

Motivation

1. **Offline black-box optimization** searches for better designs without new evaluations.
2. Optimizer **exploits surrogate overestimations** far from data manifold (out-of-distribution).
3. Can **diffusion model** serve as **calibrated forward surrogate**?

Background

Offline BBO: with only historical data

$$\mathcal{D} = \{(x_i, y_i)\}_{i=1}^N, \quad y_i = f(x_i),$$

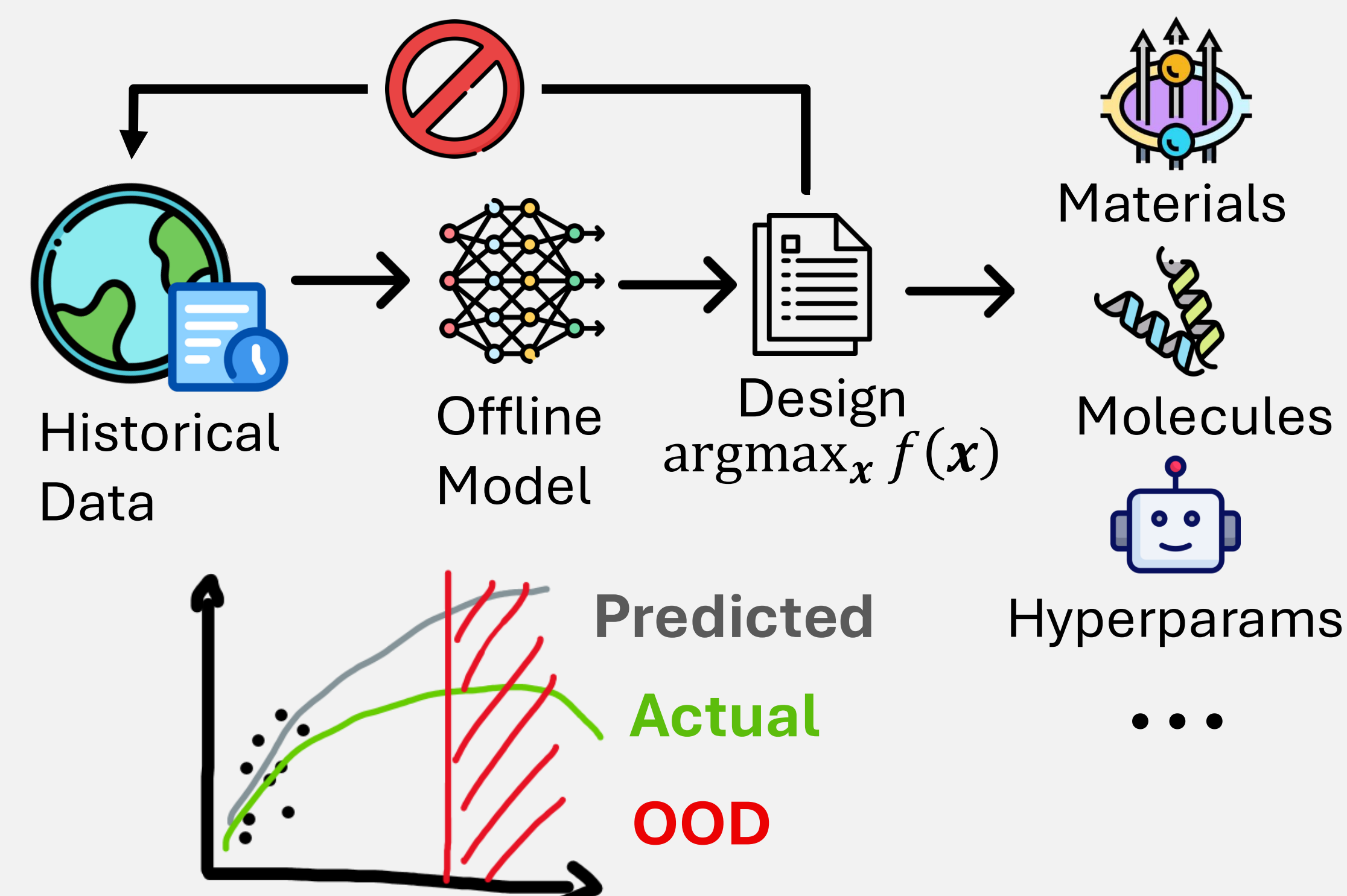
Find new design $x^* = \operatorname{argmax}_x f(x)$ without new oracle calls.

$$p(x | y) \propto p(y | x) p(x)$$

Failure mode: surrogate optimization can leave $\operatorname{supp}(\mathcal{D})$ and overestimate out-of-distribution designs.

Need: a probabilistic forward surrogate that is both

- i) Calibrated to preserve important statistics;
- ii) Support-aware to be “skeptical” of OOD designs.



Methodology

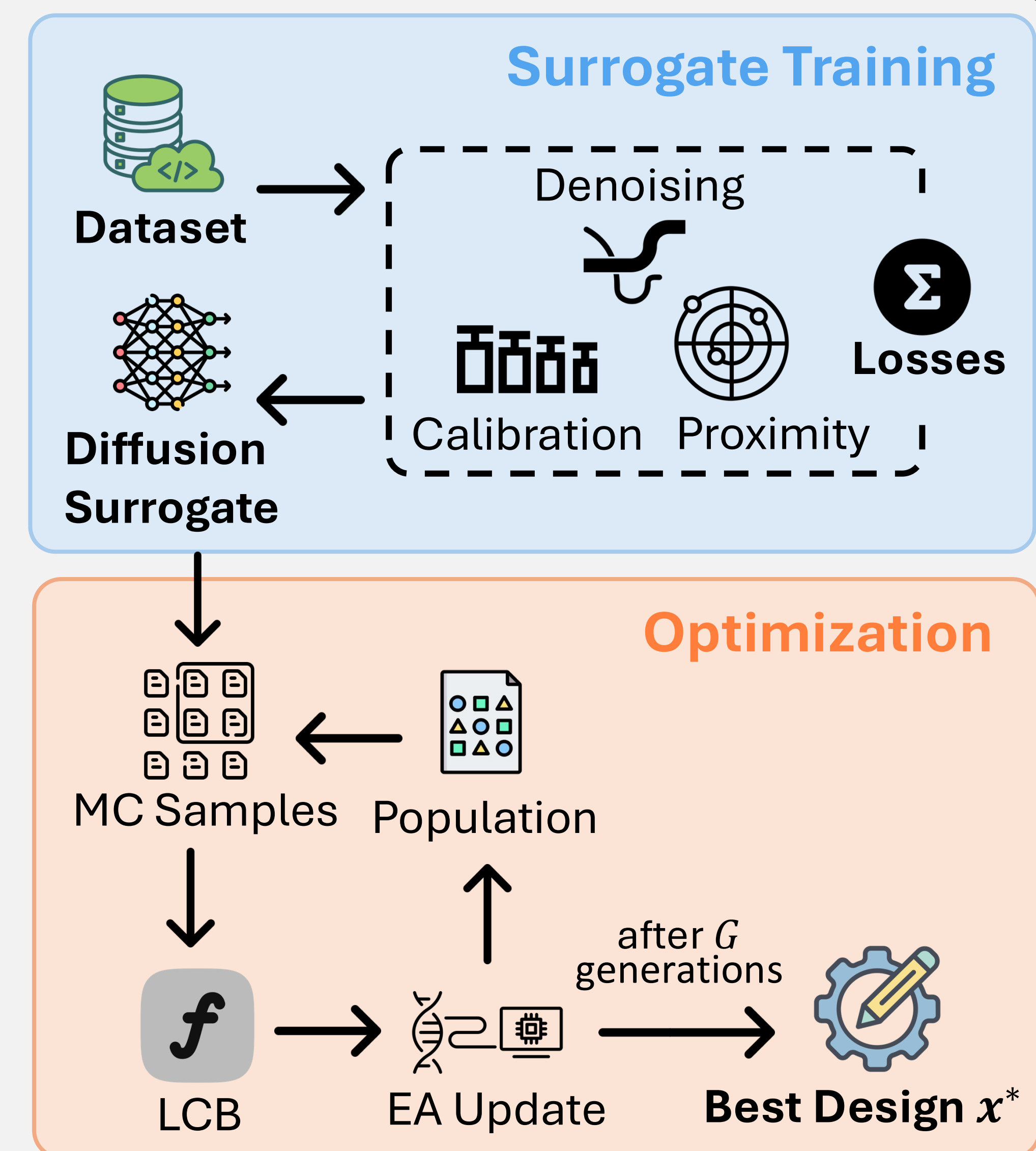
Goal: learn probabilistic forward surrogate, then search conservatively. SPADE models $p_\theta(y | x)$ with a **conditional diffusion model**. **MC samples** give a predictive mean and uncertainty.

Optimization-aware training:

$$\mathcal{L} = \mathcal{L}_{\text{diff}} + \lambda_1 \mathcal{L}_{\text{calib}} + \lambda_2 \mathcal{L}_{\text{prox}}, \quad d(x) = \log R_k(x)$$

Conservative search: candidates scored by a lower confidence bound and updated by evolutionary search. Mean decreases, uncertainty increases $\Rightarrow$ LCB decreases when far from data support.

$$y^{(m)} \sim p_\theta(y | x) \\ \text{LCB}(x) = \widehat{\mu}_\theta(x) - \beta \widehat{\sigma}_\theta(x), \\ x^* = \arg \max_x \text{LCB}(x).$$



Applications

Impact & future work on broader offline-MBO use cases:

- Proteins / DNA / molecules
- Materials / robots / chips
- Safe extrapolation
- LLM data mixtures
- Self-evolving agent
- **Many more!**

Please feel free to apply SPADE / collaborate!

Benchmark & Ablation Results

Method	SuperC	Ant	D’Kitty	LLM-DM	TF8	TF10	Mean Rank	Median Rank
$\mathcal{D}(\text{best})$	0.399	0.565	0.884	1.000	0.439	0.511	—	—
<i>Standard Optimization Baselines</i>								
CMA-ES	0.465 ± 0.024	1.561 ± 0.896	0.724 ± 0.001	0.975 ± 0.016	0.939 ± 0.039	0.692 ± 0.013	13.3/24	13.5/24
REINFORCE	0.481 ± 0.013	0.263 ± 0.026	0.573 ± 0.204	0.305 ± 0.027	0.961 ± 0.034	0.632 ± 0.012	19.8/24	23.0/24
BO-qEI	0.402 ± 0.034	0.812 ± 0.000	0.896 ± 0.000	0.953 ± 0.022	0.825 ± 0.091	0.663 ± 0.011	18.7/24	19.5/24
<i>Forward Surrogate Methods</i>								
Standard GA	0.505 ± 0.013	0.293 ± 0.029	0.860 ± 0.021	0.998 ± 0.000	0.923 ± 0.011	0.732 ± 0.041	14.0/24	14.5/24
GA on GP	0.499 ± 0.019	0.948 ± 0.013	0.946 ± 0.001	0.846 ± 0.029	0.770 ± 0.087	0.599 ± 0.004	16.0/24	17.0/24
MC-Dropout	0.535 ± 0.064	0.805 ± 0.021	0.934 ± 0.004	0.781 ± 0.251	0.911 ± 0.042	0.817 ± 0.031	13.0/24	13.0/24
COMs	0.481 ± 0.028	0.878 ± 0.031	0.929 ± 0.016	0.815 ± 0.008	0.937 ± 0.025	0.755 ± 0.017	15.3/24	15.5/24
RoMA	0.509 ± 0.015	0.592 ± 0.059	0.825 ± 0.016	0.968 ± 0.011	0.662 ± 0.000	0.801 ± 0.000	16.2/24	16.0/24
ICT	0.503 ± 0.017	0.911 ± 0.030	0.945 ± 0.011	0.839 ± 0.021	0.888 ± 0.047	0.814 ± 0.027	14.0/24	13.5/24
Tri-mentoring	0.514 ± 0.018	0.944 ± 0.033	0.950 ± 0.015	0.750 ± 0.002	0.899 ± 0.045	0.811 ± 0.039	12.2/24	9.5/24
BDI	0.513 ± 0.000	0.964 ± 0.000	0.941 ± 0.000	0.988 ± 0.025	0.973 ± 0.000	<u>0.882 ± 0.000</u>	5.7/24	4.0/24
LTR	0.514 ± 0.022	0.904 ± 0.036	0.958 ± 0.012	0.982 ± 0.041	0.973 ± 0.010	0.849 ± 0.023	6.2/24	5.0/24
MATCH-OPT	0.504 ± 0.021	0.931 ± 0.011	0.957 ± 0.014	0.850 ± 0.033	0.977 ± 0.004	0.824 ± 0.008	9.2/24	8.5/24
PGS	0.563 ± 0.058	0.949 ± 0.017	0.966 ± 0.013	0.595 ± 0.027	<u>0.981 ± 0.015</u>	0.793 ± 0.021	8.0/24	4.0/24
<i>Inverse Generative Methods</i>								
CbAS	0.503 ± 0.069	0.846 ± 0.033	0.895 ± 0.016	0.919 ± 0.043	0.903 ± 0.028	0.652 ± 0.032	16.2/24	16.0/24
MINs	0.499 ± 0.017	0.894 ± 0.022	0.939 ± 0.004	0.982 ± 0.007	0.908 ± 0.063	0.647 ± 0.012	14.0/24	14.0/24
DDOM	0.481 ± 0.015	0.926 ± 0.025	0.923 ± 0.009	0.983 ± 0.028	0.884 ± 0.042	0.708 ± 0.015	14.7/24	17.0/24
GABO	0.508 ± 0.007	0.224 ± 0.051	0.719 ± 0.001	0.975 ± 0.019	0.939 ± 0.038	0.739 ± 0.009	15.0/24	13.5/24
GTG	0.480 ± 0.055	0.865 ± 0.040	0.935 ± 0.010	0.910 ± 0.030	0.901 ± 0.039	0.801 ± 0.004	15.0/24	14.0/24
RGD	0.515 ± 0.011	0.922 ± 0.020	0.883 ± 0.014	1.004 ± 0.008	0.889 ± 0.068	0.825 ± 0.039	10.7/24	9.5/24
BONET	0.434 ± 0.021	0.948 ± 0.025	0.955 ± 0.010	0.874 ± 0.039	0.894 ± 0.086	0.796 ± 0.018	13.3/24	14.5/24
DEMO	0.520 ± 0.012	0.948 ± 0.013	0.954 ± 0.013	0.907 ± 0.005	0.808 ± 0.044	0.836 ± 0.042	9.7/24	6.5/24
ROOT	0.525 ± 0.012	0.965 ± 0.014	<u>0.972 ± 0.005</u>	0.905 ± 0.006	0.986 ± 0.007	0.833 ± 0.046	5.0/24	3.5/24
SPADE (Ours)	0.546 ± 0.013	0.978 ± 0.006	0.981 ± 0.003	1.019 ± 0.064	0.923 ± 0.015	0.915 ± 0.010	2.8/24	1.5/24

SOTA: SPADE finds high-scoring designs reliably across six offline BBO tasks.

Ablation: Diffusion + Prox + Calib = best. Calibration and support regularization both needed.

Observation on synthetic functions: SPADE recovers global structure.

