

Table S1. List of surface sediment samples collected in the Mediterranean Sea, locality, date, geographical coordinates and depth.

| Sample | Locality | Date (dd/mm/yyyy) | Latitude (DD.ddddd) | Longitude (DD.ddddd) | Depth |
|--------------|-----------------|----------------------|------------------------|-------------------------|-------|
| MS15_09 | Bari | 07/11/2015 | 41.30400 | 16.92400 | 115 |
| MS15_55 | Gargano Sud | 10/11/2015 | 41.51400 | 16.32300 | 35 |
| MS15_56 | Gargano Sud | 11/11/2015 | 41.55400 | 16.23000 | 22 |
| MS15_120 | Leuca | 16/11/2015 | 39.54720 | 18.40894 | 600 |
| MS15_121 | Leuca | 16/11/2015 | 39.57741 | 18.40133 | 515 |
| MS15_137 | Gallipoli Sud | 17/11/2015 | 39.78800 | 17.97600 | 94 |
| MS15_145 | Crotone | 18/11/2015 | 39.58355 | 16.87080 | 65 |
| MS15_147 | Crotone | 18/11/2015 | 39.48957 | 17.10846 | 155 |
| MS15_151 | Crotone | 18/11/2015 | 39.34500 | 17.19500 | 227 |
| MS15_152 | Crotone | 18/11/2015 | 39.25500 | 17.15800 | 188 |
| MS15_159 | Crotone | 18/11/2015 | 38.99725 | 17.25582 | 321 |
| MS15_161 | Crotone | 19/11/2015 | 38.90300 | 17.20400 | 225 |
| MS15_174 | Crotone | 19/11/2015 | 38.87500 | 17.14000 | 180 |
| MS15_175 | Crotone | 19/11/2015 | 38.76444 | 16.89083 | 564 |
| MS15_181 | Roccella Ionica | 20/11/2015 | 38.34500 | 16.52400 | 85 |
| MS15_196 | Augusta | 21/11/2015 | 37.24600 | 15.28200 | 81 |
| MS15_185 | Riace | 20/11/2015 | 38.34600 | 16.50300 | 51 |
| MS16_I_155 | Orosei | 28/07/2016 | 40.26010 | 9.90530 | 446 |
| MS16_I_157 | Orosei | 28/07/2016 | 40.20990 | 9.67640 | 276 |
| MS16_I_166 | Nora Canyon | 30/07/2016 | 38.80270 | 8.92530 | 123 |
| MS16_I_168 | Nora Canyon | 30/07/2016 | 38.65490 | 9.04010 | 495 |
| MS16_II_28 | Gela | 04/09/2016 | 36.91869 | 14.27464 | 70 |
| MS16_II_31 | Gela | 04/09/2016 | 36.83518 | 14.20068 | 415 |
| MS16_II_34 | Gela | 04/09/2016 | 36.75515 | 14.12661 | 766 |
| MS16_II_44 | Augusta | 04/09/2016 | 37.22468 | 15.29330 | 128 |
| MS16_II_76 | Porto Cesareo | 09/09/2016 | 40.25050 | 17.46871 | 189 |
| MS16_II_77a | Porto Cesareo | 09/09/2016 | 40.15356 | 17.40916 | 930 |
| MS16_II_77b | Porto Cesareo | 09/09/2016 | 40.15356 | 17.40916 | 930 |
| MS16_II_104a | S. Maria Leuca | 11/09/2016 | 39.80009 | 18.64332 | 145 |
| MS16_II_104b | S. Maria Leuca | 11/09/2016 | 39.80009 | 18.64332 | 145 |

| | | | | | |
|----------------|------------------------|------------|----------|----------|-----|
| MS16_II_108a | S. Maria Leuca | 11/09/2016 | 39.82152 | 18.52397 | 125 |
| MS16_II_108b | S. Maria Leuca | 11/09/2016 | 39.82152 | 18.52397 | 125 |
| MS16_II_119a | S. Cataldo | 12/09/2016 | 40.50142 | 18.60157 | 270 |
| MS16_II_119b | S. Cataldo | 12/09/2016 | 40.50142 | 18.60157 | 270 |
| MS16_II_121a | S. Cataldo | 12/09/2016 | 40.43669 | 18.41888 | 104 |
| MS16_II_121b | S. Cataldo | 12/09/2016 | 40.43669 | 18.41888 | 104 |
| MS16_III_26 | Bari | 09/12/2016 | 41.33600 | 17.19300 | 608 |
| MS16_III_34 | Bari | 09/12/2016 | 41.11800 | 17.05900 | 82 |
| MS16_III_41 | Barletta | 09/12/2016 | 41.42500 | 16.37500 | 39 |
| MS16_III_45 | Barletta | 10/12/2016 | 41.50600 | 16.66800 | 99 |
| MS16_III_49 | Vieste (DART) | 10/12/2016 | 42.24300 | 15.92300 | 119 |
| MS16_III_65 | Vieste (DART) | 10/12/2016 | 42.02200 | 15.86900 | 45 |
| MS16_III_69 | Pescara | 11/12/2016 | 42.55100 | 14.45200 | 91 |
| MS16_III_72 | Pescara | 11/12/2016 | 42.51900 | 14.40000 | 75 |
| MS16_III_75 | Pescara | 11/12/2016 | 42.47600 | 14.33400 | 26 |
| MS16_III_79 | S. Benedetto Tronto | 11/12/2016 | 42.80600 | 14.07500 | 22 |
| MS16_III_85 | S. Benedetto Tronto | 11/12/2016 | 42.88800 | 14.18600 | 73 |
| MS16_III_90 | Civitanova Marche | 11/12/2016 | 43.29100 | 14.03800 | 55 |
| MS16_III_96 | Civitanova Marche | 11/12/2016 | 43.20800 | 13.89000 | 17 |
| MS16_III_102 | Senigallia | 12/12/2016 | 43.78600 | 13.31800 | 20 |
| MS16_III_110 | Senigallia | 12/12/2016 | 43.85800 | 13.42200 | 49 |
| MS16_III_113 | Ancona | 12/12/2016 | 43.72100 | 13.77100 | 65 |
| MS16_III_122 | Ancona | 12/12/2016 | 43.61900 | 13.65000 | 1 |
| MS16_III_126 | Rimini | 14/12/2016 | 44.22400 | 12.68900 | 22 |
| MS16_III_129 | Rimini Boa E1 | 14/12/2016 | 44.14000 | 12.57000 | 10 |
| MS16_III_147 | MEDA S1 | 15/12/2016 | 44.73500 | 12.45400 | 22 |
| MS16_III_155 | Delta Po | 16/12/2016 | 44.64000 | 12.60000 | 31 |
| MS16_III_157 | Chioggia | 16/12/2016 | 44.98500 | 12.62100 | 30 |
| MS16_III_165 | Chioggia | 16/12/2016 | 45.00100 | 12.80700 | 33 |
| MS16_III_184 | Trieste | 16/12/2016 | 45.65700 | 13.65900 | 23 |
| MS17_I_83_C | Golfo di Napoli | 12/07/2017 | 40.73183 | 14.23217 | 187 |
| MS17_I_84_C R1 | Golfo di Napoli | 12/07/2017 | 40.77417 | 14.16750 | 127 |
| MS17_I_88_C R2 | Golfo di Napoli | 12/07/2017 | 40.77133 | 14.17750 | 180 |
| MS17_I_90_C R1 | Golfo di Napoli | 12/07/2017 | 40.75163 | 14.15683 | 120 |

| | | | | | |
|------------------|--------------------|------------|----------|----------|------|
| MS17_I_94_C R2 | Golfo di Napoli | 12/07/2017 | 40.75733 | 14.11500 | 95 |
| MS17_I_107_C R2 | Isole Pontine | 14/07/2017 | 40.91117 | 12.88283 | 65 |
| MS17_I_111_C | Anzio | 14/07/2017 | 41.38317 | 12.68417 | 41 |
| MS17_I_114_C R2 | Elba Nord | 15/07/2017 | 43.01733 | 10.34417 | 110 |
| MS17_I_118_C R2 | Elba Nord | 15/07/2017 | 43.02167 | 10.47183 | 58 |
| MS17_I_120_C | Livorno | 15/07/2017 | 43.46717 | 10.24817 | 66 |
| MS17_I_125_C R2 | Gorgona | 16/07/2017 | 43.40917 | 9.91250 | 111 |
| MS17_I_131_C | Capraia | 16/07/2017 | 43.00000 | 9.79050 | 110 |
| MS17_I_142_C | Anzio | 18/07/2017 | 41.37867 | 12.42800 | 139 |
| MS17_II_008_C R2 | Egadi | 22/07/2017 | 37.96283 | 12.14567 | 171 |
| MS17_II_025_C R1 | Porto Empedocle | 24/07/2017 | 37.12983 | 13.34650 | 440 |
| MS17_II_032_C R1 | Gela | 24/07/2017 | 36.75850 | 14.13350 | 781 |
| MS17_II_035_C R1 | Gela | 24/07/2017 | 36.83633 | 14.20133 | 418 |
| MS17_II_039_C R2 | Gela | 24/07/2017 | 36.92000 | 14.28000 | 70 |
| MS17_II_041_C R1 | Augusta | 25/07/2017 | 37.22983 | 15.30000 | 130 |
| MS17_II_066_C R1 | Soverato | 26/07/2017 | 38.68000 | 16.61983 | 692 |
| MS17_II_089_C R3 | Amendolara | 28/07/2017 | 39.93067 | 16.70850 | 99 |
| MS17_II_096_C R2 | Amendolara | 29/07/2017 | 39.93183 | 16.71083 | 81 |
| MS17_II_108_C R1 | Campomarino | 30/07/2017 | 40.13283 | 17.41517 | 1018 |
| MS17_II_143_C R1 | Bari | 02/08/2017 | 41.25650 | 17.14750 | 151 |
| MS17_II_145_C R1 | Bari | 02/08/2017 | 41.18833 | 17.10900 | 125 |
| MS17_II_147_C R1 | Bari | 03/08/2017 | 41.11833 | 17.06467 | 84 |
| MS17_II_167_C R2 | Brindisi | 04/08/2017 | 40.76633 | 17.93650 | 106 |
| MS17_II_169_C R1 | Brindisi | 04/08/2017 | 40.76217 | 17.94350 | 106 |
| MS17_II_172_C R1 | Brindisi | 04/08/2017 | 40.76500 | 17.93883 | 106 |
| MS17_II_176_C R2 | Brindisi | 04/08/2017 | 40.76367 | 17.94117 | 106 |
| MS17_III_33 | Lesina | 09/08/2017 | 42.03683 | 15.31517 | 68 |
| MS17_III_76 | Chioggia | 11/08/2017 | 44.99183 | 12.69167 | 32 |
| MS17_III_81 | Venezia | 11/08/2017 | 45.29100 | 12.57217 | 24 |
| MS17_III_94 | Trieste | 12/08/2017 | 45.62450 | 13.47083 | 21 |

Table S2. List of microalgal species used to test the new primer specificity.

| Species | Strain | Class | Medium |
|------------------------------------------------|----------|-------------------|-----------|
| <i>Alexandrium andersoni</i> | VGO664 | Dinophyceae | f/2 - Si |
| <i>Alexandrium mediterraneum</i> | CNR-AT4 | Dinophyceae | f/2 - Si |
| <i>Alexandrium minutum</i> | CBA57 | Dinophyceae | f/2 - Si |
| <i>Alexandrium pacificum</i> | CNR-SRA4 | Dinophyceae | f/2 - Si |
| <i>Coolia monotis</i> | CBA1 | Dinophyceae | f/4 - Si |
| <i>Gonyaulax spinifera</i> | CBA5 | Dinophyceae | f/4 - Si |
| <i>Gymnodinium impudicum</i> | GY6V | Dinophyceae | L1 - Si |
| <i>Heterocapsa</i> sp. | CBAB/D5 | Dinophyceae | f/2 - Si |
| <i>Heterocapsa triquetra</i> | RCC4813 | Dinophyceae | L1 - Si |
| <i>Lingulodinium polyedrum</i> | LPA0510 | Dinophyceae | f/10 - Si |
| <i>Ostreopsis</i> cf. <i>ovata</i> | CBA1291 | Dinophyceae | f/4 - Si |
| <i>Prorocentrum lima</i> | CBA3 | Dinophyceae | f/4 - Si |
| <i>Protoceratium reticulatum</i> | PRA0311 | Dinophyceae | f/4 - Si |
| <i>Scrippsiella trochoidea</i> | CBA3 | Dinophyceae | f/2 - Si |
| <i>Fibrocapsa japonica</i> | CBA1 | Raphidophyceae | f/2 - Si |
| <i>Heterosigma akashiwo</i> | HA2V | Raphidophyceae | L1 |
| <i>Chaetoceros socialis</i> | CBA22 | Bacillariophyceae | f/2 |
| <i>Cylindroteca</i> sp. | CBA7 | Bacillariophyceae | f/2 |
| <i>Coscinodiscus</i> sp. | CBA1 | Bacillariophyceae | f/2 |
| <i>Ditylum brightwellii</i> | CBA2 | Bacillariophyceae | f/2 |
| <i>Guinardia flaccida</i> | CBA1 | Bacillariophyceae | f/2 |
| <i>Pseudo-nitzschia</i> cf. <i>arenysensis</i> | CBA163 | Bacillariophyceae | f/2 |
| <i>Pseudo-nitzschia calliantha</i> | CBA189 | Bacillariophyceae | f/2 |
| <i>Pseudo-nitzschia pungens</i> | CBA180 | Bacillariophyceae | f/2 |
| <i>Skeletonema marinoi</i> | CBA4 | Bacillariophyceae | f/2 |
| <i>Thalassionema</i> sp. | CBA10 | Bacillariophyceae | f/2 |
| <i>Thalassiosira</i> sp. | CBA3 | Bacillariophyceae | f/2 |

Table S3. List of species-specific primer sequences of various diatom and dinoflagellate taxa targeting different rDNA regions, qPCR reagent concentrations, amplicon melting temperature (T_m) and size.

| Taxon | Primer name | Forward primer sequence (5'–3') Reverse primer sequence (5'–3') | Primer concentration [nM] | MgCl ₂ [mM] | Amplicon T _m (°C) | Amplicon size (bp) | Primer locations | Reference |
|--------------------------------------|-------------|--------------------------------------------------------------------|---------------------------|------------------------|------------------------------|--------------------|------------------|--------------------------|
| <i>Chaetoceros socialis</i> | CsocF | 5'-GGAGCGTCTGAGTATGGTCGT-3' | 300 | 3.5 | 83.98 | 100 | ITS2 (5' → 3') | This study |
| | CsocR | 5'-GCCATCCAGACGCAAAAAGTGT-3' | | | | | ITS2 (3' ← 5') | |
| <i>Ditylum brightwellii</i> | DbrighF | 5'-ACCGCGTGTGCTTATGTAT-3' | 300 | 2.5 | 83.46 | 103 | ITS1 (5' → 3') | This study |
| | DbrighR | 5'-ACCAAACGACTTTCGGGGTT-3' | | | | | ITS1 (3' ← 5') | |
| <i>Skeletonema</i> spp. | SkelsppF | 5'-CGATACACTGGTAGCGAGCC-3' | 200 | 1.5 | 86.52 | 118 | ITS1 (5' → 3') | This study |
| | SkelsppR | 5'-TTCAGTTCGGTAATGGGCGG-3' | | | | | ITS1 (3' ← 5') | |
| <i>Thalassiosira</i> spp. | ThalsppF | 5'-AGCGGGAAGAGCTCACCAT-3' | 200 | 2.5 | 84.35 | 99 | LSU (5' → 3') | This study |
| | ThalsppR | 5'-AAGAGACTTGGCCCGGGAAC-3' | | | | | LSU (3' ← 5') | |
| <i>Pseudo-nitzschia</i> spp. | Pseudo 5' | 5'-CGATACGTAATGCGAATTGCAA-3' | 300 | 1.5 | 82.10 | 111 | 5.8S (5' → 3') | Penna et al. (2007) |
| | Pseudo 3' | 5'-GTGGGATCCRCAGACACTCAGA-3' | | | | | 5.8S (3' ← 5') | |
| Bacillariophyceae | 1209f | 5'-CAGGTCTGTGATGCCCTT-3' | 300 | 3.5 | 84.95 | 189 | SSU (5' → 3') | Giovannoni et al. (1988) |
| | Diatom18SR1 | 5'-CAATGCAGWTTGATGAWCTG-3' | | | | | SSU (3' ← 5') | Godhe et al. (2008) |
| <i>Alexandrium minutum</i> | ITS1m | 5'-CATGCTGCTGTGTTGATGACC-3' | 200 | 2.5 | 82.0 | 212 | ITS1 (5' → 3') | Galluzzi et al. (2005) |
| | 5.8S 3' | 5'-GCAMACCTTCAAGMATATCCC-3' | | | | | 5.8S (3' ← 5') | |
| <i>A. pacificum</i> | ITS1c | 5'-AGCATGATTTGTTTTCAAGC-3' | 200 | 2.5 | 81.90 | 226 | ITS1 (5' → 3') | Penna et al. (2007) |
| | 5.8S-3' | 5'-GCAMACCTTCAAGMATATCCC-3' | | | | | 5.8S (3' ← 5') | |
| <i>A. tamarense/A. mediterraneum</i> | 5.8S 5' | 5'-TGTTACTTGTACCTTTGGGA-3' | 300 | 2.5 | 81.34 | 134 | 5.8S (5' → 3') | Penna et al. (2007) |
| | ITS2t | 5'-ACAACACCCAGGTTCAAT-3' | | | | | ITS2 (3' ← 5') | |
| <i>Gonyaulax spinifera</i> | GspinF_for | 5'-GAAACTCCTTCTGTGGATGC-3' | 200 | 3.5 | 82.90 | 154 | LSU (5' → 3') | Perini et al. (2019) |

| | | | | | | | | |
|--------------------------------------------|--------------|----------------------------------|-----|-----|-------|-----|----------------|----------------------|
| | GspinF_for | 5'-TCACAGTTC CCTCATGGTACT-3' | | | | | LSU (3' ← 5') | |
| <i>Gymnodinium</i> spp. | GymnosppF | 5'-CAGCGACGGATGTCTCGGTT-3' | 200 | 2.5 | 84.20 | 102 | 5.8S (5' → 3') | This study |
| | GymnosppR | 5'-TGCGTTCAAGTTTCTGTCGGT-3' | | | | | 5.8S (3' ← 5') | |
| <i>Heterocapsa</i> <i>triquetra</i> | HtriF | 5'-ATCCTCCTTGCGAGGGTTGG-3' | 200 | 2.5 | 86.27 | 73 | ITS2 (5' → 3') | This study |
| | HtriR | 5'-AATGCGTCAGGGGTGGGAAA-3' | | | | | ITS2 (3' ← 5') | |
| <i>Lingulodinium</i> <i>polyedrum</i> | Lpoly RT-for | 5'-AACTCGTTGGCGAGCATTTTT-3' | 400 | 2.5 | 81.90 | 58 | ITS2 (5' → 3') | Perini et al. (2019) |
| | Lpoly RT-rev | 5'-CGCTAGCAAAGCACTCGCTTA-3' | | | | | ITS2 (3' ← 5') | |
| <i>Protoceratium</i> <i>reticulatum</i> | Pret RT-for | 5'-GGTGCAGTGAAATGTATTAGGCATT-3' | 200 | 2.5 | 78.90 | 83 | 5.8S (5' → 3') | Perini et al. (2018) |
| | Pret RT-rev | 5'-TCCCAAAAACATAGAATACGTTCAAT-3' | | | | | 5.8S (3' ← 5') | |
| <i>S. trochoidea</i> - species complex | StcompF | 5'-GGGTGTGCTTGTGCGTCAAAA-3' | 200 | 1.5 | 83.39 | 81 | ITS2 (5' → 3') | This study |
| | StcompR | 5'-AGCAAGTTGTGCGCCAAGAG-3' | | | | | ITS2 (3' ← 5') | |
| <i>Heterosigma</i> <i>akashii</i> | HaF | 5'-CCGACGGGCGTGGTAGC-3' | 200 | 2.5 | 83.47 | 73 | ITS2 (5' → 3') | Yuan et al. (2015) |
| | HaR | 5'-TCCTCTGTCAGAACAACCGAAGT-3' | | | | | ITS2 (3' ← 5') | |
| Dinophyceae | EUK528f | 5'-CCGCGGTAATCCAGCTC-3' | 300 | 3.5 | 84.07 | 110 | SSU (5' → 3') | Elwood et al. (1985) |
| | Dino18SR1 | 5'-GAGCCAGATRCDCACCCA-3' | | | | | SSU (3' ← 5') | Lin et al. (2006) |

Table S4. List of the environmental variables used in the RDA and MRT. All variables were used to train the MRT, while only those with short names followed by an asterisk (VIF<10) were used in the RDA.

| No. | Name of the variable (mean value if not specified) | Source | Short name |
|-----|-------------------------------------------------------|------------------------------------------------------|-------------------|
| 1 | Sampling depth | Field data from sampling | Depth* |
| 2 | Bottom salinity | Boyer et al. (2005) | Botsal* |
| 3 | Bottom temperature | Boyer et al. (2005) | Bottemp* |
| 4 | Calcite concentration | Feldman & McClain (2010) Tyberghein et al. (2012) | Calcite |
| 5 | Chlorophyll <i>a</i> concentration (mean) | Feldman & McClain (2010) Tyberghein et al. (2012) | Chl |
| 6 | Chlorophyll <i>a</i> concentration (annual range) | Feldman & McClain (2010) Tyberghein et al. (2012) | Chl range |
| 7 | Euphotic depth | Feldman & McClain (2010) | Zeumean |
| 8 | Human impact to marine ecosystems | Halpern et al. (2008) | Impact* |
| 9 | Nitrate concentration | Feldman & McClain (2010) Tyberghein et al. (2012) | NO ₃ |
| 10 | Nutrient input (fertilizers) | Halpern et al. (2008) | Nutrients |
| 11 | Ocean acidification | Halpern et al. (2008) | Acid* |
| 12 | Phosphate concentration | Boyer et al. (2009) Tyberghein et al. (2012) | PO ₄ * |
| 13 | Pollutants (inorganic) | Halpern et al. (2008) | Inorg |
| 14 | Pollutants (organic) | Halpern et al. (2008) | Orgpol |
| 15 | Salinity | Boyer et al. (2009) Tyberghein et al. (2012) | Sal |
| 16 | Sea surface temperature (mean) | Feldman & McClain (2010) Tyberghein et al. (2012) | Sst |
| 17 | Sea surface temperature (annual range) | Feldman & McClain (2010) Tyberghein et al. (2012) | Sstrange |
| 18 | Shipping intensity | Halpern et al. (2008) | Shipping* |
| 19 | Silicate concentration | Boyer et al. (2009) Tyberghein et al. (2012) | Silic |
| 20 | Seafloor slope | GIS based on bathymetry | Slope* |
| 21 | Otter trawling | VMS based estimation (see Russo et al. 2014) | Fishing* |

Table S5. Mean standard curves, mean Ct cell⁻¹, rDNA copy number cell⁻¹ and efficiency obtained from diatom and dinoflagellate cultured strains by qPCR assay.

| Taxon | Mean standard curves | Mean Ct cell ⁻¹ ± SD ^a | rDNA copy number cell ⁻¹ ± SD ^b | Efficiency (%) |
|------------------------------------------|-------------------------|----------------------------------------------|-------------------------------------------------------|----------------|
| <i>Chaetoceros socialis</i> | $y = -3.4696x + 38.441$ | 34.27 ± 0.29 | 15 ± 2 | 94 |
| <i>Ditylum brightwellii</i> | $y = -3.4302x + 34.205$ | 25.48 ± 0.13 | 396 ± 140 | 96 |
| <i>Skeletonema</i> spp. | $y = -3.4549x + 34.536$ | 29.92 ± 0.10 | 25 ± 5 | 95 |
| <i>Thalassiosira</i> spp. | $y = -3.549x + 34.422$ | 25.35 ± 0.17 | 344 ± 38 | 91 |
| <i>A. tamarense/A. mediterraneum</i> | $y = -3.3218x + 36.286$ | 22.51 ± 0.03 | 14676 ± 1662 | 100 |
| <i>Gymnodinium</i> spp. | $y = -3.377x + 35.00$ | 26.41 ± 0.14 | 412 ± 58 | 98 |
| <i>Heterocapsa triquetra</i> | $y = -3.5951x + 35.62$ | 25.71 ± 0.13 | 570 ± 53 | 90 |
| <i>S. trochoidea</i> species complex | $y = -3.5916x + 38.039$ | 24.33 ± 0.13 | 5625 ± 1375 | 90 |
| <i>Heterosigma akashiwo</i> ^c | $y = -3.406x + 38.253$ | n.d. | 367 ± 12 | 97 |

^amean Ct values measured in 2 independent experiments ± standard deviation (SD)

^bmean rDNA gene copy number calculated in triplicate ± standard deviation (SD)

^cfrom Yuan et al. (2015)

n.d., not determined

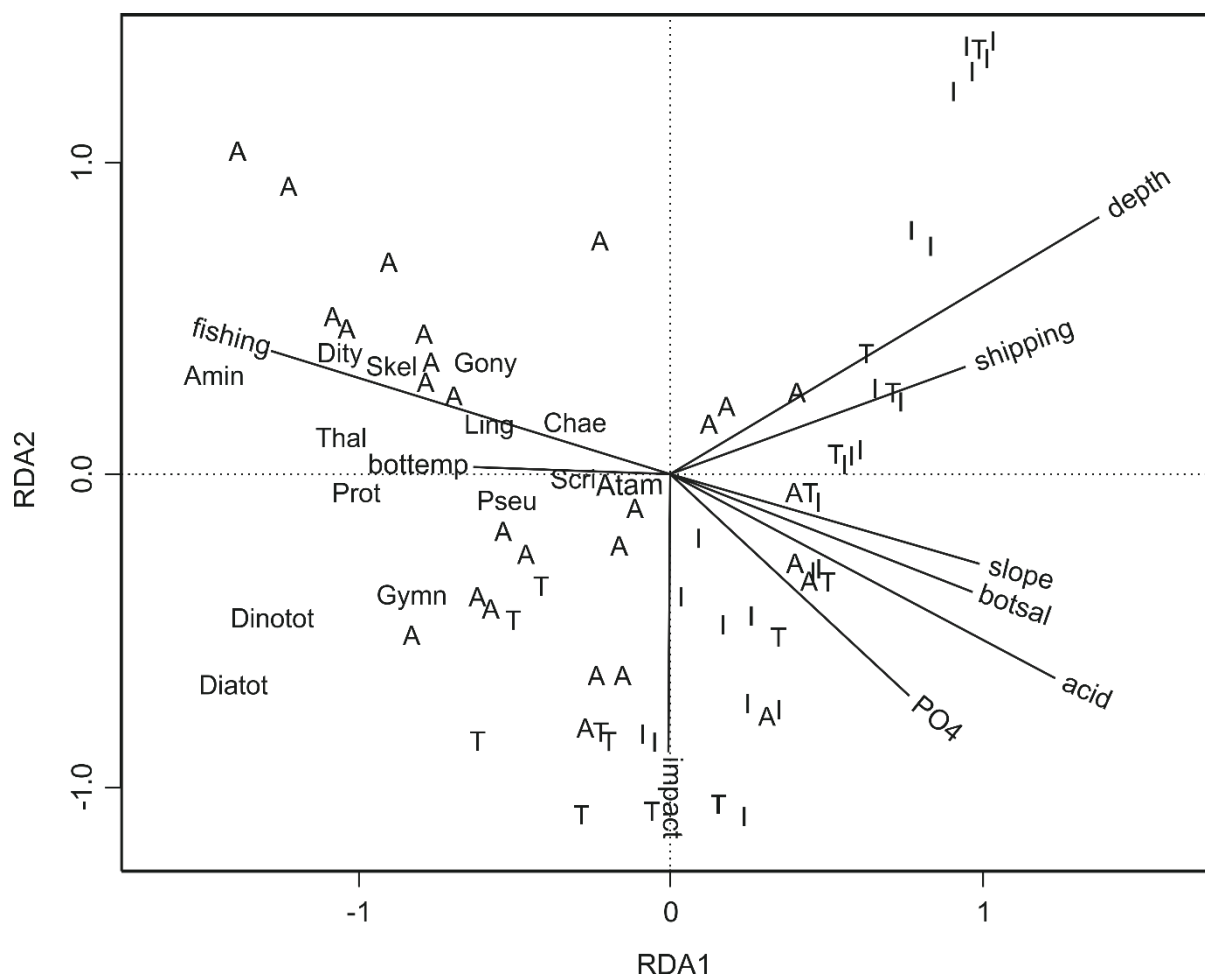


Figure S1. Redundancy Analysis triplot: samples are labelled according to the basin where they were collected (A = Adriatic Sea, I = Ionian Sea and T = Tyrrhenian Sea), while short names are shown for taxa and for environmental variables (solid lines). Full names can be found in Table 1. All the resting stages point towards the positive end of the first canonical axis (RDA1), thus showing that samples that are found in that semi-plane are the most abundant. The latter include almost all the samples from the Adriatic Sea. Short names for the resting stage taxa can be found in Table 1 next to the full names, while those for the environmental variables are shown in the rightmost column in Table S4. Three resting stage taxa (Hets, Apac and Hetc) whose coordinates were very close to the origin and therefore very poorly represented, are not shown in the ordination. The first two canonical axes, both significant to the permutation test, accounted for a very large share of variance, i.e. 49.88% (RDA1 44.27% and RDA2 5.59%).

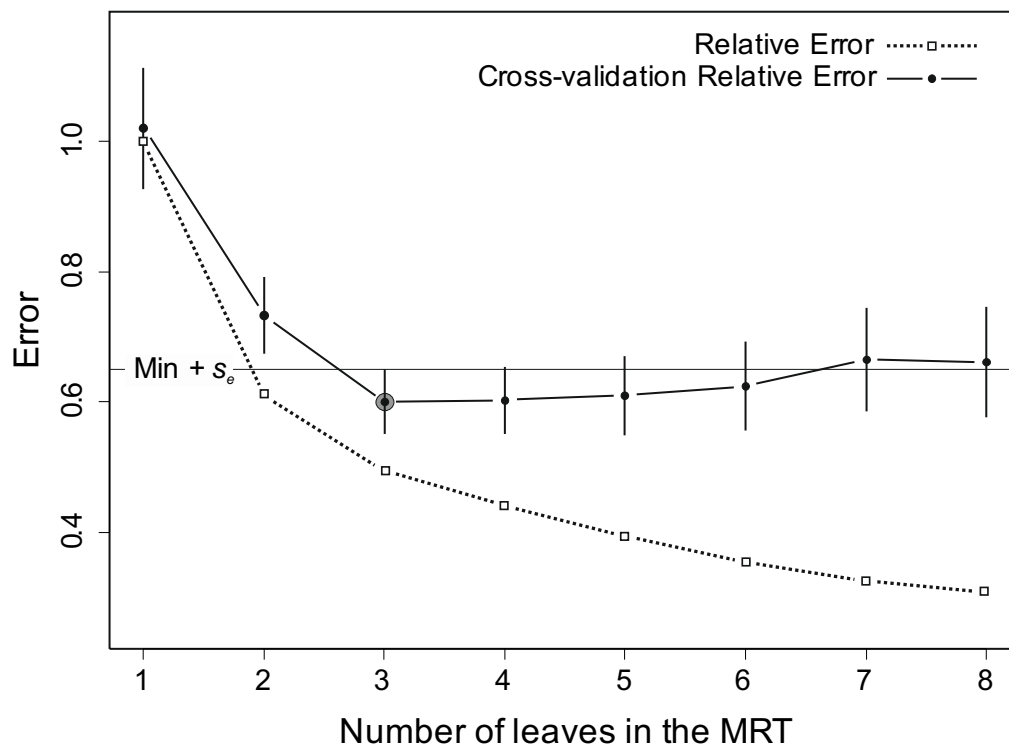


Figure S2. Relative error (dotted line) and cross-validation relative error (solid line) for the Multivariate Regression Tree (MRT). The best MRT has three leaves and corresponds both to the minimum cross-validation error and to the most parsimonious solution among those within one standard error above the optimum (i.e. below the solid horizontal line).

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