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| **S1 Table. Summary of the RNA-Seq raw data in *Tribolium castaneum*** | | | |  |
|  |  |  |  |  |
| Sample | Raw Reads | Clean Reads | Useful Reads % | GC % |
| Control | 17,307,627 | 12,524,554 | 72.36% | 48.09% |
| Hypoxia/hypercapnia | 19,866,284 | 14,193,339 | 71.44% | 50.74% |
|  |  |  |  |  |
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