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Supplemental information

Evolution of ion transporter Na⁺/K⁺-ATPase expression in the osmore-

gulatory maxillary glands of an invasive copepod

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Figure S1. NKA *in situ* expression in response to treatment salinities (0, 5, 15 PSU) for saline (SW) and freshwater (FW) populations, measured under common garden conditions, related to Figure 7. (A) Total NKA immunolabeled maxillary gland (IMG) area calculated as the sum of area (μm^2) in all cuts multiplied by the thickness of each cut (5 μ m) divided by the prosome length (μ m) of that individual. (B) The maximum NKA IMG intensity (in pixel value). Data are represented as Mean ± SE. P < 0.05 = *. Mean values and standard errors are reported in Table S3.

Table S1. Statistical results for comparisons of body size (prosome length μ m) and Whole-Animal Na⁺/K⁺-ATPase expression between saline *versus* freshwater populations and between salinity treatments, related to Figure 2 and 3 Two-way ANOVA tests were performed, followed by Tukey post hoc tests. Additional *a priori* pairwise comparisons were performed between populations at each treatment salinity using Student's t-tests. Results show the effects of the factors *Population* (saline vs. freshwater populations), *Salinity* (0, 5, 15 PSU), and the interaction of *Population* × *Salinity* on body size and Na⁺/K⁺-ATPase expression. df = degrees of freedom. Significant results are indicated as * = *P* < 0.05, *** = *P* < 0.0001.

| Comparison (greater value indicated by > and <) | | Test | Test statistic | df | P-value |
|---|---------------------------------------|------------------------|-------------------|-------|------------|
| A. Body | y Size (μm) – log transformed | | | | |
| Population | | Two-way ANOVA | F = 0.693 | 1,284 | 0.406 |
| | 0 PSU | a priori t-test | t = 0.40 | 111 | 0.60 |
| | 5 PSU | a priori t-test | <i>t</i> = -1.13 | 77 | 0.262 |
| | 15 PSU, fresh > saline population | a priori t-test | <i>t</i> = 1.95 | 85 | 0.050* |
| Salinity | | Two-way ANOVA | F = 39.67 | 2,283 | <0.0001*** |
| | Saline population, $0 < 5 PSU$ | Tukey HSD | | | <0.0001*** |
| | Saline population, 0 < 15 PSU | Tukey HSD | | | <0.0001*** |
| | Saline population, 5 vs 15 PSU | Tukey HSD | | | 0.472 |
| | Freshwater population, 0 < 5 PSU | Tukey HSD | | | 0.00319* |
| | Freshwater population, 0 < 15 PSU | Tukey HSD | | | <0.0001*** |
| | Freshwater population, 5 < 15 PSU | Tukey HSD | | | 0.0360* |
| Population x Salinity interaction | | Two-way ANOVA | F = 3.02 | 2,283 | 0.050* |
| B. Who | le-Animal NKA – cube root transformed | | | | |
| Population | | Two-way ANOVA | F = 0.805 | 1,35 | 0.376 |
| | 0 PSU | <i>a priori</i> t-test | t = -0.650 | 14 | 0.526 |
| | 5 PSU | <i>a priori</i> t-test | t = 0.929 | 9 | 0.373 |
| | 15 PSU, saline > fresh population | a priori t-test | <i>t</i> = -2.17 | 11 | 0.050* |
| Salinity | | Two-way ANOVA | F = 3.30 | 2,34 | 0.0490* |
| | Saline population, 0 vs 5 PSU | Tukey HSD | | | 0.892 |
| | Saline population, $0 < 15$ PSU | Tukey HSD | | | 0.0169* |
| | Saline population, 5 < 15 PSU | Tukey HSD | | | 0.0444* |
| | Freshwater population, 0 vs 5 PSU | Tukey HSD | | | 0.0607 |
| | Freshwater population, 0 vs 15 PSU | Tukey HSD | | | 0.361 |
| | Freshwater population, 5 vs 15 PSU | Tukey HSD | | | 0.350 |
| Population x Salinity interaction | | Two-way ANOVA | F = 2.32 | 2,34 | 0.114 |

Table S2: Statistical results for comparisons of *in situ* Na⁺/K⁺-ATPase expression in the maxillary glands between saline *versus* freshwater populations and between salinity treatments, related to Figure 7. Analysis was done using a two-way ANOVA or Kruskal-Wallis test with *a priori* pairwise Student's t-tests and *a priori* Wilcoxon ranked sum tests. Results show effects of the factors *Population* (saline vs. freshwater populations), *Salinity* (0, 5, 15 PSU), and the interaction of *Population × Salinity* on Na⁺/K⁺-ATPase expression. df = degrees of freedom. *Post-hoc* comparisons are reported if significant differences were observed at P < 0.05 = *.

| (| Comparis | son (greater value indicated by >) | Test | Test statistic | df | <i>P</i> - value |
|---|----------|--|------------------------------|-------------------|-------|---------------------|
| | A. NKA | IMG Area -square root transforme | d | | | |
| | Populati | on | Two-way ANOVA | F = 2.33 | 1, 35 | 0.137 |
| | | 0 PSU, saline > fresh | <i>a priori</i> t-test | t = -2.83 | 10 | 0.0181* |
| | | 5 PSU | <i>a priori</i> t-test | t = -0.287 | 10 | 0.780 |
| | | 15 PSU | <i>a priori</i> t-test | t = 0.600 | 11 | 0.560 |
| | Salinity | | Two-way ANOVA | F = 1.32 | 2, 34 | 0.281 |
| | | Saline population, 0 <i>vs</i> 5 PSU | <i>a priori</i> t-test | <i>t</i> = 2.10 | 10 | 0.0620 |
| | | Saline population, $0 > 15 PSU$ | <i>a priori</i> t-test | t = 2.47 | 11 | 0.0332* |
| | | Saline population, 5 vs 15 PSU | a priori t-test | t = 0.0117 | 11 | 0.991 |
| | | Freshwater population, 0 vs 5 PSU | a priori t-test | t = -0.140 | 10 | 0.891 |
| F | PSU | Freshwater population, 0 vs 15 | a priori t-test | <i>t</i> = -1.12 | 11 | 0.286 |
| | PSU | Freshwater population, 5 vs 15 | a priori t-test | <i>t</i> = -1.22 | 11 | 0.249 |
| | Populati | on x Salinity interaction | Two-way ANOVA | F = 3.17 | 2, 34 | 0.050* |
| | B. NKA | IMG Intensity | | | | |
| | Populati | on | Kruskal Wallis | H = 0.577 | 1, 35 | 0.447 |
| | | 0 PSU, saline > fresh | a priori Wilcoxon ranked sum | W = 5 | n = 6 | 0.0411* |
| | | 5 PSU | a priori Wilcoxon ranked sum | W = 20 | n = 6 | 0.818 |
| | | 15 PSU | a priori Wilcoxon ranked sum | W = 27 | n = 6 | 0.445 |
| | Salinity | | Kruskal Wallis | <i>H</i> = 1.23 | 2, 34 | 0.542 |
| | | Saline population, 0 vs 5 PSU | a priori Wilcoxon ranked sum | <i>W</i> = 30 | n = 6 | 0.0649 |
| | | Saline population, 0 vs 15 PSU | a priori Wilcoxon ranked sum | W = 27 | n = 6 | 0.179 |
| | | Saline population, 5 <i>v</i> s 15 PSU | a priori Wilcoxon ranked sum | <i>W</i> = 13 | n = 6 | 0.485 |
| | | Freshwater population, 0 vs 5 PSU | a priori Wilcoxon ranked sum | <i>W</i> = 14 | n = 6 | 0.589 |
| | | Freshwater population, 0 vs 15 PSU | a priori Wilcoxon ranked sum | <i>W</i> = 10 | n = 6 | 0.1375 |
| | PSU | Freshwater population, 5 vs 15 | a priori Wilcoxon ranked sum | <i>W</i> = 15 | n = 6 | 0.445 |
| | Populati | on x Salinity interaction | Kruskal Wallis | <i>H</i> = 7.70 | 5 | 0.174 |

Table S3. Values of body size and Na⁺/K⁺-ATPase (NKA) expression of freshwater and saline populations across three salinities (0, 5, 15 PSU), related to Figure 2, 3, and 7. Response variables include *Body Size, Whole-Animal NKA* band signal intensity, *NKA IMG Area, NKA IMG Intensity*. Mean values and standard errors of the mean (SE) are reported.

| Population | Salinity (PSU) | Mean | SE | | |
|--|---------------------|-----------------------|------------------------|--|--|
| A. Body Size | (µm) | | | | |
| Freshwater | 0 | 764.5 | 9.66 | | |
| Freshwater | 5 | 834.7 | 13.6 | | |
| Freshwater | 15 | 900.7 | 15.3 | | |
| Saline | 0 | 758.6 | 8.17 | | |
| Saline | 5 | 863.2 | 17.2 | | |
| Saline | 15 | 857.0 | 19.2 | | |
| B. Whole-Ani | mal NKA (band signa | l intensity) | | | |
| Freshwater | 0 | 8.16x10 ⁻⁵ | 1.97x10 ⁻⁵ | | |
| Freshwater | 5 | 2.79x10 ⁻⁴ | 1.11 x10 ⁻⁴ | | |
| Freshwater | 15 | 1.38x10 ⁻⁴ | 3.44 x10 ⁻⁵ | | |
| Saline | 0 | 1.44x10 ⁻⁴ | 4.90 x10 ⁻⁵ | | |
| Saline | 5 | 1.66x10 ⁻⁴ | 8.28 x10 ⁻⁵ | | |
| Saline | 15 | 4.10x10 ⁻⁴ | 1.05 x10 ⁻⁴ | | |
| C. NKA IMG A | Area (<i>µ</i> m²) | | | | |
| Freshwater | 0 | 6.46 | 1.88 | | |
| Freshwater | 5 | 6.34 | 1.618 | | |
| Freshwater | 15 | 8.49 | 1.258 | | |
| Saline | 0 | 15.8 | 2.74 | | |
| Saline | 5 | 8.11 | 2.95 | | |
| Saline | 15 | 7.60 | 2.38 | | |
| D. NKA IMG Intensity (pixel intensity) | | | | | |
| Freshwater | 0 | 2.03x10 ⁴ | 1.15 x10 ⁴ | | |
| Freshwater | 5 | 2.45x10 ⁴ | 2.40x10 ⁴ | | |
| Freshwater | 15 | 2.16x10 ⁴ | 1.08x10 ⁴ | | |
| Saline | 0 | 2.67x10 ⁴ | 2.17x10 ⁴ | | |
| Saline | 5 | 2.48x10 ⁴ | 2.09x10 ⁴ | | |
| Saline | 15 | 2.45x10 ⁴ | 1.76x10 ⁴ | | |

Comparison of Integument Permeability between Saline and Freshwater Populations, related to

Figures 3 and 7.

Methods

To compare the capacity of copepods from both the saline and freshwater populations to retain ions at low salinities, ion efflux was measured using radioactive sodium (²²Na). Individual *Eurytemora carolleeae* copepods were placed in native salinities with ²²Na for 12 hours (freshwater population n = 7 at 0 PSU, saline population n = 5 at 15 PSU). Individual copepods were then transferred to 0 PSU containing no ²²Na. Ion (²²Na) efflux was measured in the new water using a scintillation counter for 3 hours after transfer. The variable calculated was the proportion of radioactive sodium retained = [²²Na retained / (²²Na effluxed + ²²Na retained)]. This variable was tested for normality and comparisons between the two populations was performed using a Student's t-test.

Results

After transfer to 0 PSU for 3 hours, two saline copepods went comatose, and one died (Table S4). The ion efflux values for these individuals were kept for analysis because ion efflux is a passive process and does not require conscious or active excretion. Comparisons between the two populations revealed that there were significant differences in the proportion of radioactive sodium retained after three hours in 0 PSU (Student's t-test; t = 2.33, df = 9.42, P = 0.0219) with the freshwater population retaining more ions than the saline population (Figure S2). Sample sizes for this study are low, and while the findings are significant, further data collection is necessary to establish conclusive results.



Figure S2. Proportion of radioactive sodium (²²Na) retained after transfer to 0 PSU for 3 hours for both the freshwater (FW) and saline (SW) populations, related to Figures 3 and 7.

Proportion of ^{22}Na retained was calculated as [^{22}Na retained / (^{22}Na effluxed + ^{22}Na retained)]. Data are represented as Mean \pm SE.

Table S4: Ion efflux (²²Na) measurements obtained after 3 hours in 0 PSU for both saline (SW) and freshwater (FW) populations, related to Figures 3 and 7. Measures include ion efflux, ion retention, and the proportion of ions retained (ions retained / total). Copepod status reports whether a copepod went comatose or died after extended time in 0 PSU.

| Population | Efflux | Retained | Retained/Total | Copepod status |
|------------|---------|----------|-----------------------|----------------|
| FW | 7828.9 | 317.8 | 0.039 | |
| FW | 5761.9 | 354.8 | 0.058 | |
| FW | 14171.0 | 2194.9 | 0.134 | |
| FW | 23878.0 | 2523.9 | 0.096 | |
| FW | 9197.9 | 877.8 | 0.087 | |
| FW | 15282.0 | 1614.9 | 0.096 | |
| FW | 5496.9 | 803.8 | 0.128 | |
| SW | 21215.0 | 1470.9 | 0.065 | Comatose |
| SW | 5662.9 | 512.8 | 0.083 | |
| SW | 1188.9 | 55.0 | 0.044 | Died |
| SW | 2314.9 | 105.3 | 0.044 | |
| SW | 7031.9 | 308.8 | 0.042 | Comatose |