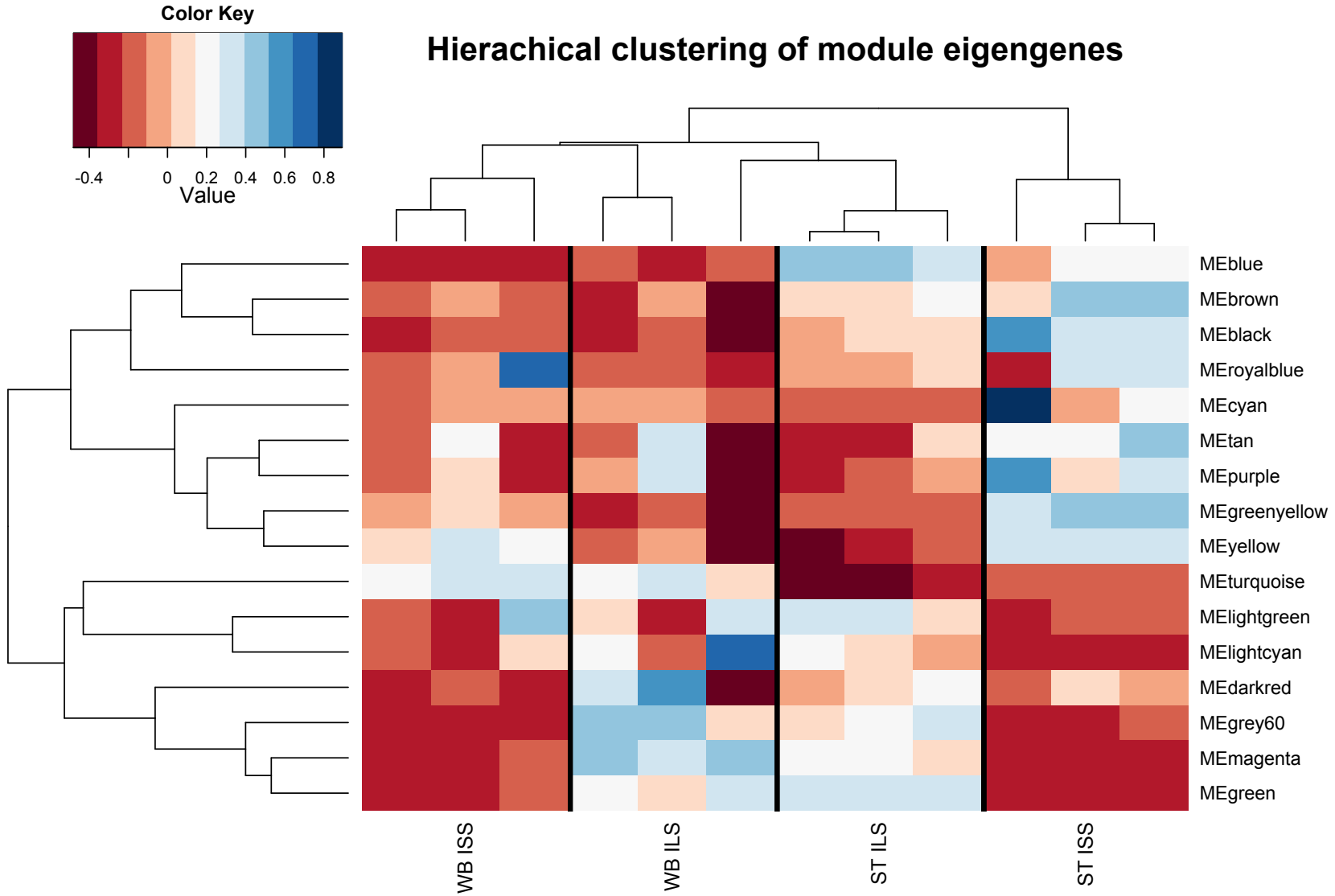

Hierarchical cluster of each module eigengene (rows) and each sample (columns). Eigengene values range from approximately -0.5 to 0.9, representing sample specific expression levels. Higher expression is denoted with blue colors, lower expression by red colors.