

Supplementary Table S.3 = Proteins unique to the recovered and control conditions.

Accession	Description	Score	Condition	Biological Process	Biological Process	SignalP	LipoP	Gene symbol	Subcellular localization	SecP
I3QZS6_CORPS	Cytochrome c nitrate reductase small subunit NrFH	1035.81	Control	Energy metabolism	Metabolism cellular			<i>nfrC</i>	MEM	0.898422
I3QXN4_CORPS	Cytochrome oxidase assembly protein	893.51	Recovered	Energy metabolism	Metabolism cellular			<i>Cp258_1127</i>	MEM	0.966085
I3QWA4_CORPS	Multidrug resistance protein norM	561.85	Recovered	Defense mechanism	Cellular processes and signaling			<i>norM</i>	MEM	0.968227
I3QY62_CORPS	Multidrug resistance protein norM	487.65	Recovered	Defense mechanism	Cellular processes and signaling			<i>norM</i>	MEM	0.971397
I3QV51_CORPS	Thioredoxin related protein	259.85	Recovered	Post-translational modification, protein turnover, chaperones	Cellular processes and signaling	X		<i>Cp258_0224</i>	SEC	0.623525
I3QVD0_CORPS	Unknown function	836.52	Recovered	Unknow function	Poorly characterized	X		<i>Cp258_0305</i>	SEC	0.951009
I3QVQ7_CORPS	Unknown function	3502.56	Recovered	Unknow function	Poorly characterized	X		<i>Cp258_0448</i>	SEC	0.628003
I3QWR5_CORPS	Unknown function	1286.94	Recovered	Unknow function	Poorly characterized	X		<i>Cp258_0810</i>	SEC	0.811667

SecP value = SecP score, value above 0.5 indicates possible secretion