**Supplementary Table S2**

|  |
| --- |
| Legend |
| \* | Manipulative study |
| ¤ | Non-manipulative study |
| ~ | Value not specified in paper, circumstantial information provided in paper informs an estimated value. Accuracy of estimations vary.  |
| gl/f | glochidia per fish |
| gl/fg | glochidia per gram of fish  |
| dpi | days post infestation |
| (W) | Wild caught fish |
| (H) | Hatchery reared fish |

|  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Mussel species | Fish species | Specific effect | Glochidia load | Against | DPI |  + -0 | p | Correlation Coefficient | n= | Paper |
| Whole body effects |
| *Margaritifera margaritifera* | *Salmo trutta* 0+ (W) | Growth during infestation period | / | Control \* | 30 - 33 dpi |  -  | 0,046 | / | 159 | Andersson, 2018 |
| Condition Factor (k) during infestation period | 0 | 0,727 |
| Growth in the field | 65 - 75 dpi | 0 | 0,17 | 62 |
| Condition Factor (k) in the field | 0 | 0,207 |
| *Salmo trutta* 0+ (H) | Condition factor (k) | 4.5 gl/f | Glochidia Load ¤ | 60 dpi | 0 | 0,887 | 0,0186 | 30 | Marwaha et la, 2019 |
| Weight | 0 | 0,857 | -0,0238 |
| Length | 0 | 0,334 | -0,127 |
| Condition factor (k) | Glochidia Load ¤ | 200 dpi | 0 | 0,481 | -0,1337 |
| Weight | 0 | 0,956 | -0,01 |
| Length | 0 | 0,7531 | 0,0599 |
| Condition factor (k) | Glochidia Load ¤ | 300 dpi |  -  | 0,0817 | -0,2051 |
| Weight |  -  | 0,0006 | -0,288 |
| Length |  -  | 0,003 | -0,256 |
| Condition factor (k) | Control \* | 300 dpi | 0 | 0,865 | / | 70 |
| Weight | 0 | 0,514 |
| Length | 0 | 0,232 |
| *Salmo trutta* (W) | Fork Length | 1 to 309 gl/f | Control ¤ | ~50 dpi | 0 | >0.1 | / | 28 | Filipsson et al, 2016 |
| Weight | 0 | >0.1 |
| *Salmo trutta* 1+ (H) | Growth | 7889 ± 390 gl/f or 138 ± 3 gl/fg | Control \* | 84 dpi |  -  | <0.094 | / | ~319 | Chowdhury et al, 2019 |
| Survival | 0 | / |
| Growth | 5473 ± 445 gl/f | Control \* | 203 dpi |  -  | <0.003 | / |
| Survival | 0 | / |
| Growth | 5202 ± 804 gl/f | Control \* | 266 dpi |  -  | <0.002 | / |
| Survival | 0 | / |
| Growth | 3125 ± 566 gl/f | Control \* | 315 dpi |  -  | <0.006 | / |
| Survival | 0 | / |
| *Salmo trutta* 1+ (W) | Fork Length | Between 0 and 11 gl/fg or ~66 gl/f | Control ¤ | ~250 dpi | 0 | >0.45 | / | 62 | Filipsson et al, 2017 |
| Weight | 0 | >0.45 |
| Condition factor (k) | 0 | >0.45 |
| *Salmo trutta* 1+ (H) | Condition factor (k) | 212.8 gl/f Infestation intensity category cutoff at 200 gl/f | Glochidia Load ¤ | 300 dpi |  +  | 0,0368 | 0,3054 | 50 | Marwaha et la, 2019 |
| High vs low infestation ¤ |  +  | 2.73e−07 | / |
| Control \* |  +  | 0,0001 |
| Weight | High glochidia load (>200) ¤ | 0 | 0,06 | 0,3968 |
| Length | 0 | 0,055 | 0,4052 |
| *Salmo salar* 0+ (H) | Growth in freshwater | ~282 gl/f | Control ¤ | ~150 dpi | 0 | / | / | ~24 | Bruno et al, 1988 |
| Survival in freshwater | 0 |
| Growth in saltwater | ~116 gl/f | Control ¤ | ~250 dpi | 0 | / | / | ~29 |
| Survival in saltwater | 0 |
| *Salmo salar* 0+ (W) | Fork Length | pooled avg of 191 gl/f at one site and 31 gl/f | Glochidia Load ¤ | ~250 dpi | 0 | >0.6 & >0.3 | 0.065 & -0.117 | ~74 | Cunjak & McGladdery, 1991 |
| condition factor (k) of all glochidia loads | 0 | <0.005 | -0,24 |
| condition factor (k) for glochidia load >200 | >200 gl/f |  -  | <0.005 | -0,56 | 24 |
| condition factor (k) | pooled avg of 191 gl/f at one site and 31 gl/f | Between field sites ¤ | ~250 dpi | 0 | >0.2 | / | ~74 |
| Survival |  -  |  / |
| *Salmo salar* (H) | Length | 818 ± 377 gl/f at 14dpi 1392 ± 641 gl/f at 42dpi 1385 ± 553 gl/f @70dpi ~81± 121 gl/f at ~105dpi <1 ± 2.3 gl/f at ~182 dpi | Control \* | ~30 dpi | 0 | 0,51 | / | 600 |  Treasurer et al, 2006 |
| Weight | 0 | 0,64 |
| Length | ~105 dpi |  -  | 0,0006 |
| Weight |  -  | 3E-05 |
| Length | ~182 dpi |  -  | 0,0004 |
| Weight | 0 | / |
| Length | ~273 dpi | 0 | 0,1172 |
| Weight | 0 | 0,13 |
| Condition factor (k) | 0 | >0.05 |
| Survival | 0 | / |
| *Salmo trutta* (W) | Specific Growth rate | High infestation > 226 ± 24 Low infestation <18±15 | Low infestation ¤ | ~100 dpi |  -  | 0,003 | / | 21 | Freitt 2016 |
| Specific growth rate (only 1+) |  -  | 0,004 | / |
| Initial condition factor | ~15 dpi | 0 | >0.05 | 18 |
| secondary condition factor | ~100 dpi | 0 | 21 |
| Change in condition factor | 0 | / |
| *Salmo salar* >1+ (W) | Post-spawning survival | 2000 to 7000 gl/f | Observation ¤ | / |  +  | / | / | / | Zuiganov et al, 2005 |
| Survival after temporary asphyxia | 65 to 400+ gl/f | Control \* | / |  +  | < 0.05 | / | / |
| Survival after gill burn |  +  |
| Survival after hook wounds |  +  |
| *Salmo salar* 1+ (H) | Survival | 66 to 219 gl/f (up to 1000 in some cases) | Control \* | ~438 dpi | 0 | >0.05 | / | 100 | Treasurer & Turnbull 2000 |
| Weight | 0 | >0.05 |
| Specific growth rate | 0 | >0.05 |
| *Margaritifera margaritifera falcata* | *Salmo trutta* 0+ (H) | Survival | 100-295 gl/f | Observation ¤ | ~59 dpi |  -  | / | / | 200 | Murphy, 1942 |
| *Oncorhynchus mykiss* 0+ (H) | Survival | 600-1200 gl/f | 1 dpi |  -  | / | / | 400 |
| Survival | 250-750 gl/f | ~59 dpi |  -  | / | / | 400 |
| Survival | 30 - 50 gl/f | ~23 dpi | 0 | / | / | 100 |
| *Margaritifera laevis* | *Oncorhynchus masou masou* 0+ (H) | Growth rate | 48 ± 29 gl/f | Control \* | 50 dpi | 0 | / | / | 75 | Ooue, et al, 2017 |
| 70 dpi |  -  |
| *Strophitus undulatus* | *Oncorhynchus mykiss* (H) | Condition factor (K) | 25.46 ± 2.64 gl/f on whole body 7.69 ± 2 on gills  | Control \* | 28 dpi | 0 | >0.05 | / | 9 | Defo et al, 2019 |
| Anodonta oregonen | *Oncorhynchus kisutch* 0+ (H) | Weight | 0 to 50 gl/f. 50+ gl/f dies within 30 days | Glochidia load ¤ | 90 dpi |  -  | < 0.05 | / | 2200 | Moles 1983 |
| Survival |  -  | / | / |
| *Anodonta anatina* | *Rhodeus amarus* (W) | Weight | 19 gl/f on whole body & 2 gl/f on gills | Control \* | 4 dpi | 0 | 0,241 | / | 48 | Methling et al, 2019 |
| *Sinanodonta woodiana* | *Squalius cephalus* 1+ (H) | Weight | 120 ± 39 gl/f | Control \* | 43 dp first infestation, 3 dp last infestation |  -  | >0.001 | / | 16 | Douda et al, 2017 |
| Length  | 0 | >0.05 |
| Condition factor (k) |  -  | >0.001 |
| Weight | 25 ± 4 gl/f | Control \* | 0 | <0.001 | / | 16 |
| Length  | 0 | >0.05 |
| Condition factor (k) |  -  | 0,08 |
| *Hyriopsis cumingii* | *Pelteobagrus fulvidraco* | Survival | 2480 ± 114 gl/f in low group and 3244 ± 248 gl/f in high group | Glochidia Load \* | 12 dpi |  -  | / |  +  | 270 | Du et al, 2015 |
| Metabolic rate |
| *Margaritifera margaritifera* | *Salmo trutta* 0+ (H) | Ventilation control | 37 ± 8 gl/f  | Glochidia load ¤ | 160 dpi |  -  | 0,012 | / | 50 | Thomas et al 2014 |
| Control \* |  -  | <0.001 |
| *Salmo trutta* 1+ (W) | Standard Metabolic rate (SMR) | Between 0 and 11 gl/fg & ~66 gl/f | Glochidia load ¤ | ~250 dpi |  -  | 0,017 |  −0.40 | 62 | Filipsson et al, 2017 |
| Max metabolic rate (MMR) | 0 | 0,81 | −0.08 |
| Standard Metabolic rate (SMR) | Control ¤ |  +  | 0,009 | / |
| Max metabolic rate (MMR) |  +  | 0,043 |
| *Sinanodonta woodiana* | *Rhodeus ocellatus* (W) | Initial MO2  | <120 gl/f on while body & ≤ 20 gl/f on gills | Control \* | <1 dpi | 0 | 0,239 | / | 24 | Methling et al 2018 |
| Change in MO2 post stressor |  -  | 0,017 |
| *Cyprinus carpio* 4+ (H) | Electrocardiogram (EMG) |  Infested at 4570 ± 1889 gl/l for 15 minutes in 15 liters | Control \* | Average of 4 and 8 dpi |  +  | <0.001 | / | 22 | Slavik et al 2017 |
|  +  | <0.001 |
| 4 dpi | ~ +  | / |
| 8 dpi | ~ 0 | / |
| Night EMG |  -  | 0,0001 |
| Day EMG | 0 | 0,0312 |
| *Hyriopsis cumingii* | *Pelteobagrus fulvidraco* | Rate of O consumption (MO2) | 2480 to 3244 gl/f | Control \* | ~12 dpi | 0 | >0.05 | / | 270 | Du et al, 2015 |
| Ammonia excretion |  +  | <0.05 |
| Ammonia excretion | Glochidia load \* | ~12 dpi |  +  | <0.05 |  +  |
| *Lampisilis reeveiana* | *Micropterus salmoides* 0+ | Ventilation quality |  632 gl/f | Glochidia Load ¤ |  ~10 dpi |  -  | <0.001 |  +  | Unclear58, 50, 35 | Kaiser, 2005 |
| ~90 dpi |  -  | <0.001 |  +  |
| Control \* |  ~10 dpi |  -  | <0.001 | / |
| ~90 dpi |  -  | <0.001 |
| Rate of O consumption (MO2) | Control \* | avg of ~90 days |  -  | <0.05 |  -  |
| Tolerance to hypoxia (critical DO level) | Glochidia Load ¤ | 2 dpi |  -  |  +  |
| 9dpi |  -  | <0.001 |
| 14 dpi |  -  |
| 18dpi |  -  |
| 49 dpi |  -  |
| 63dpi |  -  | <0.05 |
| 89 dpi |  -  |
| Aquatic surface respiration | Control \* | avg of ~90 days |  -  |   |
| *Lampsilis straminea* | *Lepomis macrochirus* | Tolerance to hypoxia (critical DO level) | 157.2 ± 35.8 gl/f | Control \* | 1dpi | 0 | >0.05 | / | 88 | Horne, 2021 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 65.7 ± 36.5 | 2dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 40.8 ± 7.3 | 3dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 40.8 ± 7.3 | 7dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 41.5 ± 17.7 | 14dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 30.7 ± 16.9 | 21dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 11.5 ± 5.5 | 35dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 37.0 ± 16.9 | 56dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| *Micropterus salmoides* | Tolerance to hypoxia (critical DO level) | 151.8 ± 20.8 | Control\* | 1dpi | 0 | 88 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 146.7 ± 11.1 | 2dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 138.3 ± 14.1 | 3dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 178.8 ± 38.9 | 7dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) |  133.8 ± 17.2 | 14dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 166.2 ± 11.0 | 21dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 138.0 ± 16.1 | 24dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 84.3 ± 18.4 | 35dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 95.5 ± 14.7 | 56dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| Tolerance to hypoxia (critical DO level) | 83.8 ± 8.8 | 77dpi | 0 |
| Metabolic rate | 0 |
| Regulation Index | 0 |
| *Venustaconcha pleasii* | *Etheostomacaeruleum* (W) | Ventilation quality | 73.5 ± 35.4 initially attached. 13.2 ± 11.8 juveniles recovered by 20 DPI | Control \* | 2 dpi |  -  | 0,005 | / | 20 | Crane et al, 2011 |
| 4 dpi | 0 | >0.05 |
| 6 dpi | 0 |
| 8 dpi | 0 |
| 10 dpi | 0 |
| 12 dpi | 0 |
| 14 dpi | 0 |
| 16 dpi | 0 |
| 18 dpi | 0 |
| 20 dpi | 0 |
| 22 dpi | 0 |
| *Ptychobranchus occidentalis* | Ventilation quality | 31.5 ± 23.7 initially attached. 13.2 ± 16.7 juveniles recovered by 22 DPI | Control \* | 2 dpi | 0 | > 0.05 | / | 36 |
| 8 dpi | 0 |
| 14 dpi |  -  | 0,009 |
| 20 dpi | 0 | > 0.05 |
| 28 dpi | 0 |
| *Anodonta anatina* | *Rhodeus amarus* (W) | Change in SMR | 19 gl/f on whole body & 2 gl/f on gills | Control \* | 1 dpi |  -  | 0,009 | / | 48 | Methling et al 2019 |
| 2 dpi | 0 |
| 3 dpi | 0 |
| 4 dpi |  +  |
| SMR | Glochidia load ¤ | 4 dpi |  -  | 0,011 |  +  |
| Change in MO2 | Control \* | 0 | 0,205 | / |
| MO2 Max | Glochidia Load ¤ |  +  | 0,048 |  +  |
| Organ Histology |
| *Margaritifera margaritifera* | *Salmo trutta* (H) | Spleen size | 100.7 (±18.62) gl/f (SD) | Control \*  | 15 dpi | 0 | > 0.05 | / | 27 | Thomas et al 2014 |
| 150.9 (±2.94) gl/f (SD) | 30 dpi |  +  | < 0.001 | 27 |
| 36.7 (±7.94) gl/f (SD) | 160 dpi | 0 | > 0.05 | 30 |
| *Sinanodonta woodiana* | *Squalius cephalus* (H) | Spleen size |  120.5 ± 39.3 gl/f (SD) | Control \*  | 43 dp first infestation, 3 dp last infestation | 0 | > 0.05 | / | 16 | Douda et al, 2017 |
|  24.6 ± 4.1 gl/f (SD) | 0 |
| *Strophitus undulatus* | *Oncorhynchus mykiss* (H) | Liver size | 26 ± 3 gl/f (SD) on whole body 8 ± 2 (SD) on gills  | Control \*  | 28 dpi | 0 | > 0.05 | / | 9 | Defo et al, 2019 |
| Toxicology |
| *Anodonta oregone* | *Oncorhynchus kiutch* <1 month old (H) | 96h LC50 of naphthalene | 130 to 10 gl/f avg of 69 gl/f | Glochidia Load \* | <1 dpi |  -  | / | / | 540 | Moles 1980 |
| Control \* |  -  |
| 96-hour LC50 toluene | 110 to 5 gl/f avg of 69 gl/f | Glochidia Load \* |  -  | 700 |
| Control \* |  -  |
|  96-hour LC50 crude oil | 100 to 10 gl/f avg of 81 gl/f | Glochidia Load \* |  -  | 420 |
| Control \* |  -  |
| *Strophitus undulatus* | *Oncorhynchus mykiss* (H) | Liver cadmium concentration at baseline cadmium levels | 26 ± 3 gl/f (SD) on whole body 8 ± 2 (SD) on gills  | Control \* | 28 dpi | 0 | >0.05 | / | 77 | Defo et al, 2019 |
| Liver cadmium concentration with excess cadmium | 0 | >0.05 | / | 78 |
| Gene Transcription |
| *Strophitus undulatus* | *Oncorhynchus mykiss* (H) | Transcriptomic changes under glochidial stress | 26 ± 3 gl/f (SD) on whole body 8 ± 2 (SD) on gills  | Control \* | 28 dpi | 0 | / | / | 77 | Defo et al, 2019 |
| Transcriptomic changes to loc110530601 under combined glochidia and high cadmium stress in liver (immune system) | Not shared with high cadmium stress \* |  +  | 78 |
| Transcriptomic changes to loc110514382 under combined glochidia and high cadmium stress in liver (immune system) |  +  |
| Transcriptomic changes to loc110504376 under combined glochidia and high cadmium stress in liver (malignant tumor) |  - |
| *Elliptio complanata* | *Perca flavescens* 0+ (W) | Transcription of HSP70 in liver | 59 to 93 gl/f | Control \* | 1 dpi |  +  | < 0.01 | / | 20 | Grendon, et al., 2019 |
| Transcription of g6pd in liver | 0 | >0.05 |
| Transcription of g6p in liver | 0 |
| Transcription of glyc in liver |  -  | < 0.0001 |
| Transcription of grlf1 in liver |  -  | < 0.001 |
| Transcription of HSP70 in gills |  +  | < 0.0001 |
| Transcription of g6pd in gills | 0 | >0.05 |
| Transcription of grlf1 in gills | 0 |
| *Lampsilis radiata* | Transcription of HSP70 in liver | 51 to 78 gl/f |  +  | < 0.01 |
| Transcription of g6pd in liver | 0 | >0.05 |
| Transcription of g6p in liver | 0 |
| Transcription of glyc in liver | 0 |
| Transcription of grlf1 in liver | 0 |
| Transcription of HSP70 in gills | 0 |
| Transcription of g6pd in gills | 0 |
| Transcription of grlf1 in gills | 0 |
| Coloration |
| *Margaritifera margaritifera* | *Salmo trutta* (W) | Dark coloration | 1 to 309 gl/f | Control ¤ | ~50 dpi |  +  | <0.001 | / | 14 | Filipsson et al, 2016 |
| *Anodonta anatina* | *Phoxinus phoxinus* (W) | Mating coloration | 5 ± 1 gl/f in high & 1 ± 0 gl/f in low | Glochidia Load \* | 14 dpi | 0 | 0,48 | / | ~90 | Kekalainen et al, 2014 |
| Mating |
| *Anodonta anatina* | *Phoxinus phoxinus* (W) | Sperm motility including all individuals | 5 ± 1 gl/f in high & 1 ± 0 gl/f in low | Glochidia Load \* | 14 dpi | 0 | 0,129 | / | ~90 | Kekalainen et al, 2014 |
| Sperm motility including only parasitized individuals |  -  | 0,011 |
| Curvature of sperm swimming trajectory |  -  | 0,039 |
| Breeding tubercle number |  -  | 0,047 |
| Molecular Changes (present in plasma unless otherwise specified) |
| *Sinanodonta woodiana* | *Cyprinus carpio* 4+ (H) | Hematocrit |  Infested at 4570 ± 1889 gl/l for 15 minutes in 15 liters | Control \* | 8 dpi | 0 | >0.01 | / | 22 | Slavik et al 2017 |
| Hemoglobin concentration | 0 | >0.01 |
| Cortisol | 0 | >0.01 |
| Aspartate aminotransferase  |  +  | < 0.01 |
| Alanine aminotransferase |  +  | < 0.01 |
| Lactate dehydrogenase | 0 | >0.01 |
| Alaline phosphatase | 0 | >0.01 |
| Calcium (Ca) | 0 | >0.01 |
| Potassium (K) |  + | < 0.01 |
| Sodium (Na) | 0 | >0.01 |
| Chloride (Cl) |  + | < 0.01 |
| *Squalius cephalus* | Potassium (K) |  Low load 24.6 ± 4.1 gl/f (SD) | Control \* | 43 dp first infestation, 3 dp last infestation |  +  | <0.001 |   |   | Douda et al, 2017 |
| Sodium (Na) | 0 | >0.05 |
| Calcium (Ca) | 0 | >0.05 |
| Chloride (Cl) |  +  | <0.001 |
| Aspartate aminotransferase  | 0 | >0.05 |
| Alanine aminotransferase | 0 | >0.05 |
| Lactate dehydrogenase | 0 | >0.05 |
| Alaline phosphatase |  +  | <0.05 |
| Cortisol | 0 | >0.05 |
| Potassium (K) | High load 120.5 ± 39.3 gl/f (SD) |  +  | <0.001 |
| Sodium (Na) | 0 | >0.05 |
| Calcium (Ca) | 0 | >0.05 |
| Chloride (Cl) |  +  | <0.001 |
| Aspartate aminotransferase  |  +  | <0.01 |
| Alanine aminotransferase |  +  | <0.01 |
| Lactate dehydrogenase |  +  | <0.01 |
| Alaline phosphatase |  +  | <0.01 |
| Cortisol | 0 | >0.05 |
| *Margaritifera margaritifera* | *Salno trutta* (H) | Hematocrit | 150.9 (±2.94) gl/f (SD) | Control \* | 30 dpi | 0 | 0,627 | / | 50 | Thomas et al 2014 |
| Glochidia Load ¤  | 0 | 0,17 | 21 |
| *Salmo trutta* 1+ (W) | Hematocrit | Between 0 and 11 gl/fg & ~66 gl/f | Control ¤ | ~250 dpi |  + | 0,038 | / | 62 | Filipsson et al, 2017 |
| *Salmo trutta* 1+ (H) | Hematocrit | 212.8 gl/f | Control \* | 300 dpi | 0 | 0,233 | / | 500 | Marwaha, et al 2019 |
| Glochidia Load ¤  |  +  | 0,029 | 0,33 |
| *Salmo salar* 1+ (H) | Chloride (Cl) | 66 to 219 gl/f (up to 1000 in some cases) | Control \* | ~288 dpi |  +  | 0,0002 | / | ~40 | Treasurer & Turnbull 2000 |
| *Hyriopsis cumingii* | *Tilapia nilotica* | Total Protein | ~ > 24141 gl/f | Control \* | ~ 7 dpi | 0 | >0.05 | / | ~ 8+ | Wen et al 2009 |
| Albumin | 0 |
| Glucose | 0 |
| Cholesterol | 0 |
| Triglyceride | 0 |
| Low Density Lipoprotein | 0 |
| High Density Lipoprotein | 0 |
| Total Protein | 0 gl/f | Control \* | ~ 13 dpi | 0 | >0.05 | / | ~ 8+ |
| Albumin | 0 |
| Glucose | 0 |
| Cholesterol | 0 |
| Triglyceride |  -  | 0,001 |
| Low Density Lipoprotein |  -  | 0,02 |
| High Density Lipoprotein | 0 | >0.05 |
| *Pelteobagrus fulvidraco* (H) | Liver Glycogen | /  | Control \*  | 2 dpi | 0 | > 0.05 | / | 120 | Ma et al 2018 |
| Liver Free Fatty Acid | 0 |
| Liver Total Protein | 0 |
| Liver Total Amino Acid | 0 |
| Muscle Glycogen | 0 |
| Muscle Free fatty Acid | 0 |
| Muscle Total Protein | 0 |
| Muscle Total Amino Acid | 0 |
| Plasma Free Fatty Acid | 0 |
| Plasma Total Protein | 0 |
| Plasma Total Amino Acid | 0 |
| Liver Glycogen | 5 dpi | 0 | > 0.05 |
| Liver Free Fatty Acid | 0 |
| Liver Total Protein | 0 |
| Liver Total Amino Acid | 0 |
| Muscle Glycogen | 0 |
| Muscle Free fatty Acid | 0 |
| Muscle Total Protein | 0 |
| Muscle Total Amino Acid | 0 |
| Plasma Free Fatty Acid | 0 |
| Plasma Total Protein | 0 |
| Plasma Total Amino Acid |  -  |  <0.05 |
| Liver Glycogen | 12 dpi | 0 | > 0.05 |
| Liver Free Fatty Acid |  -  | <0.05 |
| Liver Total Protein | 0 | > 0.05 |
| Liver Total Amino Acid |  -  |  <0.05 |
| Muscle Glycogen | 0 | > 0.05 |
| Muscle Free fatty Acid | 0 |
| Muscle Total Protein | 0 |
| Muscle Total Amino Acid | 0 |
| Plasma Free Fatty Acid | 0 |
| Plasma Total Protein | 0 |
| Plasma Total Amino Acid |  -  |  <0.05 |
| *Hyriopsis myersiana* | Average of *Cyprinus carprio* & *Oreochromis nilotica* & *Clarias macrocephalus x C. gariepinus* & *Pangasius pangasius* | Total Protein | Infested with 2662.5 to 129.4 glochidia per branchial chamber, the chambers being kept closed for 2 min | Control \* | 3 dpi |  +  | <0.01 | / | 48 | Uthaiwan, et al, 2003 |
| Triglyceride | 0 | >0.05 |
| Glucose | 0 |
| Copper (Cu) | 0 |
| Manganese (Mn) | 0 |
| Phosphorus (P) | 0 |
| Sulfur (S) | 0 |
| Sodium (Na) | 0 |
| Calcium (Ca) | 0 |
| Potassium (K) |  -  | <0.05 |
| Magnesium (Mg) | 0 | >0.05 |
| Chloride (Cl) | 0 |
| Osmolality  | 0 |
| Total Protein | 6 dpi | 0 | >0.05 |
| Triglyceride | 0 |
| Glucose | 0 |
| Copper (Cu) | 0 |
| Manganese (Mn) | 0 |
| Phosphorus (P) | 0 |
| Sulfur (S) | 0 |
| Sodium (Na) | 0 |
| Calcium (Ca) | 0 |
| Potassium (K) | 0 |
| Magnesium (Mg) |  -  | <0.01 |
| Chloride (Cl) | 0 | >0.05 |
| Osmolality  | 0 |
| Total Protein | 12 dpi |  +  | <0.01 |
| Triglyceride |  +  | <0.01 |
| Glucose | 0 | >0.05 |
| Copper (Cu) | 0 |
| Manganese (Mn) | 0 |
| Phosphorus (P) | 0 |
| Sulfur (S) | 0 |
| Sodium (Na) | 0 |
| Calcium (Ca) |  +  | 0,05 |
| Potassium (K) | 0 | >0.05 |
| Magnesium (Mg) |  -  | <0.01 |
| Chloride (Cl) |  -  | <0.05 |
| Osmolality  | 0 | >0.05 |
| *Cyprinus carprio* | Total Protein | 12 dpi | 0 | >0.05 | / | 6 |
| Glucose | 0 |
| Triglyceride | 0 |
| Copper (Cu) | 0 |
| Phosphorus (P) | 0 |
| Manganese (Mn) | 0 |
| Magnesium (Mg) | 0 |
| Calcium (Ca) | 0 |
| Sulfur (S) | 0 |
| Sodium (Na) | 0 |
| Chloride (Cl) | 0 |
| Osmolality  | 0 |
| *Oreochromis nilotica* | Total Protein | 0 | >0.05 | 6 |
| Glucose | 0 |
| Triglyceride |  -  | <0.01 |
| Copper (Cu) | 0 | >0.05 |
| Phosphorus (P) | 0 |
| Manganese (Mn) | 0 |
| Magnesium (Mg) |  +  | 0,05 |
| Calcium (Ca) | 0 | >0.05 |
| Sulfur (S) | 0 |
| Sodium (Na) | 0 |
| Chloride (Cl) | 0 |
| Osmolality  | 0 |
|  *Clarias macrocephalus x C. gariepinus* | Total Protein | Control ¤ | 0 | >0.05 | 6 |
| Glucose | 0 |
| Triglyceride |  -  | <0.01 |
| Copper (Cu) | 0 | >0.05 |
| Phosphorus (P) | 0 |
| Manganese (Mn) | 0 |
| Calcium (Ca) | 0 |
| Magnesium (Mg) | 0 |
| Potassium (K) | 0 |
| Sulfur (S) |  +  | <0.05 |
| Sodium (Na) | 0 | >0.05 |
| Chloride (Cl) | 0 |
| Osmolality  | 0 |
|  *Pangasius pangasius* | Total Protein | 0 | >0.05 | 6 |
| Glucose | 0 |
| Triglyceride |  -  | <0.05 |
| Copper (Cu) | 0 | >0.05 |
| Phosphorus (P) | 0 |
| Manganese (Mn) | 0 |
| Calcium (Ca) | 0 |
| Magnesium (Mg) | 0 |
| Potassium (K) | 0 |
| Sulfur (S) |  +  | <0.01 |
| Sodium (Na) | 0 | >0.05 |
| Chloride (Cl) |  +  | <0.01 |
| Osmolality  | 0 | >0.05 |
| *Utterbackia imbecillis* | *Lepomis macrochirus* | Hematocrit | 25 minutes in 1000 gl/l in 19l | Control \* | 1 dpi |  -  | < 0.05 | / | 24 | Dubansky et al 2011 |
| Sodium (Na) | 0 | > 0.05 |
| Glucose | 0 |
| Lactate | 0 |
| Cortisol | 0 |
| Hematocrit | 25 minutes in 2000 gl/l in 19l | 0 | > 0.05 |
| Sodium (Na) | 0 |
| Glucose | 0 |
| Lactate | 0 |
| Cortisol |  +  | < 0.05 |
| Hematocrit | 25 minutes in 4000 gl/l in 19l |  -  | < 0.05 |
| Sodium (Na) | 0 | > 0.05 |
| Glucose | 0 |
| Lactate | 0 |
| Cortisol |  +  | < 0.05 |
| Hematocrit | 25 minutes in 8000 gl/l in 19l | 0 | > 0.05 |
| Sodium (Na) | 0 |
| Glucose | 0 |
| Lactate | 0 |
| Cortisol |  +  | < 0.05 |