

Self-Supervised Pretraining for Fine-Grained Plankton Recognition

Supplementary Material

S1. Pretraining dataset class overlap

The class overlaps with different datasets are shown in Table 1. In total, there are not many overlapping classes between the dataset except with DAPlankton and SYKE-Plankton_IFCB_2022, which share most of their classes.

Table 1. Overlapping Classes Between Plankton Datasets

Dataset Pair	Overlapping Classes	Count
DAPlankton [1] & SYKE-Plankton_IFCB_2022 [5]	aphanizomenon-flosaqueae, centrales-sp, chaetoceros-sp, chaetoceros-sp-single, chlorococcales, chroococcales ciliata, cryptomonadales, cryptophyceae-teleaulax cyclotella-choctawhatcheana, dinophyceae, dinophysis-acuminata dolichospermum-anabaenopsis, dolichospermum-anabaenopsis-coiled euglenophyceae, eutreptiella-sp, gymnodiniales, gymnodinium-like, heterocapsa-rotundata, heterocapsa-triquetra heterocyte, katablepharis-remigera, melosira-arctica mesodinium-rubrum, monoraphidium-contortum, nitzschia-paleacea nodularia-spumigena, oocystis-sp peridiniella-catenata (as peridiniella-catenata-single/chain in SYKE) pseudopediella-sp, pyramimonas-sp, skeletonema-marinoi snowella-woronichinia	33
DAPlankton [1] & PMID2019 [7]	chaetoceros (chaetoceros-sp in DA), skeletonema (skeletonema-marinoi in DA)	2
SYKE-Plankton_IFCB_2022 [5] & PMID2019 [7]	chaetoceros (chaetoceros-sp in SYKE), skeletonema (skeletonema-marinoi in SYKE)	2
PMID2019 [7] & UDE Diatoms [4]	navicula	1
SYKE-Plankton_ZooScan_2024 [3] & Kaggle Plankton [2]	copepod-calanoid, copepod-cyclopoid (multiple cyclopoid in Kaggle)	2
SYKE-Plankton_ZooScan_2024 [3] & Lake Zooplankton [6]	daphnia, synchaeta (daphnia-sp, synchaeta-sp in SYKE)	2
Lake Zooplankton [6] & Kaggle Plankton [2]	diatom-chain (diatom-chain-string/tube in Kaggle)	1

S2. Results

The fine-tuning results for DAPlankton_{LAB} and DAPlankton_{SEA} are shown in Table 2 and Table 3, respectively. For DAPlankton_{SEA}, the full fine-tuning results look very similar to DAPlankton_{LAB}, and there is no real difference between the different pretraining methods.

Table 2. Mean accuracy and standard deviation (in %) for DAPlankton_{LAB} with limited training labels.

Model	IFCB			CS			FC		
	1%	5%	10%	1%	5%	10%	1%	5%	10%
ViT-L (imagenet)	78.08 ± 3.54	95.23 ± 0.25	97.51 ± 0.26	62.87 ± 7.45	91.63 ± 1.12	95.22 ± 0.76	52.04 ± 4.39	91.72 ± 0.34	94.31 ± 0.35
ViT-L (no-daplankton)	83.83 ± 2.03	95.68 ± 0.35	97.41 ± 0.23	69.90 ± 4.73	90.30 ± 0.80	93.57 ± 0.47	79.58 ± 2.58	92.87 ± 0.33	95.00 ± 0.48
ViT-L (with-daplankton)	96.92 ± 0.37	98.29 ± 0.30	98.75 ± 0.24	91.76 ± 0.89	96.26 ± 0.52	96.97 ± 0.33	83.10 ± 1.44	93.86 ± 0.65	95.72 ± 0.64

Table 3. Mean accuracy and standard deviation (in %) for DAPlankton_{SEA} with limited training labels.

Model	IFCB				CS			
	1%	5%	10%	Full	1%	5%	10%	Full
ViT-L (imagenet)	86.47 ± 1.16	96.82 ± 0.19	97.90 ± 0.13	98.91 ± 0.04	53.92 ± 6.44	85.68 ± 2.17	91.29 ± 0.59	95.80 ± 0.44
ViT-L (no-daplankton)	89.43 ± 0.16	96.35 ± 0.30	97.54 ± 0.17	98.84 ± 0.10	72.65 ± 2.72	87.32 ± 0.54	90.49 ± 0.79	95.32 ± 0.06
ViT-L (with-daplankton)	95.44 ± 0.36	97.79 ± 0.18	98.24 ± 0.14	98.85 ± 0.06	85.89 ± 0.24	91.70 ± 0.54	93.23 ± 0.49	95.57 ± 0.24

References

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