**Fas2:**

- #48449 (GMR11C08; X:4061472..4064988) has a Hb target motif with:

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| 1 putative sites were predicted with these settings (99%) in sequence named **dm6\_ct\_coords\_7288** | | | | | | | |
| Model ID | Model name | Score | Relative score | Start | End | Strand | predicted site sequence |
| MA0049.1 | hb | 12.824 | 1.00001474388335 | 3417 | 3426 | -1 | GCATAAAAAA |

**CadN:**

- #49660 (GMR31A11; 2L:17665454..17666638) has two Hb target motifs with:

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| 2 putative sites were predicted with these settings (93%) in sequence named **dm6\_dna** | | | | | | | |
| Model ID | Model name | Score | Relative score | Start | End | Strand | predicted site sequence |
| MA0049.1 | hb | 10.601 | 0.931525897820554 | 406 | 415 | 1 | GCATTAAAAA |
| MA0049.1 | hb | 10.724 | 0.935315429114177 | 1088 | 1097 | 1 | GAATAAAAAC |