S1 Table. Software and package versions used in the study.

Software/Package	Version
R software	v3.2.0
BioConductor	v3.1
clusterProfiler	v2.2.3
FactoMineR	v1.31.4
fastICA	v1.2-0
genefilter	v1.50.0
gplots	v2.17.0
hugene20sttranscriptcluster.db	v8.3.1
limma	v3.24.13
oligo	v1.32.0