



TaxaDiffusion: Progressively Trained Diffusion Model for Fine-Grained Species Generation

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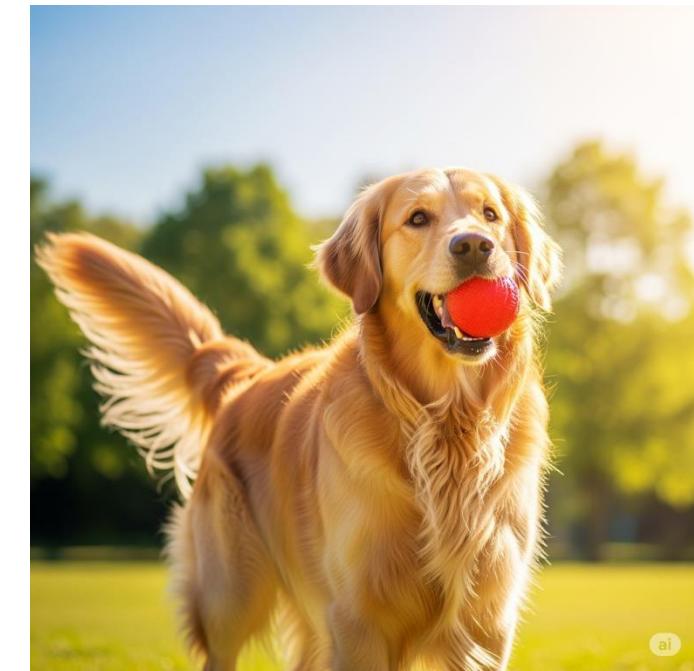
Motivation

- Gemini 2.5 Flash: Golden Retriever
 - Prompt: generate an image of Golden Retriever.

Real Image



Generated Image



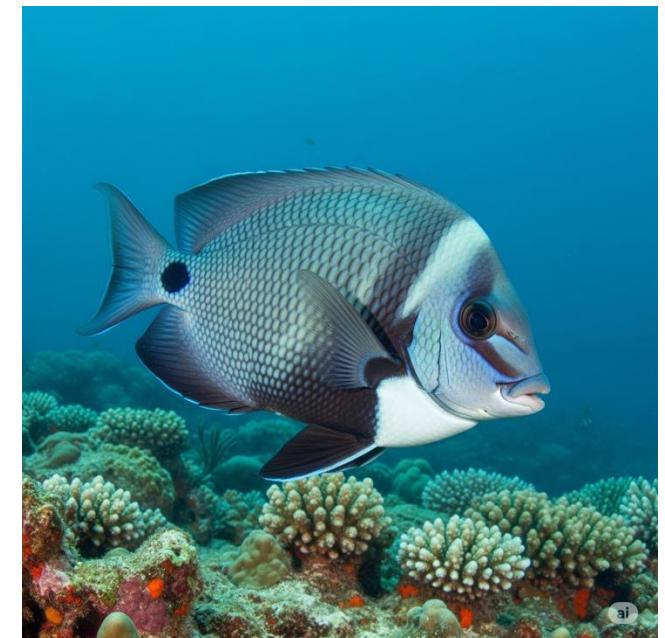
Motivation & Challenges

- Gemini 2.5 Flash: *Abudefduf abdominalis*
 - Prompt: generate an image of *Abudefduf abdominalis*.

Real Image



Generated Image

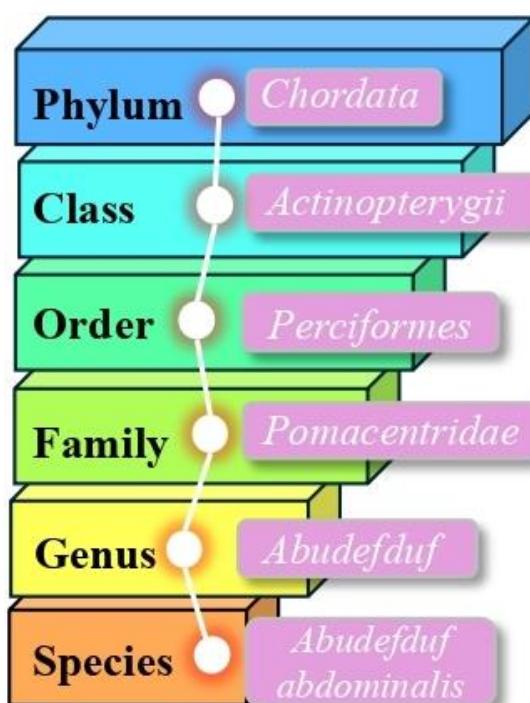


Motivation & Challenges

- Why generative models can't generate some species?
 - Limited training data per species.
 - High morphological diversity within species.
- Question:
 - How can we learn high-quality generative models for animals without a large amount of data per species?

Goal

- Generate accurate, fine-grained species images.
 - We introduced: **TaxaDiffusion**.



Real Image



Zero-Shot



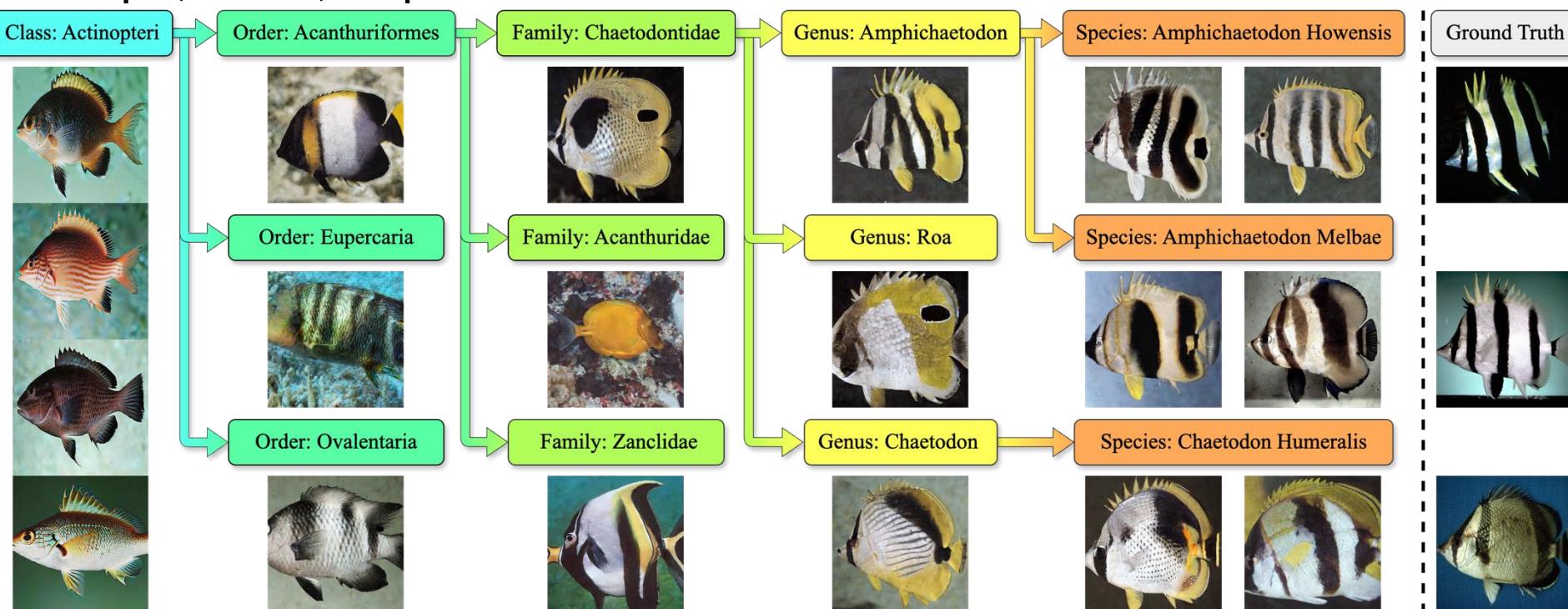
LoRA



TaxaDiffusion

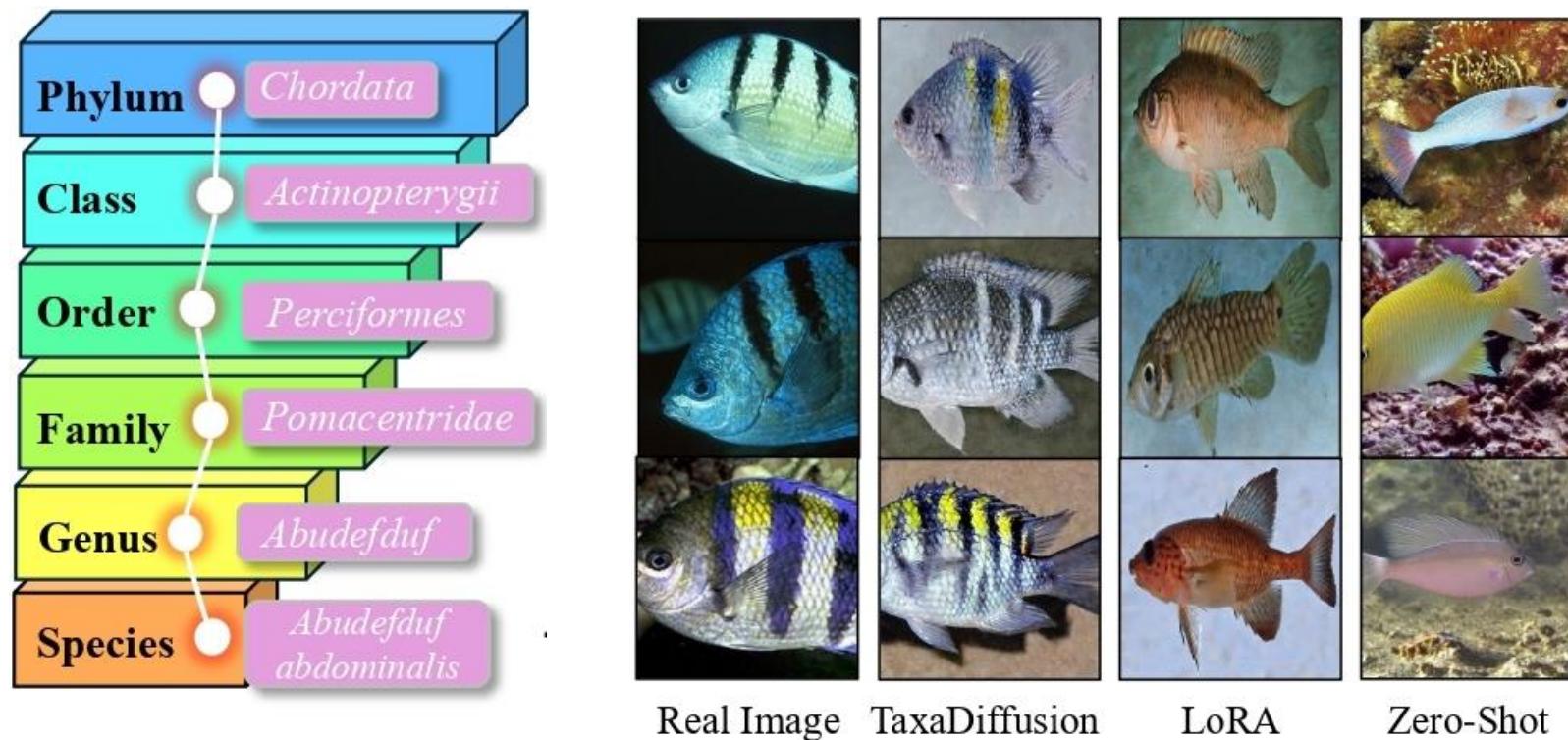
Key Insight - Leveraging Taxonomy

- **Taxonomy:**
 - Shared ancestry leads to similar characteristics in species.
 - Closely related species (same Genus/Family) differ mainly in subtle shape, color, or pattern variations.



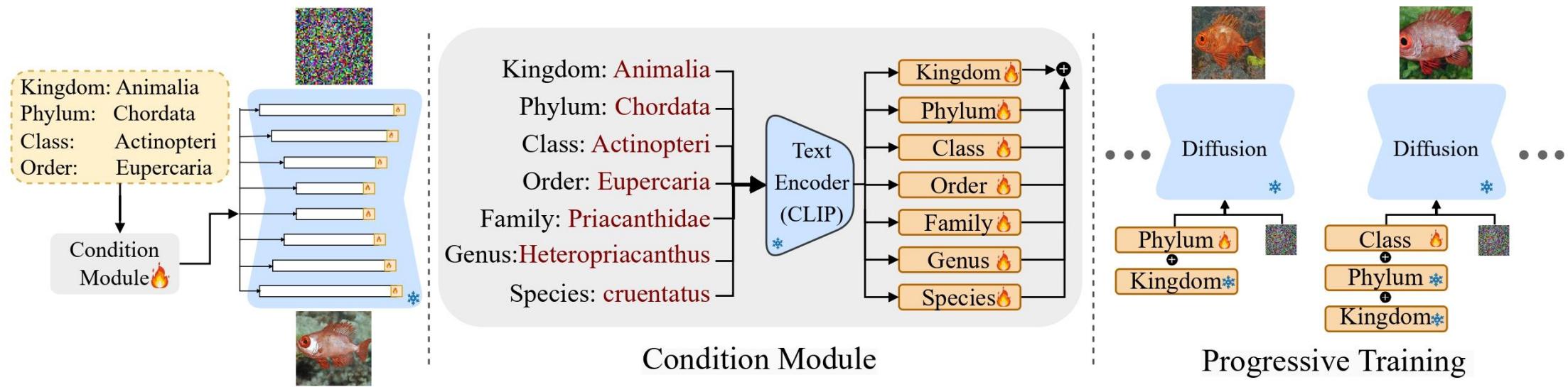
Key Insight - Leveraging Taxonomy

- Question: How can we use taxonomy in training diffusion models?



TaxaDiffusion

- Efficient domain adaptation using LoRA.
- Use of CLIP embeddings refined through transformer-based module.
- Progressive Training (Kingdom → Species).



TaxaDiffusion: Fine-grained Generation

- Classifier-Free Guidance (CFG):
 - Generation quality and diversity by refining predictions without a separate classifier.
 - Combines conditional and unconditional score estimates.

$$\tilde{\epsilon}_\theta(\mathbf{x}_t, t, \mathbf{c}) = (1 + w) \times \epsilon_\theta(\mathbf{x}_t, t, \mathbf{c}) - w \times \epsilon_\theta(\mathbf{x}_t, t),$$

- TaxaDiffusion CFG:

$$(1 + w) \times \epsilon_\theta(\mathbf{x}_t, t, \mathbf{c}^{(i)}) - w \times \epsilon_\theta(\mathbf{x}_t, t, \mathbf{c}^{(0)}),$$

Experimental Setup

Evaluation on 3 Datasets:

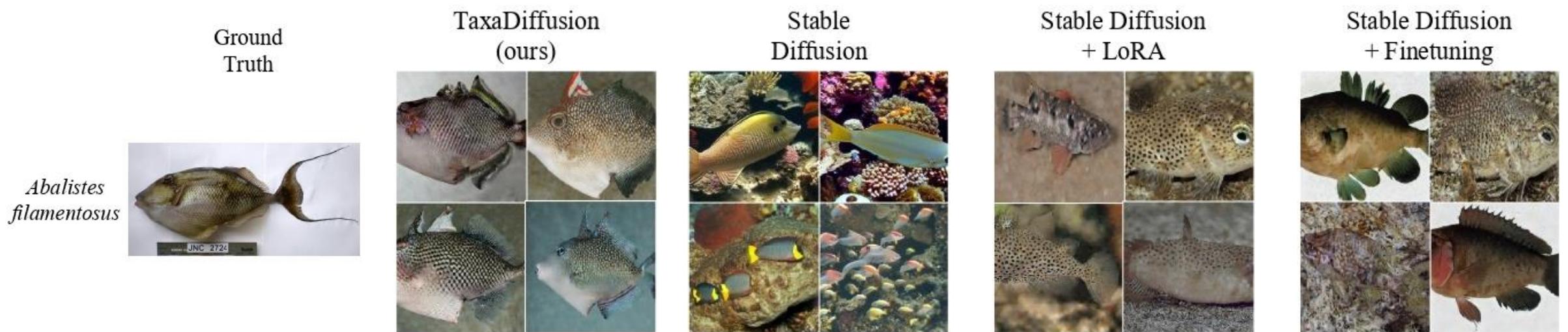
- **FishNet:** 17,357 distinct fish
- **BIOSCAN-1M:** 8,355 insect species
- **iNaturalist:** 10,000 species



Metrics:

- FID, LPIPS (image quality)
- BioCLIP (semantic alignment)

Results & Performance



Results & Performance

Table 1. **Quantitative results on FishNet [17]**. We report FID and LPIPS to assess image generation quality and BioCLIP for text-to-image alignment, where the text corresponds to the species “taxonomic name.” We generate 10 images per category at each level for evaluation, i.e., Order, Family, Genus, and Species. TaxaDiffusion outperforms the baselines for generating species-specific images.

Method	FID ↓				LPIPS ↓				BioCLIP - Score ↑			
	Order	Family	Genus	Species	Order	Family	Genus	Species	Order	Family	Genus	Species
SD [36]	74.98	67.46	65.12	61.93	0.7854	0.7859	0.7821	0.7737	7.48	8.12	8.15	3.35
SD + LoRA [14]	56.12	52.81	45.41	43.91	0.7698	0.7705	0.7617	0.7574	10.11	10.77	12.20	7.61
SD + Full [36]	50.72	47.35	41.72	39.41	0.7536	0.7616	0.7582	0.7574	11.83	11.98	13.74	8.31
TaxaDiffusion (ours)	41.92	40.16	27.35	31.87	0.7303	0.7324	0.7349	0.7319	15.06	16.67	18.42	10.43

Results & Performance

- Ablation of guidance:

TaxaDiffusion



Vanilla CFG



Results & Performance

- Importance of progressive training:

Strategy	FID ↓	LPIPS ↓	BioCLIP ↑
All	76.96	0.7273	5.66
Random	72.31	0.7178	6.68
Progressive (ours)	82.75	0.7097	7.87

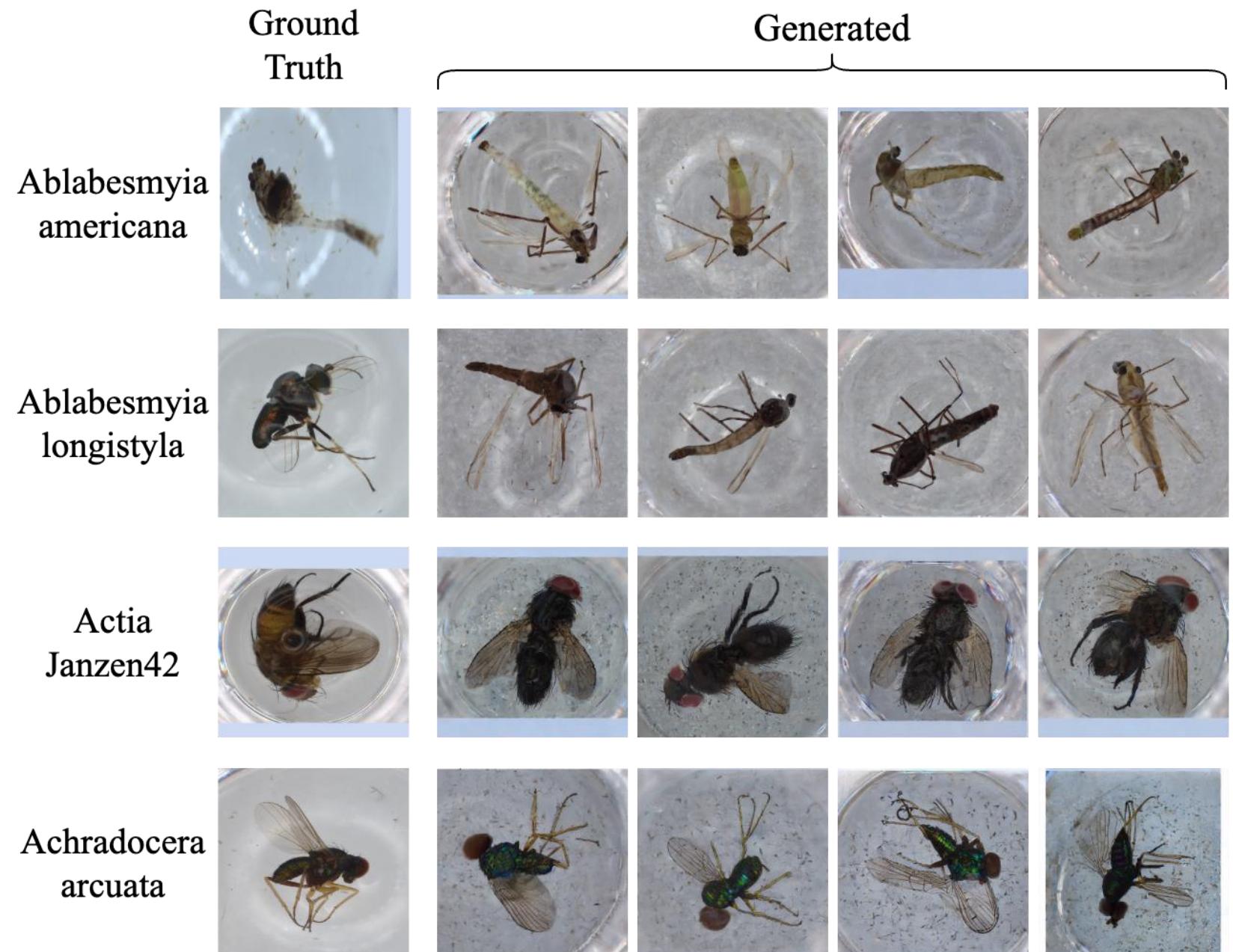
Results

- iNaturalist:



Results

- BIOSCAN-1M:



Conclusion

- **Taxonomy-Guided Training:** Learns coarse-to-fine traits by progressing through Kingdom→...→Species.
- **Rare-Species Transfer:** Generates high-fidelity images with minimal samples.

Questions

