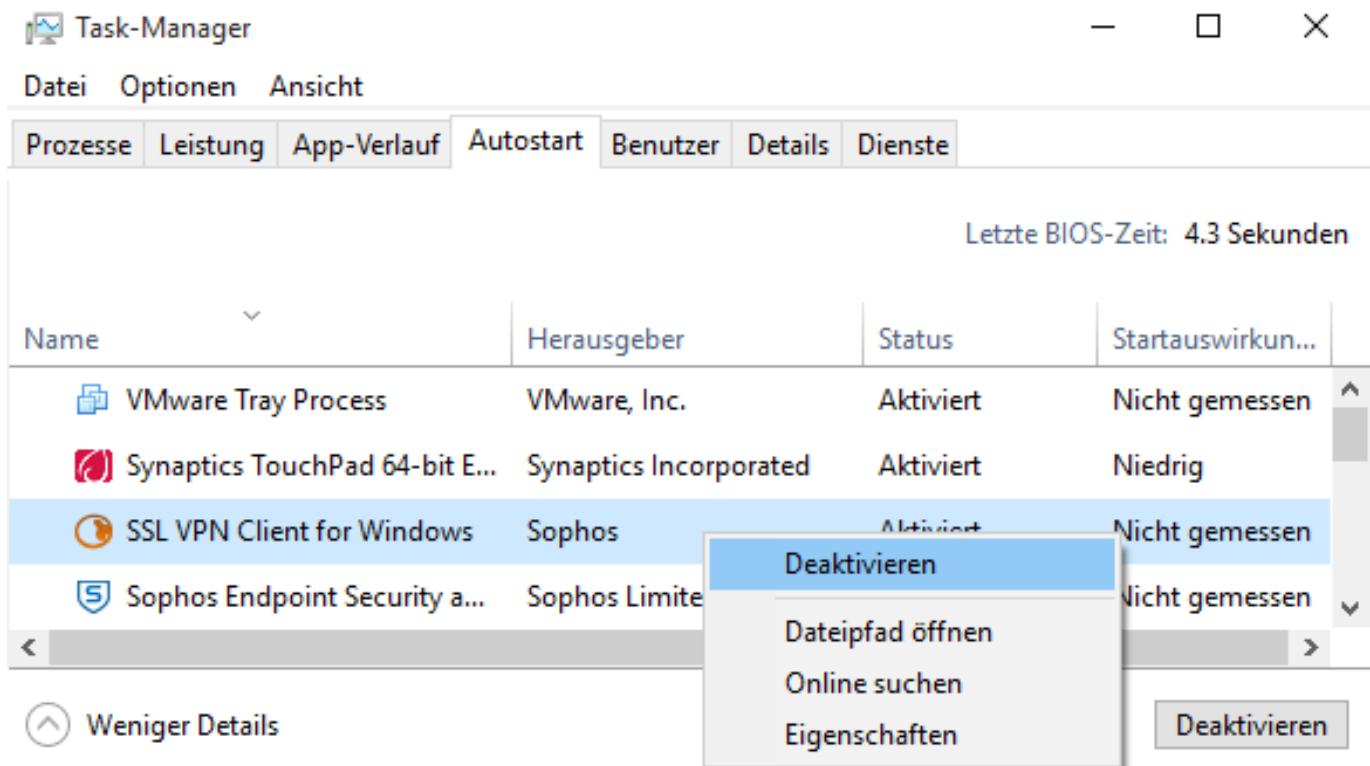


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How to fix this? A: You need to make sure the user is using at least Windows 7 or above. It looks like you're using Windows 8 and you need to upgrade or get a new computer. Q: SQLite count unique values I have a problem with counting the unique values in a table, which is in SQLite Database. I tried this: `SELECT COLUMN_NAME, COUNT(*) FROM ALL_TABLES GROUP BY COLUMN_NAME HAVING COUNT(*)>1;` but it doesn't work properly. Try this: `select column_name, count(*) from all_tables group by column_name having count(*) > 1` A summary of gene expression patterns in the venom gland transcriptome of *Anadasmus tectoricus*. The transcriptome of the venom gland of the Brazilian scorpion *Anadasmus tectoricus* (Lycosidae, Scorpiones) was analyzed by high-throughput sequencing. A total of 6,558,674 reads were generated, with an average length of 532bp. A total of 4,761,219 reads (73.9%) could be successfully aligned to the annotated scorpion genome of *A. tectoricus*, indicating that more than half of the reads were unannotated and non-coding sequences. We also identified sequences of four cDNA clones corresponding to venom peptides, characterized and deposited in GenBank: the alpha-KTx toxin 1 from the scorpion alpha-KTx1 family, the bradykinin-potentiating peptide from the tTx scorpion family and two novel

peptides: the delta-KTx4 from the tTx family and the tTX4.1 from the antimicrobial peptides (AMP) family. The latter was a novel peptide whose sequence was not reported in the databases, but whose function is related to the defense system of the scorpion, since its mRNA is expressed in other tissues, such as the claws and pereonites, and in the mature venom. The expression levels of the coding sequences identified in the venom gland were lower than those reported in the transcriptome of the stinger and venom gland, probably due to the smaller size of the scorpion venom gland compared to the rest of the tissue. No correlation was found between the expression levels and the number of ESTs and 82157476af

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