Figure 4.15. Blast2Go Analysis progress of groundnut genotype GJG-22

Figure 4.16. Data distribution Pie chart of groundnut genotype GJG-22
Figure 4.17. Number of sequences with length (X) of groundnut genotype GJG-22

Figure 4.18 Maximum blast hit of groundnut genotype GJG-22
Figure 4.19. Species distribution of groundnut genotype GJG-22

Figure 4.20. Top-Hit Species distribution of groundnut genotype GJG-22
Figure 4.21. E-value distribution of groundnut genotype GJG-22

Figure 4.22. Sequence similarity distribution of groundnut genotype GJG-22
Figure 4.23. Direct GO count (BP) of groundnut genotype GJG-22

Figure 4.24. Overall Biological process (BP) of groundnut genotype GJG-22
Figure 4.25. Direct GO count (MF) of groundnut genotype GJG-22

Figure 4.26. Overall Molecular Function (MF) of groundnut genotype GJG-22
Figure 4.27. Direct GO count (CC) of groundnut genotype GJG-22

Figure 4.28. Overall Cellular Component (CC) of groundnut genotype GJG-22
Figure 4.29. GO distribution by level (2) – Top 20 of groundnut genotype GJG-22

Figure 4.30. Enzyme code distribution of groundnut genotype GJG-22
Figure 4.31. Evidence code distribution for BLAST Hits of groundnut genotype GJG-22

Figure 4.32. Mapping database source of groundnut genotype GJG-22