Figure S1: Ratio diagram of selected taxa and variables compared with *C. lupaster* (standard - all 0).
Figure S2: Ratio diagram of selected taxa and variables compared to *C. lupaster* (standard - all 0)
Figure S3: Ratio diagram comparing male and female *C. lupaster* to *C. aureus* (standard - all 0)
Figure S4: Histogram of results of discriminant analysis of *C. lupaster* (blue) and *C. aureus* (red).
Fig S5. Median-joining network displaying the mitochondrial 12S gene variation (380 bp). Nucleotide sequences used in the network are represented based on their geographic sources as follows: ET= Ethiopia, MR=Morocco, WS=West Sahara, KN=Kenya, IN=India, IS=Israel, SW=Sweden, CH=China, US=United States of America. Sequences used in the network are given in Table S1.