

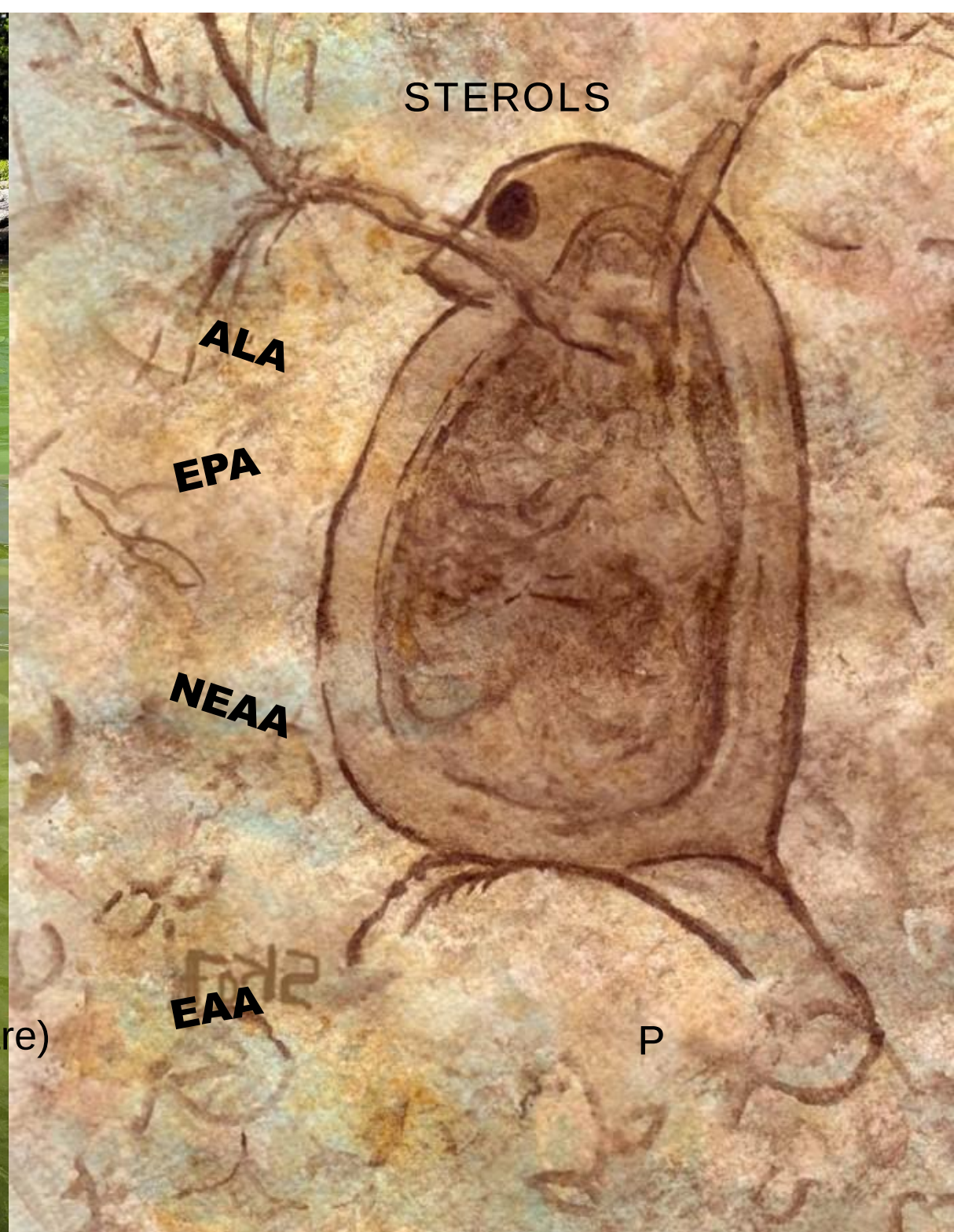


Eutrophication and lake restoration influence on nutritional quality of food webs

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Photo by Kirsi Kuoppamäki



ESSENTIAL BIOMOLECULES

- Nutritionally essential biomolecules are defined as compounds that cannot be synthesized *de novo* in animals or as that are insufficiently synthesized *de novo* for optimal growth and reproduction.

ESSENTIAL BIOMOLECULES FOR ZOOPLANKTON

CAROTENOIDS

• ASTAXANTHIN

β-CAROTENE

ECHINENONE

CANTHAXANTHIN

STEROLS

• CHOLESTEROL

PHYTOSTEROLS

HIGH EFFICIENCY

(Fucosterol, brassica-sterol, stigma-terol)

LOW EFFICIENCY

(b-sitosterol, desmosterol, chondrilla-sterol)

FATTY ACIDS

ω-3 PUFA

DHA (22:6w3)

EPA (20:5w3)

SDA (18:4w3)

ALA (18:3w3)

ω-6 PUFA

ARA (20:4w6)

DPA (22:5w6)

GLA (18:3w6)

LIN (18:2w6)

AMINO ACIDS

Arginine*

Histidine*

Isoleucine

Leucine

Lysine

Methionine

Phenylalanine

Threonine

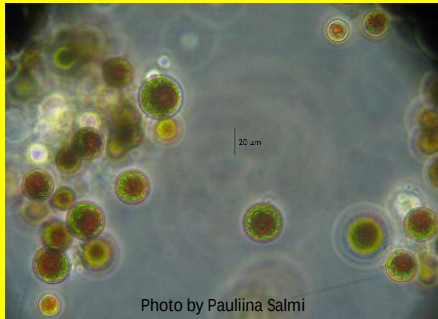
Tryptophan

Valine

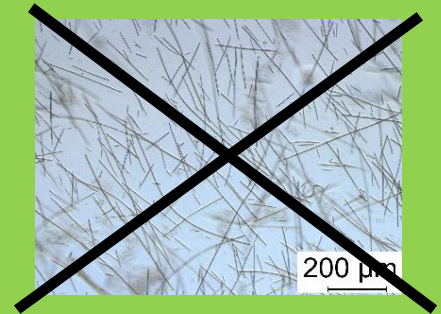
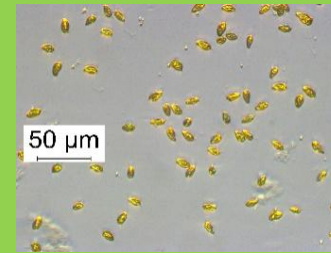
ESSENTIAL BIOMOLECULES

- In freshwater systems essential biomolecules are synthesized by phytoplankton.

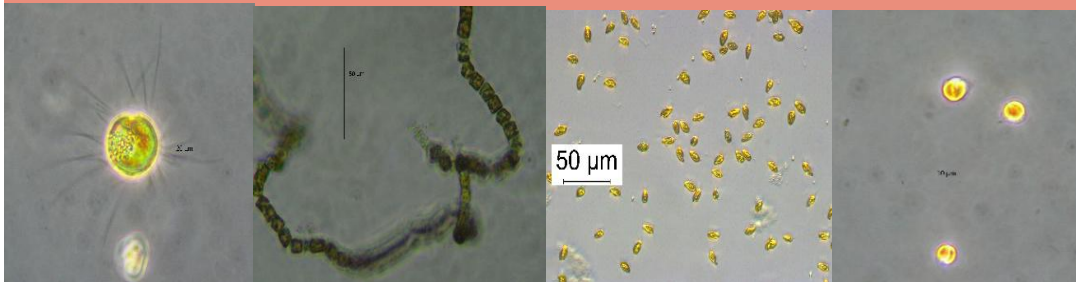
- ASTAXANTHIN IS SYNTHESIZED ONLY BY GREEN ALGAE BUT B-CAROTENE IS SYNTHESIZED BY ALL PHYTOPLANKTON TAXA.



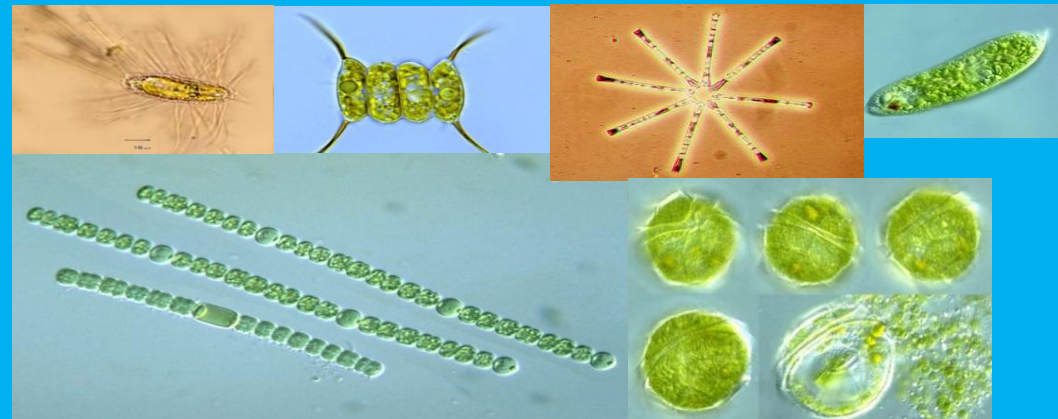
- STEROLS ARE SYNTHESIZED ALL PHYTO TAXA BUT NOT BY CYANOBACTERIA. CRYPTOPHYTES ARE BEST SOURCE FOR STEROLS.



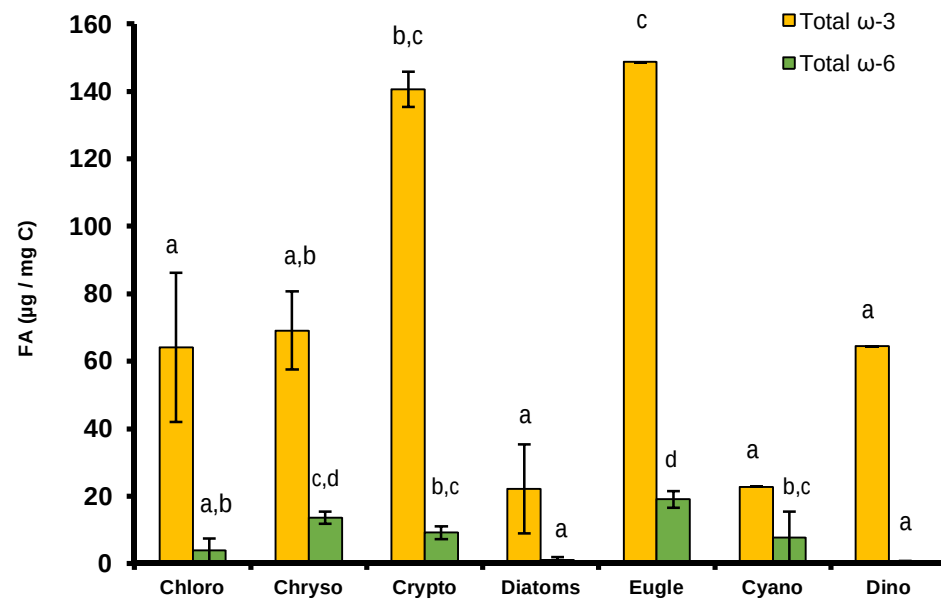
- ALA (18:3W3) AND LIN (18:2W6) ARE SYNTHESIZED BY ALL PHYTOPLANKTON TAXA, BUT EPA (20:5W3) AND DHA (22:6W3) ARE SYNTHESIZED BY ONLY FEW TAXA.



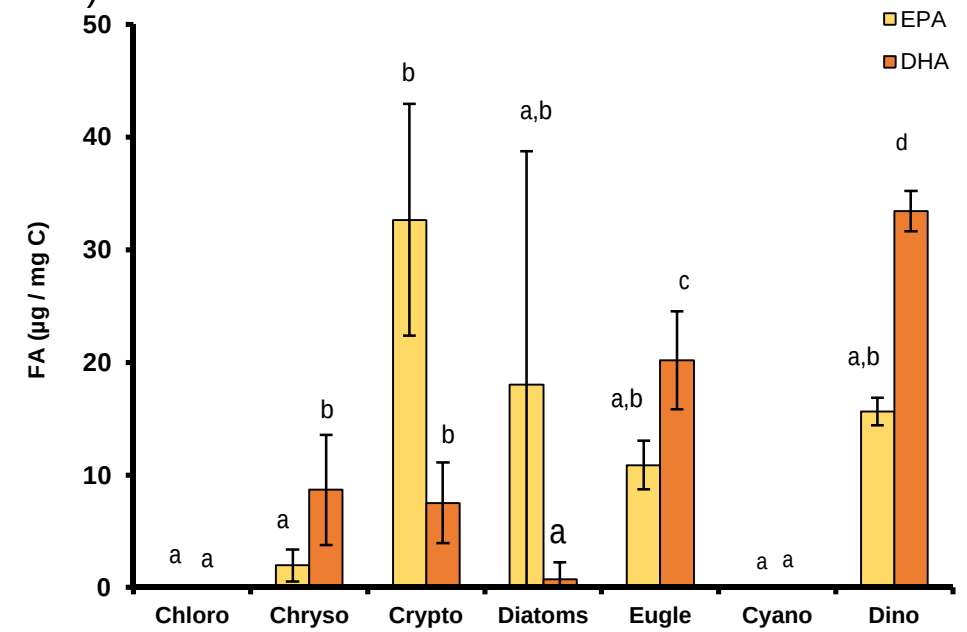
- AMINO ACIDS ARE SYNTHESIZED BY ALL PHYTOPLANKTON TAXA AS WELL BY BACTERIA, BUT CONCENTRATIONS VARY.



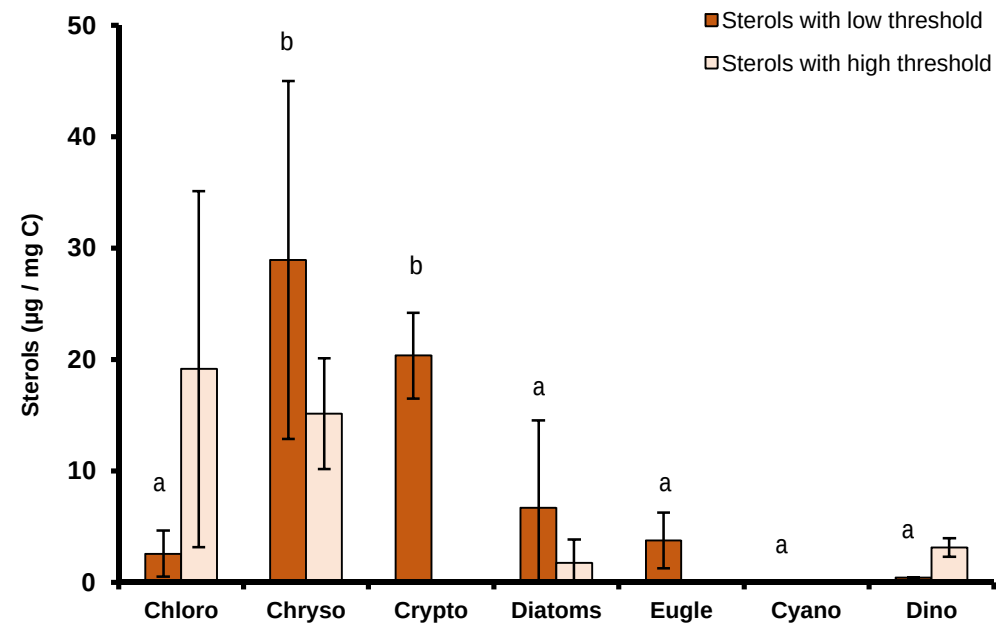
A)

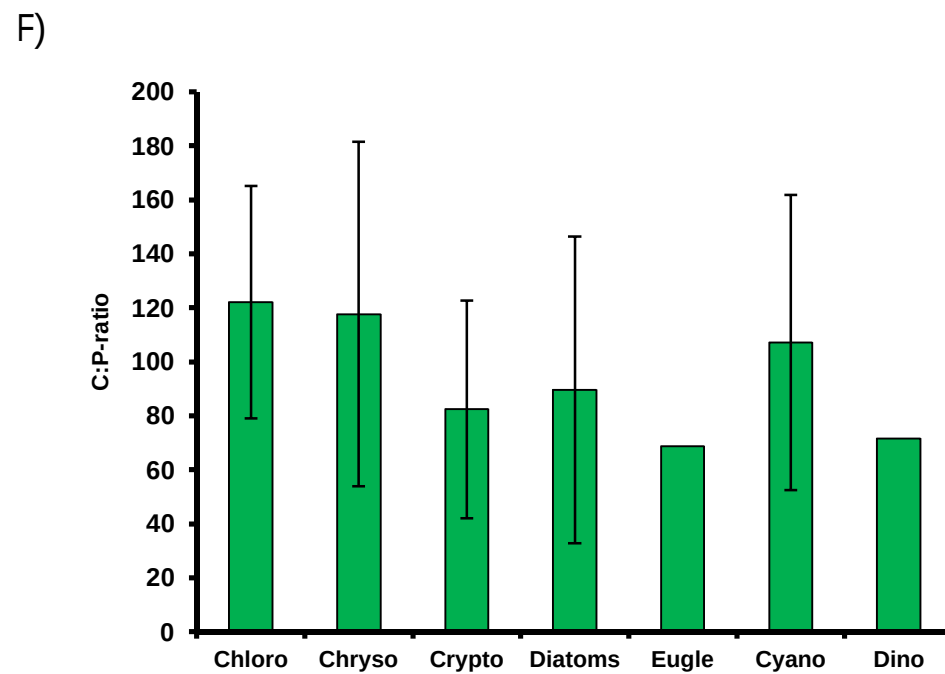
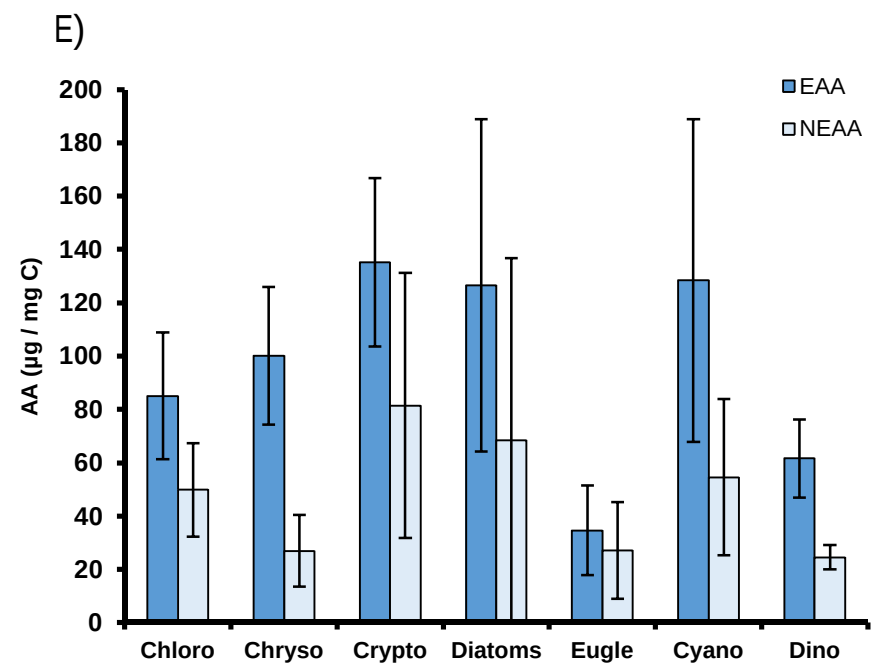
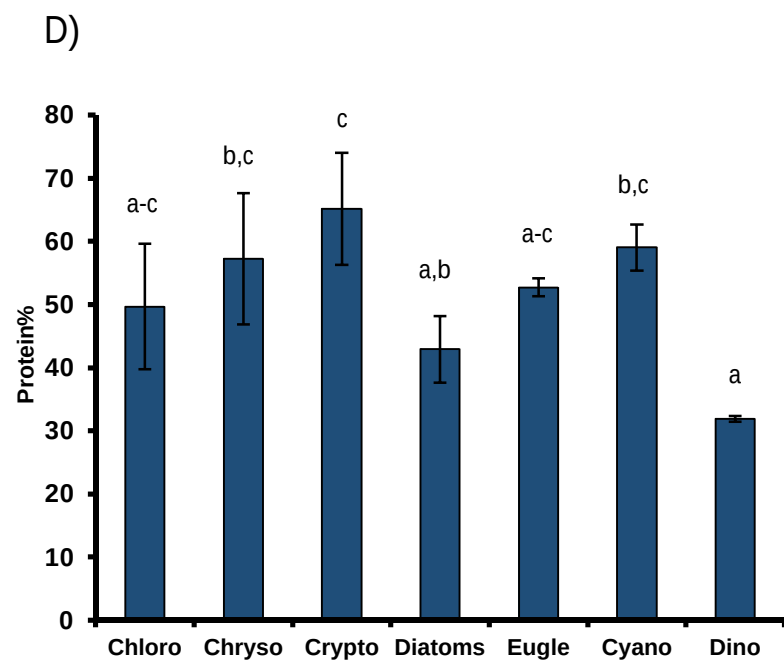


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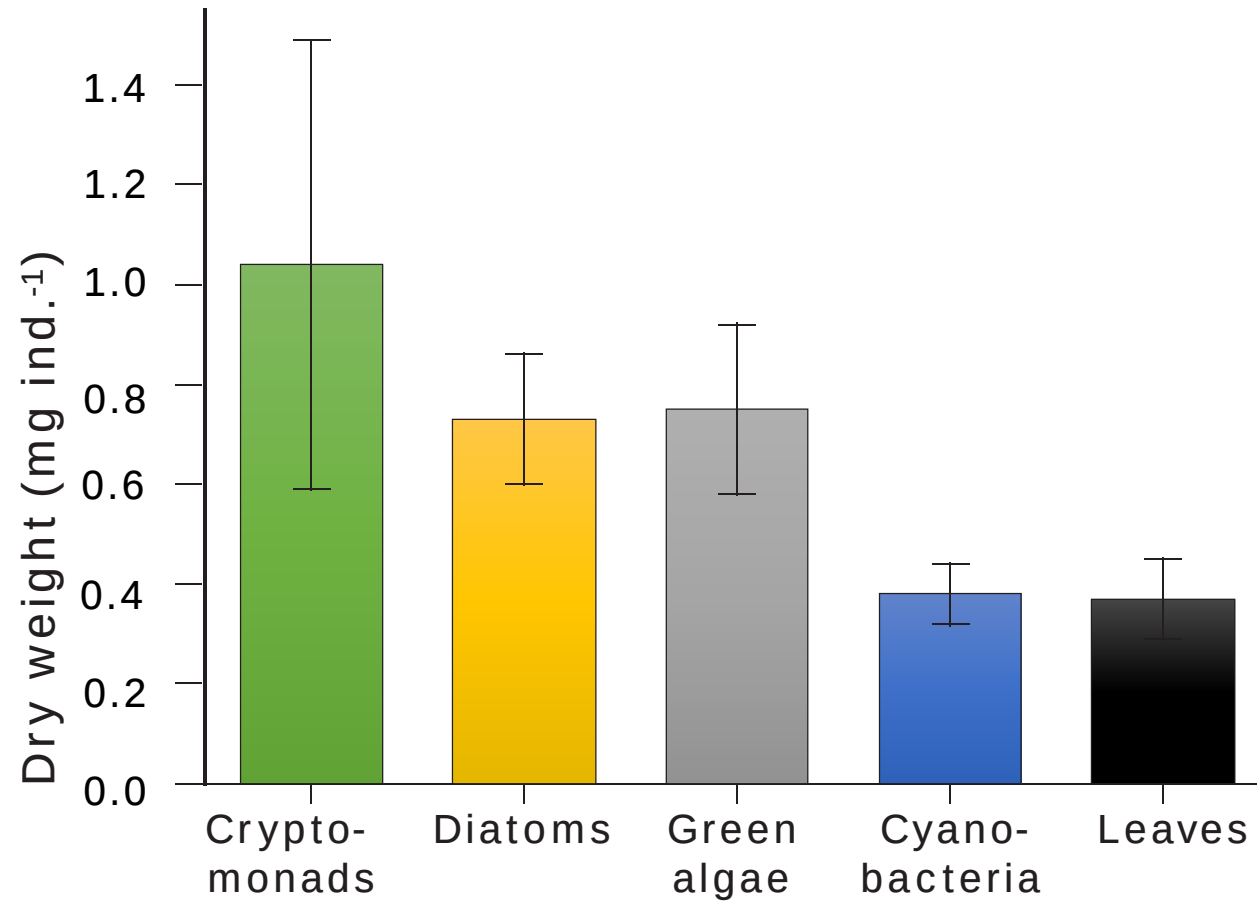


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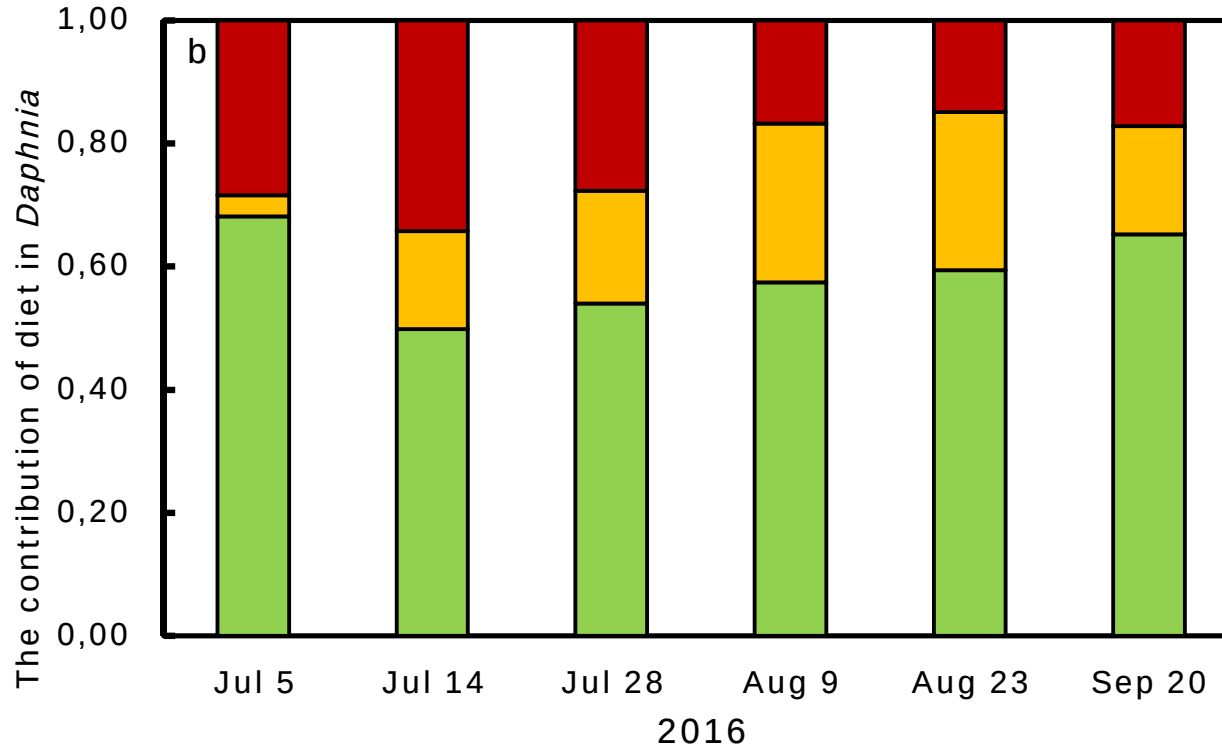
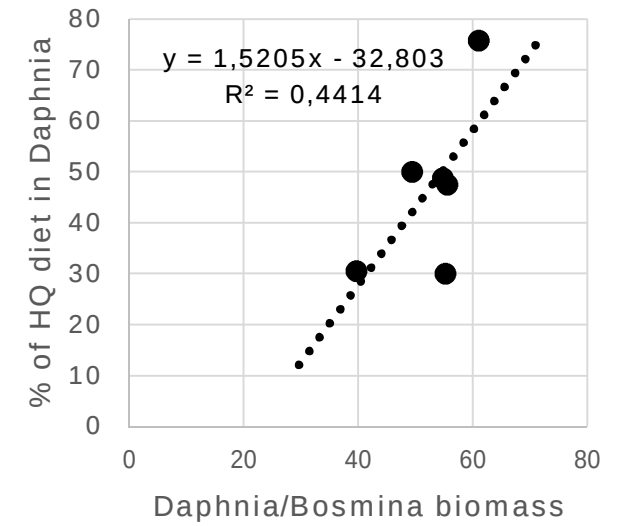
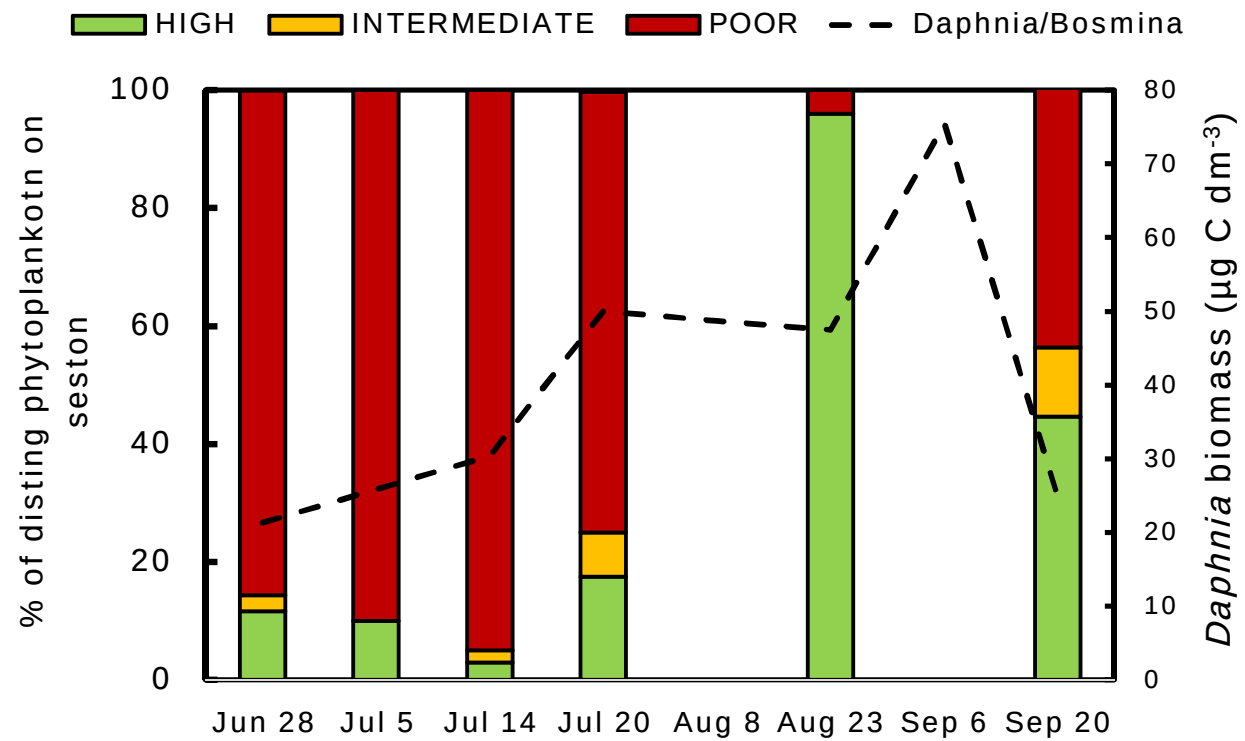


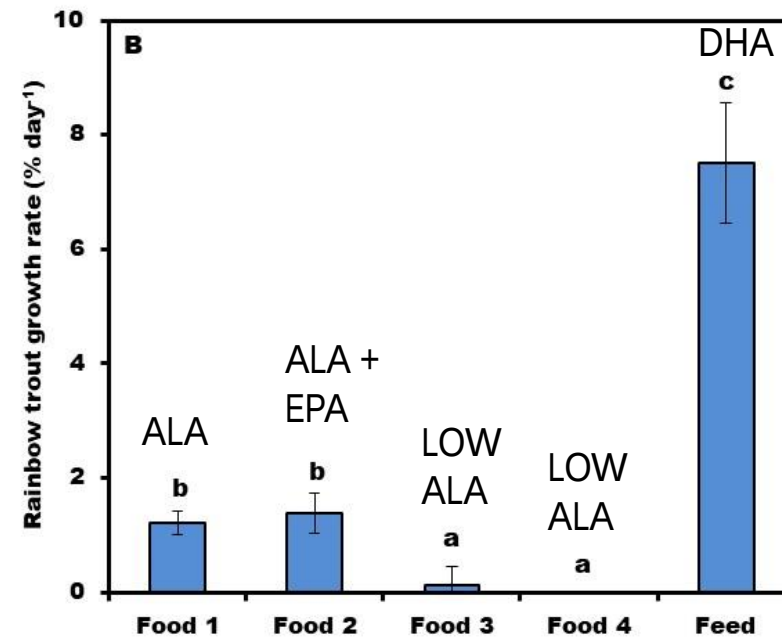
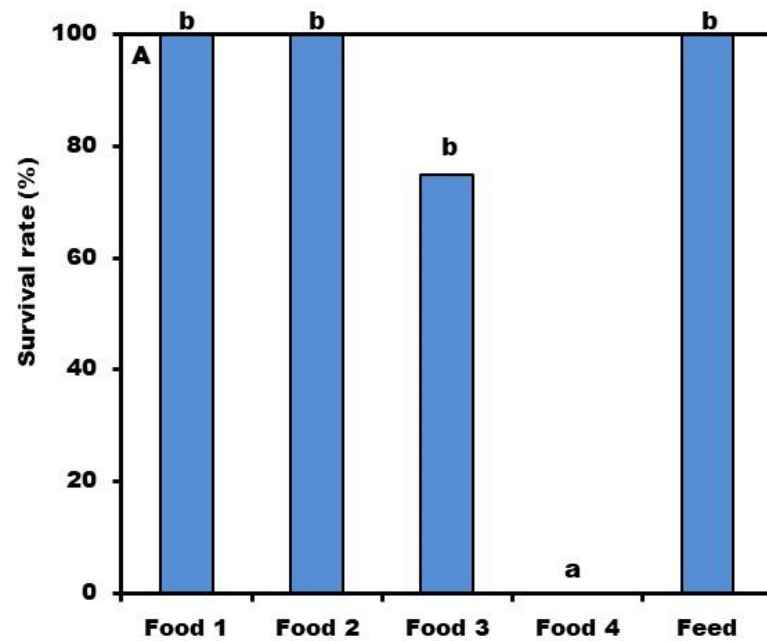


Zooplankton growth is regulated by w-3 and sterol content of phytoplankton.

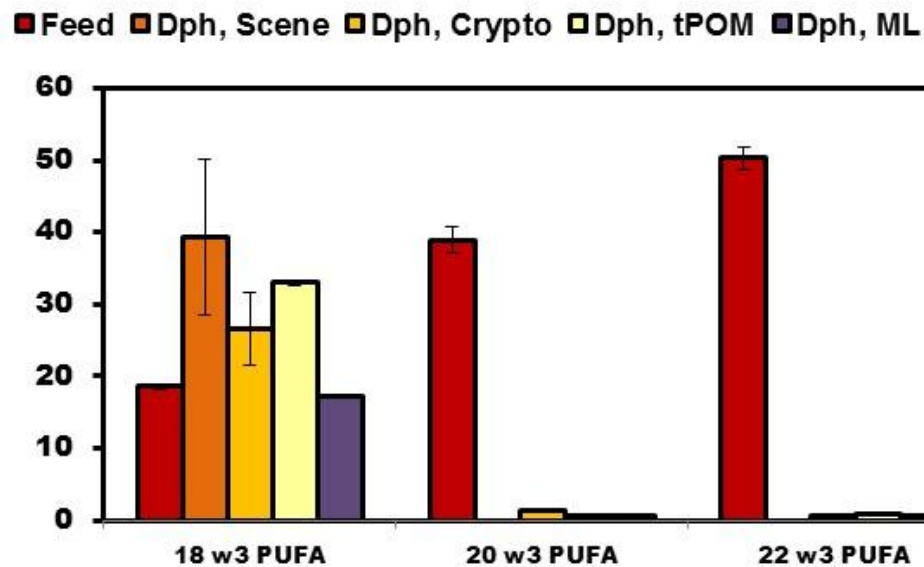


Brett et al. 2009. PNAS vol. 106 no. 50 21197-21201.

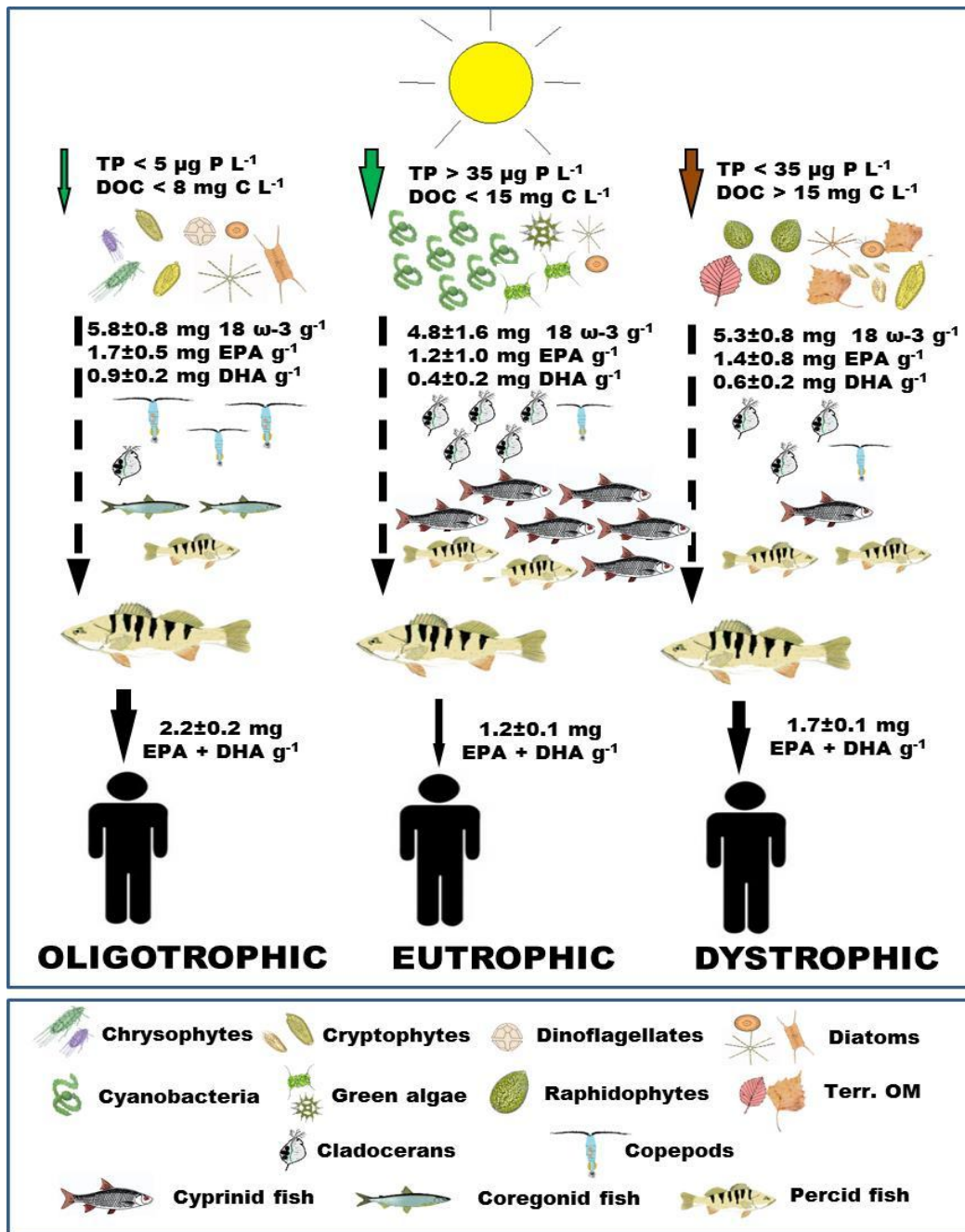




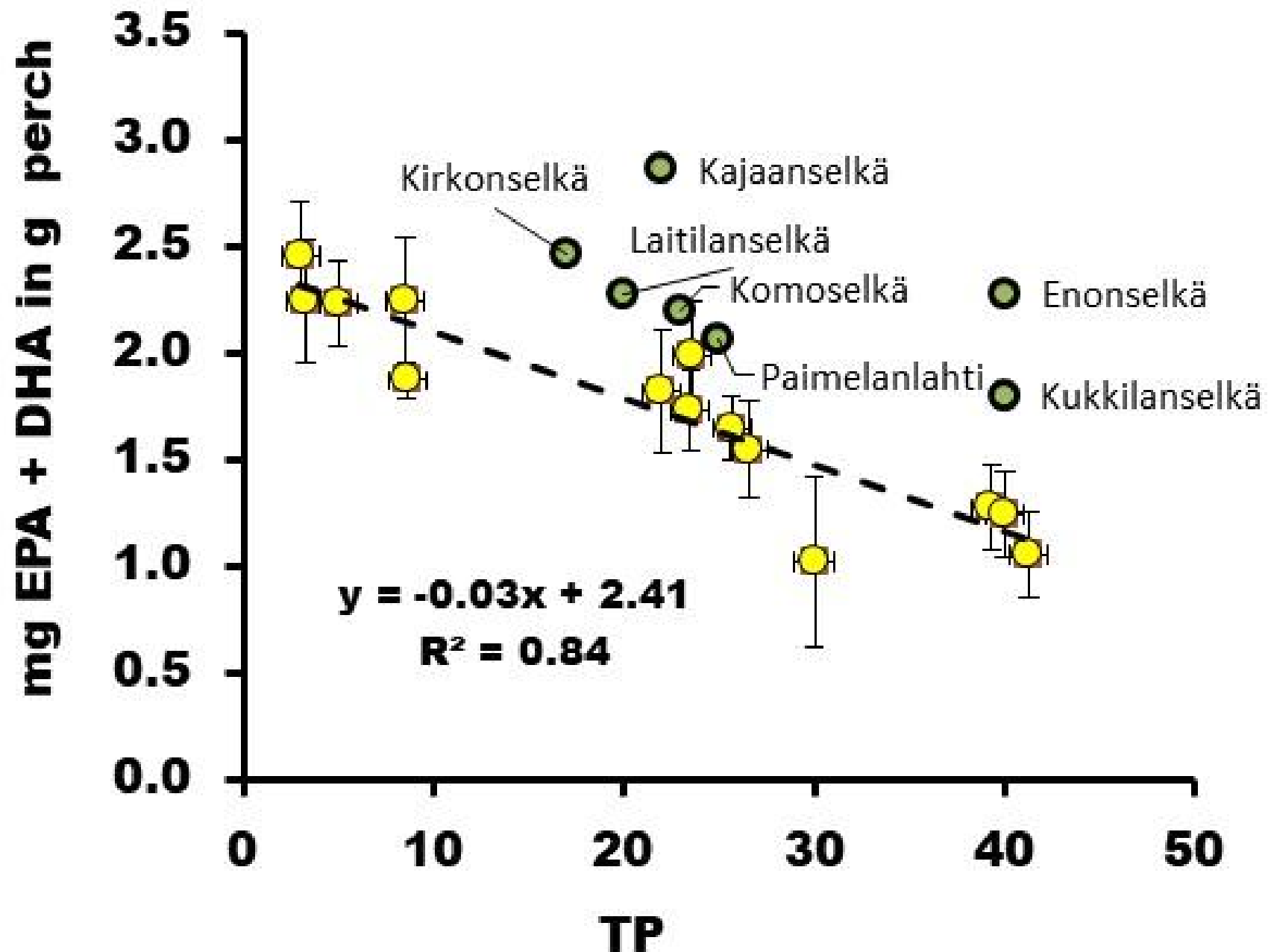
Taipale et al. Ecol and Evol 2018.



- Fish require DHA (and ARA) for disease resistance, neural tissue and eye development, pigmentation and reproduction (Sargent et al. 1999).
- Juvenile rainbow trout were not able to grow on *Daphnia* with low ALA content and grew moderately with high ALA.



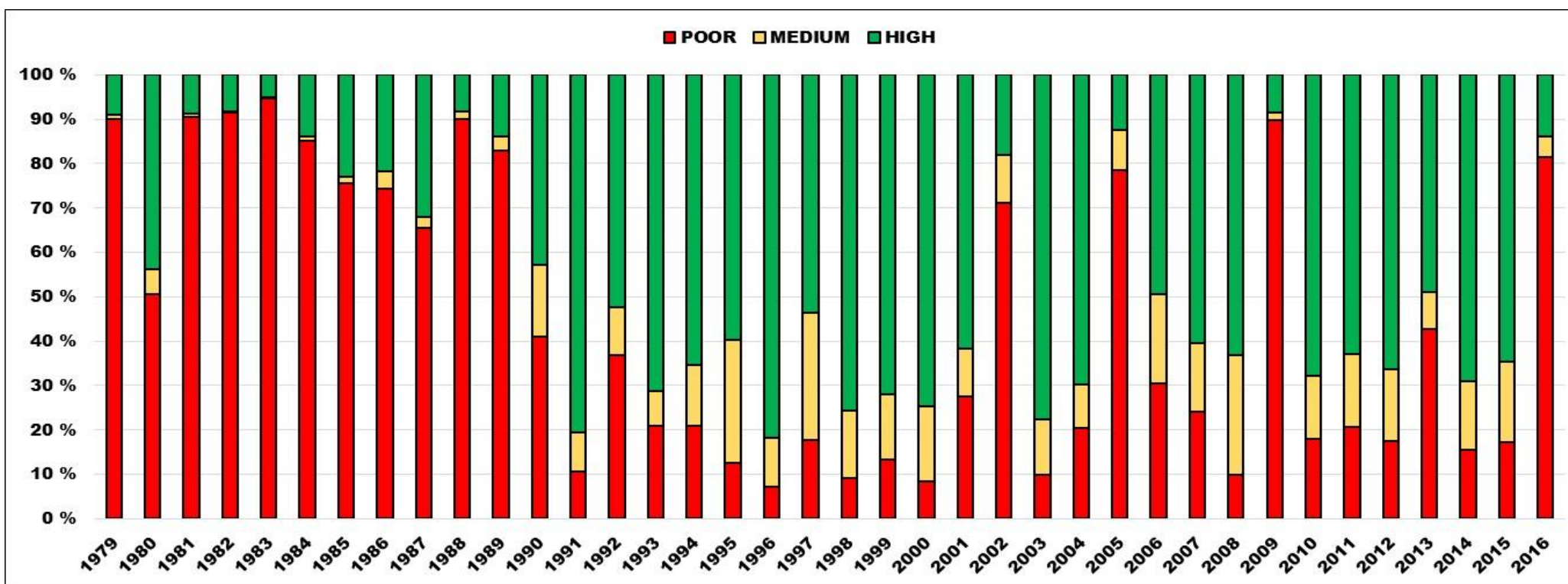
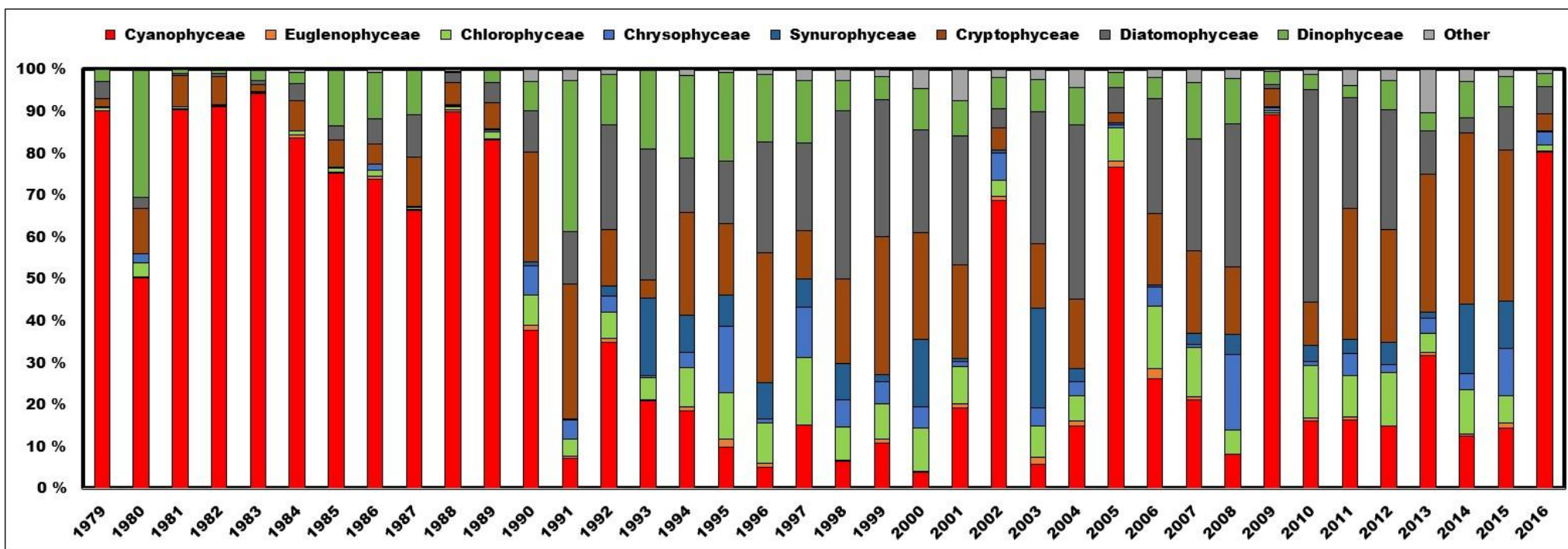
- The ω-3 FA is transferred via herbivorous zooplankton and planktivorous fish to piscivorous fish (in this case large perch).
- TP and DOC concentration influence phytoplankton structure and synthesis of ω-3 FA .
- The EPA and DHA content in individual perch is the highest in the oligo-mesotrophic lakes. Thus, a person should eat 1.9 and 1.5 times more perch from eutrophic and dystrophic lakes compared with those from oligotrophic lakes to achieve the daily recommended intake of EPA and DHA.



1. HOW BIOMANIPULATION AND HYPOLIMENTIC AEROTATION INFLUENCE PHYTOPLANKTON COMPOSITION?

2. DOES THEY ENHANCE EPA- AND DHA-SYNTHESIZING PHYTOPLANKTON TAXA?

- Phytoplankton composition from Lake Vesijärvi 1979-2016
- Using modern statistical test: non-metric multidimensional scaling (NMS), permutational multivariate analysis of variance (PERMANOVA), similarity percentages (SIMPER) and test of functional similarity (PERMDISP).
- The sestonic w-3, EPA, DHA and sterol content was calculated based on phytoplankton biomass and laboratory cultures of distinct phytoplankton taxa.



Non-metric Multi-Dimensional Scaling

Phytoplankton composition

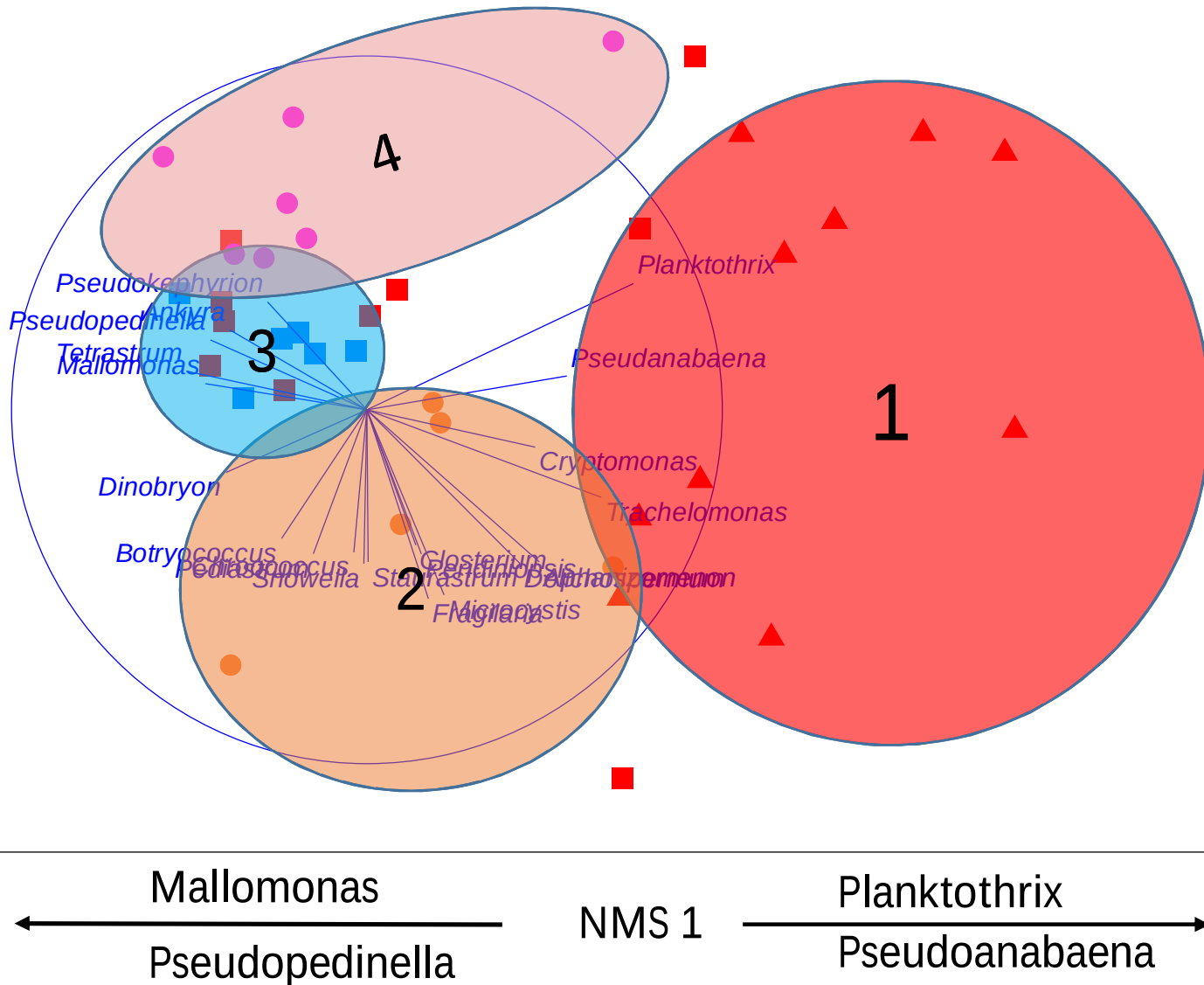
Resemblance: S17 Bray Curtis similarity

2D Stress: 0.14

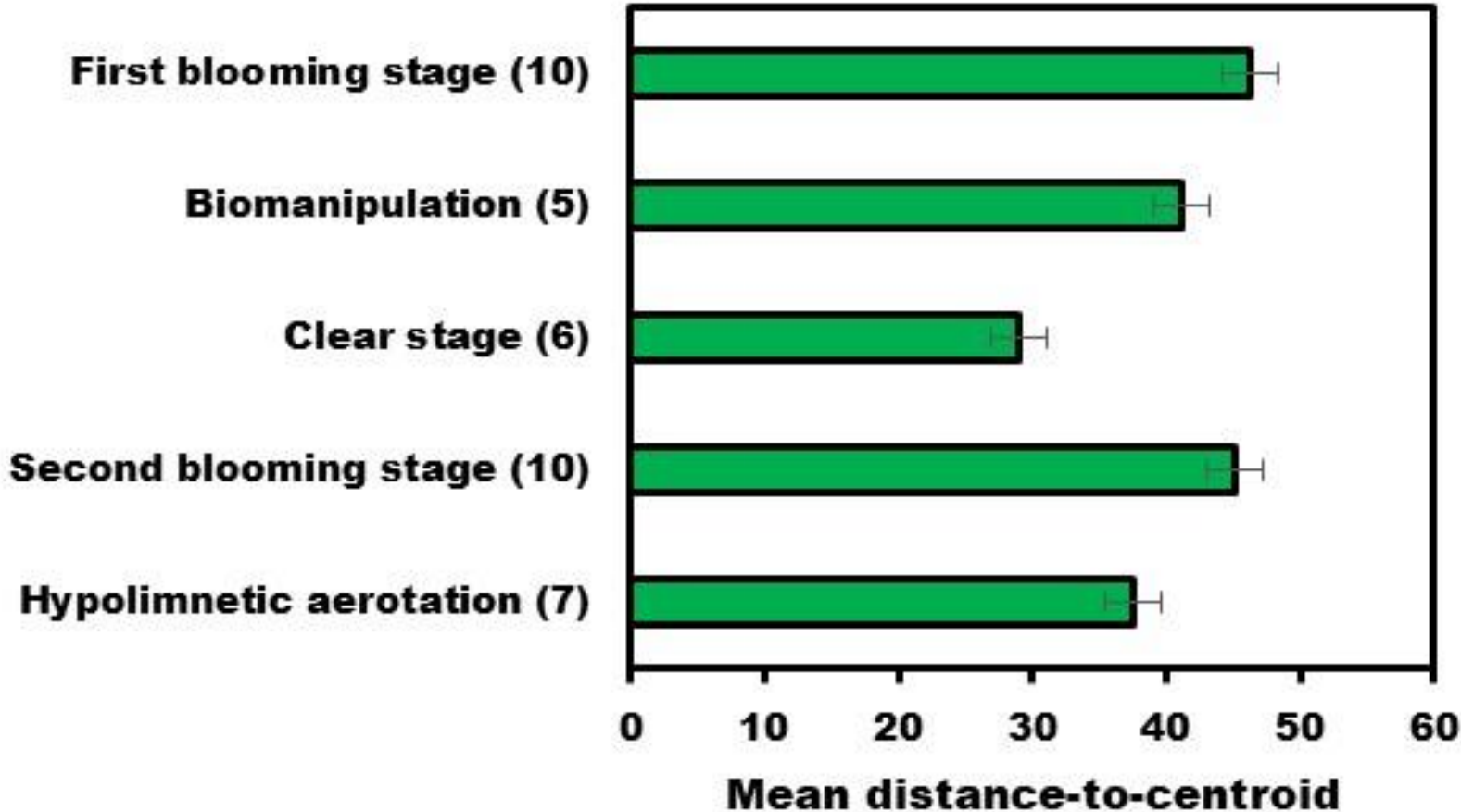
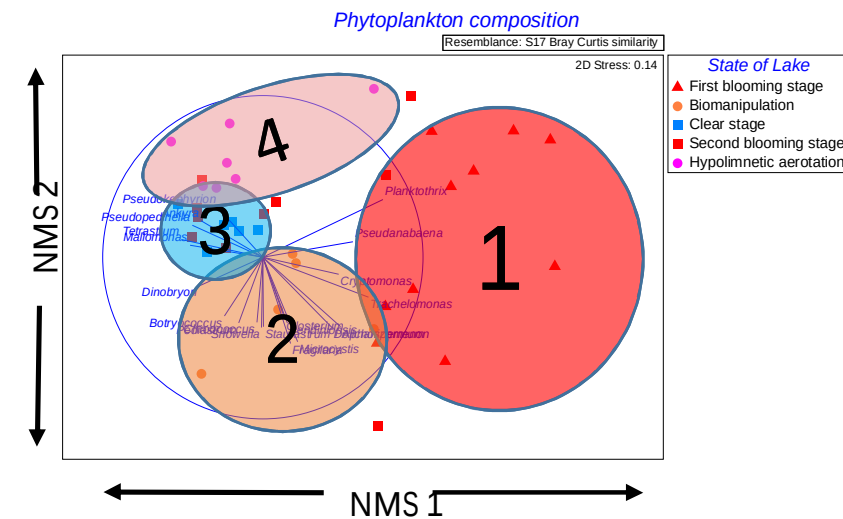
Planktothrix
Kirchneriella
NMS 2
Microcystis
Fragilaria

State of Lake

- ▲ First blooming stage
- Biomanipulation
- Clear stage
- Second blooming stage
- Hypolimnetic aerotation



Permutational Analysis of Multivariate Dispersions (PERMIDISP) was used to evaluate dispersion within each stage.

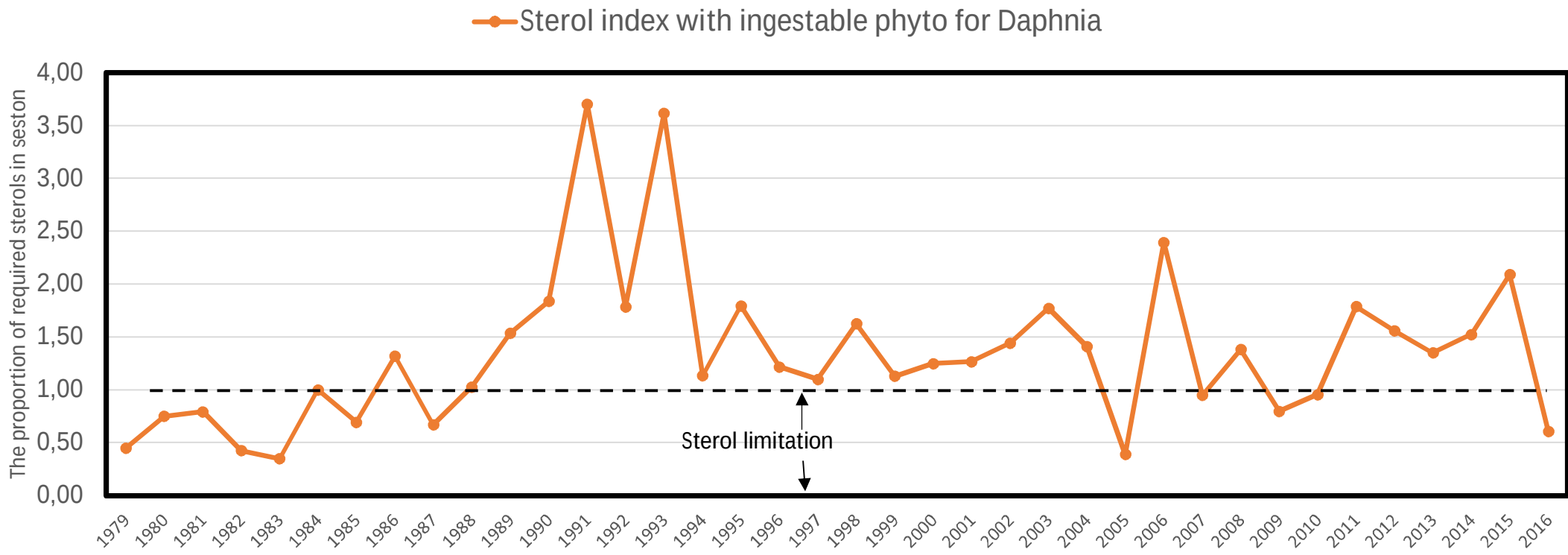
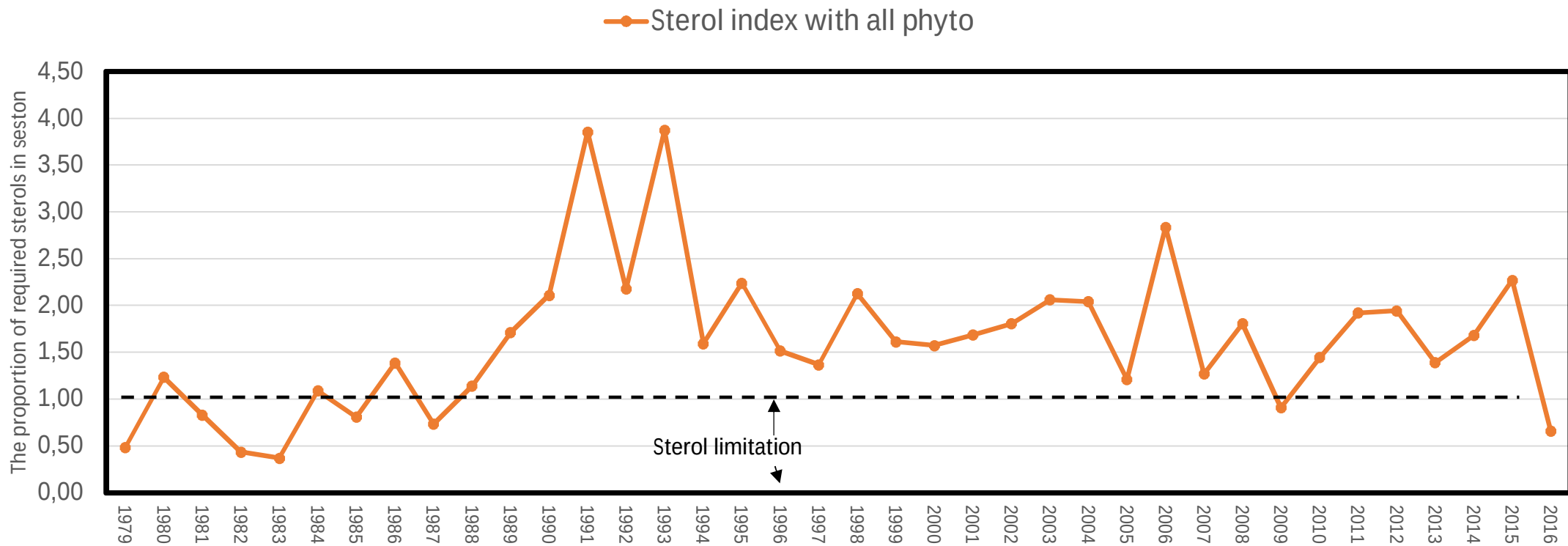


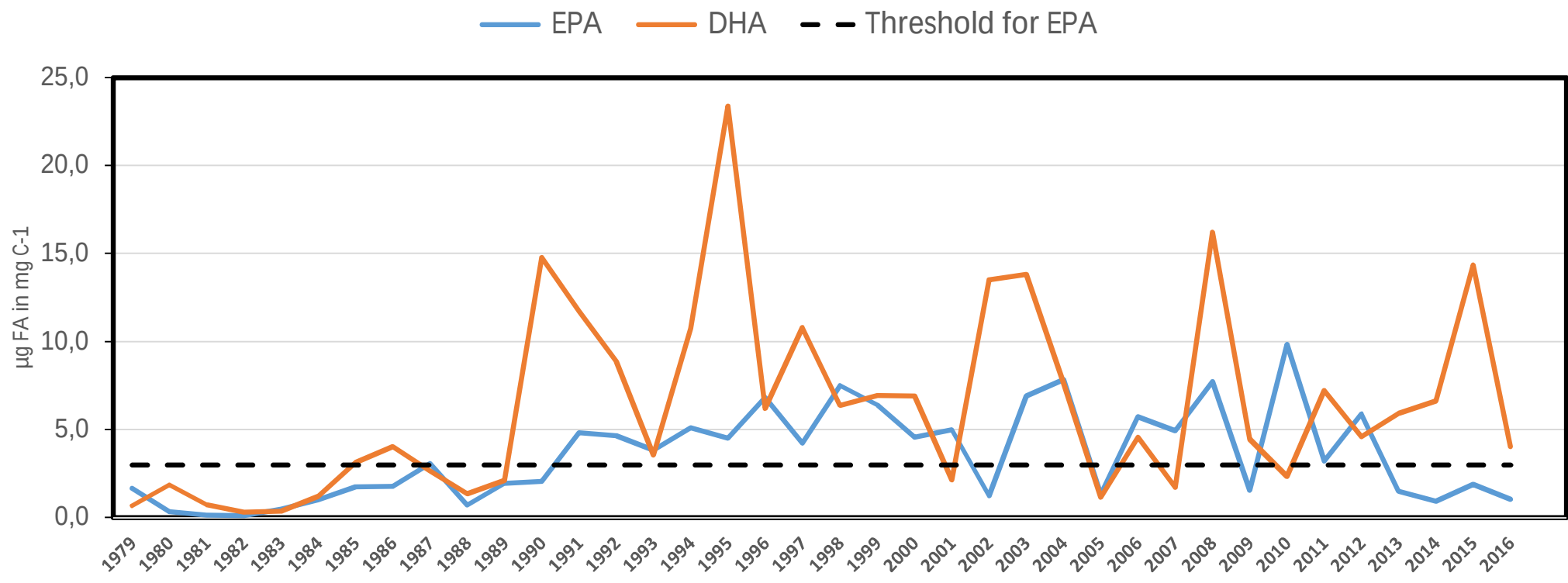
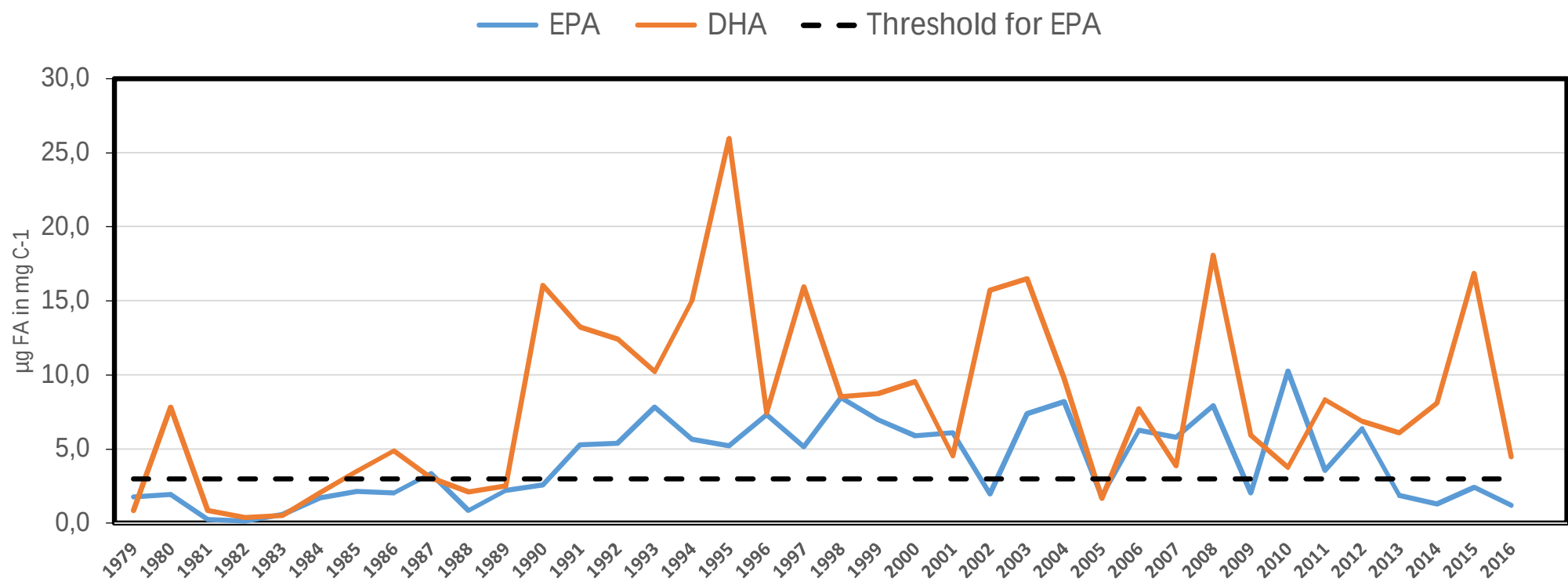
Similarity Percentage Analysis (SIMPER)

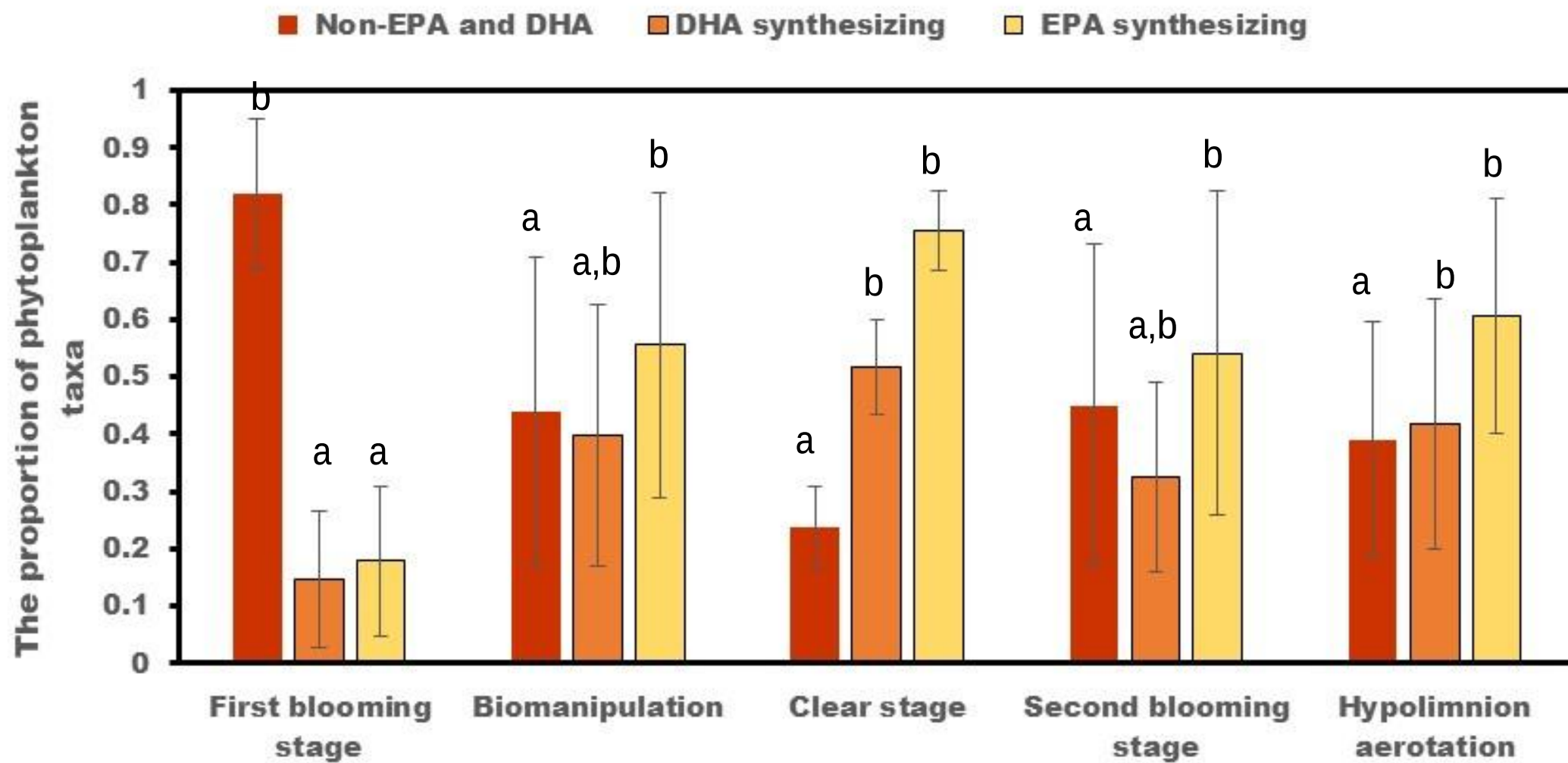
Group First blooming stage						Group Biomanipulation					
Average similarity: 32.50						Average similarity: 34.23					
Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%	Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Planktothrix	307.0	11.9	0.8	36.5	36.5	Cryptomonas	40.58	7.66	1.38	22.37	22.37
Aphanizomenon	151.7	6.1	0.7	18.7	55.1	Dolichospermum	31.96	4.5	0.63	13.15	35.52
Cryptomonas	40.7	3.9	2.1	12.1	67.2	Microcystis	28.71	3.94	1.36	11.52	47.03
Dolichospermum	101.3	3.1	0.8	9.4	76.6	Tabellaria	13.38	2.24	1.06	6.54	53.57
Pseudanabaena	152.7	2.1	0.4	6.3	83.0	Fragilaria	7.49	2.13	2.88	6.22	59.79
Microcystis	19.5	0.9	0.6	2.8	85.8	Snowella	6.72	1.61	1.77	4.72	64.51
Peridiniopsis	21.2	0.8	0.4	2.6	88.4	Ceratium	9.2	1.6	1.39	4.67	69.18
Rhodomonas	8.4	0.6	1.8	1.9	90.3	Rhodomonas	7.63	1.41	1.04	4.12	73.3
Group Clear stage						Group Hypolimnion aerotation					
Average similarity: 55.45						Average similarity: 42.43					
Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%	Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Tabellaria	22.59	10.61	1.49	19.14	19.14	Rhodomonas	14.4	10.99	1.66	25.91	25.91
Cryptomonas	19.82	10.3	1.89	18.58	37.71	Cryptomonas	8.81	6.89	1.73	16.23	42.14
Rhodomonas	11.03	5.06	1.83	9.12	46.83	Dolichospermum	7.36	4.58	1.75	10.81	52.94
Mallomonas	7.69	3.74	1.94	6.74	53.57	Asterionella	6.16	2.67	0.88	6.3	59.24
Ceratium	5.53	3.73	2.84	6.73	60.31	Mallomonas	4.46	2.39	0.99	5.64	64.88
Peridiniopsis	10.96	3.29	0.68	5.92	66.23	Planktothrix	20.44	2.11	1.01	4.98	69.86
Snowella	5.72	2.95	1.85	5.32	71.55	Ceratium	2.29	1.37	1.12	3.23	73.09
Asterionella	3.46	1.81	1.44	3.26	74.81	Tabellaria	11.27	1.37	0.59	3.22	76.31
Uroglena	5.86	1.7	0.83	3.06	77.87	Gymnodinium	2.09	1.19	0.9	2.8	79.11
Dolichospermum	5.82	1.64	1.23	2.96	80.83	Uroglena	2.72	1.02	1.05	2.41	81.51
Fragilaria	2.79	1.4	2.33	2.53	83.36	Aulacoseira	2.09	0.96	1.04	2.25	83.76
Stephanodiscus	1.98	1	1.17	1.8	85.16	Stephanodiscus	1.16	0.73	1.17	1.73	85.49
Peridinium	1.56	0.86	2.81	1.56	86.72	Fragilaria	1.16	0.51	1.46	1.2	86.69
Dinobryon	1.57	0.75	1.43	1.35	88.07	Woronichinia	0.98	0.47	1.3	1.1	87.79
Oocystis	1.46	0.71	2.35	1.29	89.36	Pseudopedinella	0.53	0.44	2.23	1.04	88.83
Synura	2.07	0.6	0.78	1.09	90.45	Pediastrum	0.71	0.41	1.34	0.98	89.8
						Microcystis	0.75	0.38	1.77	0.9	90.7

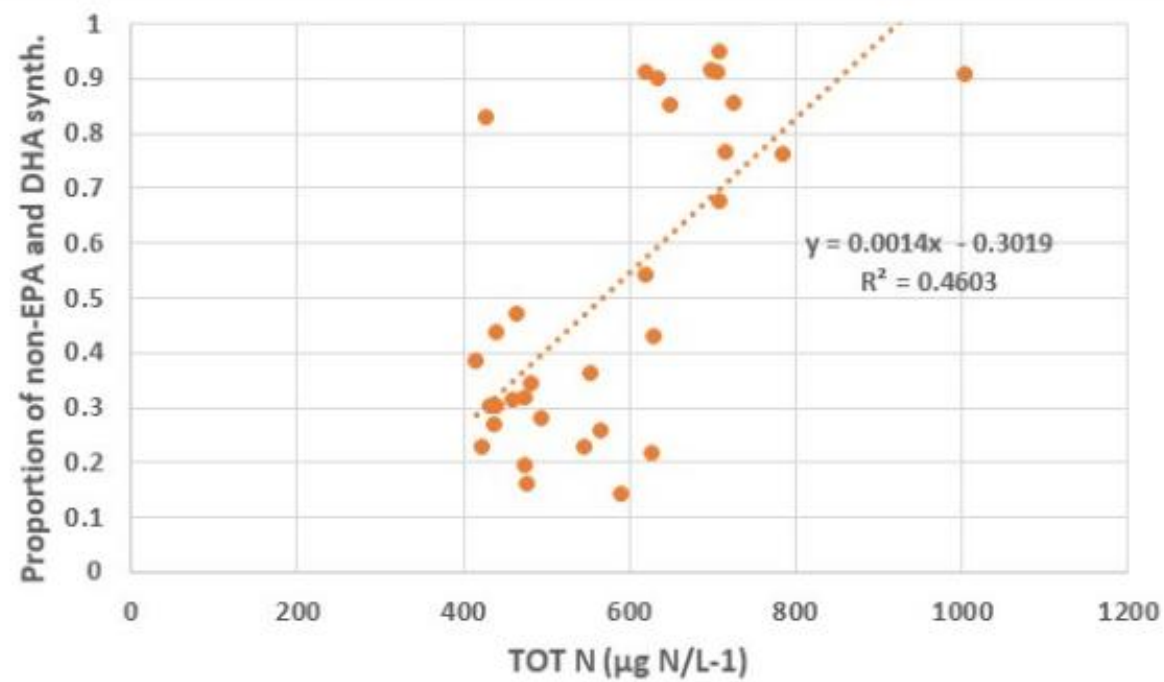
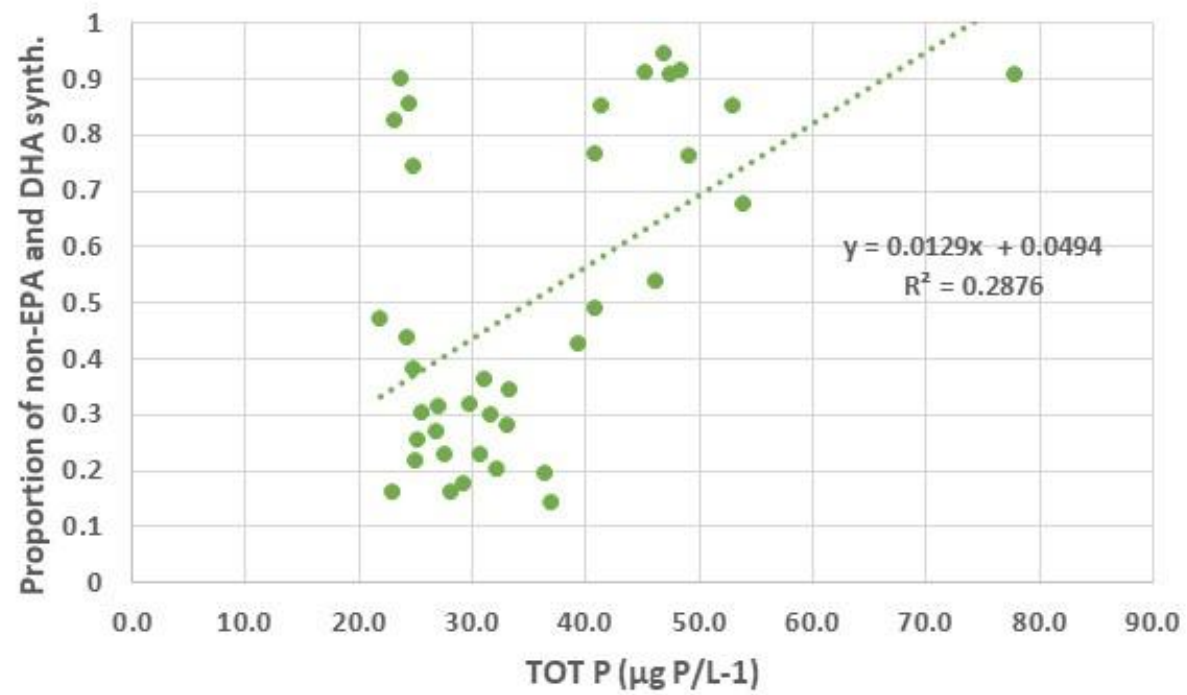
Groups First blooming stage & Clear stage						
Average dissimilarity = 85.86						
	Group First blooming stage	Group Clear stage				
Species	Av.Abund	Av.Abund	Av.Diss	Diss/SD	Contrib%	Cum.%
Planktothrix	307	0	27.7	1.08	32.26	32.26
Aphanizomenon	152	0	16.71	0.81	19.46	51.72
Pseudanabaena	153	0	10.45	0.71	12.18	63.89
Dolichospermum	101	5.82	7.91	0.86	9.22	73.11
Peridiniopsis	21	10.96	2.93	0.93	3.41	76.52
Microcystis	20	0.71	2.74	0.69	3.19	79.71
Ceratium	19	5.53	2.61	0.37	3.04	82.76
Cryptomonas	41	19.82	2.53	1.3	2.94	85.7
Tabellaria	2	22.59	2.5	1.31	2.91	88.61
Woronichinia	8	0.63	1.09	0.58	1.27	89.87
Aulacoseira	7	0.69	0.85	0.74	0.99	90.87
Groups Biomanipulation & Clear stage						
Average dissimilarity = 64.76						
	Group Biomanipulation	Group Clear stage				
Species	Av.Abund	Av.Abund	Av.Diss	Diss/SD	Contrib%	Cum.%
Aphanizomenon	45.9	0	7.81	0.5	12.05	12.05
Dolichospermum	31.96	5.82	6.97	1.35	10.76	22.81
Cryptomonas	40.58	19.82	6.58	1.13	10.16	32.98
Peridiniopsis	24.71	10.96	6.48	0.74	10.01	42.99
Microcystis	28.71	0.71	5.98	1.3	9.23	52.22
Tabellaria	13.38	22.59	3.72	1.18	5.74	57.96
Aulacoseira	8.64	0.69	2.55	0.52	3.94	61.9
Synura	6.91	2.07	2.3	0.61	3.55	65.45
Peridinium	6.86	1.56	1.76	0.54	2.72	68.17
Rhodomonas	7.63	11.03	1.73	1.24	2.67	70.84
Uroglena	7.4	5.86	1.72	1.24	2.66	73.5
Woronichinia	7.03	0.63	1.53	0.84	2.37	75.87
Ceratium	9.2	5.53	1.34	0.8	2.08	77.95
Mallomonas	3	7.69	1.27	1.45	1.95	79.9
Fragilaria	7.49	2.79	1.18	1.77	1.82	81.72

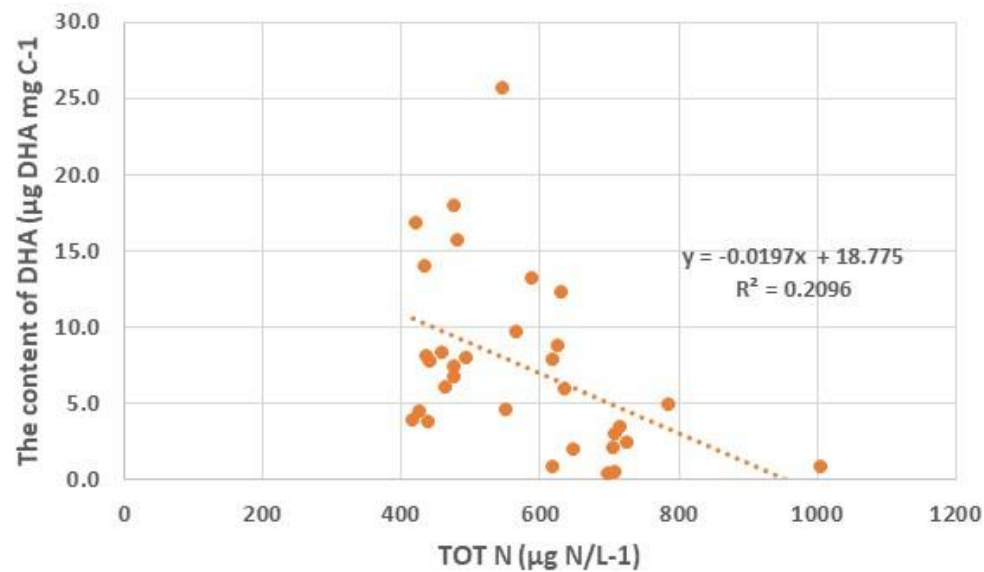
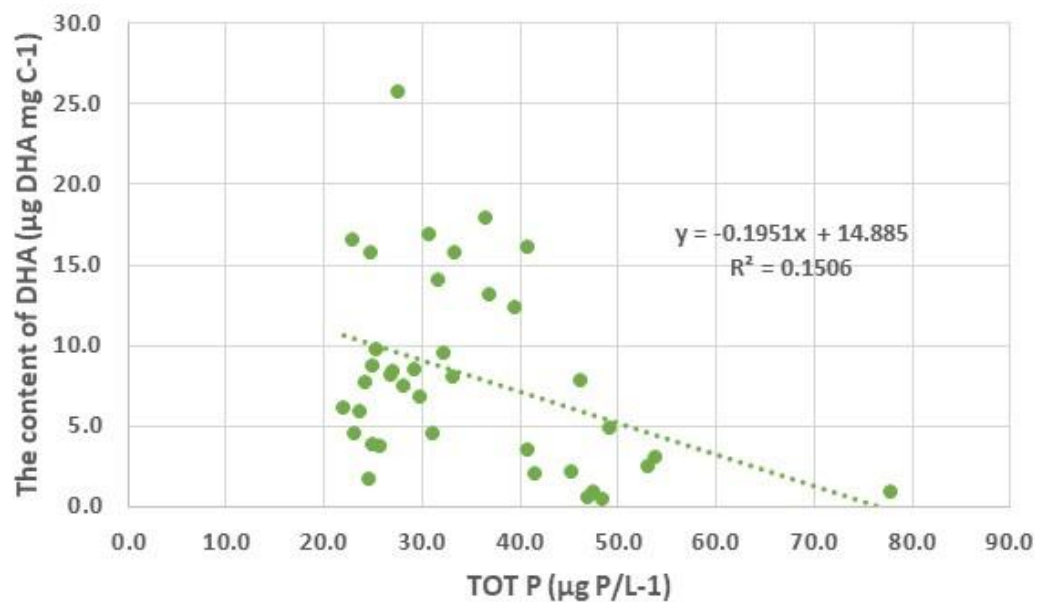
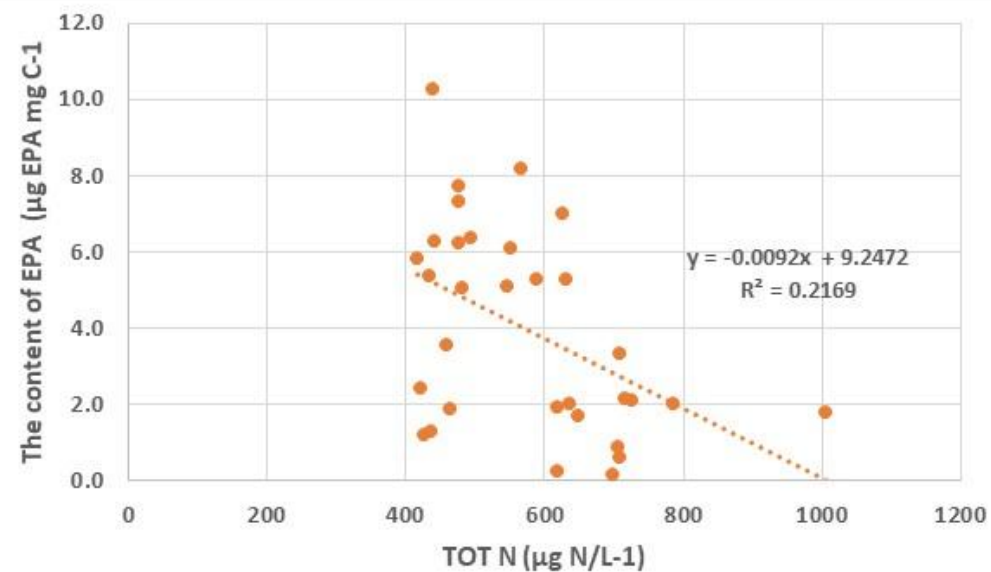
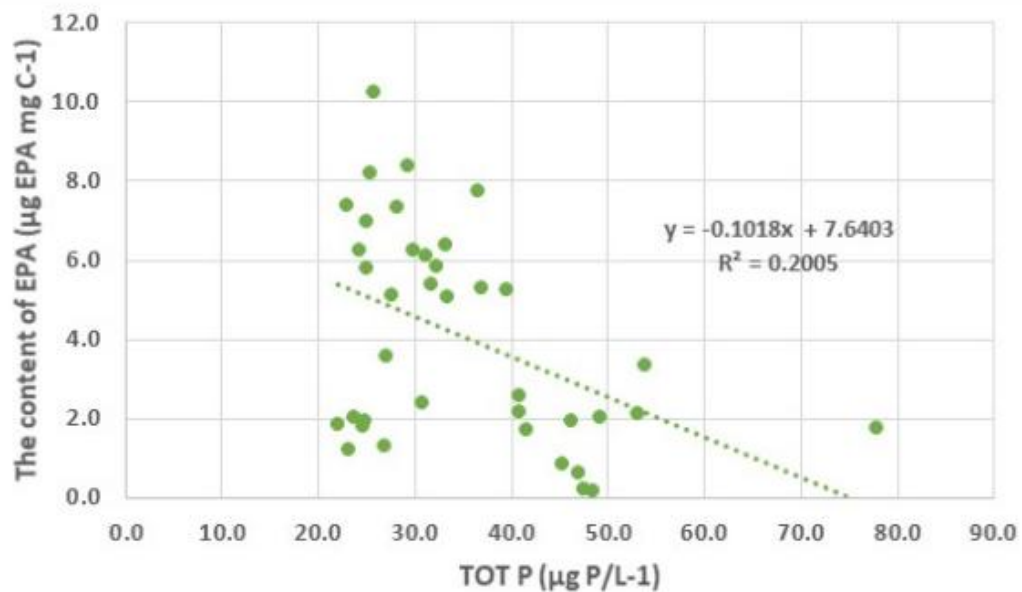
Groups Clear stage & Hypolimnion aerotation						
Average dissimilarity = 59.55						
	Group Clear stage	Group Hypolimnion aerotation				
Species	Av.Abund	Av.Abund	Av.Diss	Diss/SD	Contrib%	Cum.%
Tabellaria	22.59	11.27	9.98	1.51	16.77	16.77
Planktothrix	0	20.44	7.21	0.52	12.11	28.87
Cryptomonas	19.82	8.81	5.44	1.32	9.13	38
Peridiniopsis	10.96	0	4.7	1.13	7.89	45.9
Rhodomonas	11.03	14.4	3.88	1.23	6.51	52.4
Dolichospermum	5.82	7.36	2.87	1.17	4.82	57.23
Uroglena	5.86	2.72	2.47	0.97	4.16	61.38
Snowella	5.72	0.61	2.41	1.39	4.04	65.43
Mallomonas	7.69	4.46	2.26	1.47	3.8	69.23
Asterionella	3.46	6.16	2.21	1.27	3.71	72.94
Ceratium	5.53	2.29	1.57	1.58	2.63	75.57
Fragilaria	2.79	1.16	0.94	1.24	1.58	77.15
Synura	2.07	0.49	0.91	0.99	1.53	78.68
Eudorina	1.89	0.99	0.82	1.08	1.38	80.05
Sphaerocystis	1.45	0	0.78	0.53	1.31	81.36
Aulacoseira	0.69	2.09	0.74	1.42	1.24	82.6
Gymnodinium	1.13	2.09	0.72	1.69	1.21	83.81
Stephanodiscus	1.98	1.16	0.64	1.3	1.08	84.89
Aphanizomenon	0	1.49	0.61	0.77	1.03	85.91
Dinobryon	1.57	0.57	0.57	1.07	0.95	86.87
Peridinium	1.56	0.63	0.52	1.38	0.87	87.74
Oocystis	1.46	0.36	0.5	1.29	0.83	88.57
Trachelomonas	1	0.31	0.4	1.15	0.68	89.24
Woronichinia	0.63	0.98	0.38	1	0.64	89.88
Dictyosphaerium	0.56	0.39	0.35	0.82	0.59	90.47







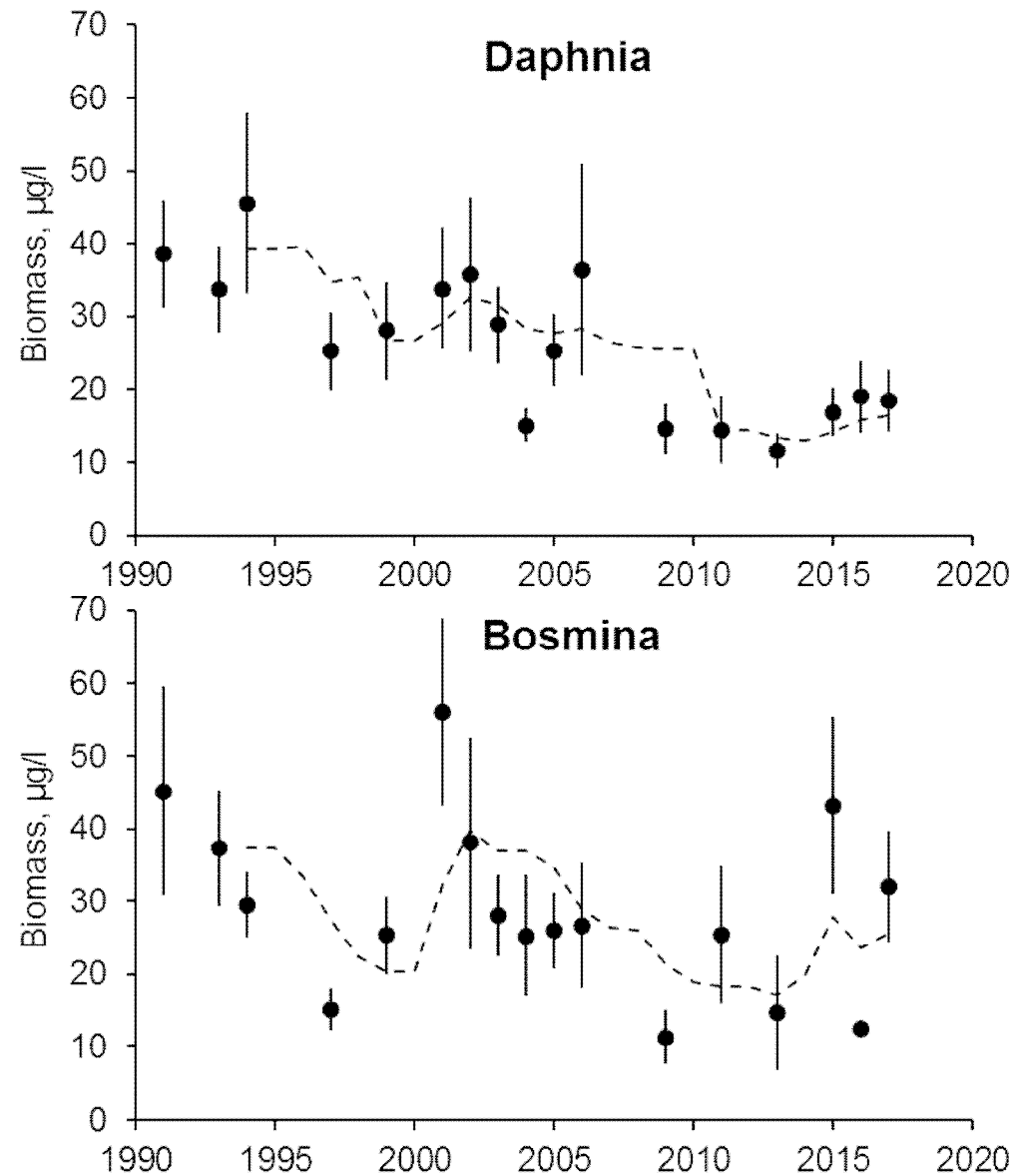






CLADOCERA

June-August average ($\pm$ SE) biomass with 5-year moving average



EUDIAPTOMUS

June-August average ($\pm$ SE) biomass
with 5-year moving average

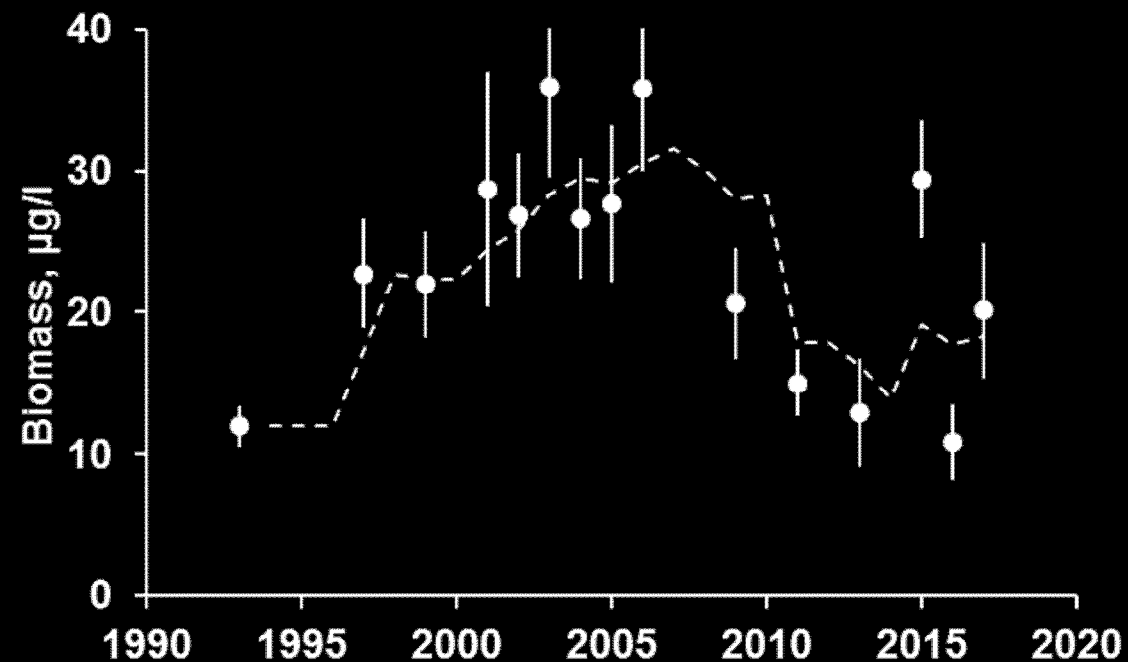
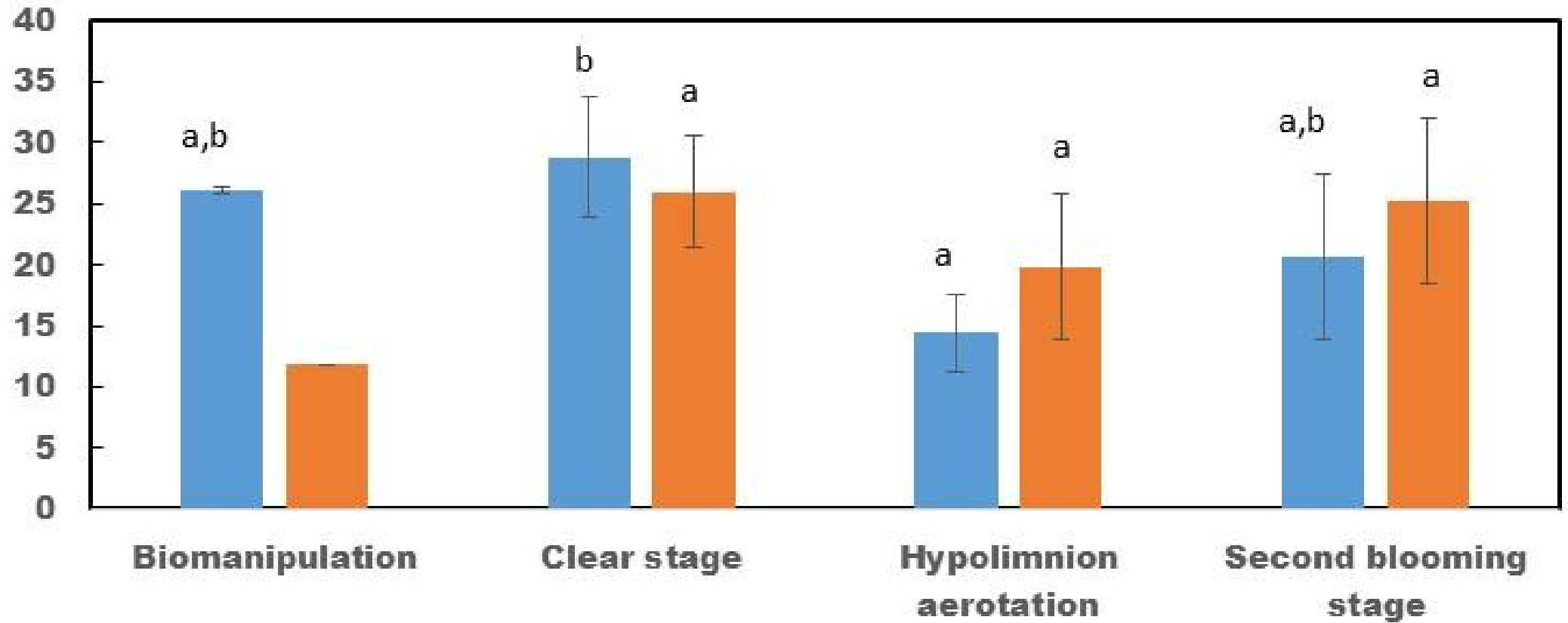


Photo: Kirsi Kuoppamäki



■ **Daphnia** ■ **Eudiaptomus**



CONCLUSIONS

- AFTER BIOMANIPULATION (CLEAR STAGE) PHYTOPLANKTON COMMUNITY WAS STABLE AND CHARACTERIZED BY OLIGOTROPHIC SPECIES
- DURING HYPOLIMNETIC AEROTATION PHYTOPLANKTON COMMUNITY DIFFERED FROM ANY PREVIOUS STAGE HAVING DIVERSE AFFECT ON PHYTOPLANKTON COMMUNITY: CYANOBACTERIA AND RHODOMONAS INCREASED
- THE PROPORTION OF NON-EPA AND NON-DHA SYNTHESIZING PHYTOPLANKTON TAXA WERE MOST ABUNDANT ON FIRST BLOOMING STAGE AND LOWEST IN CLEAR-STAGE.
- THE PROPORTION OF EPA AND DHA -SYNTHESIZING PHYTOPLANKTON TAXA WERE LOWEST IN FIRST BLOOMING STAGE AND MOST ABUNDANT IN CLEAR-STAGE, HOWEVER, THERE WAS NOT STATISTICAL DIFFERENCE BETWEEN CLEAR-STAGE, SECOND BLOOMING STAGE AND HYPOLIMNETIC AEROTATION.
- BIOMASS OF DAPHNIA WAS HIGHEST IN CLEAR STAGE AND LOWEST DURING HYPOLIMNETIC AEROTATION, BUT BIOMASS OF EUDIPTOMUS DID NOT DIFFER BETWEEN STAGES.