Table S1

Number and proportion of reads conserved after each step of bioinformatic treatment of Brazilian (BRE) and Guadeloupean (GH2) strains of *Schistosoma mansoni*.

|  |  |  |
| --- | --- | --- |
|  | BRE | GH2 |
| Raw Clusters  | 540,196,370 | 508,893,394 |
| Reads after quality filters  | 203,373,434 | 204,453,418 |
| Reads successfully aligned  | 95,474,777 | 122,346,146 |