

Table 4 Basic features of assembled RNA transcripts and functional annotation in *A. niger*

Total genomic loci	9939
Loci having one transcript	2913 (29.3%)
Loci having more than one transcript	7026 (70.7%)
Mapped to gene model loci	8347
New genomic loci	1592
Total unique transcripts	66007
Average transcript length (bp)	3735
Transcripts match to gene model transcripts	27566 (41.8 %)
BLASTX match against Swiss-Prot dataset	22162 (33.6%)
Total predicted ORFs	61153 (92.6%)
Average ORF length (amino acids)	285
Total ORFs with a Pfam match	19177 (31.4%)