Table S10. List of KEGG Orthology (KO) genes differentially (FDR<0.05) expressed in EGDe *L. monocytogenes* strain throughout the comparison of control samples (non-exposed to lactate and non-pressurized) to samples exposed to lactate and pressurized. Positive Log2 fold change indicate genes more abundant in samples exposed to lactate and pressurized.

Log2 Fold Change	FDR	KEGG annotation at level 1	KEGG annotation at level 2	KEGG pathway	KEGG Orthology (KO) genes
2.164	3.40E-03	Unclassified	Unclassified: metabolism	Enzymes with EC numbers	K01531 - P-type Mg2+ transporter [EC:7.2.2.14], mgtA, mgtB