|  |  |
| --- | --- |
| Annotation parameter | Number or percent of genes |
| Number of gene models | 13,107 |
| Percent complete (with start and stop codons) | 88% |
| Percent of genes with homology support | 74% |
| Percent of genes with Pfam domains | 49% |
| Percent of genes with EST support | 30% |