|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  |  | **Concentration of cDNA****(ng/µl)** | **Coverage depth****(mean)** | **Number of Sequenced Reads** | **Coverage per genome (Percentage)** | **CT****(N2 Gene)** |
| **d1** | Mean | 29.64 | 519.91 | 42582.01 | 89.38 |   |
| std | 20.97 | 563.50 | 42879.65 | 19.53 |   |
| Min | 1.00 | 0.11 | 9.00 | 0.00 |   |
| 0.25 | 15.00 | 161.18 | 16865.00 | 91.19 |   |
| Median | 24.20 | 380.15 | 32540.00 | 96.02 |   |
| 0.75 | 39.00 | 679.00 | 53896.00 | 98.68 |   |
| Max | 120.00 | 6152.87 | 477778.00 | 100.00 |   |
| **d2** | Mean | 29.94 | 510.19 | 42806.61 | 90.94 | 16.13 |
| std | 21.03 | 453.22 | 34931.28 | 16.34 | 4.95 |
| Min | 2.28 | 0.12 | 17.00 | 0.00 | 0.00 |
| 0.25 | 15.80 | 182.43 | 18320.50 | 91.35 | 13.00 |
| Median | 23.80 | 412.30 | 34582.00 | 95.87 | 15.70 |
| 0.75 | 36.40 | 682.63 | 54796.00 | 98.68 | 19.05 |
| Max | 120.00 | 2998.20 | 233031.00 | 100.00 | 33.20 |