Table S1: Comparison of Prochlorococcus, Synechococcus and  $2\mu m$  beads abundances measured with the benchtop FACSCalibur flow cytometer and the CytoSense flow cytometer on five replicates.

Sample	Prochlorococcus		Synechococcus		2μm beads	
	(cell.cm <sup>-3</sup> )		(cell.cm <sup>-3</sup> )		(beads.cm <sup>-3</sup> )	
	FACS	CS	FACS	CS	FACS	CS
#1	18436	18236	27766	27906	5646	5376
#2	17762	17764	27153	26758	5371	5329
#3	17469	17102	27117	27564	5362	5613
#4	17759	17953	27797	27852	5553	5679
#5	18017	17291	28214	27050	5734	5443
Mean	17888	17669	27610	27426	5533	5488
StdDev	270	357	380	418	133	126
	t.test p=0,43	26	t.test p=0,568		t.test p=0,573	

Figure S1: Temperature (°C), salinity, Chl-*a* (μg.dm<sup>-3</sup>, converted from fluorescence) and density profiles (kg.m<sup>-3</sup>) at 6 stations.

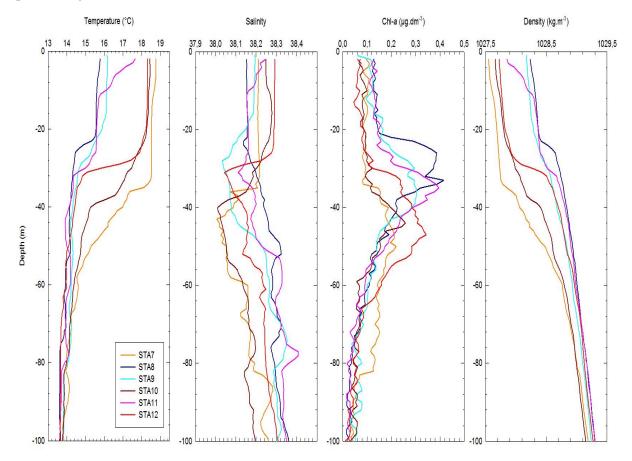


Figure S2: Temperature (°C), salinity, fluorescence (a.u.) and PAR (μE.m<sup>-2</sup>.d<sup>-1</sup>) profiles recorded by the CTD-Rosette, with associated *Prochlorococcus*, *Synechococcus* and picoeukaryotes abundances (cells.cm<sup>-3</sup>) and nitrate, nitrite, silicate and phosphate concentrations (μmol.dm<sup>-3</sup>) at Station 11.

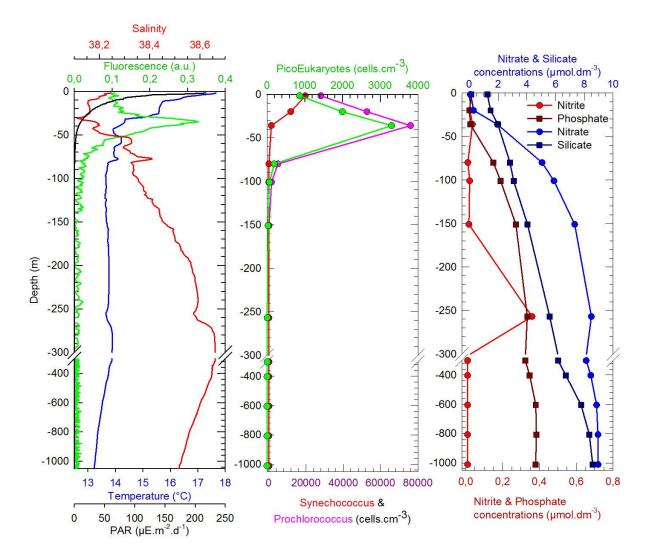


Figure S3: Mean abundances and standard deviation (green circles and error bars) during and at the vicinity of fixed stations 1 to 11 of *Prochlorococcus*, *Synechococcus* and picoeukaryotes (cells.cm<sup>-3</sup>) by the CytoSense AFCM in surface waters compared to abundances recorded by benchtop flow cytometry at the 2 first depths (from 1 to 5 m depth) during PASTIS-HVR vertical samplings (yellow triangles).

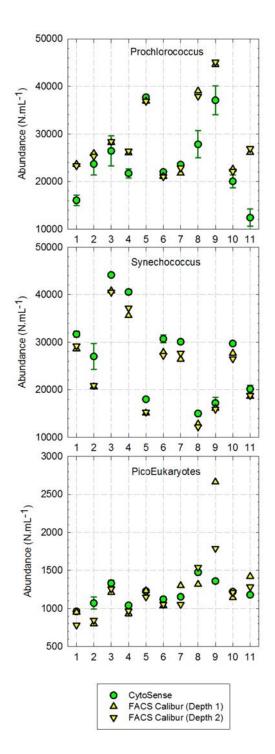


Figure S4: Mean SWS and FLR of *Prochlorococcus* and *Synechococcus* recorded by benchtop flow cytometry at fixed station 6 to 11 from PASTIS-HVR vertical samplings and at Station 11 (SWS: black circles, FLR: with squares).

