

Supplementary Information

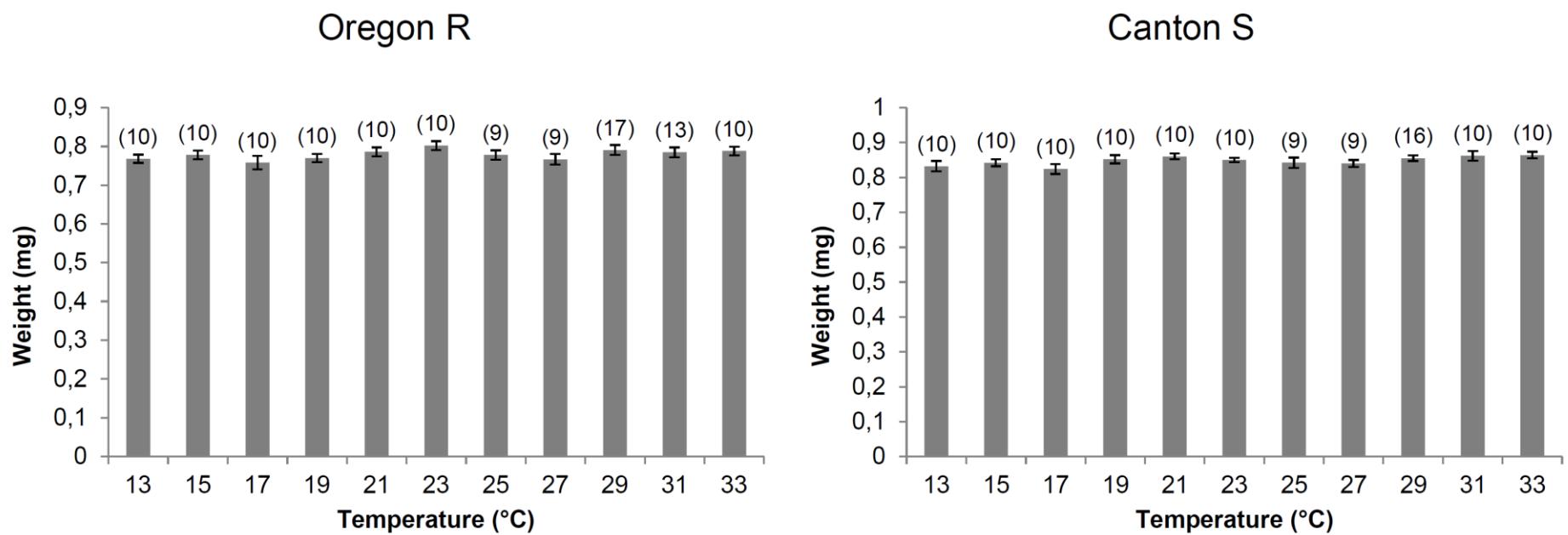
Temperature induces changes in *Drosophila* energy stores

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Supplementary Figures and Tables

Supplementary Figure S1. Body weight. Comparison of individual body weights of the flies used for the metabolic rate measurements. Data are presented as a mean body weight per individual \pm s.e.m. Sample size is reported in parenthesis. Data analyses by one-way ANOVA (with Tukey's HSD (honestly significant difference) post-hoc tests with $\alpha = 0.05$) did not reveal any significant differences among samples (Oregon: $F_{10,107} = 0.98$, P -value = 0.46; Canton: $F_{10,103} = 1.26$, P -value = 0.27).

Supplementary Figure S1.



Supplementary Table S1. The Akaike Information Criterion (AICc) and coefficient of determination (R^2) for different functions fitted to data on the temperature dependence of different traits. See Materials and methods for further details.

| Trait | Genotype | Linear | | Quadratic | | Cubic | | Exponential | |
|---|----------|---------------|-------------|---------------|-------------|--------|-------|---------------|-------------|
| | | AICc | R^2 | AICc | R^2 | AICc | R^2 | AICc | R^2 |
| Fat content | Oregon R | 92.64 | 0.21 | 70.37 | 0.94 | 77.08 | 0.94 | - | - |
| | Colony S | 84.90 | 0.02 | 64.74 | 0.93 | 72.19 | 0.94 | - | - |
| | combined | 172.12 | 0.14 | 127.15 | 0.93 | 136.85 | 0.94 | - | - |
| Relative changes in the fat content | Oregon R | 17.82 | 0.38 | 3.49 | 0.90 | 10.21 | 0.90 | - | - |
| | Colony S | 26.17 | 0.08 | 12.88 | 0.87 | 17.86 | 0.91 | - | - |
| | combined | 40.44 | 0.25 | 9.61 | 0.88 | 15.77 | 0.91 | - | - |
| Glycogen content | Oregon R | 78.52 | 0.40 | 76.38 | 0.69 | 81.84 | 0.74 | - | - |
| | Colony S | 81.95 | 0.04 | 78.00 | 0.65 | 86.89 | 0.65 | - | - |
| | combined | 157.18 | 0.20 | 147.23 | 0.67 | 157.55 | 0.69 | - | - |
| Metabolic rate | Oregon R | -5.90 | 0.99 | -1.77 | 0.99 | - | - | 11.77 | 0.94 |
| | Colony S | -8.86 | 0.99 | -11.51 | 0.99 | - | - | 10.19 | 0.94 |
| | combined | -19.67 | 0.99 | -18.53 | 0.99 | - | - | 16.90 | 0.94 |
| Food intake | Oregon R | -40.98 | 0.99 | -37.36 | 0.99 | - | - | -22.69 | 0.96 |
| | Colony S | -37.94 | 0.99 | -33.46 | 0.99 | - | - | -24.37 | 0.96 |
| | combined | -83.82 | 0.99 | -77.81 | 0.99 | - | - | -52.10 | 0.96 |
| Starvation survival time | Oregon R | 89.78 | 0.92 | - | - | - | - | 63.68 | 0.99 |
| | Colony S | 83.87 | 0.94 | - | - | - | - | 55.84 | 0.99 |
| | combined | 169.33 | 0.93 | - | - | - | - | 115.79 | 0.99 |
| Starvation survival time (vs inverse temperature) | Oregon R | 60.33 | 0.99 | 64.74 | 0.99 | - | - | - | - |
| | Colony S | 43.53 | 0.99 | 48.55 | 0.99 | - | - | - | - |
| | combined | 104.62 | 0.99 | 111.44 | 0.99 | - | - | - | - |

Supplementary Table S2. Multiple nonlinear (quadratic) regression analyses for the changes in the fat content (μg glycerides per fly) and the glycogen content (μg glycogen per fly). See Materials and methods for further details. df - degrees of freedom; SSQ - the sum of squares for each source of variation.

| Trait | Source of variation | df | SSQ | F-ratio | P-value |
|------------------|--|----|---------|---------|---------|
| Fat content | Genotype | 1 | 1.79 | 0.15 | 0.70 |
| | Initial fat content | 1 | 79.03 | 6.63 | 0.013 |
| | Temperature | 1 | 5503.73 | 461.84 | <0.0001 |
| | $(\text{Temperature})^2$ | 1 | 6158.22 | 516.76 | <0.0001 |
| | Genotype \times Temperature | 1 | 90.08 | 7.56 | 0.008 |
| | Genotype \times $(\text{Temperature})^2$ | 1 | 93.04 | 7.81 | 0.007 |
| | Error | 56 | 667.36 | - | - |
| Glycogen content | Genotype | 1 | 1.67 | 0.05 | 0.83 |
| | Temperature | 1 | 1810.38 | 50.30 | <0.0001 |
| | $(\text{Temperature})^2$ | 1 | 1942.77 | 53.98 | <0.0001 |
| | Genotype \times Temperature | 1 | 333.10 | 9.26 | 0.004 |
| | Genotype \times $(\text{Temperature})^2$ | 1 | 267.44 | 7.43 | 0.009 |
| | Error | 57 | 2051.44 | - | - |

Supplementary Table S3. Linear regression analyses for metabolic and consumption rate. *df*

- degrees of freedom; SSQ - the sum of squares for each source of variation.

| Trait | Source of variation | df | SSQ | F-ratio | P-value |
|------------------|------------------------|-----|--------|---------|----------|
| Metabolic rate | Genotype | 1 | 0.006 | 0.03 | 0.87 |
| | Temperature | 1 | 271.09 | 1152.19 | < 0.0001 |
| | Genotype × Temperature | 1 | 0.54 | 2.27 | 0.13 |
| | Error | 248 | 58.35 | - | - |
| Consumption rate | Genotype | 1 | 1.76 | 105.65 | < 0.0001 |
| | Temperature | 1 | 37.21 | 2238.56 | < 0.0001 |
| | Genotype × Temperature | 1 | 0.0004 | 0.02 | 0.88 |
| | Error | 503 | 8.36 | - | - |

Supplementary Table S4. Comparison of temperature dependence of metabolic and consumption rates (based on the mean values) by linear regression analyses for individual genotypes. *df* - degrees of freedom; SSQ - the sum of squares for each source of variation.

| Genotype | Source of variation | <i>df</i> | SSQ | <i>F</i> -ratio | <i>P</i> -value |
|----------|---------------------|-----------|-------|-----------------|-----------------|
| Oregon R | Trait | 1 | 0.007 | 4.28 | 0.053 |
| | Temperature | 1 | 2.98 | 1803.41 | < 0.0001 |
| | Trait × Temperature | 1 | 0.005 | 2.75 | 0.12 |
| | Error | 18 | 0.03 | - | - |
| Canton S | Trait | 1 | 0.02 | 14.57 | 0.0013 |
| | Temperature | 1 | 2.35 | 1685.11 | < 0.0001 |
| | Trait × Temperature | 1 | 0.005 | 3.71 | 0.07 |
| | Error | 18 | 0.03 | - | - |