|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| GeneID | Genename |  | Top BLASTx hit |  |  |  |  |  |
|  |  |  | Description | E-value | %ID | Species | Acc.no |  |
| Unigene5201 | GR1 |  | Gustatory receptor 3 | 0.0 | 73% | *Helicoverpa assulta* | KJ542712.1 |  |
| CL1105.Contig1 | GR2 |  | Gustatory receptor 4 | 0.0 | 74% | *Conogethes punctiferalis* | KX084520.1 |  |
| CL1021.Contig1 | GR3 |  | Gustatory receptor(GR9)  | 4e-123 | 70% | *Helicoverpa armigera* | JX970522.1 |  |
| Unigene11342 | SNMP1 |  | Sensory neuron membrane protein1 | 0e+00 | 70% | *Cnaphalocrocis medinalis* | JN867063.1 |  |
| CL3030.Contig3 | SNMP2 |  | SNMP2 | 0e+00 | 75% | *Ostrinia furnacalis* | [LC017802.1](https://www.ncbi.nlm.nih.gov/nucleotide/814544249?report=genbank&log$=nucltop&blast_rank=2&RID=V3V59J6W014) |  |