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Supplement of

Organic matter contents and degradation in a highly trawled area during fresh particle inputs (Gulf of Castellammare, southwestern Mediterranean)

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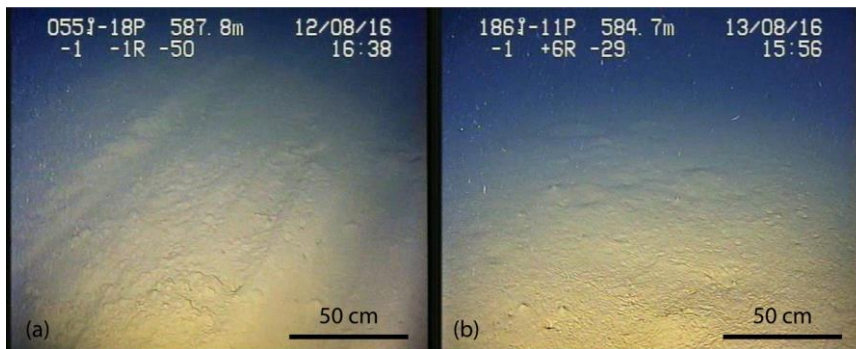
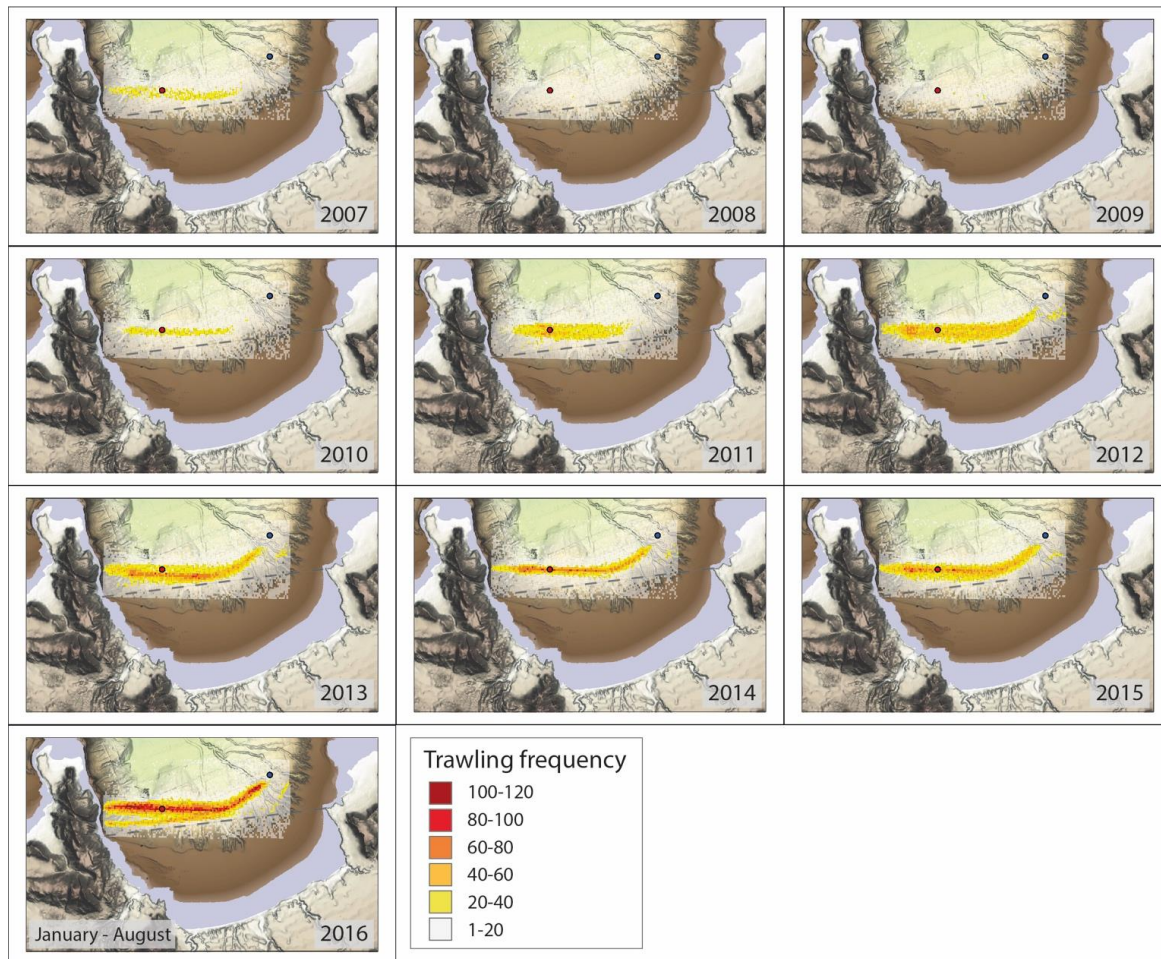


Figure S1. ROV-seafloor pictures from the trawled (a) and untrawled (b) sites in the Gulf of Castellammare. The trawled site presents linear marks and furrows on eroded sediments produced by the intense trawling activity in the area, whereas the untrawled site shows mounds and tracks characteristic of bioturbation activity.



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Figure S2. Evolution of trawling intensity. Number of times trawled per year and per grid cell (200 x 200 m) in the Gulf of Castellammare. Note the increasing trawling frequency since 2007, with the exception of 2008 and 2009, and that computation of trawling frequency in 2016 is limited to VMS data prior to the sampling period, August 10th. Blue and red circles indicate the sampling sites of the untrawled, and trawled site, respectively.

Table S1. Results of the PERMANOVA test for differences in organic matter quantity (protein, carbohydrate, lipid, biopolymeric C, and phytopigment) between trawled and untrawled sites, depth layers, and their interaction. The percentage variance explained by each source of variation is also reported. df = degrees of freedom; MS = mean square; Pseudo-F = permutational F; significance of test obtained from Monte Carlo simulations: * = p (MC) < 0.05; ** = p (MC) < 0.01; *** = p (MC) < 0.001; n.s.= not significant.

Variable	Source	df	MS	Pseudo-F	p (MC)	Explained variance (%)
Protein (mgC·g ⁻¹)	Site	1	4.62	13.2	**	22
	Depth	4	3.51	10.1	***	40
	Site x Depth	4	0.82	2.4	n.s.	12
	Residual	20	0.35			27
	Total	29				
Carbohydrate (mgC·g ⁻¹)	Site	1	11.29	47.9	***	44
	Depth	4	1.68	7.1	**	15
	Site x Depth	4	1.57	6.6	**	27
	Residual	20	0.23			14
	Total	29				
Lipid (mgC·g ⁻¹)	Site	1	4.68	20.4	***	23
	Depth	4	4.73	20.6	***	59
	Site x Depth	4	0.20	0.9	n.s.	0
	Residual	20	0.23			18
	Total	29				
Biopolymeric C (mgC·g ⁻¹)	Site	1	13.02	36.9	***	55
	Depth	4	1.36	3.9	*	11
	Site x Depth	4	0.87	2.5	n.s.	11
	Residual	20	0.35			23
	Total	29				
Phytopigment (μgC·g ⁻¹)	Site	1	11.25	93.0	***	50
	Depth	4	3.53	29.1	***	38
	Site x Depth	4	0.29	2.4	n.s.	4
	Residual	20	0.12			8
	Total	29				

Table S2. Results of the pairwise comparison testing for differences in the quantity of each investigated variable between trawled and untrawled sites at each depth. df = degrees of freedom; MS = mean square; Pseudo-F = permutational F; significance of test obtained from Monte Carlo simulations: * = p (MC) < 0.05; ** = p (MC) < 0.01; *** = p (MC) < 0.001; n.s.= not significant.

Variable	Depth (cm)	Pair-wise comparisons	
		t	p (MC)
Protein (mgC·g ⁻¹)	0-1	0.502	n.s.
	1-3	1.236	n.s.
	3-5	1.871	n.s.
	5-7	2.94	*
	7-9	1.799	n.s.
Carbohydrate (mgC·g ⁻¹)	0-1	3.648	*
	1-3	9.139	**
	3-5	2.997	*
	5-7	0.643	n.s.
	7-9	1.06	n.s.
Lipid (mgC·g ⁻¹)	0-1	0.666	n.s.
	1-3	3.753	*
	3-5	1.044	n.s.
	5-7	2.921	*
	7-9	5.63	**
Biopolymeric C (mgC·g ⁻¹)	0-1	2.058	n.s.
	1-3	4.197	*
	3-5	2.737	n.s.
	5-7	3.105	*
	7-9	3.783	*
Phytopigment (µgC·g ⁻¹)	0-1	1.846	n.s.
	1-3	6.128	**
	3-5	11.417	***
	5-7	4.572	*
	7-9	3.251	*

5 **Table S3.** Results of the PERMANOVA testing for differences in biochemical composition (protein, carbohydrate, lipid and phytopigment contents) between trawled and untrawled sites, depth layers, and their interaction. The percentage variance explained by each source of variation is also reported. df = degrees of freedom; MS = mean square; Pseudo-F = permutational F; significance of test obtained from Monte Carlo simulations: * = p (MC) < 0.05; ** = p (MC) < 0.01; *** = p (MC) < 0.001; n.s.= not significant.

	Source	df	MS	Pseudo-F	p (MC)	Explained variance (%)
Biochemical composition	Site	1	31.9	34.0	***	36
	Depth	1	13.5	14.4	***	36
	Site x Depth	4	2.9	3.0	**	11
	Residual	20	0.9			16
	Total	26				

Table S4. Results of the pairwise comparison testing for differences in the biochemical composition (in terms of protein, carbohydrate, lipid and phytopigment contents) between trawled and untrawled sites at each depth. df = degrees of freedom; MS = mean square; Pseudo-F = permutational F; significance of test obtained from Monte Carlo simulations: * = p (MC) < 0.05; ** = p (MC) < 0.01; *** = p (MC) < 0.001; n.s.= not significant.

	Depth (cm)	Pair-wise comparisons	
		t	p (MC)
Biochemical composition	0-1	2.237	n.s.
	1-3	5.353	**
	3-5	2.456	*
	5-7	2.565	*
	7-9	2.875	*

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Table S5. Results of the PERMANOVA testing for differences in (a) relative contribution of phytopigments to biopolymeric C and (b) protein C turnover rates between examined factors and their interaction. The pairwise comparison testing for differences of each investigated variable between trawled and untrawled sites at each depth is also included. df = degrees of freedom; MS = mean square; Pseudo-F = permutational F; significance of test obtained from Monte Carlo simulations: * = p (MC) < 0.05; ** = p (MC) < 0.01; *** = p (MC) < 0.001; n.s.= not significant.

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(a)

Source	df	MS	Pseudo-F	p (MC)	Explained variance (%)	Pair-wise comparisons			
						Depth (cm)	t	p (MC)	
Phytopigment/BPC (%)	Site	1	3.6	59.1	***	18	0-1	1.438	n.s.
	Depth	4	6	99.5	***	77	1-3	3.440	*
	Site x Depth	4	0.1	1.3	n.s.	0	3-5	6.698	**
	Residual	20	0.1			5	5-7	4.143	*
	Total	29				18	7-9	4.588	**

(b)

Source	df	MS	Pseudo-F	p (MC)	Explained variance (%)	Pair-wise comparisons			
						Depth (cm)	t	p (MC)	
Turnover ($10^{-3} d^{-1}$)	Site	1	1.2	26	***	5	0-1	5.400	***
	Depth	4	5.5	120.4	***	64	1-3	5.452	**
	Site x Depth	4	1.2	26.4	***	27	3-5	4.242	*
	Residual	20	0			3	5-7	1.234	n.s.
	Total	29				5	7-9	2.418	n.s.