

**Table S2. Identification of differentially expressed proteins in inner membrane fraction of *Acinetobacter baumannii* low resistant strain, RS259 with reference to ATCC 19606. Differential expression is shown as fold change (minimum 2 fold, p-value  $\leq 0.05$ ).**

Master No.	Fold change	Mw kDa	pI	Identification and Accession No.	Score	Role in Resistance
806	3.05	43.1	5.21	Elongation factor Tu OS=Acinetobacter baumannii, EFTU_ACIBT	74	Associated with efficient translation.
1003	3.01	51.5	5.43	3-isopropyl malate dehydratase large subunit OS=Acinetobacter baumannii LEUC_ACIBT	40	Involved in amino-acid biosynthesis, branched-chain amino acid biosynthesis, leucine biosynthesis.