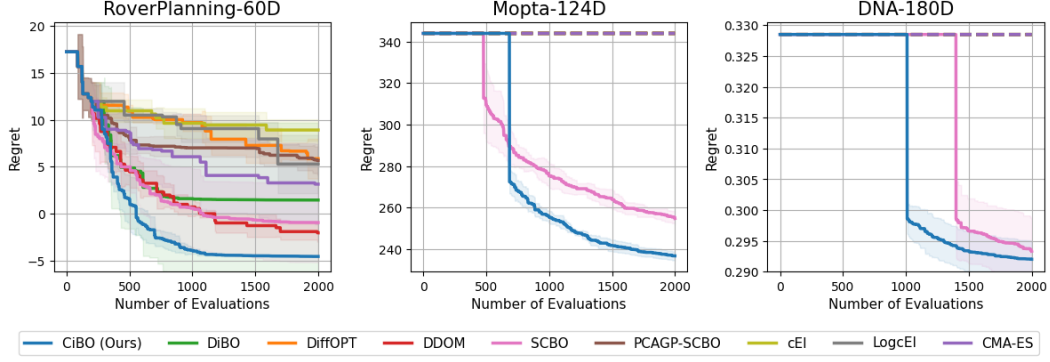


Figure 4: Comparison between our method and baselines in real-world tasks. Experiments are conducted with four random seeds, and the mean and one standard deviation are reported. A dashed line means that no feasible solutions were found.

5.2 Synthetic Experiments

We first conduct experiments on three synthetic functions, Rastrigin-200D, Ackley-200D, and Rosenbrock-200D. For each function, we utilize two inequality constraints proposed by SCBO [8]: $\sum_{d=1}^{200} x_d \leq 0$ and $\|\mathbf{x}\|_2^2 \leq 30$. We conduct all experiments with an initial dataset size of $|\mathcal{D}_0| = 200$, using a batch size of $B = 100$ and a maximum evaluation limit of 10,000. In the indicator constraints scenarios, as it is too challenging to find an initial feasible solution across all baselines, we sample 10 points within feasible regions during initialization.

As shown in the Figure 3, our method outperforms all baselines across different synthetic tasks, both in the standard and indicator constraints. Generative model-based methods, including DiffOPT and DDOM, struggle to find a feasible solution and fail to improve on indicator constraints. While DiBO achieves better feasibility, its finetuning-based approach suffers from mode collapse and tends to converge to suboptimal solutions. These results show that employing an outsourced diffusion sampler significantly enhances performance in constrained black-box optimization by effectively capturing multi-modal and expansive flat target distributions.

Constrained BO methods (SCBO, PCAGP-SCBO, and LogcEI) successfully identify feasible points but show limited sample efficiency compared to our method across all tasks. The evolutionary search algorithm CMA-ES performs modestly in general but fails to find a feasible solution for some tasks. These results underscore that our approach effectively captures both high-scoring and feasible regions in a sample-efficient manner. Furthermore, compared to other baselines, our method consistently finds feasible solutions throughout the optimization process, which is illustrated in Appendix E.1.

5.3 Real World Experiments

To validate the robustness of our approach, we evaluate our method on three challenging real-world benchmark problems: (1) Rover Planning in 60 dimensions with 15 infeasible square-shaped regions, (2) Mopta in 124 dimensions with 68 constraints, and (3) Lasso DNA in 180 dimensions with 5 constraints. For all experiments, we initialize with $|\mathcal{D}_0| = 200$ data points and limit evaluations to 2,000. We use a batch size of $B = 50$ for Rover Planning and Lasso DNA, and $B = 20$ for Mopta, as no baseline methods could identify feasible solutions with $B = 50$.

As illustrated in Figure 4, our approach consistently identifies high-quality feasible solutions with superior sample efficiency across all tasks. We observe that the performance gap between our method and other baselines becomes larger on real-world problems and most baselines failed to find any feasible solutions for the challenging Mopta-124D and DNA-180D tasks. While SCBO is the only competing method to achieve feasibility alongside our approach, it exhibits lower sample efficiency. This highlights the robustness of our approach for scalable constrained black-box optimization.

5.4 Additional Analysis

In this section, we conduct a comprehensive analysis of each component of our proposed method through ablation experiments on Rastrigin 200D and Rover Planning 60D.

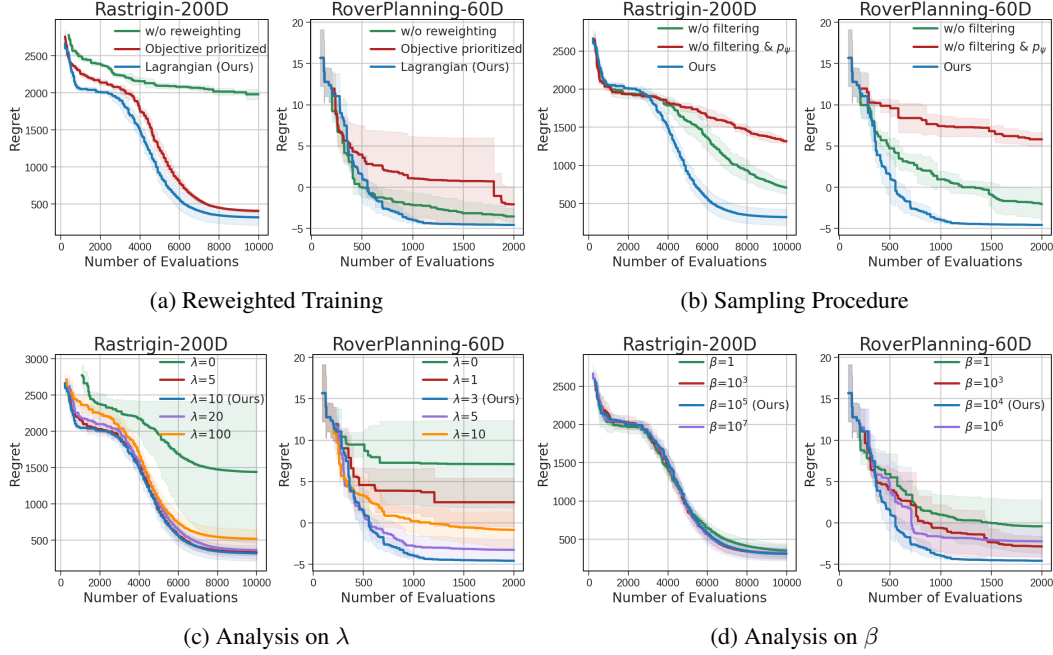


Figure 5: Additional analysis for various components of CiBO. Experiments are conducted with four random seeds, and the mean and one standard deviation are reported.

Rewighted Training. To investigate the effectiveness of our reweighted training approach suggested in Equation (7), we conduct a comparative analysis of two variants: training without reweighting, applying weights based on the objective values (Objective-prioritized). As shown in Figure 5a, variants without reweighting or using objective-prioritized reweighting exhibit low sample efficiency.

Sampling Procedure. We analyze the effect of each component in candidate sampling. We conduct experiments with two variants: removing filtering, and removing both filtering and the diffusion sampler, thus sampling candidates directly from the prior p_θ . As depicted in Figure 5b, there is a significant performance gap between our method and other variants, validating the effectiveness of each proposed component. We also experiment with the filtering coefficient N in Appendix E.2.

Lagrangian Multiplier λ . We introduce the Lagrangian multiplier λ . As shown in Figure 5c, setting $\lambda = 0$ (eliminating the constraint penalty) significantly degrades performance on both tasks, as it only focuses on high objective values and neglects the feasibility of solutions. Conversely, excessively high λ values diminish the influence of the objective function, resulting in reduced sample efficiency.

Inverse Temperature β . The inverse temperature controls the balance between the prior $p_\theta(\mathbf{x})$ and the reward function $r(\mathbf{x})$. We conduct experiments by varying β values. As shown in Figure 5d, using a moderately high β generally helps to improve sample efficiency. However, if β is too high, the performance is heavily dependent on the accuracy of surrogate models, leading to slow convergence. This validates that incorporating prior distribution is crucial for scalability (Section 4.2).

Further Analysis. To further understand our method, we analyze the impact of the buffer size L (Appendix E.3), batch size B , and initial dataset size $|\mathcal{D}_0|$ (Appendix E.5). We also investigate the effect of off-policy training (Appendix E.4) and runtime scalability of our method (Appendix F).

6 Conclusion

We introduced CiBO, a generative model-based framework for scalable constrained black-box optimization. Our approach formulates candidate selection as posterior inference, leveraging flow-based models to capture the data distribution and surrogate models to predict both objectives and constraints. By amortizing posterior sampling in the latent space with outsourced diffusion samplers, our method effectively addresses the challenges posed by highly multi-modal and flat posterior distributions that arise from incorporating constraints. Extensive experiments across synthetic and real-world benchmarks demonstrate the superiority of our proposed method.

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484 Appendix

485 A Task Details

486 A.1 Synthetic Functions

We evaluate three synthetic functions in our constrained black-box optimization experiments: Rastrigin, Ackley, and Rosenbrock. The Rastrigin and Ackley functions are highly multi-modal functions with numerous local minima, whereas the Rosenbrock function features a narrow valley that makes convergence to the global minimum notoriously difficult [52]. Following [53, 54], we define the search domains as Rastrigin: $[-5, 5]^D$, Ackley: $[-5, 10]^D$, and Rosenbrock: $[-5, 10]^D$. All functions are subject to two constraints:

$$\sum_{d=1}^{200} x_d \leq 0 \quad \text{and} \quad \|\mathbf{x}\|_2^2 \leq 30$$

487 Although prior work enforced the tighter bound $\|\mathbf{x}\|_2^2 \leq 5$, we relax this constraint in our high-
 488 dimensional setting. For the indicator constraint experiments, we sample initial feasible points by
 489 hit-and-run MCMC [55].

490 A.2 Rover Trajectory Planning

Rover Trajectory Planning is a trajectory optimization task in a 2D environment introduced by [49]. The objective is to optimize the rover’s trajectory, where its trajectory is represented by 30 points defining a B-Spline. We place 15 impassable obstacles o_i and impose collision-avoidance constraints $c_i(\mathbf{x})$ as in [8]:

$$c_i(\mathbf{x}) = \begin{cases} -d(o_i, \gamma(\mathbf{x})) & \text{if } \gamma(\mathbf{x}) \cap o_i = \emptyset, \\ \max_{\alpha \in \gamma(\mathbf{x}) \cap o_i} \min_{\beta \in \partial o_i} d(\alpha, \beta) & \text{otherwise.} \end{cases}$$

491 where $\gamma(\mathbf{x})$ denotes final trajectory, o_i is the region of the obstacle and ∂o_i denotes the boundary of o_i .
 492 A trajectory is feasible if and only if it does not intersect any obstacle. We follow the implementation
 493 from [49], but since there is no released code for the constraints, we implement the violation metric
 494 ourselves. Below is an example of the trajectory found by our method.

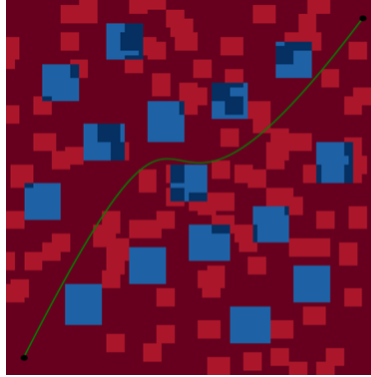


Figure 6: Trajectory found by CiBO, achieving regret of -4.59.

A.3 Vehicle Design with 68 Constraints (MOPTA)

MOPTA is the high-dimensional real-world problem of large-scale multidisciplinary mass optimization [6]. The objective is to minimize a vehicle’s mass, which incorporates decisions about materials, gauges, and vehicle shape with 68 performance constraints. The best-known optimum mass is approximately 222.74. We followed the implementation from [56].¹

A.4 LassoBench

LassoBench [50]² is a high-dimensional benchmark for hyperparameter optimization, specifically designed to tune the hyperparameters of the Weighted LASSO (Least Absolute Shrinkage and Selection Operator) regression model. It includes both synthetic tasks (simple, medium, high, and hard) and real-world tasks (Breast cancer, Diabetes, Leukemia, DNA, and RCV1). In this work, we focus on the DNA task, a microbiology classification problem. It computes the average validation error across all cross-validation folds as an unconstrained objective. We reformulate the problem by retaining the mean validation error as the objective while introducing constraints that the validation error on each fold must not exceed 0.32.

B Baselines Details

In this section, we provide a thorough description of our baseline implementation details and specify the hyperparameter settings used across all experiments.

DiBO [29]: We use the original code³, and adapt DiBO to handle constrained optimization by reformulating the objective as a Lagrangian, setting the same λ value as our methods for fair comparison.

DiffOPT [15]: As there is no publicly available code, we re-implement this baseline on our own. To approximate the data distribution, we use diffusion models with a similar architecture to our method. To enable accurate sampling from the target distribution, we implement Langevin dynamics as the energy function, which can be constructed by surrogate models in our setting, is differentiable.

DDOM [27]: Building on the original implementation⁴, we reconstruct this baseline with network architecture matching our flow-based model. While maintaining the method’s specific parameters as specified in the original work, we incorporate a Lagrangian framework and set the same λ as ours.

SCBO [8]: We follow the tutorial code for SCBO provided by `botorch`⁵ to reproduce the results.

PCAGP-SCBO [7]: To reproduce PCAGP-SCBO, we follow the code for SCBO and then apply `torch_pca`⁶ to project high-dimensional data into a reduced latent space with dimension l before fitting GP surrogates for constraints. For all synthetic tasks, we use $l = 2$ and for real-world tasks, we conduct a hyperparameter search on $[2, \lfloor D/2 \rfloor]$ and report the best one.

cEI [23]: We implement cEI acquisition function by using `qExpectedImprovement()` in `botorch` library. We train a GP surrogate model independently for the objective and each constraint.

LogcEI [24]: We implement logcEI acquisition function by using `qLogExpectedImprovement()` in `botorch` library. We train a GP surrogate model independently for the objective and each constraint.

CMA-ES [25]: We employ the `pycma`⁷ library [57]. For constraint handling, we formulate the problem using the same Lagrangian approach with the same λ value as ours for each task.

¹<https://github.com/LeoIV/BxUS>

²<https://github.com/ksehic/LassoBench>

³<https://github.com/umkiyoung/DiBO>

⁴<https://github.com/siddarthk97/ddom>

⁵https://botorch.org/docs/tutorials/scalable_constrained_bo/

⁶https://github.com/valentingol/torch_pca

⁷<https://github.com/CMA-ES/pycma>

Algorithm 1 CiBO

```

1: Input: Initial dataset  $\mathcal{D}_0$ ; Max rounds  $R$ ; Batch size  $B$ ; Buffer size  $L$ ; Number of constraints  $M$ ;
   Flow model  $p_\theta$ ; Diffusion sampler  $p_\psi$ ; Proxies  $f_{\phi_1}, \dots, f_{\phi_K}, g_{\phi}^{(1)}, \dots, g_{\phi}^{(M)}$ ;
2: for  $r = 0, \dots, R - 1$  do
3:   Initialize  $p_\theta, p_\psi, f_{\phi_1}, \dots, f_{\phi_K}, g_{\phi}^{(1)}, \dots, g_{\phi}^{(M)}$ 
4:
5:   Phase 1. Training Models
6:   Compute weights  $w(y, \mathbf{c}, \mathcal{D}_r)$  with Equation (7)
7:   Train  $p_\theta$  with Equation (8)
8:   Train  $f_{\phi_1}, \dots, f_{\phi_K}, g_{\phi}^{(1)}, \dots, g_{\phi}^{(M)}$  with Equation (9)
9:
10:  Phase 2. Sampling Candidates
11:  Train  $p_\psi$  with Equation (14) using prior  $p_\theta$  and  $\mathbf{z} \sim N(0, \mathbf{I})$ 
12:  Sample latent noise with  $\{\mathbf{z}_i\}_{i=1}^{NB} \sim p_\psi(\mathbf{z})$ 
13:  Projection to data space with learned mapping  $\mathbf{x}_i = f_\theta(\mathbf{z}_i) \quad \forall i \in \{1, \dots, NB\}$ 
14:
15:  Filtering
16:  Select top- $B$  samples  $\{\mathbf{x}_b\}_{b=1}^B$  with respect to:
      $r_\phi(\mathbf{x}_i) - \lambda \sum_{m=1}^M \max(0, g_{\phi}^{(m)}(\mathbf{x}_i)) \quad \forall i = \{1, \dots, NB\}$ 
17:
18:  Evaluation and Moving Dataset
19:  Evaluate  $y_b = f(\mathbf{x}_b), \quad c_b^m = g^{(m)}(\mathbf{x}_b) \quad \forall m = \{1, \dots, M\} \quad \forall b = \{1, \dots, B\}$ 
20:  Update  $\mathcal{D}_{r+1} \leftarrow \mathcal{D}_r \cup \{(\mathbf{x}_b, y_b, \mathbf{c}_b)\}_{b=1}^B$ 
21:  if  $|\mathcal{D}_{r+1}| > L$  then
22:    Remove last  $|\mathcal{D}_{r+1}| - L$  samples from  $\mathcal{D}_{r+1}$  with respect to:  $y - \lambda \sum_{m=1}^M \max(0, c^m)$ 
23:  end if
24: end for

```

D Implementation Details

In this section, we introduce the implementation details of our method **CiBO**. Specifically, model architectures, the training processes employed, the hyperparameter configurations used, and the computational resources required.

D.1 Training Models

D.1.1 Training Proxies

We employ an ensemble of five proxies to model the objective function and a single proxy for each constraint. Each proxy is implemented as a MLP with three hidden layers of 1024 units, using GELU [58] activations. Proxies are trained with the Adam optimizer [59] for 100 epochs per round at a learning rate of 1×10^{-3} and a batch size of 256. All hyperparameters related to the proxy are listed in Table 1.

Table 1: Hyperparameters for Training Proxy

	Parameters	Values
Architecture	Num Ensembles	5
	Number of Layers	3
	Num Units	1024
Training	Batch size	256
	Optimizer	Adam
	Learning Rate	1×10^{-3}
	Training Epochs	100

D.1.2 Training Flow-based Models

We adopt the architecture of [60] for our flow model, comprising three hidden layers with 512 units each. Training is performed using Adam optimizer for 500 epochs per round, with a learning rate of 1×10^{-3} and a batch size of 256. For ODE integration during sampling, we employ the Runge-Kutta 4 method with `torchdiffeq` [61], and set the integration steps as 250. All flow-model hyperparameters are detailed in Table 2.

Table 2: Hyperparameters for Training Flow-based Model

	Parameters	Values
Architecture	Number of Layers	3
	Num Units	512
Training	Batch size	256
	Optimizer	Adam
	Learning Rate	1×10^{-3}
	Training Epochs	500

551 D.2 Sampling Candidates

552 D.2.1 Training Diffusion Sampler

553 Various approaches have been developed to draw samples from a distribution when only an unnormalized probability density or energy function is available. Traditional methods include Markov Chain Monte Carlo (MCMC) techniques [62, 63, 64, 65, 66, 67], though their computational cost increases dramatically in high-dimensional spaces. More recently, amortized variational inference methods, particularly those based on training diffusion samplers [68, 69, 70, 71, 72, 73, 74], have gained widespread adoption as they offer improved scalability for high-dimensional problems.

559 Following the [21], we adopt [42] to train diffusion sampler to sample from the target:

$$p_{\text{post}}(\mathbf{z}) \propto p(\mathbf{z}) \exp \left(\beta \cdot \left[r_{\phi}(f_{\theta}(\mathbf{z})) - \lambda \sum_{m=1}^M \max \left(0, g_{\phi}^{(m)}(f_{\theta}(\mathbf{z})) \right) \right] \right) \quad (15)$$

560 Here, the right-hand-side term serves as an unnormalized probability density, which the diffusion sampler amortizes the sampling cost by approximating it.

Off-policy Training of Diffusion Sampler As mentioned in the Section 4.2, we use the Trajectory Balance objective to train the diffusion sampler.

$$\mathcal{L}_{\text{TB}}(\mathbf{z}_{0:1}; \psi) = \left(\log \frac{Z_{\psi} p(\mathbf{z}_0) \prod_{i=0}^{T-1} p_F(\mathbf{z}_{(i+1)\Delta t} | \mathbf{z}_{i\Delta t}; \psi)}{p(\mathbf{z}_1) r(f_{\theta}(\mathbf{z}_1)) \prod_{i=1}^T p_B(\mathbf{z}_{(i-1)\Delta t} | \mathbf{z}_{i\Delta t})} \right)^2$$

562 The primary advantage of the TB loss is off-policy training [42, 75]. We can train our model not only from the on-policy trajectories through the reverse SDE $\{\mathbf{z}_0, \dots, \mathbf{z}_1\} = \tau \sim p_F(\tau)$ but also from the trajectories through the forward SDE conditioned on the generated samples $\tau \sim p_B(\tau | \mathbf{z}_1)$. This proves its effectiveness on mode coverage and credit assignment [42].

566 Specifically, we repeat two processes. First, we sample trajectories on-policy $\tau \sim p_F(\tau)$, train the model with Equation (14), and collect the samples \mathbf{z}_1 into the buffer. Second, from the collected samples \mathbf{z}_1 , we generate off-policy trajectories through $\tau \sim p_B(\tau | \mathbf{z}_1)$, then train with the Equation (14). During the off-policy training, we prioritize the samples with low energy: $\mathcal{E}(\mathbf{z}_1) = -\log(p(\mathbf{z}_1) r(f_{\theta}(\mathbf{z}_1)))$ following [42] to make our model focus on the low energy samples. These techniques improve the overall performance of our framework (Appendix E.4).

572 We use the original code⁸ released from [42] for implementation. We also set method-specific hyperparameters with Path Integral Sampler (PIS) [68] architecture, zero initialization, and t-scale to 1 to make sure the initialized $p_F(\mathbf{z}_1)$ starts from the standard normal distribution. Detailed hyperparameters for training the diffusion sampler can be found in Table 3.

Table 3: Hyperparameters for Training Diffusion Sampler

	Parameters	Values
Architecture	Number of Layers	2
	Num Units	256
	Diffusion Time Steps	50
Training	Batch size	256
	Optimizer	Adam
	Learning Rate	1×10^{-3}
	Training Epochs	50

576 **Computational Resources.** Our experiments were conducted using NVIDIA RTX 3090 and A6000 GPUs. These resources were sufficient to train our models within a reasonable time for all reported experiments. Details of computational time can be found at Appendix F.

⁸<https://github.com/GFN0rg/gfn-diffusion>

579 D.3 Hyperparameters

580 In our formulation of constrained black-box problems, we introduce λ for Lagrangian augmentation.
 581 We draw $N \times B$ samples from the posterior distribution, then select B samples during filtering. After
 582 evaluation, we update the training set by keeping the top L highest-scoring samples subject to the
 583 Lagrangian objective. Table 4 summarizes all hyperparameter values used in candidate selection,
 584 and we include additional analysis to assess how each parameter affects overall performance in
 585 Section 5.4 and Appendix E.

Table 4: Hyperparameters during sampling candidates

	Lambda λ	Inverse Temperature β	Buffer Size L	Filtering Coefficient N
Ackley 200D	10	10^5	3000	10
Rastrigin 200D	10	10^5	2000	10
Rosenbrock 200D	10	10^5	2000	10
RoverPlanning 60D	3	10^5	1000	10
Mopta 124D	3	10^3	500	10
DNA 180D	5	10^3	1000	15

E Further Analysis

In this section, we provide further analysis on different components of our method that are not included in the main manuscript due to the page limit.

E.1 Analysis on Feasibility Ratio

To further analyze our method’s ability to effectively handle constraints, we report the feasibility ratio across optimization batches for the Rastrigin 200D task. Here, the feasibility ratio denotes the number of feasible samples over queried samples.

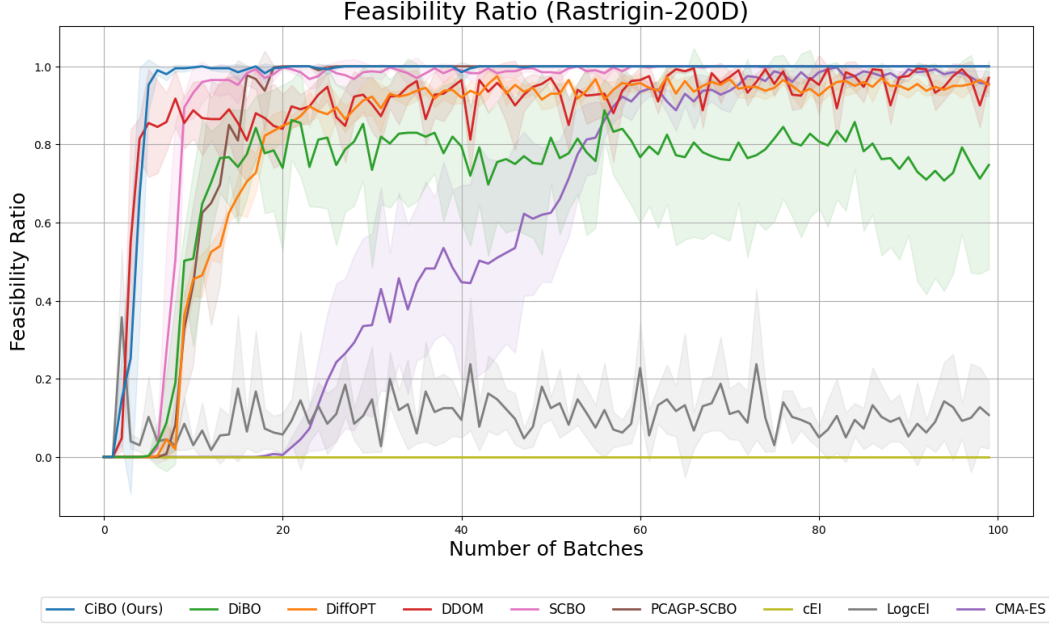


Figure 7: Feasibility ratio over all baselines. Experiments are conducted with four random seeds, and the mean and one standard deviation are reported.

As shown in Figure 7, CiBO demonstrates superior performance by rapidly achieving the highest feasibility ratio within the first 5-10 batches, significantly faster than all competing methods. While some baselines (SCBO, PCAGP-SCBO) eventually reach high feasibility ratios, they require approximately twice as many batches to achieve comparable performance. Other methods like DiBO and DiffOPT take even longer (around 20 batches), and CMA-ES struggles substantially, only reaching moderate feasibility ratios after 50 batches. Notably, CiBO not only reaches the high feasibility ratio faster but also maintains it consistently throughout the optimization process, demonstrating its robust constraint-handling capabilities in high-dimensional spaces.

601 E.2 Analysis on Filtering coefficient N

602 To improve the sample efficiency of our method, we introduce filtering, where we sample $N \times B$
 603 candidates from the posterior distribution, then select the highest B samples with respect to the
 604 Lagrangian-relaxed objective function. To analyze the impact of the filtering coefficient N , we
 605 experiment with varying N values, including our default $N = 100$.

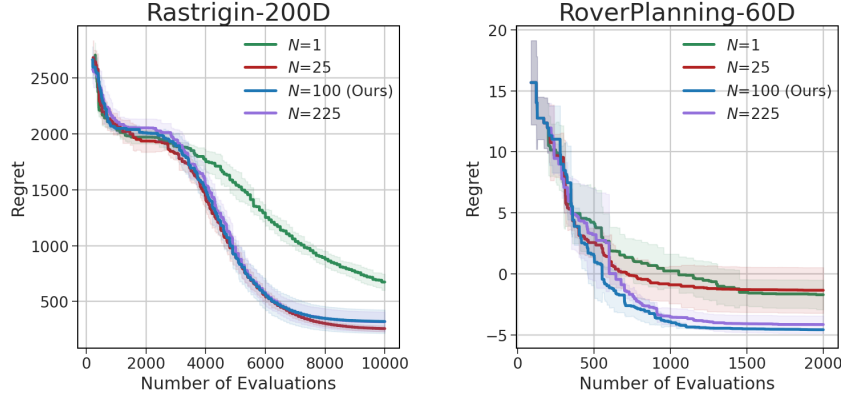


Figure 8: Performance of CiBO in Rastrigin-200D and Rover Planning-60D with varying N . Experiments are conducted with four random seeds, and the mean and one standard deviation are reported.

606 As shown in Figure 8, increasing the filtering coefficient improves sample efficiency by concentrating
 607 candidate selection in both high objective values and feasible regions. If the coefficient is set too low,
 608 we lose its exploitation capability, leading to slower convergence.

609 E.3 Analysis on Buffer Size L

610 In each round, we retain the L top-scoring samples with respect to the Lagrangian-relaxed objective
 611 function for computational efficiency. To analyze the effect of the buffer size L , we conduct
 612 experiments by varying L . As demonstrated in Figure 9, using too small L occasionally gets stuck in
 613 a sub-optimal solution while using too large L exhibits a slow convergence rate.

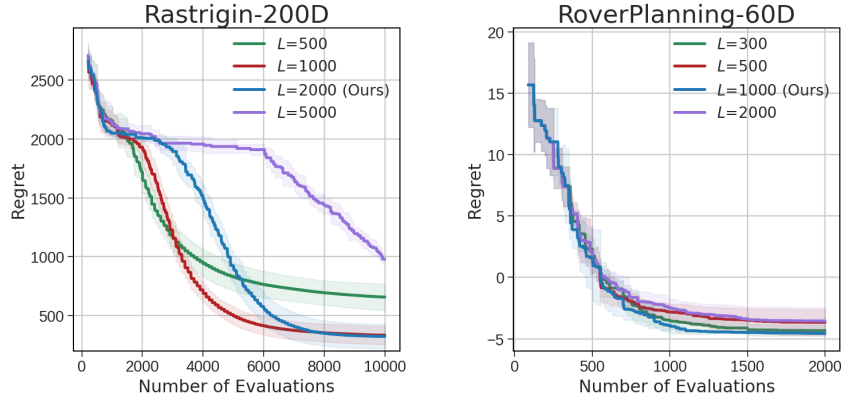


Figure 9: Performance of CiBO in Rastrigin-200D and Rover Planning-60D with varying L . Experiments are conducted with four random seeds, and the mean and one standard deviation are reported.

614 E.4 Effect of Off-policy Training in Amortized Inference

615 We employ off-policy training with the TB loss to train the diffusion sampler as detailed in Section 4.2.
 616 To analyze the impact of off-policy training on performance, we conduct ablation studies on different
 617 training schemes. As shown in Figure 10, off-policy training consistently outperforms on-policy
 618 methods, and the performance gap widens as the number of constraints grows (15 constraints in
 619 Rover Planning versus only 2 in Rastrigin). It underlines that training with off-policy samples is
 620 crucial for amortizing the posterior distribution with multiple modes and a large plateau.

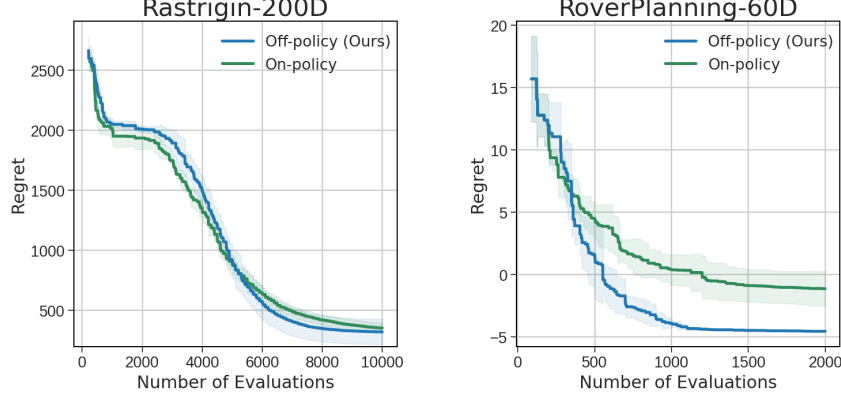


Figure 10: Comparison between off-policy and on-policy in Rastrigin-200D and Rover Planning-60D. Experiments are conducted with four random seeds, and the mean and one standard deviation are reported.

621 E.5 Analysis on Initial Dataset size $|D_0|$ and Batch size B

622 The size of the initial dataset, $|D_0|$, and batch size B play a critical role in the performance of
 623 black-box optimization algorithms. When $|D_0|$ is small and B is large, the algorithm must optimize
 624 using very limited information, making the search significantly more challenging. To this end, we
 625 conduct experiments varying $|D_0|$ and B to demonstrate the robustness of our method on initial data
 626 configurations. As shown in Figure 11, our method demonstrates robustness regarding both the initial
 627 dataset size $|D_0|$ and the batch size B .

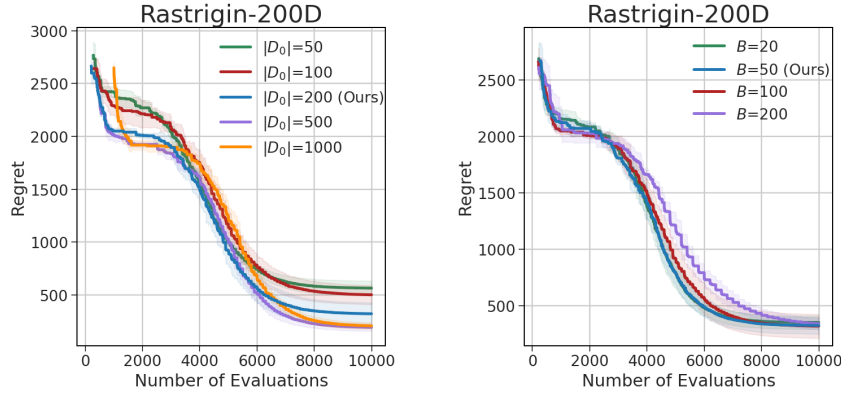


Figure 11: Performance of CiBO in Rastrigin-200D with varying $|D_0|$ and B . Experiments are conducted with four random seeds, and the mean and one standard deviation are reported.

F Runtimes

We report the running time of each method in Table 5. To measure the runtime, we conduct experiments on a single NVIDIA RTX 3090 GPU and Intel Xeon Platinum CPU @ 2.90 GHz. As shown in the table, the running time of our method is similar to other generative model-based approaches, and mostly faster than BO-based methods.

Table 5: Average time (in seconds) for each round in each method.

	Rastrigin-200D	Ackley-200D	Rosenbrock-200D	RoverPlanning-60D	Mopta-124D	DNA-180D
cEI	336.96 \pm 47.48	133.12 \pm 6.66	489.86 \pm 81.38	111.13 \pm 5.83	205.66 \pm 5.35	133.98 \pm 9.16
LogcEI	720.45 \pm 56.76	158.38 \pm 10.13	593.39 \pm 97.93	113.28 \pm 3.97	324.27 \pm 9.91	161.08 \pm 8.21
SCBO	322.81 \pm 47.38	117.81 \pm 6.50	475.02 \pm 80.11	87.83 \pm 4.42	270.30 \pm 5.19	147.83 \pm 12.57
PCAGP-SCBO	327.48 \pm 51.20	122.67 \pm 3.50	479.06 \pm 82.41	17.55 \pm 3.05	20.69 \pm 0.39	81.34 \pm 9.19
CMA-ES	0.08 \pm 0.00	0.09 \pm 0.00	0.10 \pm 0.01	0.61 \pm 0.00	5.33 \pm 0.15	46.58 \pm 3.16
DDOM	26.87 \pm 0.28	27.00 \pm 0.32	26.96 \pm 0.12	3.56 \pm 0.02	8.63 \pm 0.37	50.99 \pm 0.81
DiffOPT	91.00 \pm 5.07	111.04 \pm 1.51	89.37 \pm 8.27	29.48 \pm 1.21	105.61 \pm 2.71	60.64 \pm 1.34
DiBO	73.97 \pm 0.56	68.89 \pm 0.98	73.83 \pm 1.11	29.43 \pm 1.51	66.61 \pm 4.16	71.85 \pm 2.48
CiBO	73.39 \pm 2.09	103.43 \pm 4.84	82.24 \pm 6.50	53.58 \pm 5.14	105.43 \pm 2.24	81.77 \pm 2.37

633 **G Limitations and Future Work**

634 We are interested in improving our method further. First, as we need to train all models with the
635 updated dataset in every round, presenting a framework that can efficiently reuse the trained models
636 from the previous rounds would be beneficial. Furthermore, there are several advancements in the
637 literature on flow-based model training [76] and diffusion samplers [77, 78], which could potentially
638 yield substantial performance gains. We leave them as future work.

639 **H Broader Impact**

640 Advances in real-world design optimization have the potential to drive major innovations, but they
641 also come with potential risks and unintended consequences. For example, optimization techniques
642 in biochemical design may uncover novel compounds with therapeutic potential, but similar methods
643 could also be misused to discover harmful substances. It is essential for researchers to act responsibly
644 and ensure their work serves the public good.