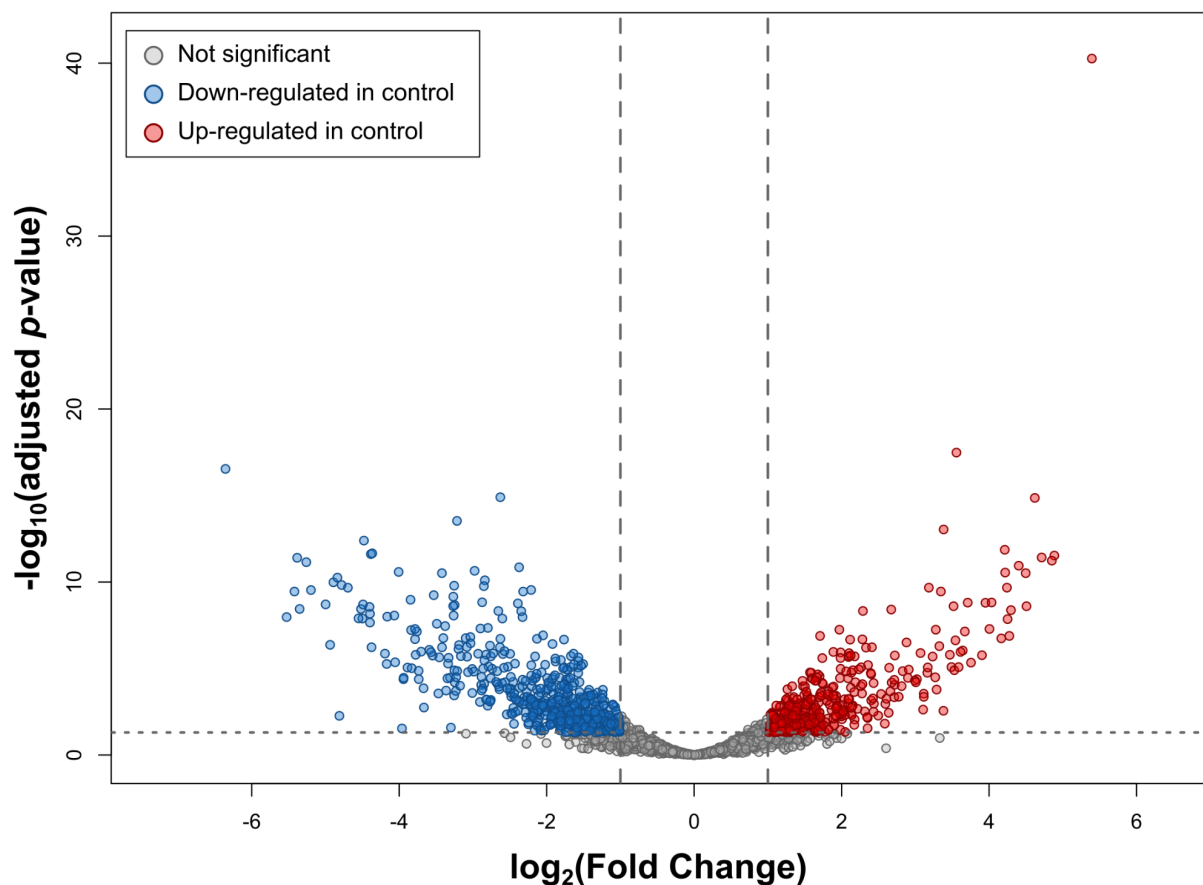


Copper tolerance and transcriptomic response after prolonged Cu^{2+} exposure in the Antarctic psychrophile *Rhodoglobus galactosidasius* strain CPHL_1^T

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Supplementary Fig. S6 Volcano plot summarizing the differential transcript abundance analysis between control and copper-exposed samples. Each point represents an individual gene, including ones coding for hypothetical proteins. Genes meeting the significance criteria (adjusted p -value < 0.05 and $|\log_2$ fold change| > 1) are highlighted. Genes with higher transcript abundance under control condition relative to the copper-exposed samples are shown in red, while genes with lower transcript abundance are shown in blue