



Building a Flexible Framework for Automated White Shark Re-Identification

Fabrice Kurmann¹, Connor Pryor¹, Charles Dickens¹, Alexandra E. DiGiacomo²,
Samantha Andrzejczek², Eriq Augustine¹, Barbara A. Block², Lise Getoor¹
University of California Santa Cruz¹, Stanford University²



Introduction and Motivation

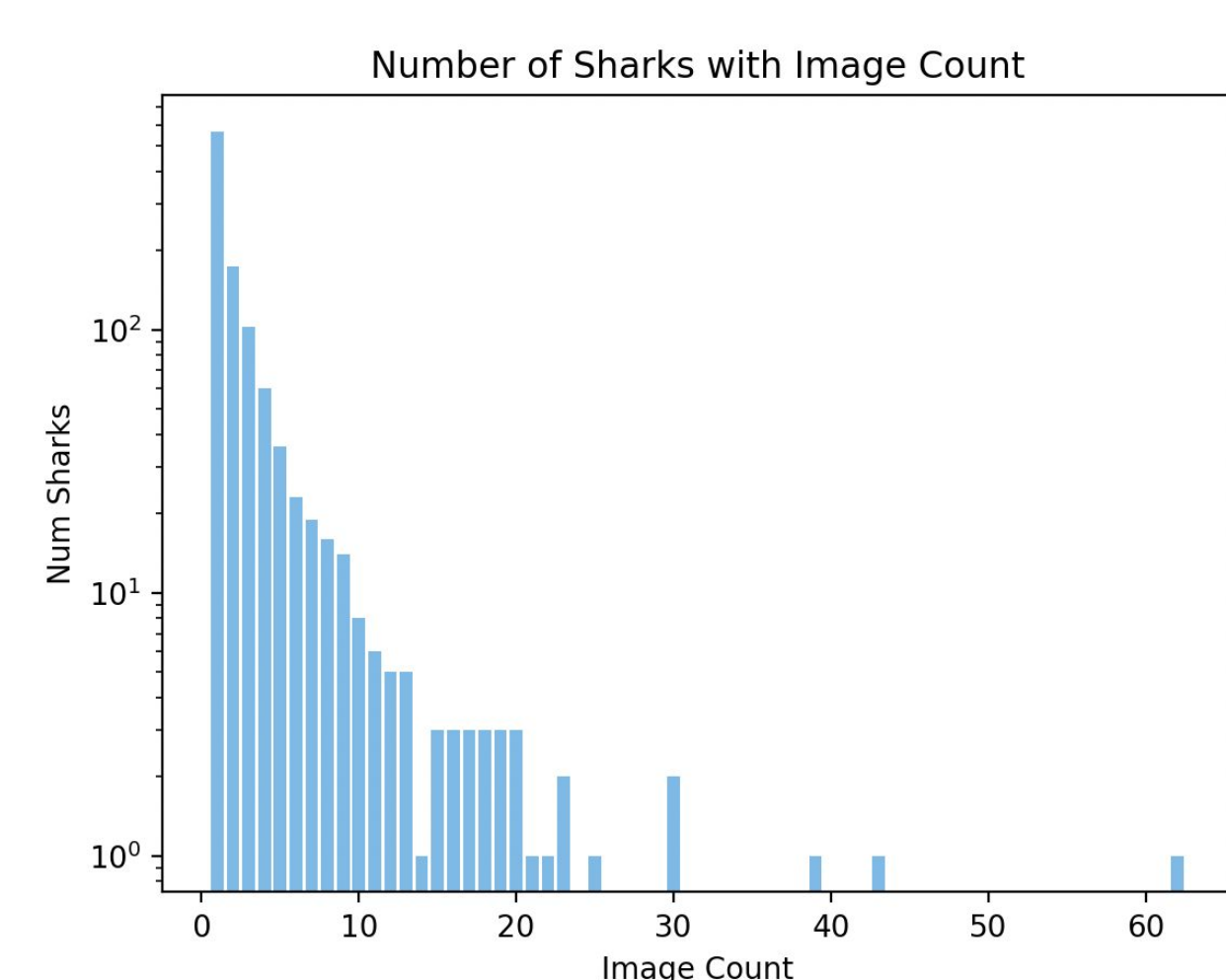
Animal Re-Identification (Re-ID), matching new observations against a catalog of known individuals, is essential for wildlife conservation, however manual re-identification is time-consuming and error-prone.

We introduce an automated white shark Re-ID framework designed to accelerate and improve this process.

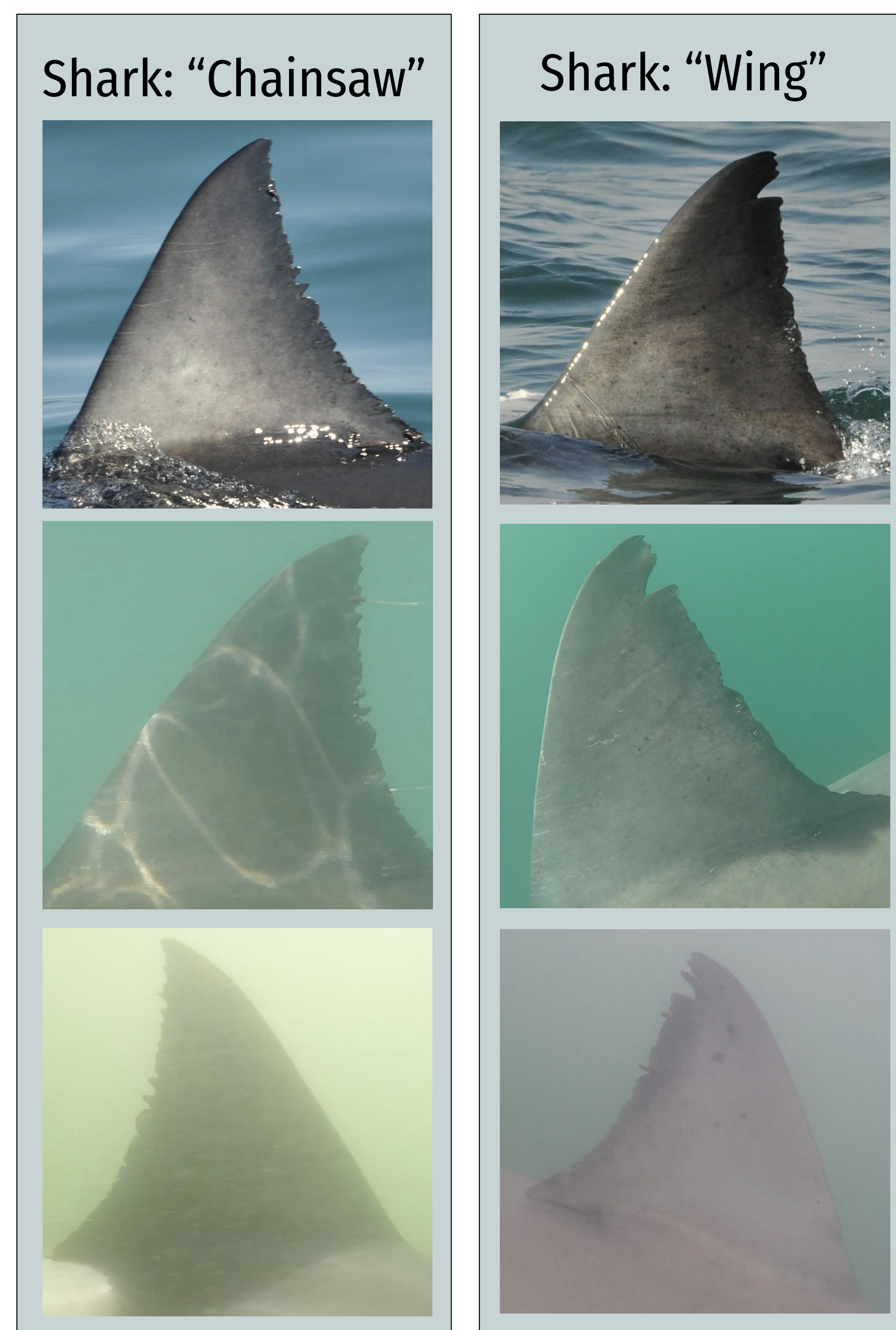
Our system leverages visual features of dorsal fins [1] while keeping humans in the loop for validation, enabling efficiency and reliability.

Dataset

- Their rarity, vast habitat, and difficulty to photograph creates a sparsely populated database of many sharks, each represented by few images
 - 3083 dorsal fin images
 - 1031 unique white sharks
 - 20+ year timespan



- Images include numerous confounding factors
 - Fin orientation, rotation, angle
 - Background color/lighting
 - Water surface, reflection, splash



Methods

- Image encoder maps images to embedding space**
 - Google/vit-large-patch16-384 feature extraction backbone [2]
 - Multilayer perceptron (MLP) projection head
 - Trained for:
 - Sensitivity to biomarkers on the dorsal fin (notches and pigmentation)
 - Invariance on confounding variables (pose, lighting, image quality)
- Retrieval algorithm searches embeddings space**
 - Well-trained model organizes embeddings into distinct clusters for each shark
 - Matching sharks are retrieved by searching the embedding space for nearest neighbors to a query image

Shark Re-Identification Framework

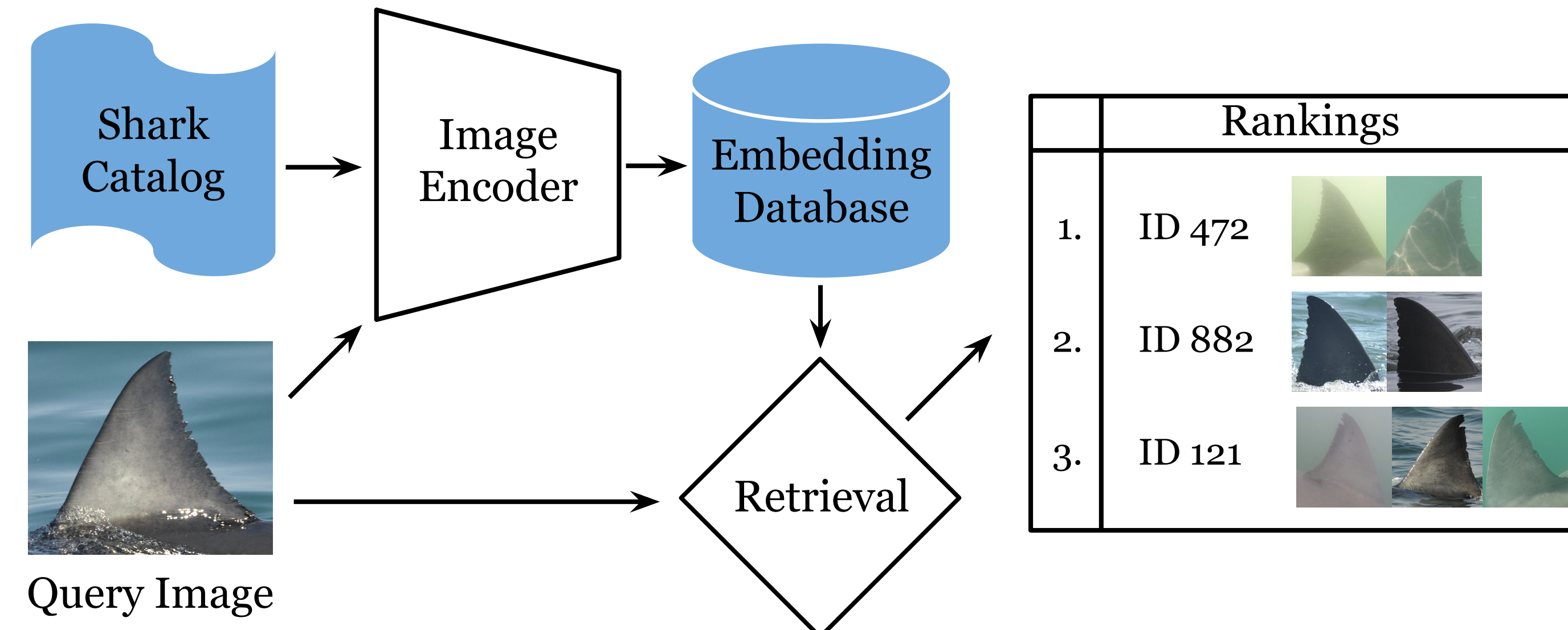


Image Encoder Training

- Training Approach:**
 - Triplet loss function* [3]:

$$\mathcal{L}(A, P, N) = \max(d(A, P) - d(A, N) + \alpha, 0)$$
 - Fine-tuning with low rank adaptation* [4]
 - Parameter efficient method allows convergence in 72 hours on single GPU
- Training Enhancements:**
 - Class-aware triplet sampling*
 - Sparse dataset: many training batches lack anchor-positive samples
 - Explicitly sample two positive samples for each anchor image
 - Training image augmentation*
 - Augmentations to perspective, rotation, scale, and color during training
 - Increase invariance to confounding traits in images
 - Reduce overfitting

Retrieval Techniques

Nearest Neighbor (NN)

- Rank all identities by their closest image embedding to the query

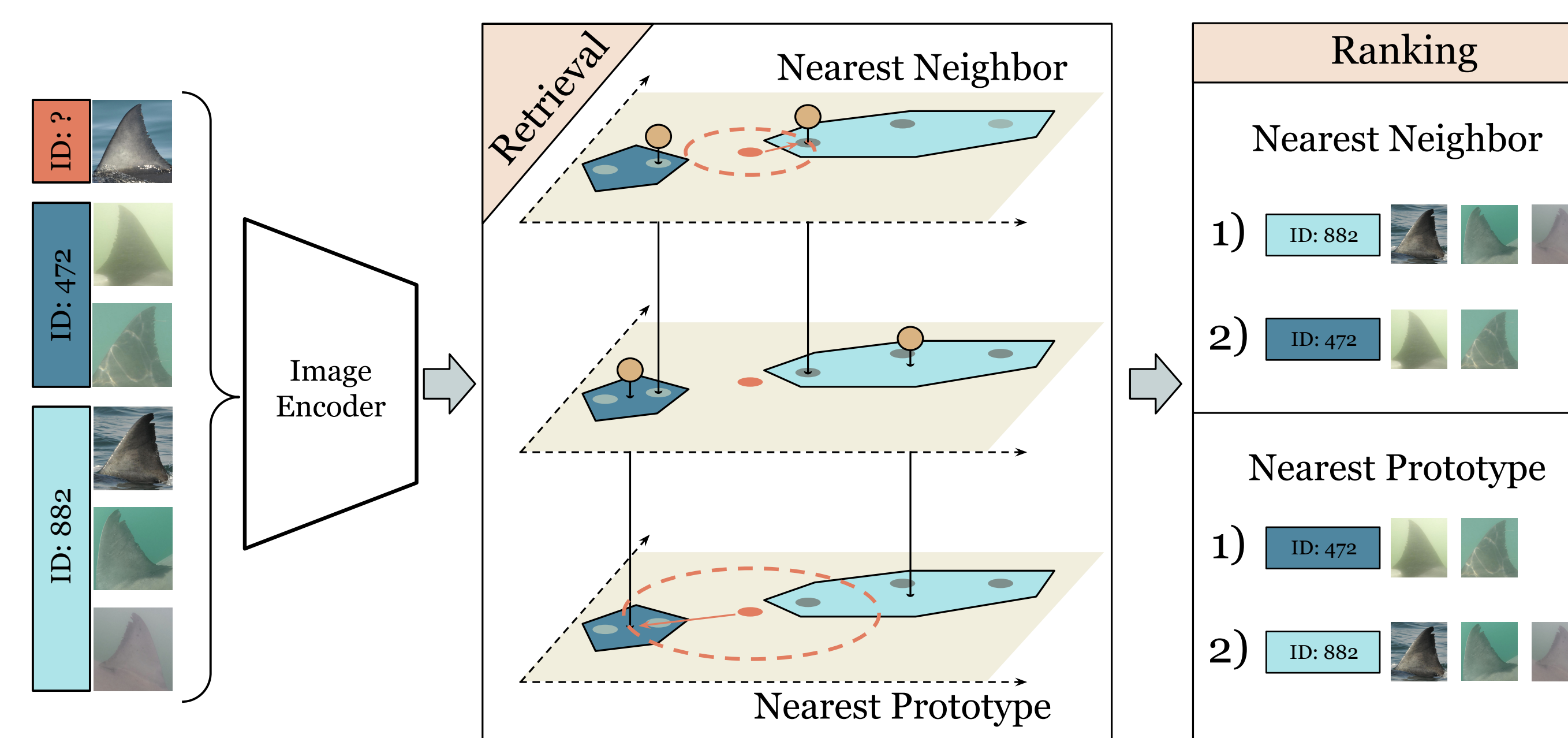
$$\hat{\mathcal{Y}}_{\text{NN}} = \text{rank} \left(y \in \mathcal{Y} \mid \min_{z \in \mathcal{Z}_y} d(z_q, z) \right)$$

Nearest Prototype (NP)

- Prototype embedding for each ID [5]

$$\mu_y = \frac{1}{|Z_y|} \sum_{z \in Z_y} z$$
- Rank by the closest prototype to the query

$$\hat{\mathcal{Y}}_{\text{NP}} = \text{rank} \left(y \in \mathcal{Y} \mid d(z_q, \mu_y) \right)$$



Results

- Hits@K scores
 - Proportion of queries where correct individual is among first k retrieved
 - K=50 represents a practical upper limit for human review
- Test set of 1 randomly selected image from each individual with > 1 image
 - 497 train, 2,586 test images

Model	Nearest Neighbor Retrieval			
	Hits@1	Hits@5	Hits@25	Hits@50
CNN	0.02	0.05	0.14	0.20
VIT	0.03	0.09	0.23	0.33
VIT + LoRA	0.09	0.23	0.40	0.53
VIT + LoRA + CAS	0.31	0.55	0.74	0.83
VIT + LoRA + CAS + Aug	0.48	0.67	0.85	0.90

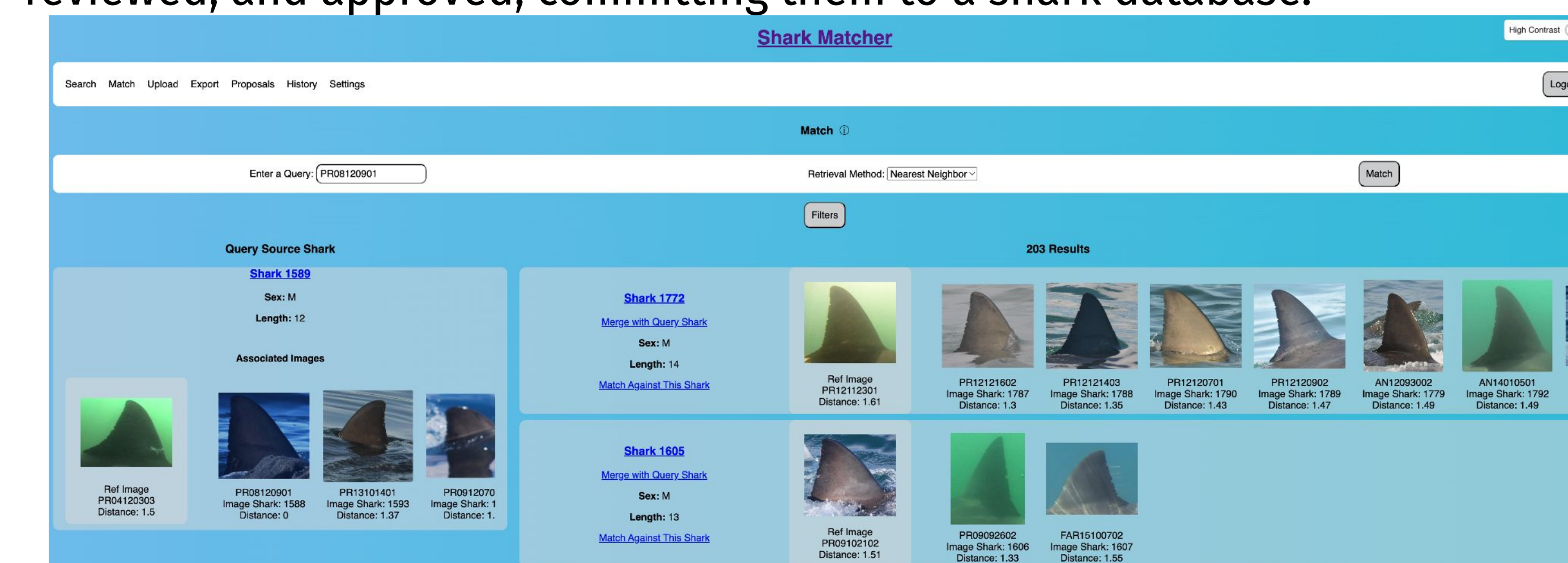
Model	Prototype Retrieval			
	Hits@1	Hits@5	Hits@25	Hits@50
CNN	0.02	0.07	0.15	0.22
VIT	0.05	0.13	0.30	0.38
VIT + LoRA	0.11	0.23	0.43	0.53
VIT + LoRA + CAS	0.33	0.56	0.75	0.82
VIT + LoRA + CAS + Aug	0.51	0.68	0.84	0.91

- Ablation shows breakdown of augmentation technique.
 - All techniques aside from random erasing improve results

Augmentations	Nearest Neighbor Retrieval			
	Hits@1	Hits@5	Hits@25	Hits@50
None	0.31	0.55	0.74	0.83
Geometric	0.45	0.65	0.83	0.89
Geo. + Color	0.48	0.67	0.85	0.90
Geo. + Color + Erase	0.35	0.57	0.74	0.83

Shark Matcher Human-in-the-Loop UI

Collaborating with marine biologists, we developed a UI tailored for their labeling workflow. Our model's match results can be browsed, filtered, reviewed, and approved, committing them to a shark database.



Conclusion and Future Work

We develop a framework for white shark Re-ID, presenting optimizations to model training and retrieval technique, showing their benefits to retrieval accuracy. Paired with our UI, this work has become a valuable tool for dataset de-duplication and matching newly captured shark images for our marine biologist collaborators.

As future work, we aim to:

- Evaluate over different animal species to better understand generalization
- Improve model adaptation with online learning of newly added data
- Develop techniques for training accurate models on noisy and mislabeled real-world data

Acknowledgements

This work was partially supported by the NSF grant CCF-2023495.

[1] Nowacek, Christiansen, Bejder, Goldbogen, Friedlaender, Studying cetacean behaviour: new technological approaches and conservation apps. (2016).
 [2] Dosovitskiy, Beyer, Kolesnikov, Weissenborn, Zhai, Unterthiner, Dehghani, Minderer, Heigold, Gelly, Uszkoreit, and Houslay. An image is worth 16x16 words: Transformers for image recognition at scale. arXiv (2021).
 [3] Schroff, Kalenichenko, and Philbin. Facenet: A unified embedding for face recognition and clustering. CVPR (2015).
 [4] Hu, Shen, Wallis, Zhu, Li, Wang, Wang, and Chen. Lora: Low-rank adaptation of large language models. arXiv (2021).
 [5] Rocchio. Relevance feedback in information retrieval. (1971).