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| **Table S2 | Sequencing data of novel miRNA mapping during acute and chronic infection with *T. gondii* oocysts.**  |
|  | 11 days post infection |  | 33 days post infection |
| Mouse groups | Control samples |  | Infected samples |  | Control samples |  | Infected samples  |
| Mapped mature | 41 | 73 | 79 |  | 69 | 73 | 50 |  | 60 | 71 | 50 |  | 51 | 57 | 68 |
| Mapped star | 5 | 9 | 6 |  | 8 | 7 | 5 |  | 6 | 6 | 6 |  | 5 | 6 | 4 |
| Mapped hairpin | 48 | 77 | 84 |  | 73 | 78 | 54 |  | 62 | 73 | 54 |  | 54 | 61 | 69 |
| Mapped unique sRNA | 64 | 109 | 114 |  | 95 | 107 | 74 |  | 94 | 108 | 76 |  | 82 | 82 | 92 |
| Mapped total sRNA | 153 | 292 | 298 |  | 299 | 230 | 149 |  | 206 | 329 | 208 |  | 170 | 187 | 231 |