



File S2. Heatmap of metabolite signals. A number of 161 metabolite signals were generated by $^1\text{H-NMR}$ and 148 signals by UPLC-HRMS (77 by positive and 71 by negative mode) for twelve replicate (1-12) of chicory flour (F) and decoction (D). Metabolite pick areas were expressed in percentage (100 % correspond at the mean for chicory flour) in the heatmap hierarchical clustering with the Ward's method.