

Background

- A pre-trained conditional diffusion model excels at modeling $p(x|c)$.
 - For example, in Stable Diffusion, $c \in C$ is a prompt, and $x \in X$ is the image generated according to this prompt.
 - Many tailored DMs are able to generate biological sequences (e.g., DDSM).
- In practice, we are often interested in adding new controls into pre-trained diffusion models, e.g.
 - Stable Diffusion.
 - existing condition: prompts
 - new condition: certain layouts or backgrounds.
 - DDSM that generates 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 $p^{pre}(x|c): C \mapsto \Delta(X)$.
- Our goal is to add new conditional controls $y \in Y$ such that we can sample from $p(x|c, y)$.
- Assume we can access to offline data:

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

We denote the conditional distribution by $p^\phi(y|x, c)$.

Target Distribution

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

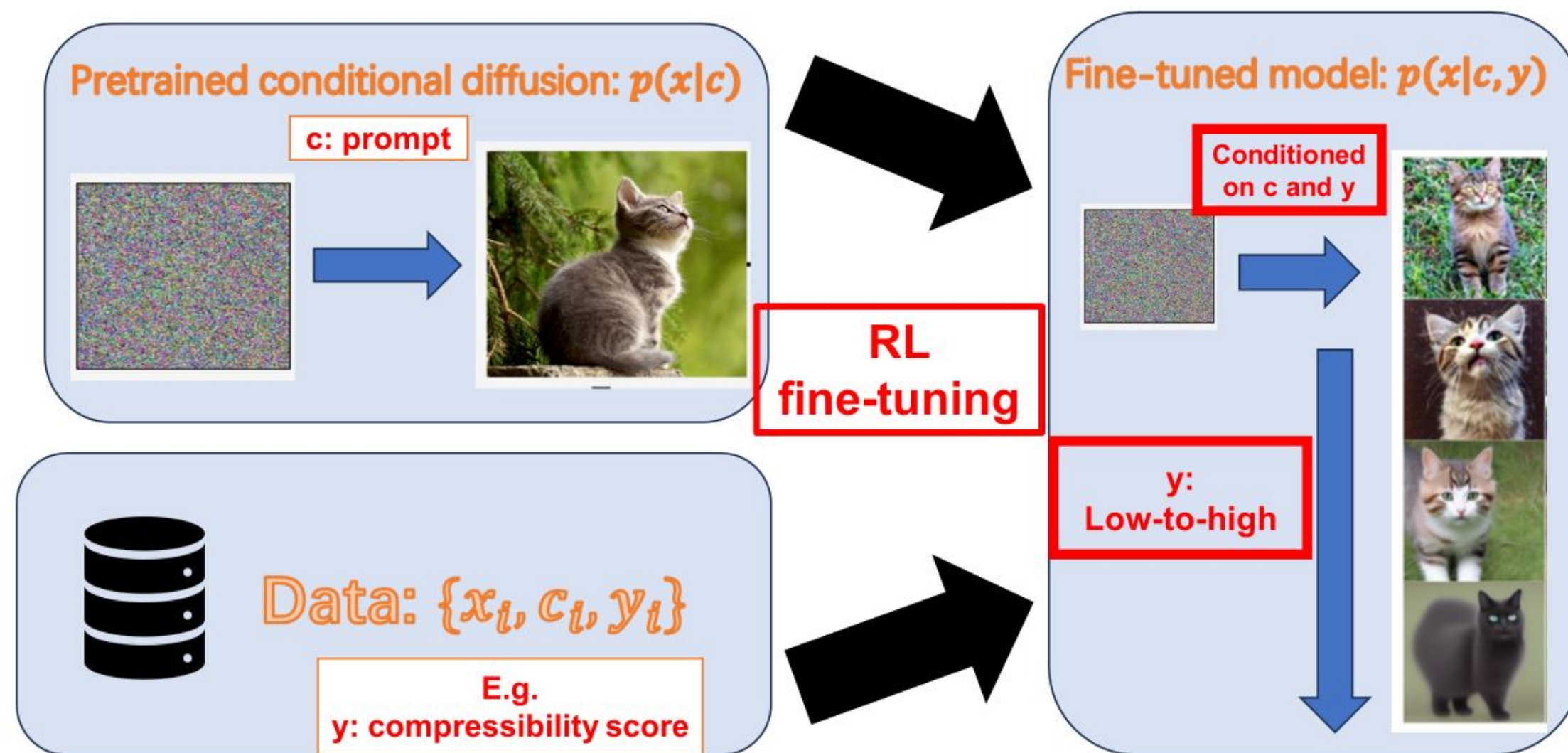
$$p_\gamma(\cdot|c, y) \propto (p^\phi(y|\cdot, c))^\gamma p^{pre}(\cdot|c)$$

where γ represents the strength of the additional guidance.

Methodology & Results

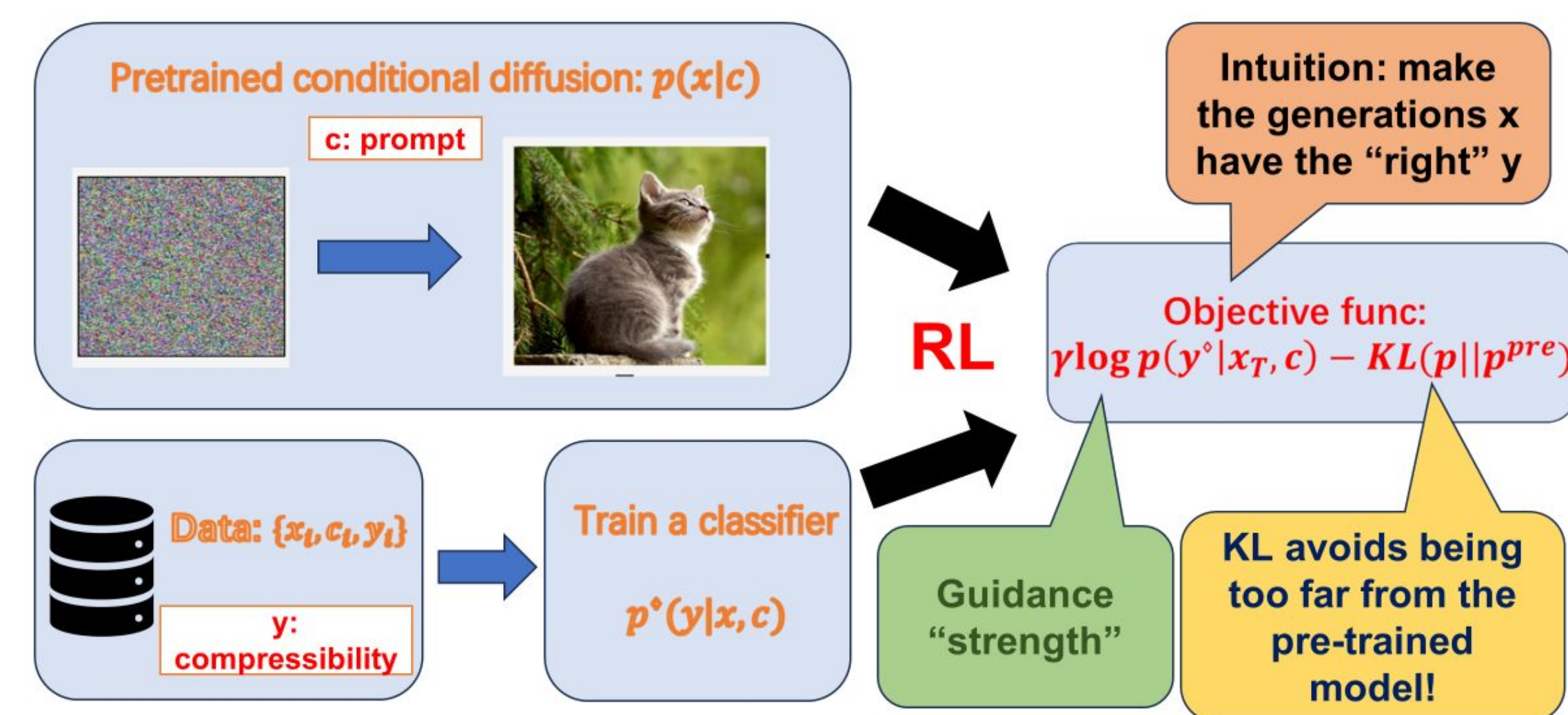
Roadmap

Our goal: adding control via fine-tuning



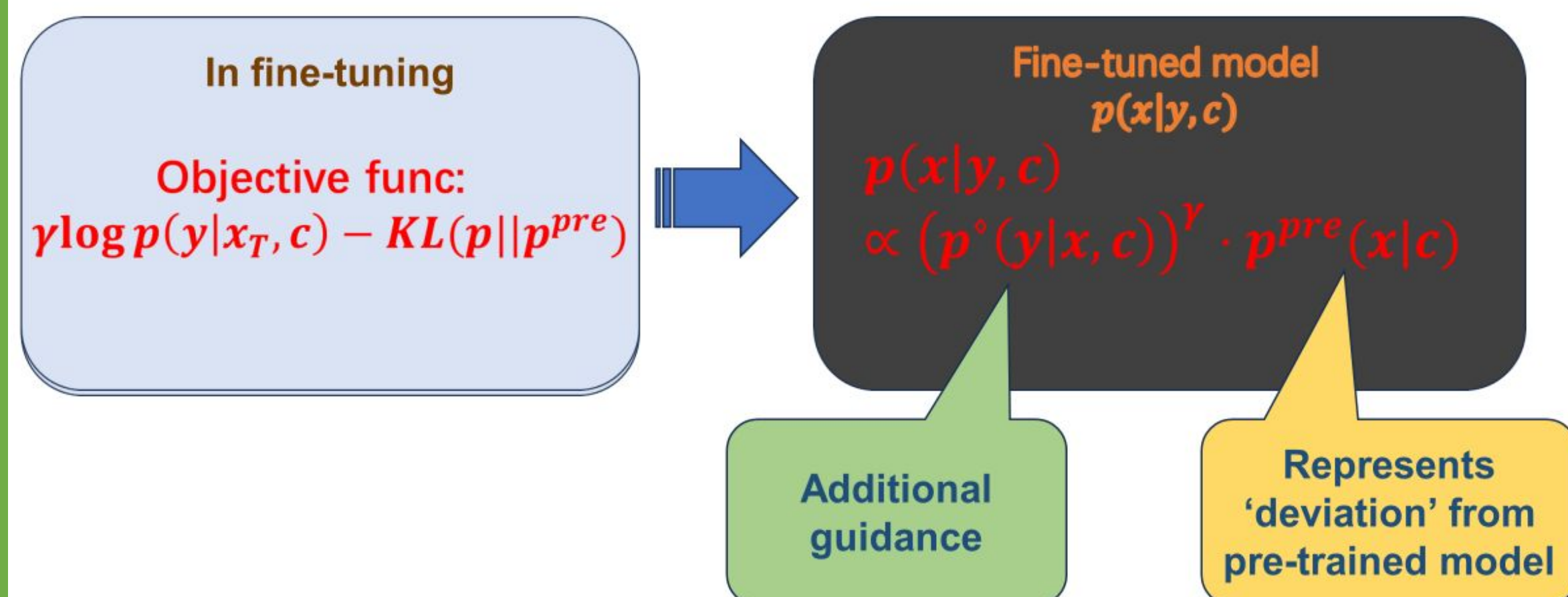
RL-based Fine-tuning

Methodology

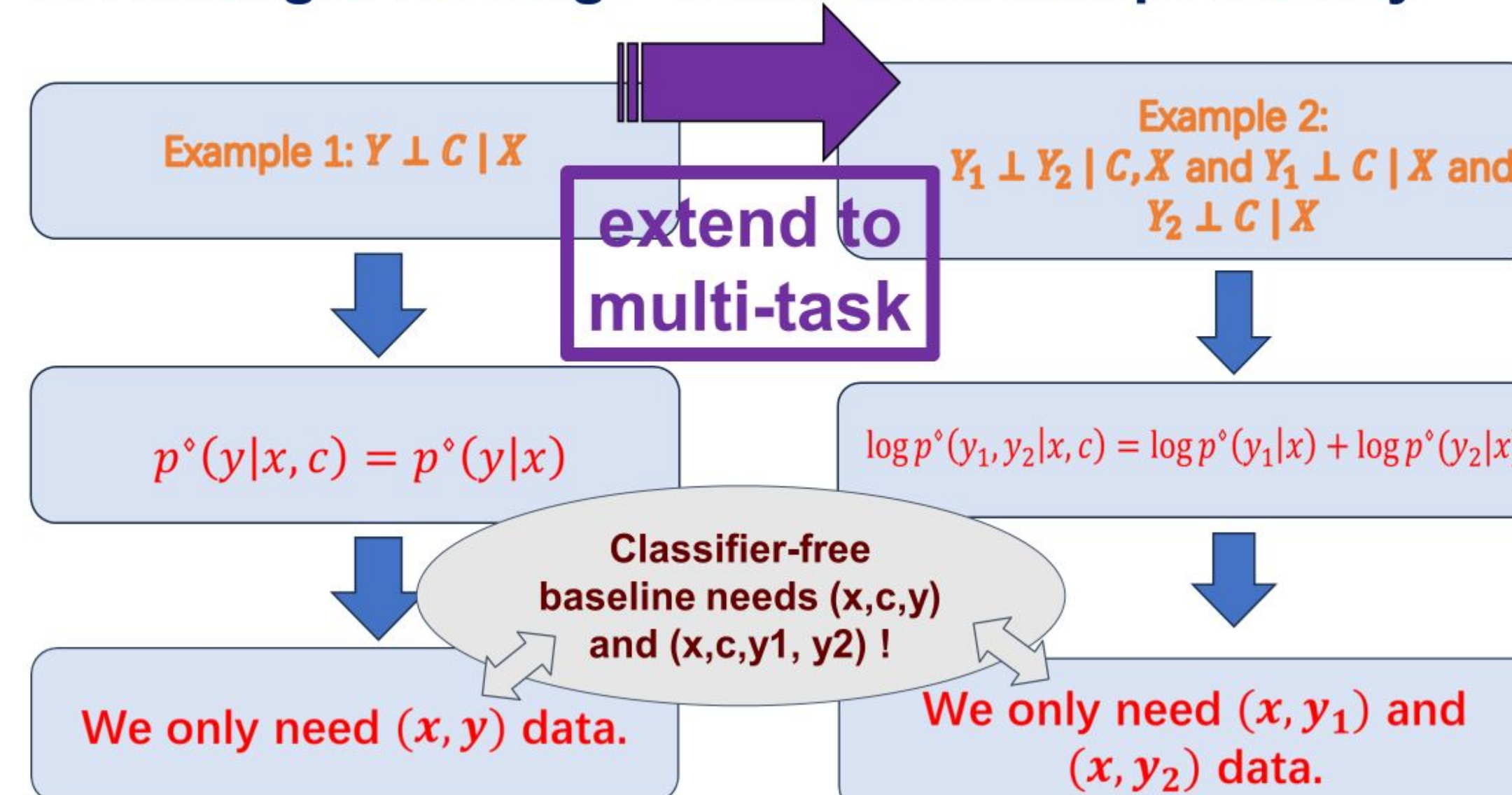


Benefits of Our Approach

Theoretical justification (incomplete)

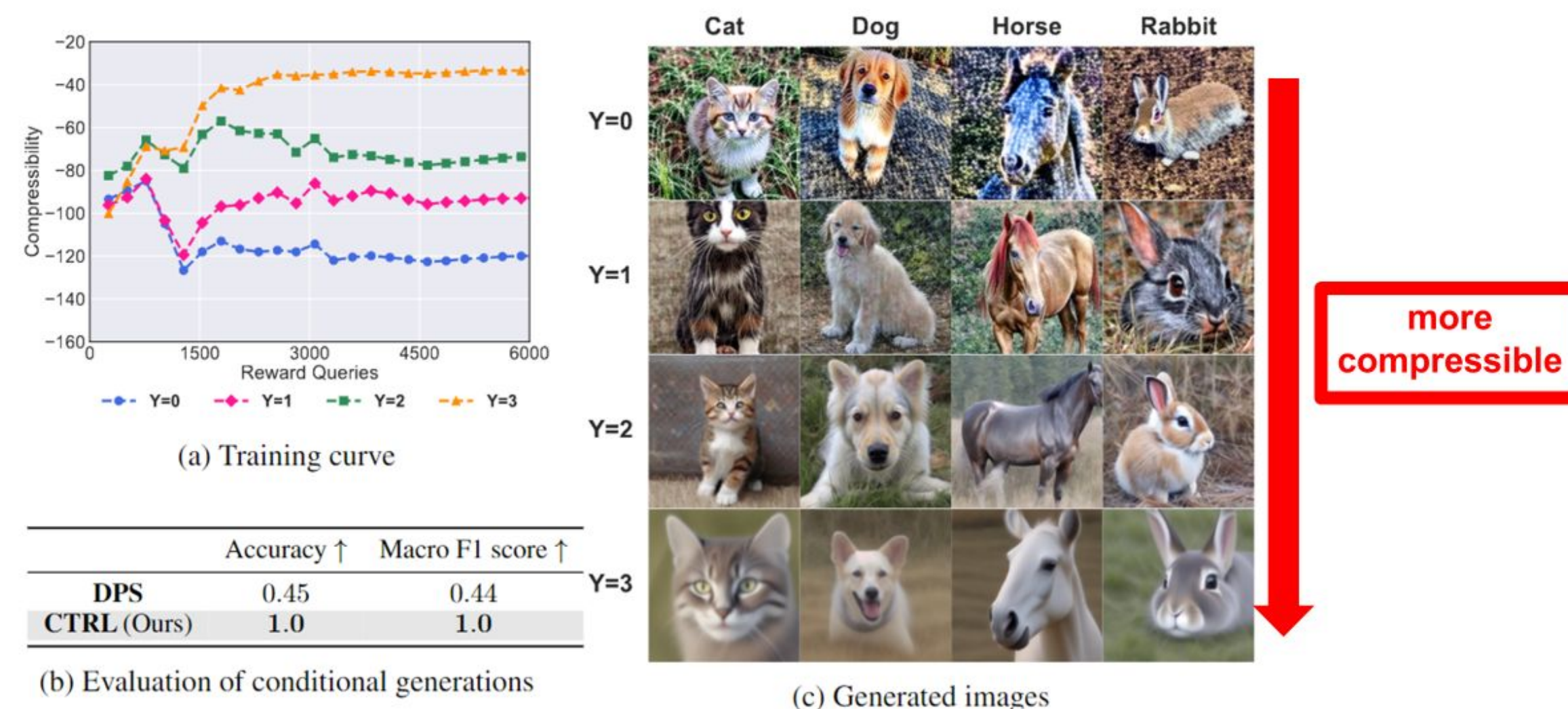


Advantage: leverage conditional independency

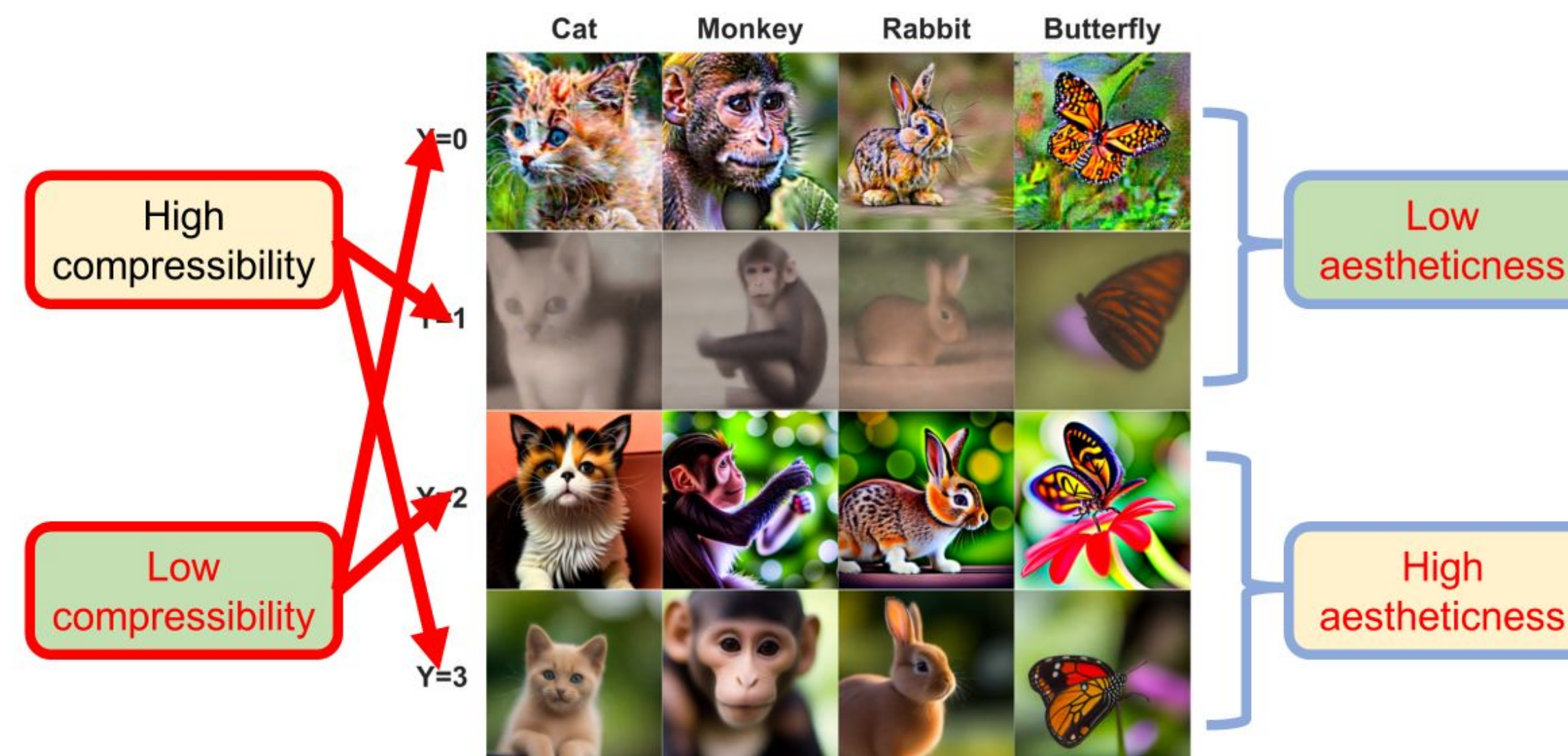


Experimental Results

Example 1: Compressibility



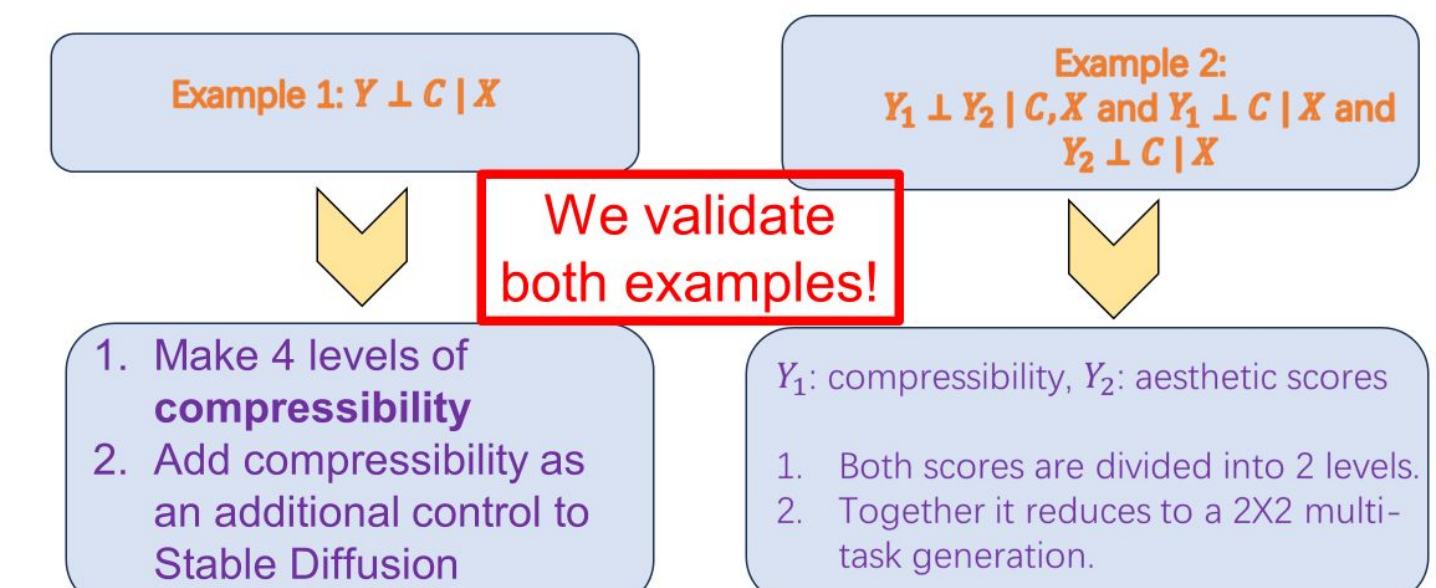
Example 2: Compressibility & Aestheticness



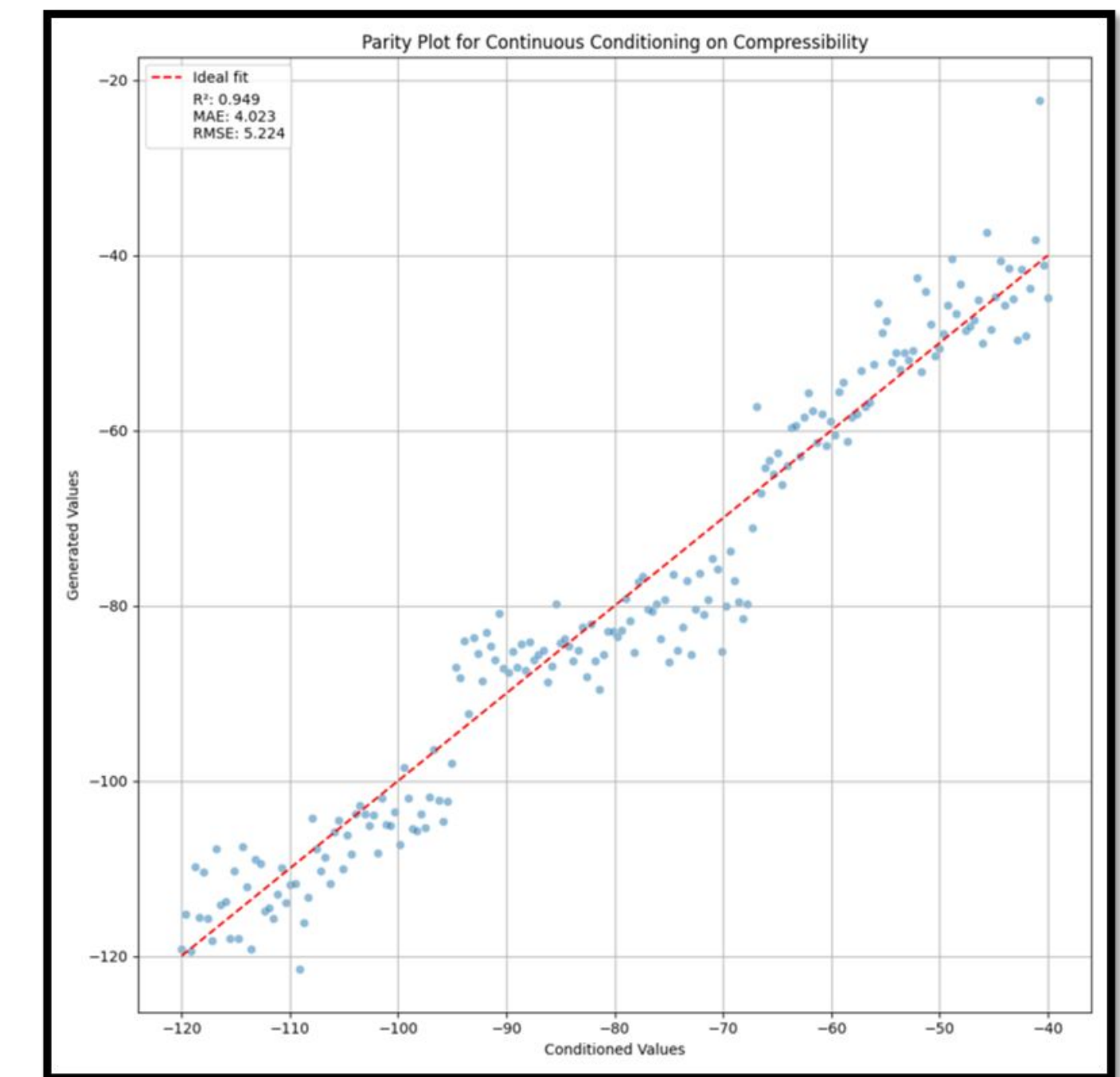
Experimental Details

Settings

Experiments



Extension: continuous condition



Conclusions & Future Work

- 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**.
- We are working on extending this work to **DNA enhancers** and **RNA 5'UTR design**.
- The goal is achieve **cell-specific promoters design**!

ArXiv

