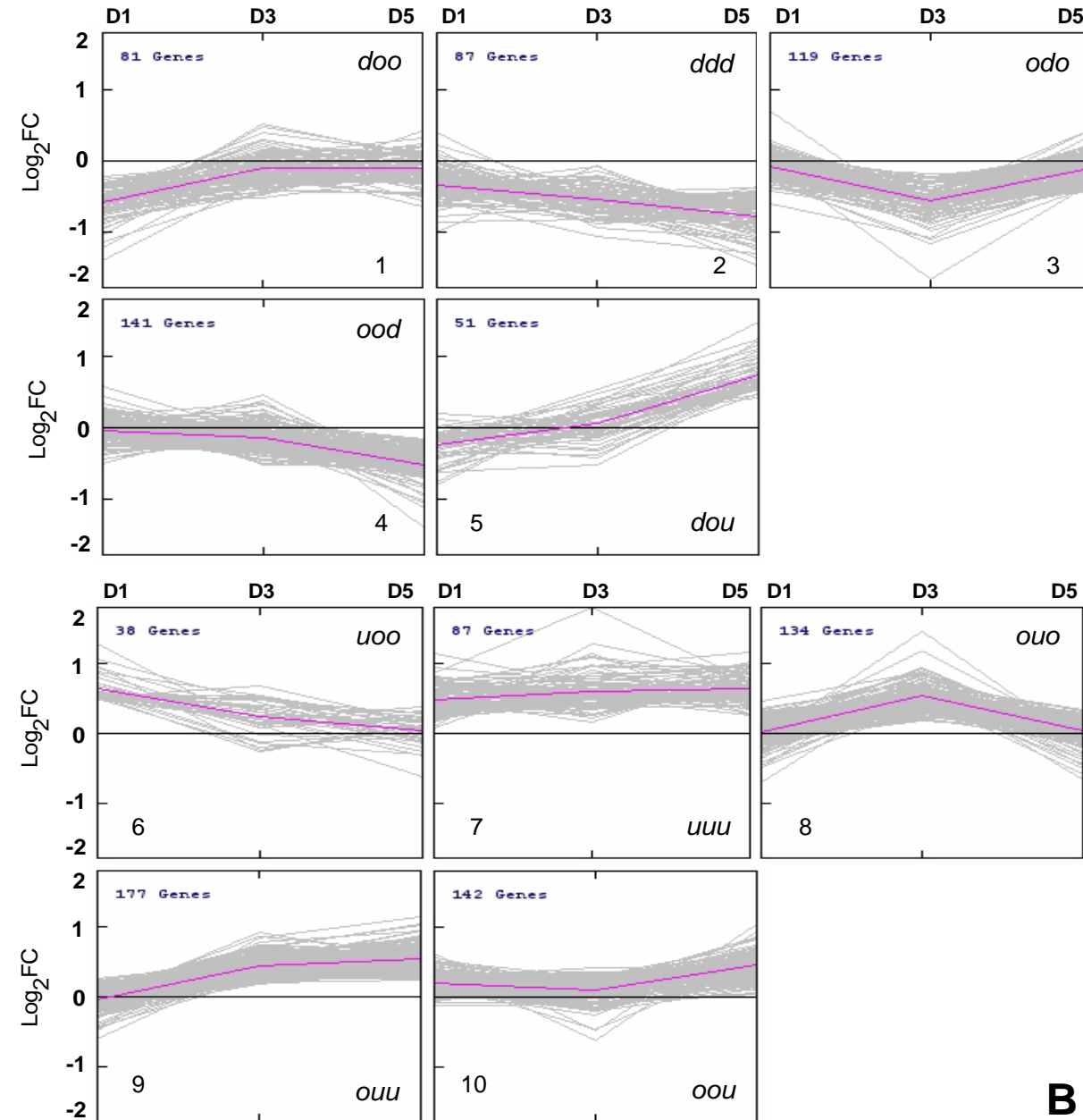


A

Additional file 4 – Clusters of NAA-responsive genes with average values (pink line) and standard deviation (grey area) of the expression levels of the selected genes are presented. In these diagrams, “y” axis represents log₂-fold change and “x” axis represents the different time points for sampling. The cluster names are assigned upregulated (u), unchanged (o) or downregulated (d) for each time point.



Additional file 4 – Clusters of shading-responsive genes with average values (pink line) and standard deviation (grey area) of the expression levels of the selected genes are presented. In these diagrams, “y” axis represents log_2 -fold change and “x” axis represents the different time points for sampling. The cluster names are assigned upregulated (u), unchanged (o) or downregulated (d) for each time point.

B