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Research goal

Can we find a way to identify multiple key elements and pathways of corn resistance to Aspergillus flavus by doing major data integration analysis?

Importance

Aspergillus flavus is a fungus with many traits. This fungus colonizes crops such as corn, cottonseed, peanut and more. Some strains produce highly toxic carcinogenic and aflatoxin. chemicals such as resistance aflatoxin Corn to accumulation is polygenic a trait.

Objectives and experiments

Use GWAS **Obj.** 1. phenotypic data to identify genes and pathways associated with corn resistance to aflatoxin accumulation

Obj. 2. Perform post-GWAS using RNA-seq kernel data from (KSA) screening assay OŤ TZAR102 (Resistant line) and Va35 (Susceptible line)



Flavonoids play a key role in resistance to accumulation of aflatoxin in corn







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