Loss of ncm⁵ and mcm⁵ wobble uridine side chains results in an altered metabolic profile

Tony Karlsborn¹, A K M Firoj Mahmud^{1†}, Hasan Tükenmez^{1†} and Anders S. Byström^{1,*}

- 1) Department of Molecular Biology, Umeå University, 901 87 Umeå, Sweden
- † These authors contributed equally
- * Corresponding author, Phone (+46)-90-785 67 64; Fax (+46)-90-77 26 30

E-mail address, Anders.Bystrom@molbiol.umu.se

Metabolomics-Springer

Online Resource 7. Summary of the PLS-DA models used in the study.

Culturing Temp.	Strain	Obs.	Tot. Obs.	Comp.	Eigenv.	R2Y	Q2	CV- ANOVA
30°C	elp3∆-l.cempty	9	18	3	5.18	0.87	0.8	3,38E-05
30°C	elp3∆-l.cELP3	9	10	3	3.16	0.87	0.8	3,36E-05
30°C	elp3∆-h.cempty	9	18	1	2.48	0.803	0.35	0,0394747
30°C	elp3∆-h.ctKQE	9	10	1	2.40	0.003	0.33	0,0354747
30°C	elp3∆-h.cempty	9	18	3	6.22	0.765	0.643	2,26E-05
30°C	WT-h.cempty	9	10		0.22	077 02	0.0.0	
34°C	elp3∆-l.cempty	9	18	2	5.82	0.951	0.903	2,67E-07
34°C	elp3∆-l.cELP3	9						,
34°C	elp3∆-h.cempty	9	17*	4	7.39	0.498	0.383	0,0479456
34°C	<i>elp3∆</i> -h.ctKQE	8*						

Obs. Indicates the number of observations from each strain used in the PLS-DA model (three biological replicates with three technical replicates for each biological replicate).

Tot. Obs. Indicates the total number of observations used in the PLS-DA model.

Comp. The number of components in the PLS-DA model.

Eigenv. Eigenvalue describes the length of the vector making up the first component of the PLS-DA model.

R2Y Fraction of Y variation modeled in the first component of the PLS-DA model.

Q2 Cross-validation of the R2-value for the first component in the PLS-DA model.

CV-ANOVA P-value from the CV-ANOVA significance test of the PLS-DA model. The 95% statistical significance was set to P < 0.05 and the 99.9% statistical significance was set to P < 0.001.

^{*} One technical replicate was defined as an outlier in the PLS-DA model and was therefore excluded.