

Figure S1:

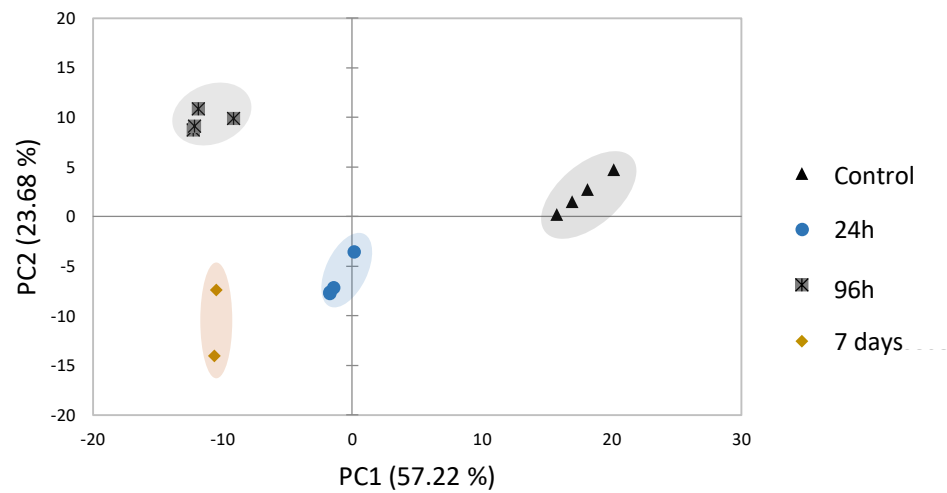


Figure S1: Total lipidome separation by PCA (including TAG). PCA explains 80% of the distribution of the data, and shows a clear separation between induced and non-induced cells suggesting redistribution of the lipidome in response to N deprivation. Data used – normalized peak area.