

Supplementary Diagram 1. Flow diagram of analytical pipeline for this study integrating SEM and residual GWAS approaches.

Integrating growth physiology and calcium status to reveal the genetic basis of lettuce tipburn via GWAS

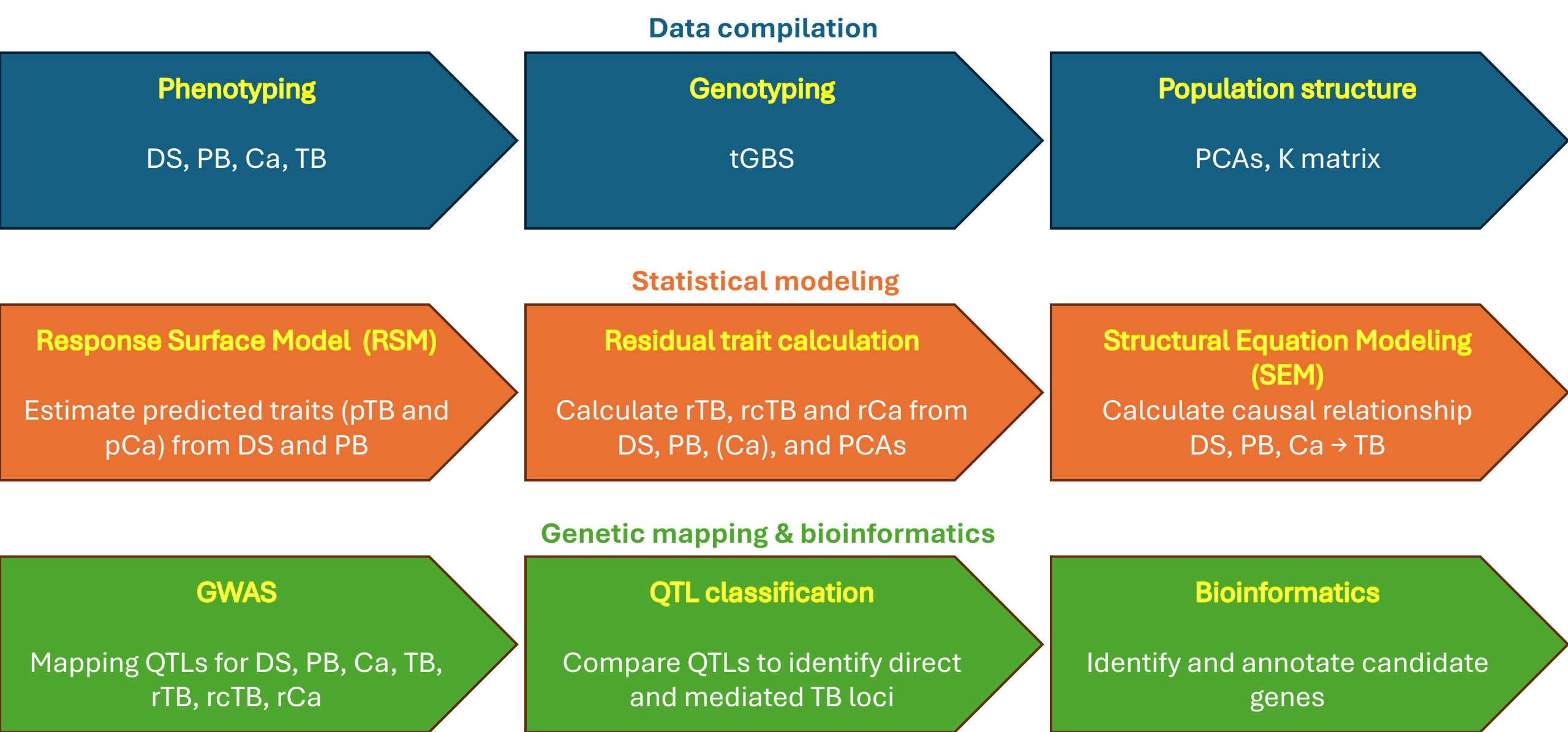
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Running title: Lettuce tipburn resistance



Legend
 DS: developmental score, PB: plant biomass, TB: tipburn incidence, Ca: bulk leaf calcium content, tGBS: tunable genotyping-by-sequencing, PCAs: three principal components, K matrix: kinship matrix, pTB & pCa: traits predicted from growth covariates (DS & PB), rTB : residual TB adjusted for DS, PB, & PCAs, rcTB: residual TB adjusted for DS, PB, Ca, & PCAs, rCa: residual Ca adjusted for DS, PB, & PCAs.