

A

| Samples | Number of reads | Avg. length (bp) | Total bp (Mbp) | Subtotals (reads) | Total contigs | Unique contigs |
|------------------------|------------------------|-------------------------|-----------------------|--------------------------|----------------------|-----------------------|
| AZ-0h-30DAP | 46,048 | 413 | 19 | 657,819 | 36,583 | 4,669 |
| AZ-3h-30DAP | 164,920 | 398.2 | 65 | | | |
| AZ-6h-30DAP | 120,367 | 422 | 50 | | | |
| AZ-9h-30DAP | 326,484 | 425 | 138.6 | 553,805 | 33,740 | 2,856 |
| AZ-0h-150DAP | 164,027 | 437 | 71.6 | | | |
| AZ-3h-150DAP | 86,217 | 371.7 | 32 | | | |
| AZ-6h-150DAP | 121,711 | 431.5 | 52 | | | |
| AZ-9h-150DAP | 181,850 | 399 | 72 | 1,099,385 | 40,897 | 6,930 |
| P-0h-150DAP | 339,657 | 427 | 145 | | | |
| P-3h-150DAP | 278,933 | 407.8 | 113 | | | |
| P-6h-150DAP | 416,935 | 450 | 187.8 | | | |
| P-9h-150DAP | 63,860 | 395 | 25 | | | |
| Totals/Averages | 2,311,009 | 415 | 971 | | | |

B

| 454 Sequencing data | Numbers |
|------------------------------------|----------------|
| Reads Assembled | 1,563,975 |
| Reads not assembled (singletons) | 747,034 |
| Percentage singletons | 32% |
| Total non-redundant contigs | 51,804 |