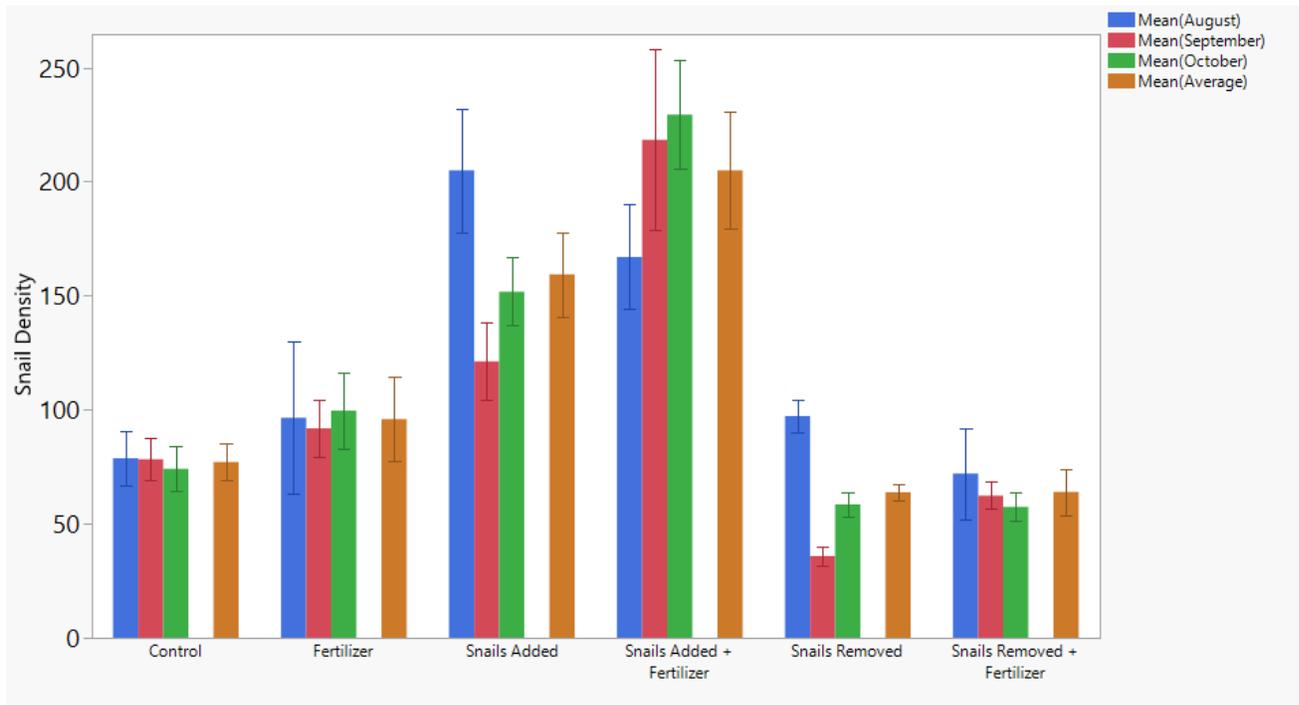


## Supplement



**Figure S1.** Densities of *Melampus bidentatus* throughout treatment plots in August, September, and October as well as the average.

**Table S1.** Results from linear mixed models examining the differences in fungal ASV richness, Pielou’s evenness, and Shannon’s diversity between timepoints, treatments, and fertilized/unfertilized plots, respectively.

<b>Response Variable</b>	<b>Predictor</b>	<b>DF Num</b>	<b>DF Den</b>	<b>F Ratio</b>	<b>P Value</b>	<b>Post hoc</b>
<b>Fungi</b> ASV Richness	<b>Time</b>	<b>1</b>	<b>48</b>	<b>7.54</b>	<b>0.0086</b>	<b>Sept &gt; Nov</b>
	Fertilizer	1	48	0.02	0.96	
	Snails	2	48	0.72	0.49	
	Fertilizer:Snails	2	48	1.44	0.25	
<b>Pielou’s Evenness</b>	Time	1	48	0.41	0.53	<b>Unfert.Sept &gt; Fert.Sept</b>
	<b>Fertilizer</b>	<b>1</b>	<b>48</b>	<b>9.84</b>	<b>0.003</b>	
	Snails	2	48	0.83	0.83	
	Fertilizer:Snails	2	48	1.93	1.93	
<b>Shannon’s Diversity</b>	Time	<b>1</b>	<b>48</b>	<b>12.83</b>	<b>&lt;0.001</b>	<b>Sept &gt; Nov Unfert&gt;Fert</b>
	Fertilizer	1	48	6.02	0.018	
	Snails	2	48	0.36	0.70	
	Fertilizer:Snails	2	48	2.34	0.11	

**Table S2.** Results from linear mixed models examining the differences in algal ASV richness, Pielou’s evenness, and Shannon’s diversity between timepoints, snail treatments, and fertilizer treatments, respectively.

<b>Response Variable</b>	<b>Predictor</b>	<b>DF Num</b>	<b>DF Den</b>	<b>F Ratio</b>	<b>P Value</b>	<b>Post hoc</b>	
<b>Algae</b>	ASV Richness	<b>Time</b>	<b>1</b>	<b>48</b>	<b>4.92</b>	<b>0.032</b>	<b>Sept &gt; Nov</b>
		Fertilizer	1	48	1.42	0.24	
		Snails	2	48	0.54	0.59	
		Fertilizer:Snails	2	48	1.64	0.21	
	Pielou’s Evenness	<b>Time</b>	<b>1</b>	<b>48</b>	<b>43.37</b>	<b>&lt;0.001</b>	<b>Sept &gt; Nov</b>
		Fertilizer	1	48	0.18	0.90	
		Snails	2	48	0.41	0.41	
		Fertilizer:Snails	2	48	0.72	0.5	
	Shannon’s Diversity	<b>Time</b>	<b>1</b>	<b>48</b>	<b>53.7</b>	<b>&lt;0.001</b>	<b>Sept &gt; Nov</b>
		Fertilizer	1	48	0.94	0.33	
		Snails	2	48	0.89	0.42	
		Fertilizer:Snails	2	48	0.93	0.38	

**Table S3.** Ranked models from linear mixed model selection on fungal alpha diversity metrics. Bold rows depict best-fit models outcompeted null models containing only random variables. Asterisks: \*\*\* =  $p < 0.001$ , \*\* =  $p < 0.01$ , \* =  $p < 0.05$ .

Response Variable	Model Rank	AIC	$\chi^2$	Predictor variables and standardized $\beta$ estimates ( $\pm$ se) of best-fit model
<b><i>Litter Bundle 1</i></b>				
<b>Fungal Richness</b>	<b>1</b>	<b>284.6</b>	<b>5.9*</b>	<b>Amphipods**</b> (-0.53 $\pm$ 0.15)
	Null	291.4		<i>Null Model</i>
<b>Fungal Evenness</b>	<b>1</b>	<b>-74.8</b>	<b>23.1***</b>	<b>Snails*</b> (0.46 $\pm$ 0.18) + <b>Isopods**</b> (0.51 $\pm$ 0.18) + <b>Live Biomass***</b> (-1.23 $\pm$ 0.23)
	2	-73.7	-	Live Biomass + Isopods
	3	-73.6	-	Live Biomass
	Null	-63.9	-	<i>Null Model</i>
<b>Fungal Diversity</b>	<b>1</b>	<b>29.4</b>	<b>18.5***</b>	<b>Live Biomass***</b> (-1.19 $\pm$ 0.24)
	2	29.8	-	Live Biomass + Amphipods
	3	30.7	-	Live Biomass + Snails
	Null	41.9	-	<i>Null Model</i>
<b><i>Litter Bundle 2</i></b>				
Fungal Richness	Null	113.7	-	<i>Null Model</i>
	1	116	-	Salinity
<b>Fungal Evenness</b>	<b>1</b>	<b>31.9</b>	<b>6.7*</b>	<b>Sediment NH<sub>4</sub><sup>+</sup>*</b> (0.51 $\pm$ 0.18) + <b>Litter %N*</b> (0.55 $\pm$ 0.25) + <b>Sediment pH</b> (0.39 $\pm$ 0.22)
	2	-32.5	-	Sediment NH <sub>4</sub> <sup>+</sup> + Litter %N
	Null	36.7	-	<i>Null Model</i>
<b>Fungal Diversity</b>	Null	17.6	-	<i>Null model</i>
	1	20.5	-	Humidity

**Table S4.** Ranked models from linear mixed model selection on algal alpha diversity metrics. Bold rows depict best-fit models outcompeted null models containing only random variables. Asterisks: \*\*\* =  $p < 0.001$ , \*\* =  $p < 0.01$ , \* =  $p < 0.05$ .

Response Variable	Model Rank	AIC	$\chi^2$	Predictor variables and standardized $\beta$ estimates ( $\pm$ se) of best-fit model
<b><i>Litter Bundle 1</i></b>				
<b>Algal Richness</b>	<b>1</b>	<b>93.7</b>	<b>6.1*</b>	<b>Amphipods*</b> (-0.44 $\pm$ 0.16)
	2	94.9	-	Amphipods – Litter %N
	Null	97.9	-	<i>Null Model</i>
<b>Algal Evenness</b>	<b>1</b>	<b>91.4</b>	<b>4.6*</b>	<b>Litter %N*</b> (0.33 $\pm$ 0.16)
	2	92.5	-	Litter %N + Live Biomass
	3	92.9	-	Live Biomass
	Null	95	-	<i>Null Model</i>
<b>Algal Diversity</b>	<b>1</b>	<b>89.3</b>	<b>6.9**</b>	<b>Amphipods*</b> (-0.44 $\pm$ 0.16)
	Null	94.2	-	<i>Null Model</i>
<b><i>Litter Bundle 2</i></b>				
<b>Algal Richness</b>	<b>1</b>	<b>61</b>	<b>4.3*</b>	<b>Salinity*</b> (0.36 $\pm$ 0.18)
	2	62.2	-	Salinity - Live Biomass
	Null	63.3	-	<i>Null Model</i>
Algal Evenness	Null	65.9	-	<i>Null Model</i>
Algal Diversity	1	64.22	-	Sediment pH
	Null	67	-	<i>Null Model</i>
	1	66.4	-	Salinity

**Table S5.** Results from PERMANOVA analysis contrasting unweighted UniFrac distances of fungal and algal communities between timepoints, treatments, and fertilized/unfertilized plots, respectively.

<b>Response Variable</b>	<b>Predictor</b>	<b>Df</b>	<b>Total Df</b>	<b>F Value</b>	<b>R<sup>2</sup></b>	<b>P Value</b>
<b><i>Collection Period</i></b>						
Unweighted Unifrac (Fungi)	<b>Time</b>	<b>1</b>	<b>53</b>	<b>2.34</b>	<b>0.043</b>	<b>0.02</b>
Unweighted Unifrac (Algae)	<b>Time</b>	<b>1</b>	<b>53</b>	<b>6.43</b>	<b>0.1</b>	<b>&lt;0.001</b>
<b><i>Litter Bundle 1</i></b>						
Unweighted Unifrac (Fungi)	<b>Fertilizer</b>	1	32	<b>2.35</b>	<b>0.061</b>	<b>0.025</b>
	Snails	2	32	0.59	0.034	0.92
	Fertilizer:Snails	2	32	1.36	0.013	0.44
Unweighted Unifrac (Algae)	Fertilizer	1	32	0.8	0.27	0.53
	Snails	2	32	0.24	0.016	0.99
	Fertilizer:Snails	2	32	0.44	0.029	0.90
<b><i>Litter Bundle 2</i></b>						
Unweighted Unifrac (Fungi)	Fertilizer	1	20	1.31	0.07	0.25
	Snails	2	20	0.90	0.084	0.58
	Fertilizer:Snails	2	20	0.66	0.069	0.89
Unweighted Unifrac (Algae)	Fertilizer	1	20	2.08	0.08	0.058
	Snails	2	20	0.88	0.068	0.55
	Fertilizer:Snails	2	20	1.27	0.097	0.28

**Table S6.** Results from PERMANOVA analysis contrasting Bray-Curtis distances of fungal and algal communities between timepoints, treatments, and fertilized/unfertilized plots, respectively.

<b>Response Variable</b>	<b>Predictor</b>	<b>Df</b>	<b>Total Df</b>	<b>F Value</b>	<b>R<sup>2</sup></b>	<b>P Value</b>
<i>Collection Period</i>						
<b>Bray-Curtis (Fungi)</b>	<b>Time</b>	<b>1</b>	<b>54</b>	<b>2.9</b>	<b>0.051</b>	<b>0.011</b>
<b>Bray-Curtis (Algae)</b>	<b>Time</b>	<b>1</b>	<b>54</b>	<b>16.4</b>	<b>0.24</b>	<b>&lt;0.001</b>
<i>Litter Bundle 1</i>						
<b>Bray-Curtis (Fungi)</b>	<b>Fertilizer</b>	<b>1</b>	<b>32</b>	<b>4.10</b>	<b>0.12</b>	<b>0.003</b>
	Snails	2	32	0.65	0.039	0.81
	Fertilizer:Snails	2	32	0.71	0.042	0.76
Bray-Curtis (Algae)	Fertilizer	1	32	0.74	0.024	0.54
	Snails	2	32	0.50	0.033	0.86
	Fertilizer:Snails	2	32	0.62	0.04	0.76
<i>Litter Bundle 2</i>						
<b>Bray-Curtis (Fungi)</b>	<b>Fertilizer</b>	<b>1</b>	<b>32</b>	<b>2.90</b>	<b>0.13</b>	<b>0.004</b>
	Snails	2	32	0.89	0.079	0.58
	Fertilizer:Snails	2	32	1.03	0.090	0.41
Bray-Curtis (Algae)	Fertilizer	1	32	1.04	0.043	0.34
	Snails	2	32	0.89	0.069	0.44
	Fertilizer:Snails	2	32	1.34	0.08	0.24

**Table S7.** Results from SIMPER analysis contrasting Bray-Curtis distances of fungal communities between timepoints, treatments, and fertilized/unfertilized plots, respectively.

Contrast	Order and/or Finest Taxonomy	Cont. %	Cum. %	Avg. % Abundances	
				1	2
<b>Timepoints 1 &amp; 2</b>	Microascales: <i>Natantispora retorquens</i>	30.0	40.1	44.8	34.7
	Unassigned Ascomycota	12.9	58.7	12.7	22.1
	Pleosporales: <i>Phaesphaeria halima</i>	7.8	69.7	13.2	8.0
	Unassigned Fungi	6.0	78.1	7.0	10.8
	Capnodiales: Neodevriesiaceae	3.6	83.1	6.7	1.9
	Lulworthiales: <i>Lulworthia spp</i>	3.1	87.5	3.5	6.0
	Unassigned Pleosporales	2.0	92.4	3.9	5.7
	Eurotiales: <i>Talaromyces helicus</i>	1.6	96.2	2.7	2.9
	Pleosporales: Phaeosphaeriaceae	1.6	97.0	2.7	3.2
	Microscales: <i>Lignincola laevis</i>	1.1	97.4	1.5	1.9
	<b>Unfert/Fert (Bundle 1)</b>				<b>Unfert</b>
Microascales: <i>Natantispora retorquens</i>		34.2	50.3	32.5	56.3
Unassigned Ascomycota		8.3	62.4	23.0	3.0
Pleosporales: <i>Phaesphaeria halima</i>		8.1	74.4	14.7	11.6
Capnodiales: Neodevriesiaceae		4.1	80.4	4.6	8.6
Unassigned Fungi		3.9	86.2	9.2	4.9
Lulworthiales: <i>Lulworthia spp</i>		1.9	89.0	5.5	1.7
Unassigned Pleosporales		1.9	91.8	4.0	3.8
Pleosporales: Phaeosphaeriaceae		1.7	94.3	1.9	3.4
Microscales: <i>Lignincola laevis</i>		1.0	95.8	1.1	1.9
<b>Unfert/Fert (Bundle 2)</b>					<b>Unfert</b>
	Unassigned Ascomycota	23.3	30.3	38.8	8.1
	Microascales: <i>Natantispora retorquens</i>	22.5	60	13.0	52.8
	Unassigned Fungi	8.7	70.8	13.6	8.4
	Pleosporales: <i>Phaesphaeria halima</i>	5.6	78.1	11.2	5.3
	Eurotiales: <i>Talaromyces helicus</i>	4.4	83.4	6.5	~0
	Lulworthiales: <i>Lulworthia spp</i>	4.4	89.5	4.8	7.0
	Unassigned Pleosporales	2.0	92.0	2.9	7.9
	Pleosporales: Phaeosphaeriaceae	1.3	93.7	3.0	3.4
	Microscales: <i>Lignincola laevis</i>	1.2	95.3	0.9	2.8
	Capnodiales: Neodevriesiaceae	1.2	96.9	2.0	1.8

**Table S8.** Results from SIMPER analysis contrasting Bray-Curtis distances of algal communities between timepoints, treatments, and fertilized/unfertilized plots, respectively.

Contrast	Order and/or Finest Taxonomy	Cont. %	Cum. %	Avg. % Abundances	
				1	2
<b>Timepoints 1 &amp; 2</b>	Ulotrichales: <i>Chlorothrix</i> spp.	29.8	40.3	0.7	50.5
	Ulvales: <i>Pseudendoclonium submarinum</i>	26.9	76.7	75.9	27.4
	Unassigned Ulvales	6.2	86.0	12.9	9.1
	Ulotrichales: <i>Acrosiphonia arcta</i>	4.2	90.7	~0.0	7.0
	Unassigned Chlorophyta	2.8	92.9	5.8	2.3
	Unassigned Ulotrichales	1.5	95.0	0.2	2.6
	Unassigned Chlorophyceae	1.0	96.8	1.6	0.5
	Ulvales: <i>Ulva compressa</i>	1.0	98.1	2.6	0.4
<b>Unfert/Fert (Bundle 1)</b>				<b>Unfert</b>	<b>Fert</b>
	Ulvales: <i>Pseudendoclonium submarinum</i>	38.7	72.0	73.7	77.7
	Unassigned Ulvales	6.8	84.7	15.1	10.8
	Unassigned Chlorophyta	3.8	92.1	6.8	4.9
	Ulvales: <i>Ulva compressa</i>	1.7	95.1	2.4	2.3
	Unassigned Chlorophyceae	1.6	98.0	0.6	2.6
<b>Unfert/Fert (Bundle 2)</b>				<b>Unfert</b>	<b>Fert</b>
	Ulotrichales: <i>Chlorothrix</i> spp.	27.7	49.3	53.7	47.7
	Ulvales: <i>Pseudendoclonium submarinum</i>	11.5	69.9	21.4	32.4
	Unassigned Ulvales	6.8	82.0	12.7	6.0
	Ulotrichales: <i>Acrosiphonia arcta</i>	5.0	90.9	6.3	7.5
	Unassigned Ulotrichales	1.7	94.0	1.8	3.2
	Unassigned Chlorophyta	1.6	95.0	2.6	4.1