

Global patterns of phosphatase activity in natural soils

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SUPPLEMENTARY INFORMATION

Competing financial interests

The author(s) declare no competing financial interests

Fig S1 Histogram

Histogram of Acid phosphatase measurements compiled in these database (left, n=329). We log-transformed Acid phosphatase values for the modeling to ensure that the residuals were approximately normally distributed (right, n=329).

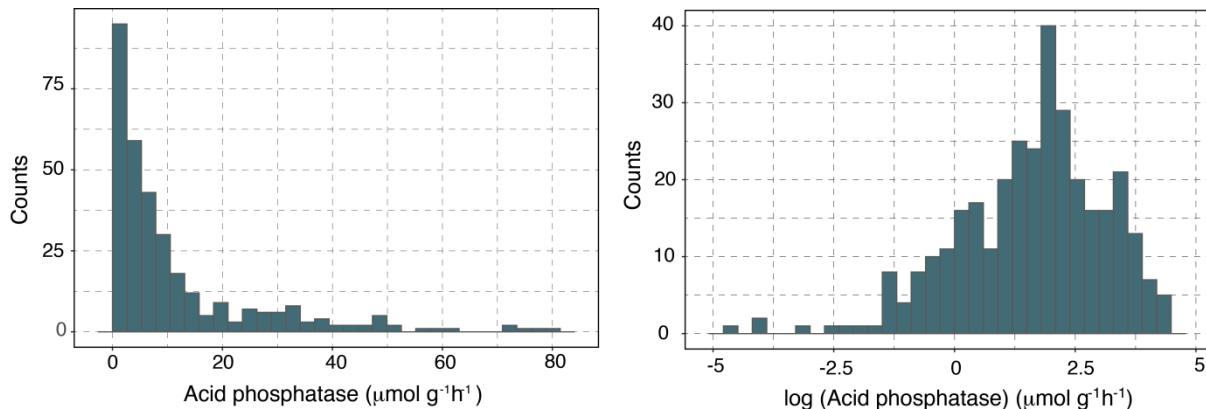
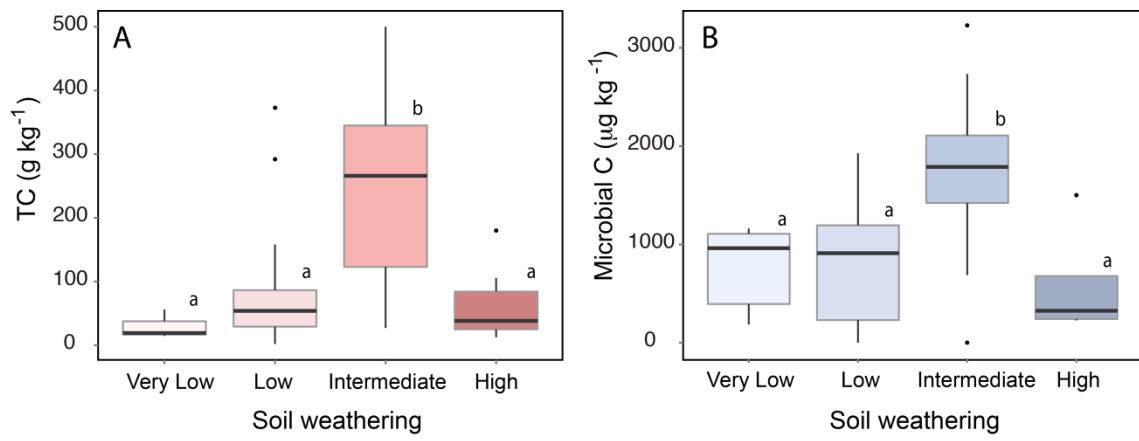


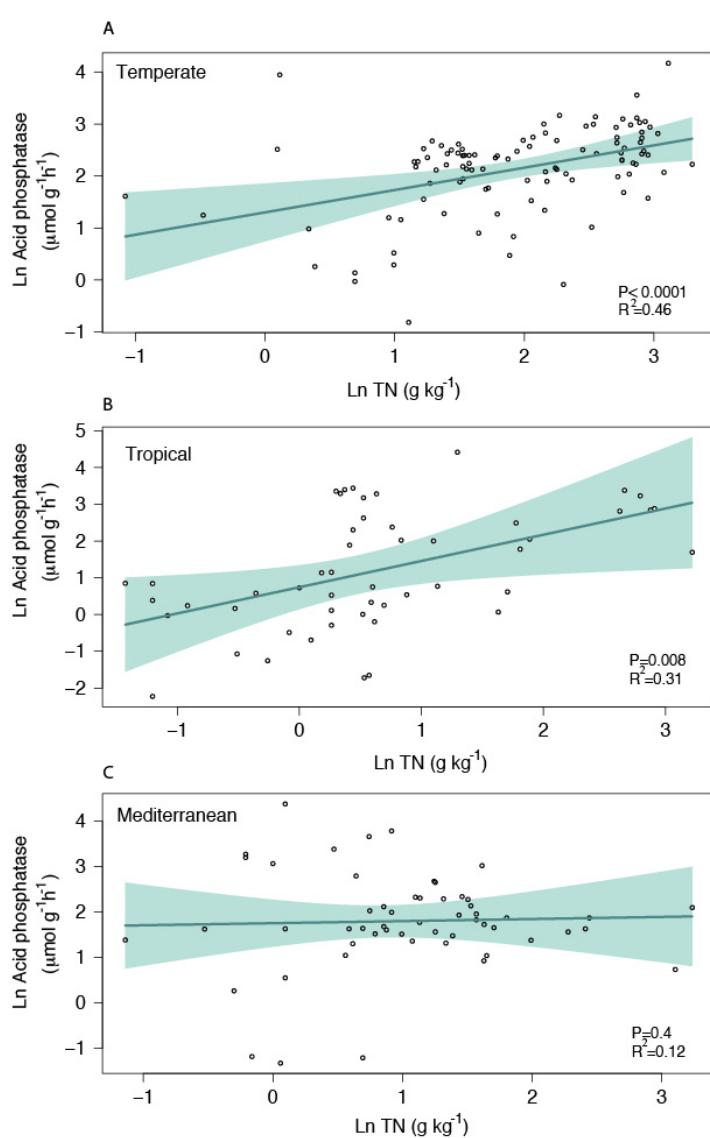
Fig S2 Soil weathering stages traits

Dependence of TC (A) and microbial C (B) on the amount of soil weathering (very low, low, intermediate and high). Boxplot show median values (solid horizontal line), 50th percentile values (box outline), 90th percentile values (whiskers), and outlier values. Letters represent the results of Tukey's post-hoc comparisons of group means.



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Fig S3 Effect of TN across biomes47
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Partial residual plot of the variability of global Ln phosphatase activity ($\mu\text{mol g}^{-1} \text{h}^{-1}$) explained by Ln TN (g kg^{-1})
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(visreg R package) for temperate (A), tropical (B) and mediterranean (C) biomes.

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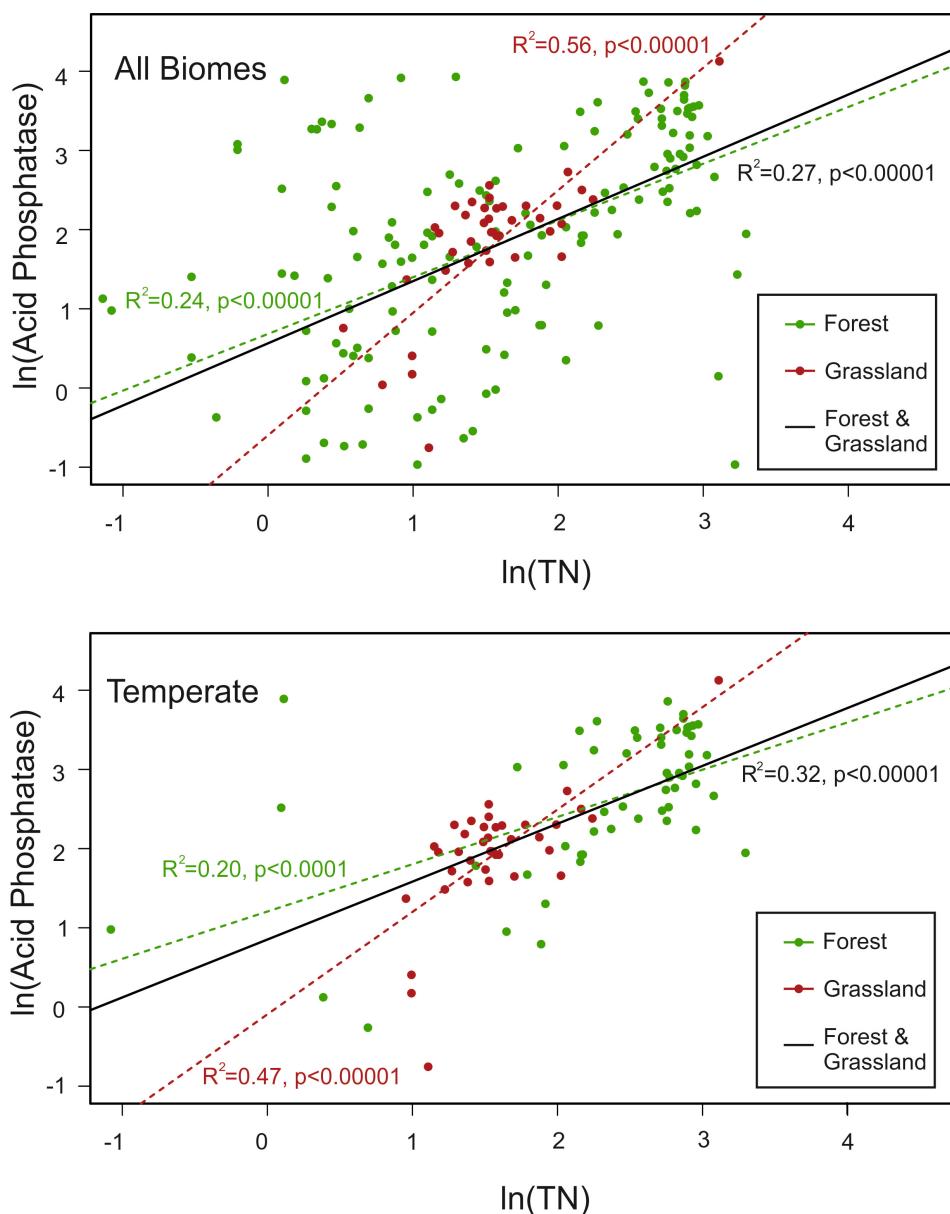
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Fig S3 Effect of TN across vegetation type

Scatterplot showing the relationship between Acid Phosphatase and TN. Variables were Ln-transformed. Forest vs. Grassland communities are highlighted. Grassland category includes grasslands and pastures.



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Table S1. Linear model with and without interactions combining TN, MAT, MAP and TN, MAT, MAP, TC and AMP as explanatory variables of acid phosphatase activity. Signif. codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

| | Estimate | SE | t | P | Sig. Code | β Coeff |
|---|------------|-----------|--------|----------|-----------|---------------|
| Model 1: TN, MAP, MAT with interactions, Multiple R2: 0.27, adjusted R2: 0.26 $F: 29.04$ on 3 and 235 DF, $P: 5.253e^{-16}$, No. observations in the fit: 239 | | | | | | |
| Intercept | -52.681 | 26.39 | -1.97 | 0.0471 | * | |
| Ln TN | 0.765 | 0.091 | 8.398 | 4.31E-15 | *** | 0.553 |
| Ln MAP | 0.079 | 0.166 | 0.477 | 0.634 | | 0.03 |
| Ln MAT | 9.323 | 4.73 | 1.97 | 0.05 | . | 0.133 |
| Model 2: TN, MAP, MAT with interactions, Multiple R2: 0.30, adjusted R2: 0.28 $F: 16.4$ on 6 and 232 DF, $P: 9.472e^{-16}$, No. observations in the fit: 239 | | | | | | |
| Intercept | -754 | 249 | -3.03 | 0.0027 | ** | - |
| Ln TN | 0.8 | 20.9 | 0.038 | 0.9695 | | - |
| Ln MAP | 100.43 | 35.9 | 2.8 | 0.0056 | ** | - |
| Ln MAT | 133.2 | 43.9 | 3.04 | 0.0027 | ** | - |
| Ln TN:Ln MAP | 0.053 | 0.14 | 0.38 | 0.71 | | - |
| Ln TN:Ln MAP | -0.076 | 3.75 | -0.02 | 0.98 | | - |
| Ln MAT:Ln MAP | -17.7 | 6.34 | -2.8 | 0.0056 | ** | - |
| Model 3 : TN, MAP, MAT, AMP, TC without interactions, Multiple R2: 0.50, adjusted R2: 0.49 $F: 21.02$ on 5 and 104 DF, $P: <1.798e^{-14}$ No. observations in the fit: 110 | | | | | | |
| Intercept | -52.681 | 26.39 | -1.97 | 0.0471 | ** | |
| Ln TN | 0.765 | 0.091 | 8.398 | 4.31E-15 | *** | 0.697 |
| Ln MAP | -0.19 | 0.329 | -0.578 | 0.564 | | -0.051 |
| Ln MAT | 27.53 | 7.988 | 3.446 | 0.0008 | *** | 0.338 |
| Ln TC | -0.012 | 0.1454 | -0.083 | 0.933 | | -0.009 |
| Ln Amp | -1.075 | 0.659 | -1.633 | 0.1056 | | -0.161 |
| Model 4 : TN, MAP, MAT, AMP, TC with interactions, Multiple R2: 0.67, adjusted R2: 0.62 $F: 12.81$ on 15 and 94 DF, $P: <2.2e^{-16}$ No. observations in the fit: 110 | | | | | | |
| Intercept | 2888.3565 | 1199.5565 | 2.408 | 0.017999 | * | - |
| Ln TN | 133.01929 | 93.06713 | 1.429 | 0.156238 | | - |
| Ln MAP | -210.50512 | 137.94625 | -1.526 | 0.130369 | | - |
| Ln MAT | -507.95906 | 209.42456 | -2.425 | 0.017197 | * | - |
| Ln Amp | -56.13413 | 71.09077 | -0.79 | 0.431742 | | - |
| Ln TC | -523.8962 | 135.09173 | -3.878 | 0.000195 | *** | - |
| Ln TN:Ln MAP | 1.04137 | 0.60578 | 1.719 | 0.088896 | . | - |
| Ln TN:Ln MAP | -23.62698 | 15.47049 | -1.527 | 0.130062 | | - |
| Ln MAT:Ln MAP | 37.7051 | 23.88633 | 1.579 | 0.117805 | | - |
| Ln TN:Ln AMP | -0.09707 | 0.14196 | -0.684 | 0.495789 | | - |
| Ln MAP:Ln AMP | -0.07585 | 0.36584 | -0.207 | 0.836205 | | - |
| Ln MAT:Ln AMP | 8.37083 | 11.9161 | 0.702 | 0.484115 | | - |
| Ln TN:Ln TC | -1.73328 | 1.30457 | -1.329 | 0.187189 | | - |
| Ln MAP:Ln TC | -1.17335 | 1.16879 | -1.004 | 0.318003 | | - |
| Ln MAT:Ln TC | 92.04352 | 24.20831 | 3.802 | 0.000255 | *** | - |
| Ln Amp:Ln TC | 3.12774 | 1.03766 | 3.014 | 0.003312 | ** | - |

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Table S2. Linear model with and without interactions combining TN, MAT, MAP and TN, MAT, MAP, TC and AMP as
explanatory variables of acid phosphatase activity for only temperate sites. Signif. codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’
0.1 ‘ ’ 1

| | Estimate | SE | t | P | Sig. Code | β Coeff |
|---|------------|----------|--------|----------|-----------|---------------|
| Model 1: TN, MAP, MAT with interactions, Multiple R2: 0.41, adjusted R2: 0.39 F: 22.98 on 3 and 100 DF, P: 2.135e ⁻¹¹ , No. observations in the fit: 104 | | | | | | |
| Intercept | -131.89 | 52.032 | -2.535 | 0.0128 | * | |
| Ln TN | 0.514 | 0.1081 | 4.751 | 6.78E-06 | *** | 0.416 |
| Ln MAP | 0.6506 | 0.2279 | 2.855 | 0.00523 | ** | 0.239 |
| Ln MAT | 22.77 | 9.21 | 2.47 | 0.015 | * | 0.199 |
| Model 2: TN, MAP, MAT with interactions, Multiple R2: 0.46, adjusted R2: 0.42 F: 13.57 on 6 and 97 DF, P: 3.998e ⁻¹¹ No. observations in the fit: 104 | | | | | | |
| Intercept | -552 | 769 | -0.717 | 0.475 | | - |
| Ln TN | -90 | 61.1 | -1.47 | 0.144 | | - |
| Ln MAP | 78.7 | 103 | 0.77 | 0.446 | | - |
| Ln MAT | 98 | 137 | 0.718 | 0.475 | | - |
| Ln TN:Ln MAP | 0.4329 | 0.34 | 1.26 | 0.21 | | - |
| Ln TN:Ln MAT | 15.5 | 11 | 1.4 | 0.164 | | - |
| Ln MAT:Ln MAP | -13.9 | 18.3 | -0.764 | 0.447 | | - |
| Model 3: TN, MAP, MAT, AMP, TC without interactions, Multiple R2: 0.50, adjusted R2: 0.56 F: 21.22 on 5 and 73 DF, P: <4.853e ⁻¹³ No. observations in the fit: 79 | | | | | | |
| Intercept | -191.5875 | 79.3329 | -2.415 | 0.0182 | * | |
| Ln TN | 0.6765 | 0.162 | 4.176 | 8.11E-05 | *** | 0.517 |
| Ln MAP | 0.8366 | 0.3351 | 2.496 | 0.0148 | * | 0.237 |
| Ln MAT | 33.79 | 13.94 | 2.424 | 0.017 | * | 0.2955 |
| Ln TC | -0.1858 | 0.1472 | -1.262 | 0.2108 | | -0.1947 |
| Ln Amp | -1.0891 | 0.672 | -1.621 | 0.1094 | | -0.1463 |
| Model 4: TN, MAP, MAT, AMP, TC with interactions, Multiple R2: 0.84, adjusted R2: 0.80 F: 21.45 on 15 and 63 DF, P: <2.2e ⁻¹⁶ , No. observations in the fit: 79 | | | | | | |
| Intercept | 3834.4786 | 2779.078 | 1.38 | 0.17254 | *** | - |
| Ln TN | 6.3961 | 137.9777 | 0.046 | 0.96317 | *** | - |
| Ln MAP | 3.3291 | 315.9972 | 0.011 | 0.99163 | | - |
| Ln MAT | -666.9843 | 492.8147 | -1.353 | 0.18076 | | - |
| Ln Amp | -120.9173 | 92.7636 | -1.303 | 0.19715 | | - |
| Ln TC | -1150.3699 | 344.781 | -3.337 | 0.00143 | ** | - |
| Ln TN:Ln MAP | 0.2948 | 0.9338 | 0.316 | 0.75325 | | - |
| Ln TN:Ln MAT | -1.3264 | 23.8159 | -0.056 | 0.95576 | | - |
| Ln MAT:Ln MAP | -2.3051 | 55.4016 | -0.042 | 0.96694 | | - |
| Ln TN:Ln AMP | -0.4588 | 0.1509 | -3.041 | 0.00343 | ** | - |
| Ln MAP:Ln AMP | 0.6284 | 0.809 | 0.777 | 0.4402 | | - |
| Ln MAT:Ln AMP | 20.9023 | 15.639 | 1.337 | 0.18618 | | - |
| Ln TN:Ln TC | 0.6802 | 1.6586 | 0.41 | 0.68315 | | - |
| Ln MAP:Ln TC | 2.2726 | 3.3639 | 0.676 | 0.50178 | | - |
| Ln MAT:Ln TC | 200.6906 | 62.6777 | 3.202 | 0.00214 | ** | - |
| Ln Amp:Ln TC | -0.2184 | 1.4342 | -0.152 | 0.87947 | | - |

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72 **Table S3.** Linear model with interactions and including TN, MAT, MAP and pH as explanatory variables of alkaline
73 phosphatase activity. Signif. codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

| | Estimate | SE | t | P | Sig. Code |
|---|----------|--------|-------|----------|-----------|
| Model alkaline: TN, MAP, MAT and pH with interactions, Multiple R2: 0.56, adjusted R2: 0.42 <i>F</i> : 4.246 on 15 and 51 DF, <i>P</i> : 5.099e ⁻⁰⁵ , No. observations in the fit: 67 | | | | | |
| Intercept | 21061.3 | 5784.8 | 3.64 | 0.000635 | *** |
| Ln TN | -8225.5 | 5729.4 | -1.44 | 0.157205 | |
| Ln MAP | -2817.6 | 784.7 | -3.59 | 0.000741 | *** |
| Ln MAT | -3717 | 1022.5 | -3.64 | 0.000645 | *** |
| Ln pH | -10376.5 | 2839.8 | -3.65 | 0.00061 | *** |
| Ln TN:Ln MAP | 1283.4 | 801 | 1.60 | 0.115289 | |
| Ln TN:Ln MAT | 1437.5 | 1012.1 | 1.42 | 0.161562 | |
| Ln MAT:Ln MAP | 497.1 | 138.6 | 3.59 | 0.00075 | *** |
| Ln TN:Ln pH | 5271.1 | 2966.6 | 1.78 | 0.081567 | . |
| Ln MAP:Ln pH | 1391.9 | 391.6 | 3.55 | 0.000828 | *** |
| Ln MAT:Ln pH | 1831.6 | 501.9 | 3.65 | 0.000619 | *** |
| Ln TN:Ln MAP:Ln MAT | -224.5 | 141.4 | -1.59 | 0.118556 | |
| Ln TN:Ln MAP:Ln pH | -810.3 | 422.8 | -1.92 | 0.060913 | . |
| Ln TN:Ln MAT:Ln pH | -922.4 | 523 | -1.76 | 0.084282 | . |
| Ln MAP:Ln MAT:Ln pH | -245.5 | 69.1 | -3.55 | 0.000838 | *** |
| Ln TN: Ln MAP:Ln MAT:Ln pH | 141.9 | 74.6 | 1.90 | 0.062919 | . |

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78 **Database literature**
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