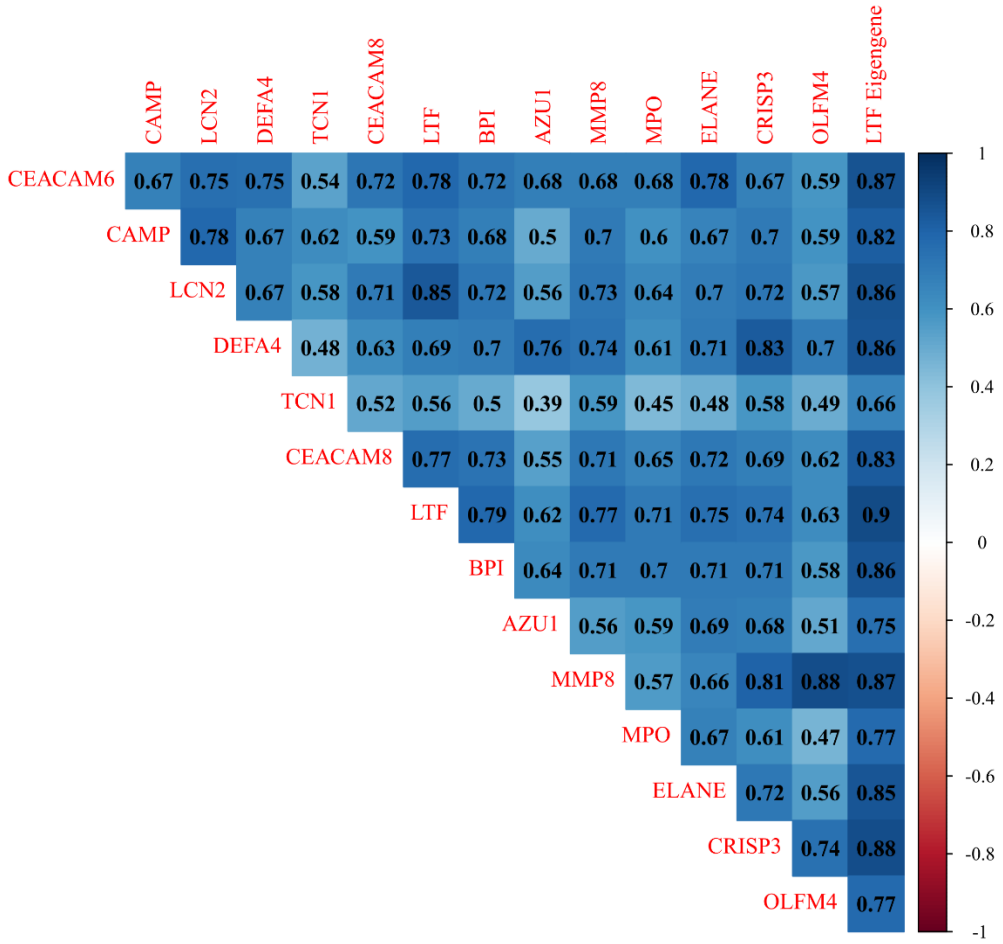
Appendix 7. Heatmap representing the correlation between 14 genes representing the LTF module and the LTF module eigengene

Description*: Heatmap representing the change in gene expression between the IA-2 and IA-1 visits adjusted for age and sex. Numeric values are Pearson correlation coefficients representing pairwise correlations between genes as well as pairwise correlations between each gene and the LTF eigengene (column on far-right side of the figure). Individual genes are ordered along diagonal based on strength of association between change in expression and group (group*visit interaction p value from the longitudinal model), ranging from CEACAM6 (p=0.0010) to OLFM4 (p=0.3210), see Table 4.*