

Table 1: Functional Classification of salivary gland transcriptome of adult female *Anopheles gambiae* mosquitoes, based on transcript similarity to gene or adjacent untranslated region (UTR)

Transcript class	Based on Proteome match		Based on Proteome + UTR match	
	Number of Contigs	Number of Sequences	Number of Contigs	Number of sequences
Housekeeping	357	554	548	762
Unknown	337	382	132	157
Secreted	155	1940	166	1954
Bacterial	9	32	11	34
TE	3	3	4	4