Module−−trait relationships in RHA

Row labels (top to bottom):
MElightcyan, MEyellowgreen, MEredsquare4, MEfloralwhite, MEdarkorange, MEdarkmagenta, MElightcyan1, MEblue2, MEwhite, MEorangered4, MEbrush4, MEivory, MEnavy, MElsteelblue, MEnavajowhite4, MEsaddlebrown, MElightcoral, MElightcyan1, MEturquoise, MEdarkgrey, MEskyblue2, MEpaleviolette..., MEfirebrick4, MEorangered2, MEdarkviolet, MEdarkorange3, MEpurple, MEbrown3, MElavenderblush3, MEmediumorchid, MEdarkturquoise, MEmaroon, MEmediumpurple3, MEpaleturquoise, MEbrown4, MEsalmon2, MEcyan, MEfirebrick4, MElilac, MEplum4, MEskyblue1, MEgreenyellow, MEmediumorchid, MEhoneydew, MEantiquewhite4, MEroyalblue, MEred, MEdarkseagreen3, MEdarkolivegreen, MEgreen, MEdarkseagreen4, MEdarkcyan, MEblack, MEdarkslateblue, MElightsteelblue, MEbrown, MEmagenta, MEdarkorange2, MEdarkorange, MElightgreen, MEsteelblue, MElightsteelblue1, MEsalmon, MEnavajowhite2, MEblue, MEskyblue3, MEgrey

Column labels (left to right):
Bench, Osmocote, Salt, Treatment, Accession, Group, Transplant_Date, Chlorophyll, Tot_AG, Tot_BG, Tot_Biomass, Reproductive, SampleDay

Legend scale: 1, 0.5, 0, −0.5, −1