## **Supplementary Materials**

## Distribution of nitrogen-cycling genes in an oxygen-depleted cyclonic eddy in the Alfonso Basin, Gulf of California

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## Table S1. qPCR primers, thermal cycling and reaction conditions used in this study

| Gene              | Primers            | Sequence (5'-3')                       | Cycling conditions   | Reaction conditions                 | Efficiency | Reference  |
|-------------------|--------------------|--|--|-------------------------------------|------------|--|
| Archaeal          | Arch-amoAF         | STAATGGTCTGGCTTAGACG                   | 95°C 5 min, 40 x (95°C 45  | 0.2 μM                              |            | Francis <i>et</i>  |
| amoA              | Arch-amoAR         | GCGGCCATCCATCTGTATGT                   | s, 53°C 1 min, 72° 1 min),<br>72°C 5 min   | primers, 0.3<br>µg/µl BSA           | 84%        | al. (2005)   |
| D ( 1             | amoA1F             | GGGGTTTCTACTGGTGGT                     | 95°C 5 min, 40 x (95°C 45  | 02 14                               |            | Rotthauwe  |
| Bacterial<br>amoA | amoA2R             | CCCCTCKGSAAAGCCTTCTTC                  | s, 55°C 45 s, 72° 45 s),<br>72°C 5 min   | 0.3 μM<br>primers                   | 111.35%    | <i>et al.</i><br>(1997)  |
|                   | nirS-cd3aF         | GTSAACGTSAAGGARACSGG                   | Touchdown (-1°C/cycle)<br>95°C 5 min, 6 x (95°C 30<br>s, 62°C 20 s, 72°C 40 s),    | 0.3 μΜ                              |            | Michotey   |
| nirS              | nirS-R3cd          | GASTTCGGRTGSGTCTTGA                    | 35 x (95°C 30 s, 60°C 20 s,<br>72°C 40 s), 72°C 5 min                              | primers, 0.3<br>µg/µl BSA           | 82%        | <i>et al.</i><br>(2000)  |
|                   | nirK-876F          | ATYGGCGGVAYGGCGA                       | Touchdown (-1°C/cycle)<br>95°C 5 min, 6 x (95°C 15                                 |                                     |            |  |
| nirK              | nirK-1040R         | GCCTCGATCAGRTTRTGGTT                   | s, 65°C 30 s, 72°C 30 s),<br>35 x (95°C 15 s, 60°C 30 s,<br>72°C 30 s), 72°C 5 min | 0.5 μM<br>primers, 0.3<br>μg/μl BSA | 96%        | Henry <i>et</i><br><i>al.</i> (2004)                                   |
| nrfA              | nrfAf2aw<br>nrfAR1 | CARTGYCAYGTBGARTA<br>TWNGGCATRTGRCARTC | 95°C 5 min, 40 x (95°C 15<br>s, 52°C 30 s, 72°C 30 s),<br>72°C 5 min               | 0.5 μM<br>primers, 0.2<br>μg/μl BSA | 93.47%     | Mohan <i>et</i><br><i>al.</i> (2004),<br>Welsh et<br><i>al.</i> (2014) |
| hzo               | hzoF1              | TGTGCATGGTCAATTGAAAG                   | 95°C 5 min, 40 x (95°C 45  | $0.4 \mu M$                         |            | Listal   |
|                   | hzoR1              | CAACCTCTTCWGCAGGTGCATG                 | min), 72°C 5 min   | µg/µl BSA                           | 88.36%     | (2010)   |

Table S2. Multiple linear regression models and their parameters for the N cycling genes Significance codes: \*\*\* = 0.001, \*\* = 0.01, \* = 0.05, . = 0.1.

| Gene   | Linear models                            |
|--------|--|
| A-amoA | lm (A-amoA ~ Oxygen + NO3)               |
| B-amoA | lm (B-amoA ~ Oxygen + NO2)               |
| nirK   | lm (nirK ~ Oxygen + Chlorophyll + NO3)   |
| nirS   | lm (nirS ~ Salinity + NO3)               |
| nrfA   | $lm (nrfA \sim Chlorophyll + NH4 + NO3)$ |
| hzo    | lm (hzo ~ Chlorophyll + NH4 + NO3)       |

| Coefficients          | Estimate | Std. Error | t value | <b>Pr(&gt; t )</b> | vif      |
|-----------------------|----------|------------|---------|--------------------|----------|
| A-amoA                |          |            |         |                    |          |
| Intercept             | 9.35132  | 0.48887    | 19.128  | 2.52E-14***        |          |
| Oxygen                | 0.15871  | 0.07966    | 1.992   | 0.0602 .           | 1.75753  |
| NO <sub>3</sub> -     | 0.14927  | 0.10739    | 1.39    | 0.1798             | 1.75753  |
| B-amoA                |          |            |         |                    |          |
| Intercept             | 1.4727   | 0.6357     | 2.316   | 0.0313*            |          |
| Oxygen                | 0.3439   | 0.1421     | 2.42    | 0.0252*            | 1.04273  |
| $\mathbf{NO}_2^-$     | -8.6419  | 3.7636     | -2.296  | 0.0326*            | 1.04273  |
| nirK                  |          |            |         |                    |          |
| Intercept             | 5.6996   | 0.7304     | 7.804   | 0.000000242***     |          |
| Oxygen                | -0.3507  | 0.136      | -2.579  | 0.01838*           | 2.404638 |
| Chlorophyll           | -0.8957  | 0.3061     | -2.927  | 0.00866**          | 1.4009   |
| $NO_3^-$              | 0.6143   | 0.162      | 3.792   | 0.00123**          | 1.87807  |
| nirS                  |          |            |         |                    |          |
| Intercept             | 121.6614 | 26.7753    | 4.544   | 0.000198***        |          |
| Salinity              | -33.1173 | 7.4669     | -4.435  | 0.000254***        | 1.01473  |
| $NO_3^-$              | 0.2307   | 0.0648     | 3.559   | 0.001965**         | 1.01473  |
| nrfA                  |          |            |         |                    |          |
| Intercept             | 2.0502   | 0.6635     | 3.09    | 0.00602**          |          |
| Chlorophyll           | -3.6969  | 0.4627     | -7.99   | 1.71E07***         | 1.0322   |
| $\mathbf{NH_4}^+$     | 1.9819   | 0.5269     | 3.762   | 0.00132**          | 1.0285   |
| $NO_3^-$              | 1.3724   | 0.2135     | 6.426   | 3.67E06***         | 1.04834  |
| hzo                   |          |            |         |                    |          |
| Intercept             | 2.0432   | 0.7658     | 2.668   | 0.015204*          |          |
| Chlorophyll           | -3.9209  | 0.5341     | -7.341  | 5.86E-7***         | 1.0322   |
| $\mathbf{NH}_{4}^{+}$ | 2.4124   | 0.6082     | 3.967   | 0.000827***        | 1.0285   |
| $NO_3^-$              | 1.5256   | 0.2465     | 6.189   | 6.03E-6***         | 1.04834  |

|        |          | Residu  | als     |        |        | Residual<br>std.<br>error | Mult. $R^2$ | Adj.<br>R <sup>2</sup> | <i>F</i> -statistic | <i>p</i> -<br>value |
|--------|----------|---------|---------|--------|--------|---------------------------|-------------|------------------------|---------------------|---------------------|
| Gene   | Min      | 1Q      | Median  | 3Q     | Max    | 20 d.f.                   |             |                        |                     |                     |
| A-amoA | -0.83204 | -0.2905 | -0.0682 | 0.2206 | 0.7726 | 0.4402                    | 0.166       | 0.0826                 | 1.99                | 0.1638              |
| B-amoA | -1.7464  | -0.713  | -0.2175 | 0.6764 | 2.2665 | 1.02                      | 0.316       | 0.2481                 | 4.629               | 0.0223              |
| nirK   | -0.8509  | -0.6006 | 0.1562  | 0.4334 | 1.052  | 0.6424                    | 0.842       | 0.8171                 | 33.76               | 8.13e <sup>-8</sup> |
| nirS   | -0.6903  | -0.2242 | -0.0411 | 0.1808 | 0.6452 | 0.3496                    | 0.647       | 0.6118                 | 18.34               | 2.99e <sup>-5</sup> |
| nrfA   | -1.53598 | -0.8664 | -0.05   | 0.7162 | 1.9986 | 1.133                     | 0.89        | 0.8726                 | 51.23               | 2.69e <sup>-9</sup> |
| hzo    | -1.7465  | -0.9536 | 0.1803  | 0.7046 | 2.8353 | 1.308                     | 0.881       | 0.8617                 | 46.68               | 5.85e <sup>-9</sup> |

| Shapiro-Wilk normality test |              |                 |  |  |  |  |
|-----------------------------|--------------|-----------------|--|--|--|--|
| Gene                        | $\mathbf{W}$ | <i>p</i> -value |  |  |  |  |
| A-amoA                      | 0.95872      | 0.4379          |  |  |  |  |
| B-amoA                      | 0.94847      | 0.2719          |  |  |  |  |
| nirK                        | 0.92372      | 0.08008         |  |  |  |  |
| nirS                        | 0.9665       | 0.6059          |  |  |  |  |
| nrfA                        | 0.9383       | 0.1652          |  |  |  |  |
| hzo                         | 0.94595      | 0.2407          |  |  |  |  |



**Fig. S1.** T-S diagram of the water masses in the Bay of La Paz. TSW, Tropical Surface Water; GCW, Gulf of California Water, StSsW, Subtropical Subsurface Water.



**Fig. S2.** Depth profiles of ammonium, nitrite, nitrate, phosphate, and silicon in the northern (up) and southern (down) sections.

10

15

5

400

0



**Fig. S3.** Nitrogen deficit (N\*) in the northern (up) and southern (down) sections.