**Table S1. Proteins identified in fluid-filled reservoirs by LC/MS.**

|  |  |  |
| --- | --- | --- |
| protein match | # of peptides | SpectrumMill score |
| GA\_JG443736 (alpha-amylase homology)1 | 3 | 37.6 |
|
| tubulin alpha chain2 | 5 | 63.4 |
| tubulin beta chain2 | 2 | 40.9 |
| actin 42A2 | 5 | 68.2 |
| actin A22 | 2 | 23.8 |

1from *Apotrechus in silico* translated cDNA library

2from Genbank’s non-redundant protein database