



**Figure 2. 1-D gel profile of *S. crista* and *H. erinaceum* fruiting body proteins.** Total protein was separated on 12.5% SDS-PAGE followed by excision of gel into six gel sections. In-gel trypsin-digested peptides were subjected to nESI-LC-MS/MS. Proteins were identified using the MASCOT search engine and NCBI nr protein database.