# Genentech

A Member of the Roche Group

# Adding Conditional Control to Diffusion Models with Reinforcement Learning

Yulai Zhao\*<sup>12</sup>, Masatoshi Uehara<sup>2</sup>\*, Gabriele Scalia<sup>2</sup>, Tommaso Biancalani<sup>2</sup>, Sergey Levine<sup>3</sup>, Ehsan Hajiramezanali<sup>2</sup>

<sup>1</sup>Princeton University, <sup>2</sup>Genentech, <sup>3</sup>University of California, Berkeley



### Background

- A pre-trained conditional diffusion model excels at modeling p(x|c).
- For example, in Stable
   Diffusion, c∈C is a prompt,
   and x∈X is the image
   generated according to this
   prompt.
- Many tailored DMs are able to generate biological sequences (e.g., DDSM).
- In practice, we often want to add additional controls into pre-trained diffusion models, e.g.
  - Stable Diffusion.
  - existing condition: prompts
  - new condition: certain layouts or backgrounds.
  - DDSM tailored for generating DNA enhancers.
    - existing condition: activity level in HepG2
    - new condition: activity level in other cell lines such as K562.

## Settings

- Given the pre-trained model, which enables us to sample from p<sup>pre</sup> (x|c): C → ∆(X).
- Goal: add new conditional controls y∈Y such that we can sample from p(x|c,y).
- Assume we can access to offline data:

 $D = \{(c^{(i)}, x^{(i)}, y^{(i)})\}_{i=1}^{n}$ 

where conditional distribution is denoted by p<sup>o</sup>(y|x,c).

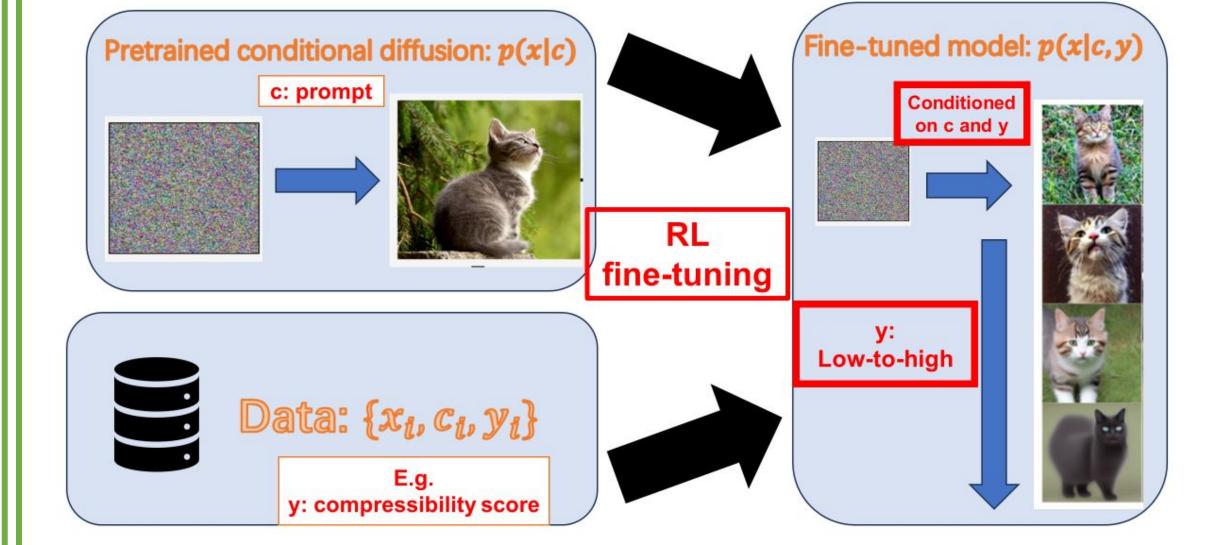
#### **Target Distribution**

our goal is to obtain a diffusion model such that we can sample from

 $p_{\gamma}(\cdot|c,y) \propto (p^{\prime}(y|\cdot,c))^{\gamma} p^{pre}(\cdot|c)$ where  $\gamma$  represents the strength of the additional guidance.

### Methodology & Results

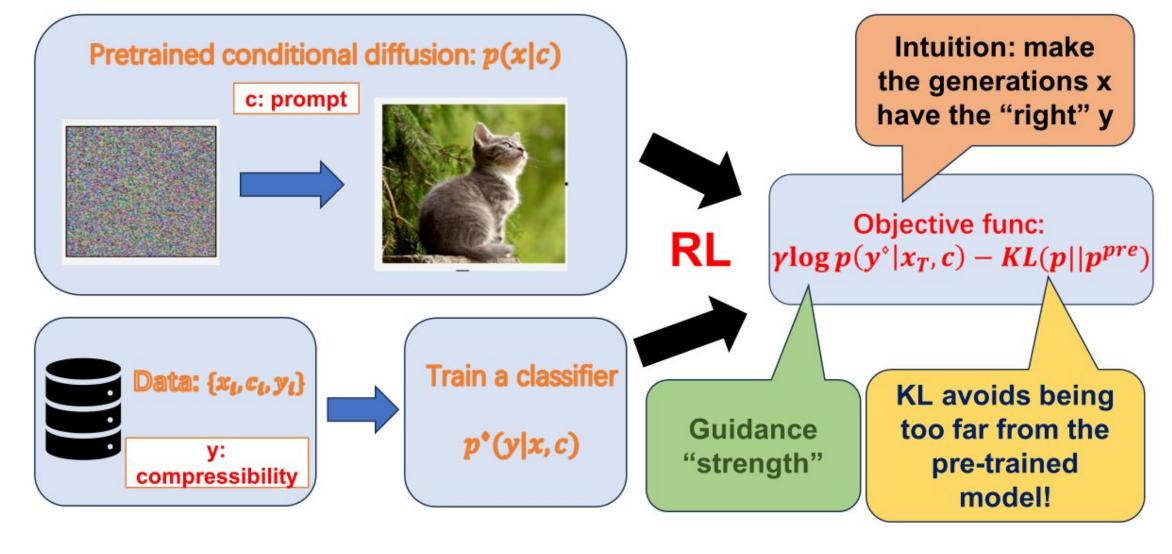
# Our goal: adding control via fine-tuning



Roadmap

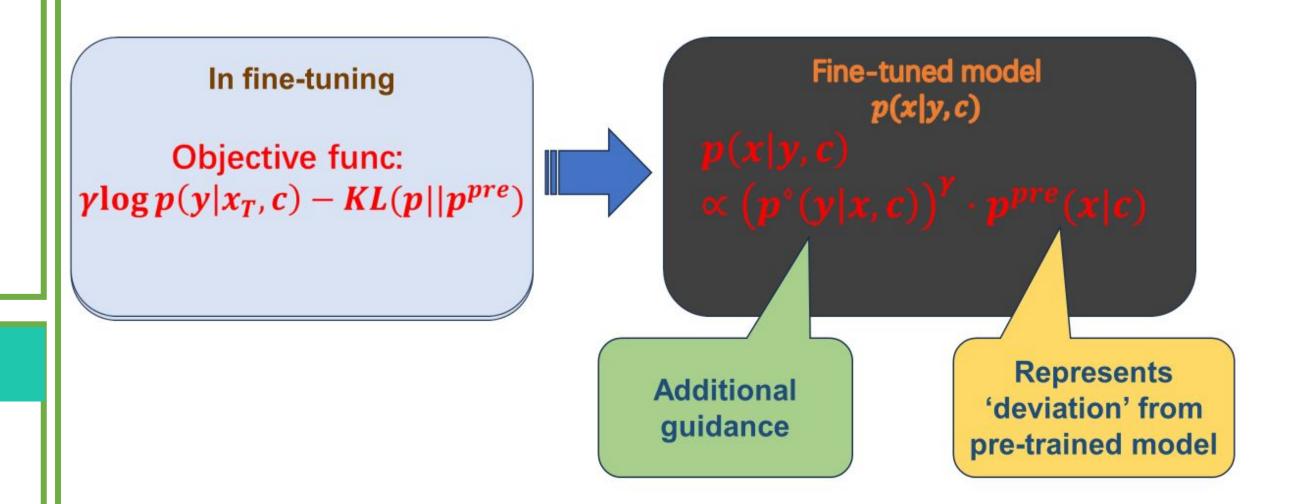
### **RL-based Fine-tuning**

## Methodology



### **Advantages of Our Approach**

# Theoretical justification (incomplete)



### **★** Compared to classifier-free guidance

Our method demonstrates superiority by leveraging the conditional independence.

• Example 1: Y  $\perp$  C | X. Then  $p^{\flat}(y|x,c) = p^{\flat}(y|x)$ 

 This means we only need (x,y) sample pairs to train the classifier, rather than triplets (c,x,y).

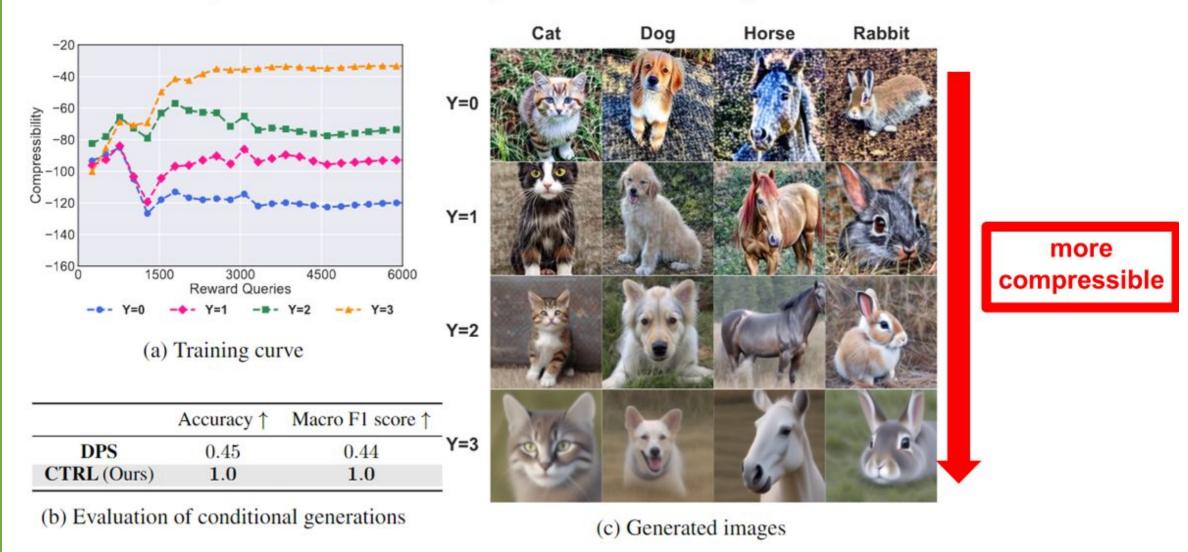
Example 2 (multi-task): Y₁ ⊥ Y₂ | C, X, Y₁ ⊥ C | X and Y₂ ⊥ C | X. Then

$$\log p^{\flat}(y_1, y_2 | x, c) = \log p^{\flat}(y_1 | x) + \log p^{\flat}(y_2 | x)$$

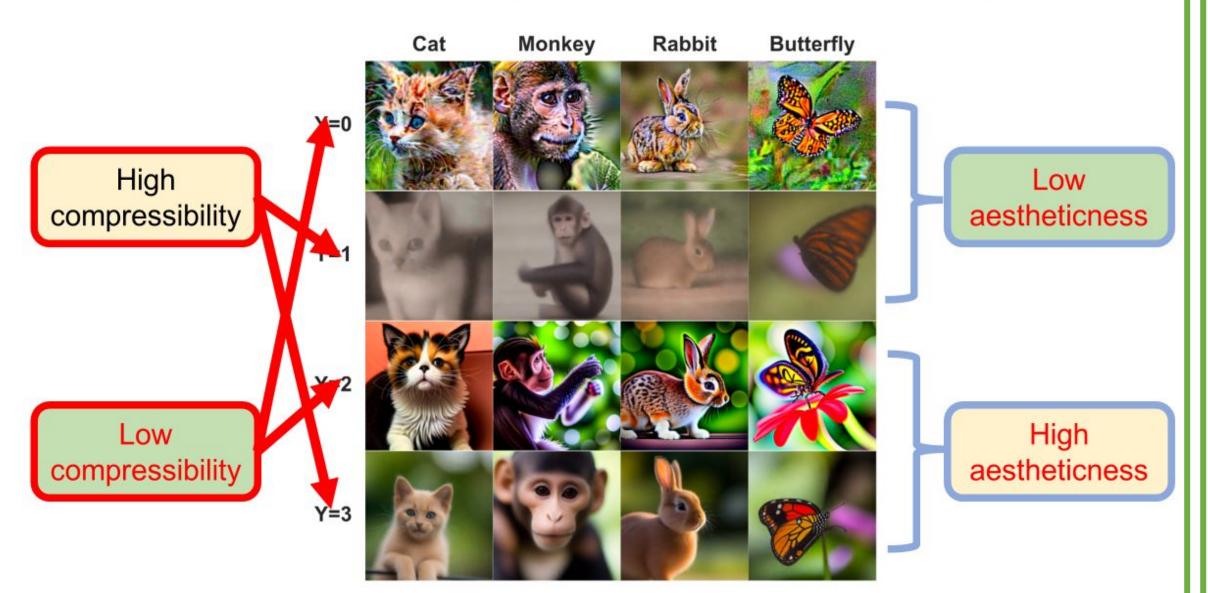
- This means we only need (x,y<sub>1</sub>) and (x,y<sub>2</sub>) pairs.
- Significantly simplifying dataset construction: classifier free guidance must require quadruples (c,x,y<sub>1</sub>,y<sub>2</sub>)!

#### Results

# **Example 1: Compressibility**



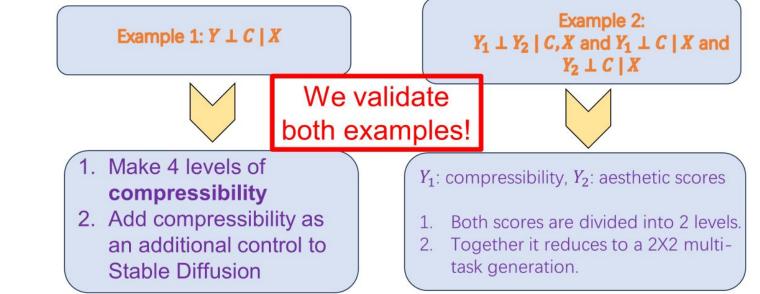
# Example 2: Compressibility & Aestheticness



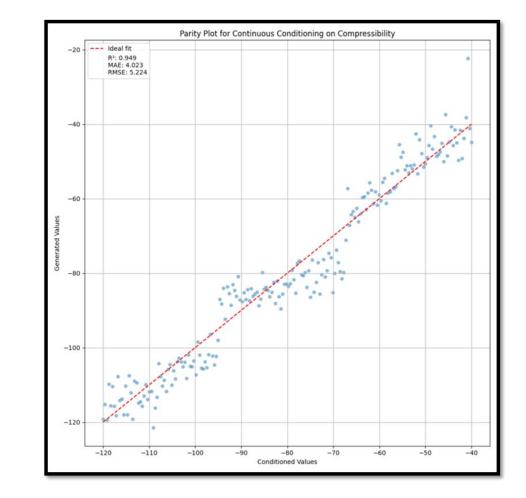
### **Experimental Details**

#### Settings

#### **Experiments**



### **Extension: conditioned on continuous y**



- We can use the model fine-tuned on discretized y and interpolate discrete class embeddings for continuous y.
- This naive approach can achieve R<sup>2</sup>=0.95

#### Conclusions

- We introduce an RL-based fine-tuning approach for conditioning pre-trained diffusion models on new additional labels.
- Compared to classifier-free guidance, our proposed method allows for leveraging the conditional independence, thereby greatly simplifying the construction of the offline dataset.
- We also theoretically justify our approach and build the connection with classifier-based guidance.
- Future work includes applying this method to design DNA enhancers and 5'UTR.
- The goal is achieve cell-specific promoters design!



