

SFW*Rel.*

SDW*Rel.*

RFW*Rel.*

RDW*Rel.*

Supplementary Figure 4. Pair plot of mixed model estimated genotypic means for eleven traits (Emergence = emergence rate; RFW*Rel.*, RDW*Rel.*, SFW*Rel.*, and SDW*Rel.* = relative root and shoot fresh and dry weight, respectively; NOD = nodulation index; RRI = root rot index; WIL = wilted nodes; CDI = cortex decay index; SLI = shoot lesion index; DIS = disease progress; see Material and Methods for a detailed description). Panels on the diagonal show the distribution of the genotypic means for each trait. Panels below the diagonal show pairwise correlations with a linear regression line. Panels above the diagonal display correlation coefficients (Spearman's rho) and associated significance levels (•= *p*-value < 0.1; \* = < 0.5; \*\* = < 0.01; \*\*\* = < 0.001).