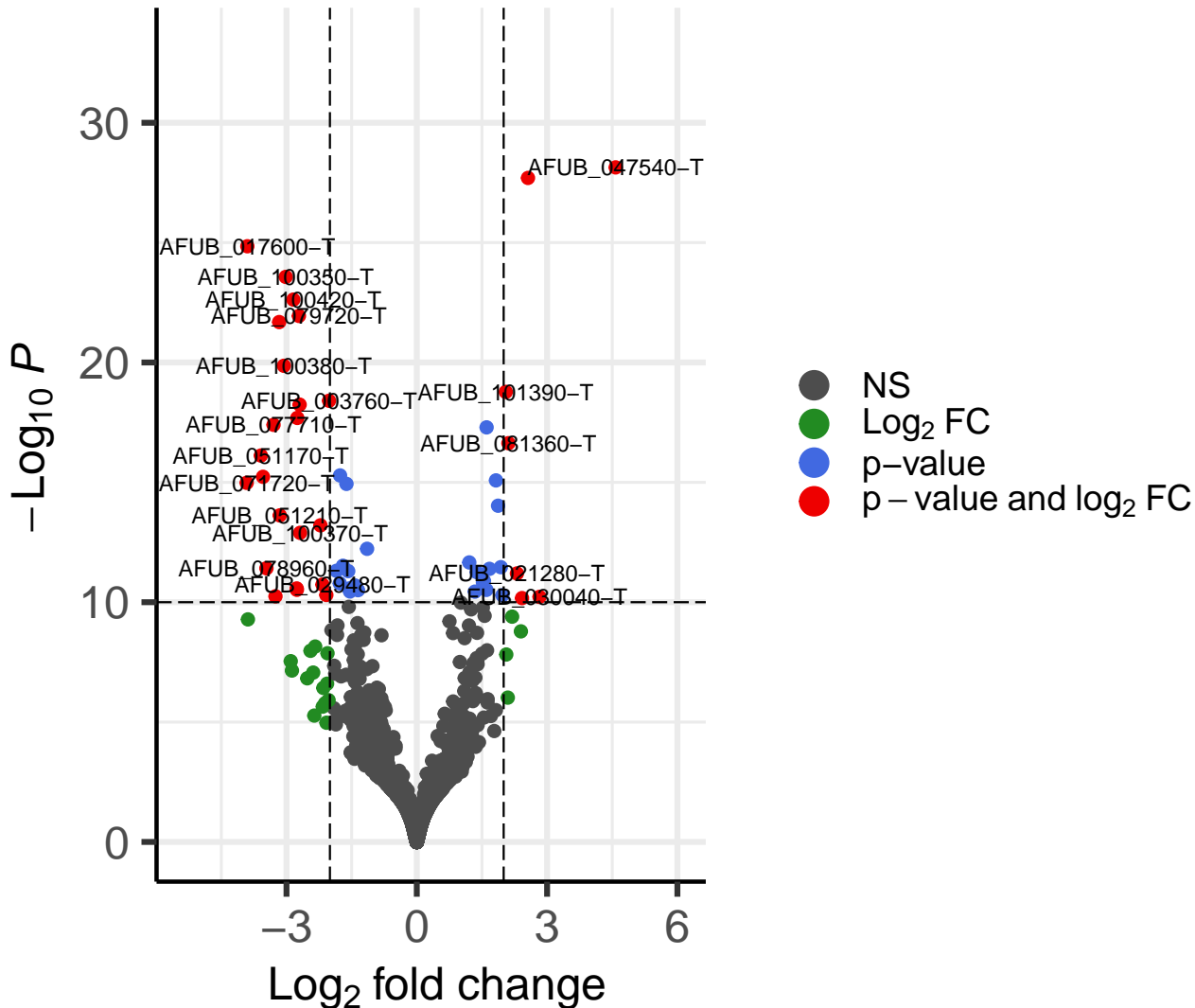
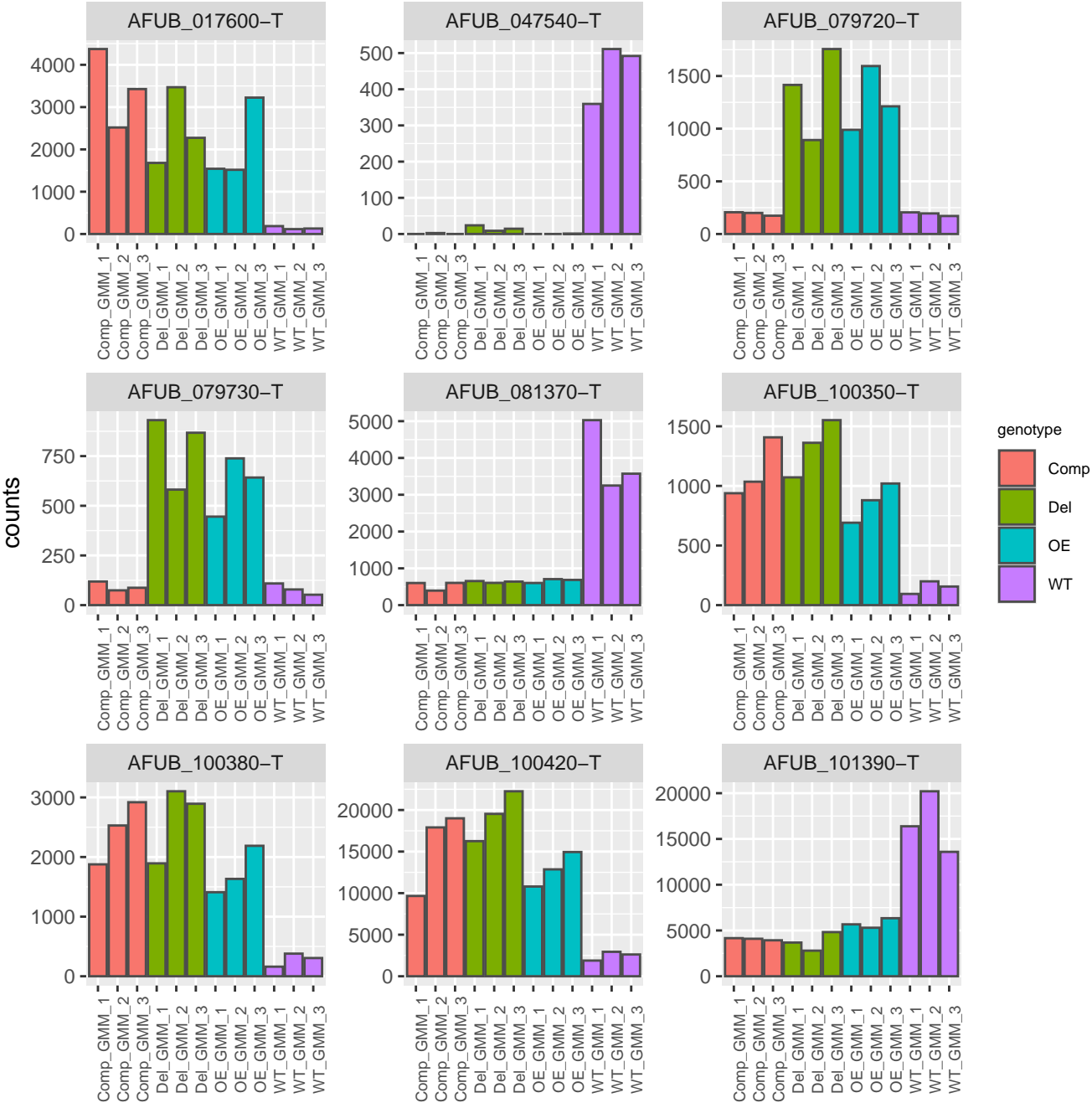


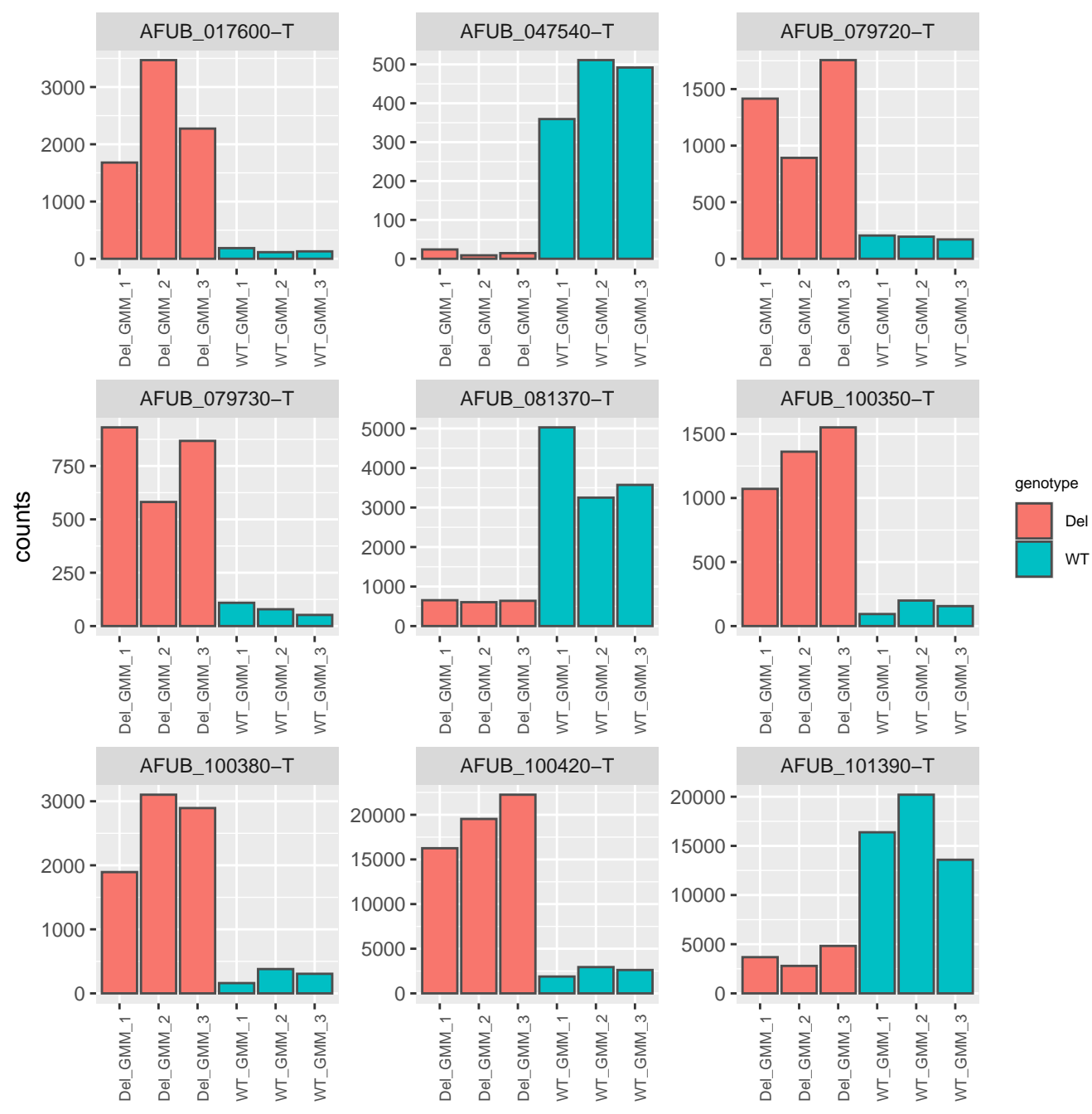
# WT versus Delta

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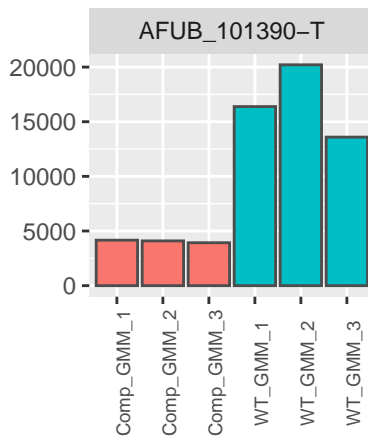
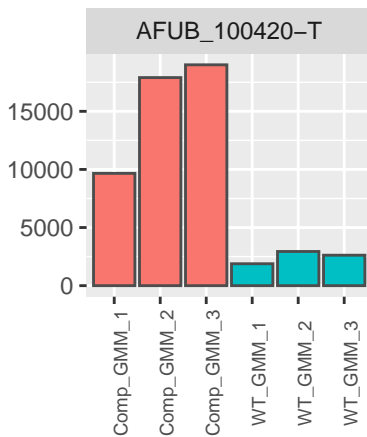
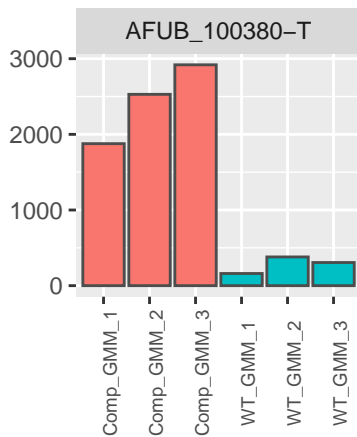
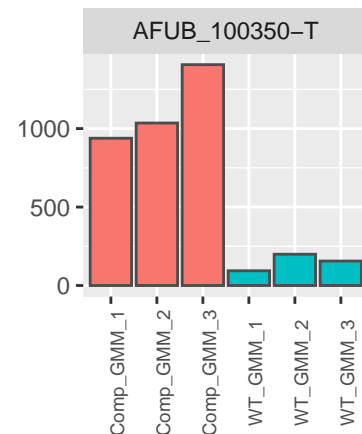
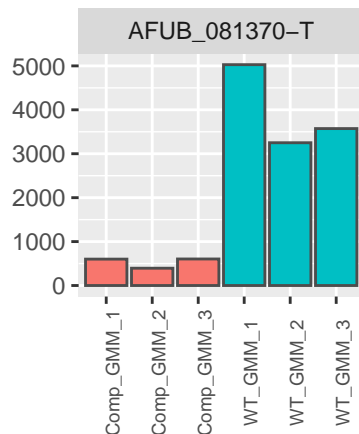
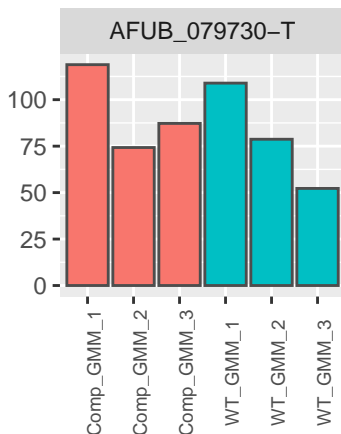
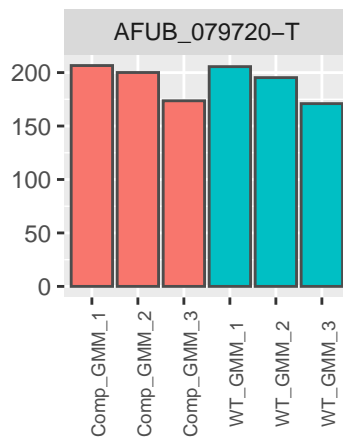
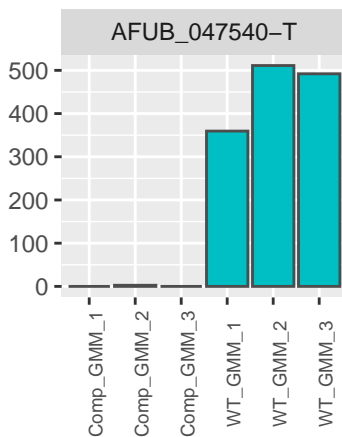
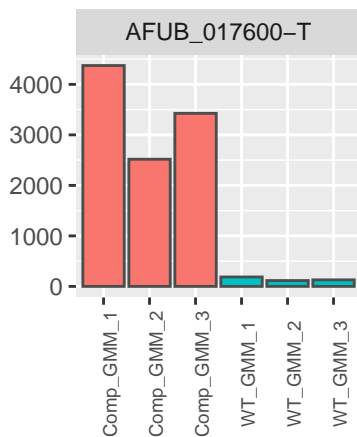


total = 10048 variables



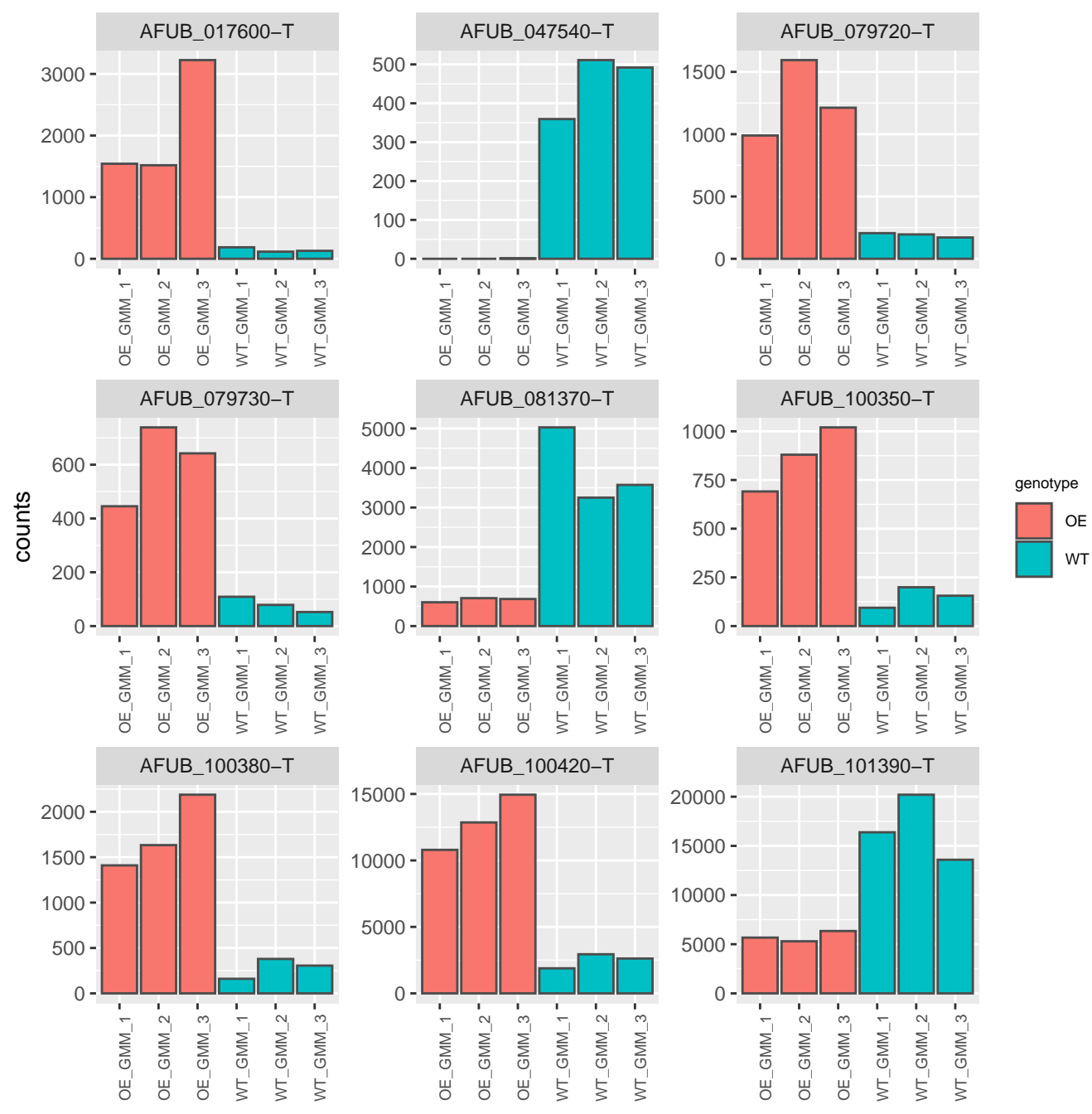


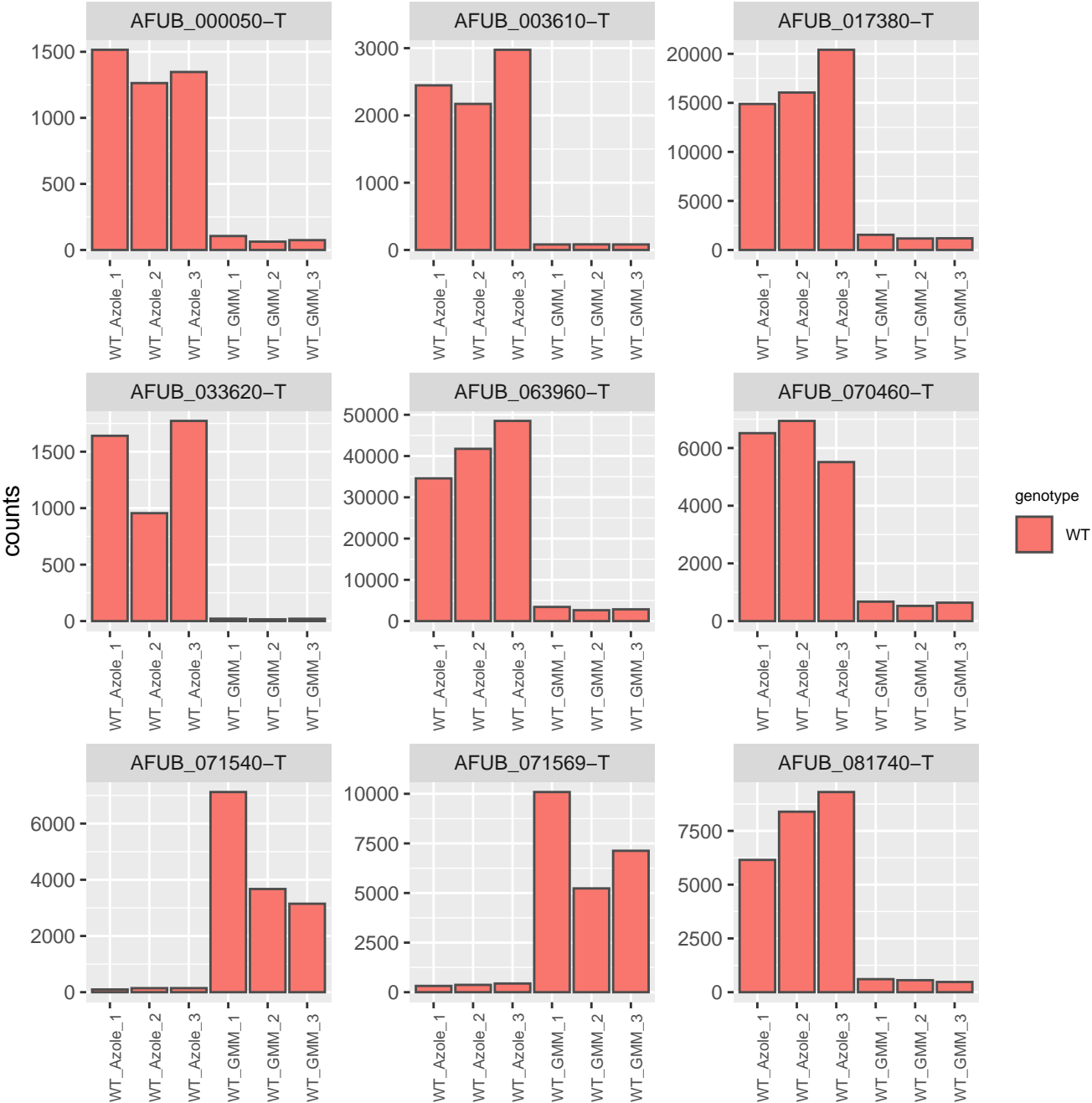
counts

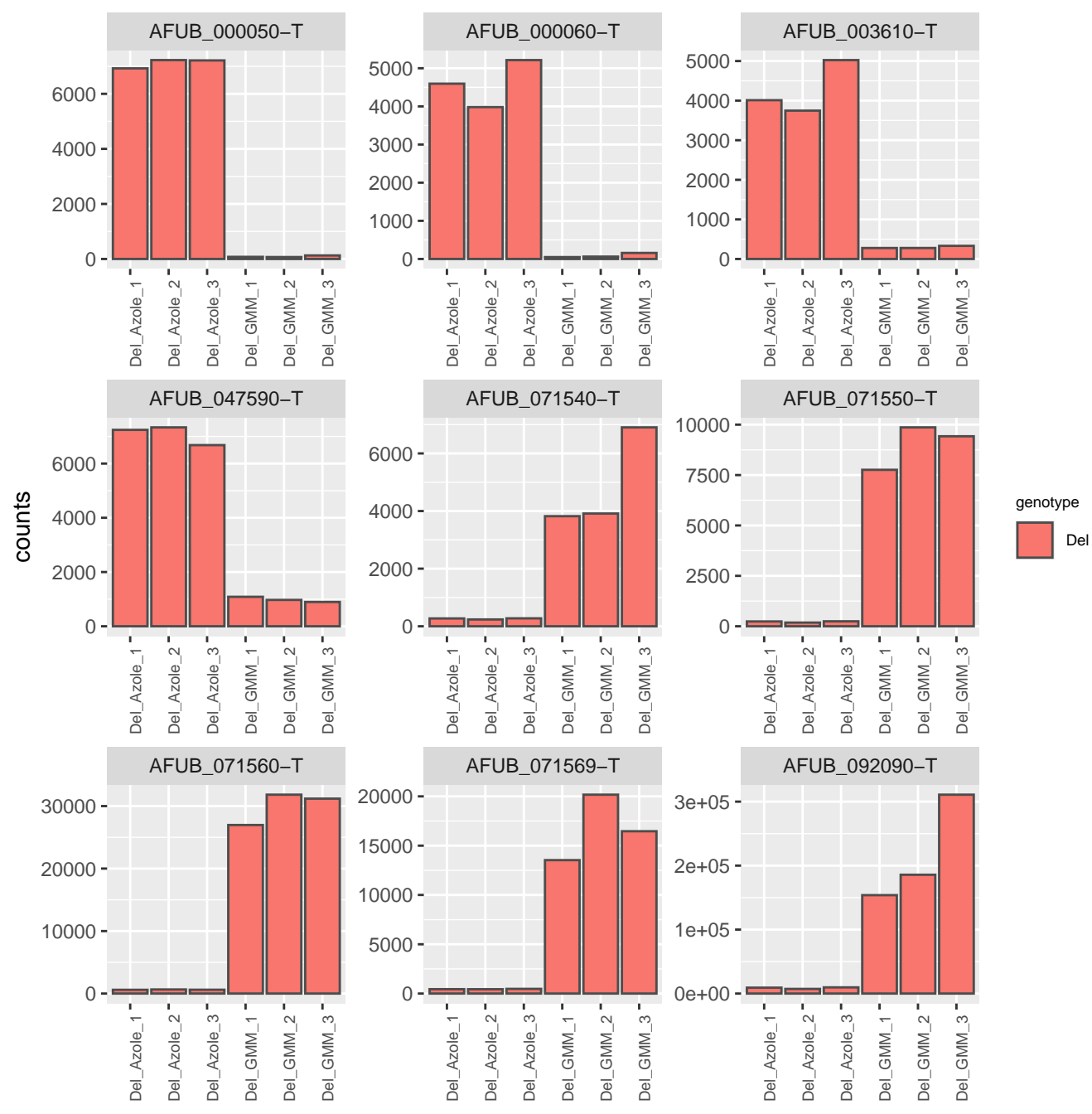


genotype



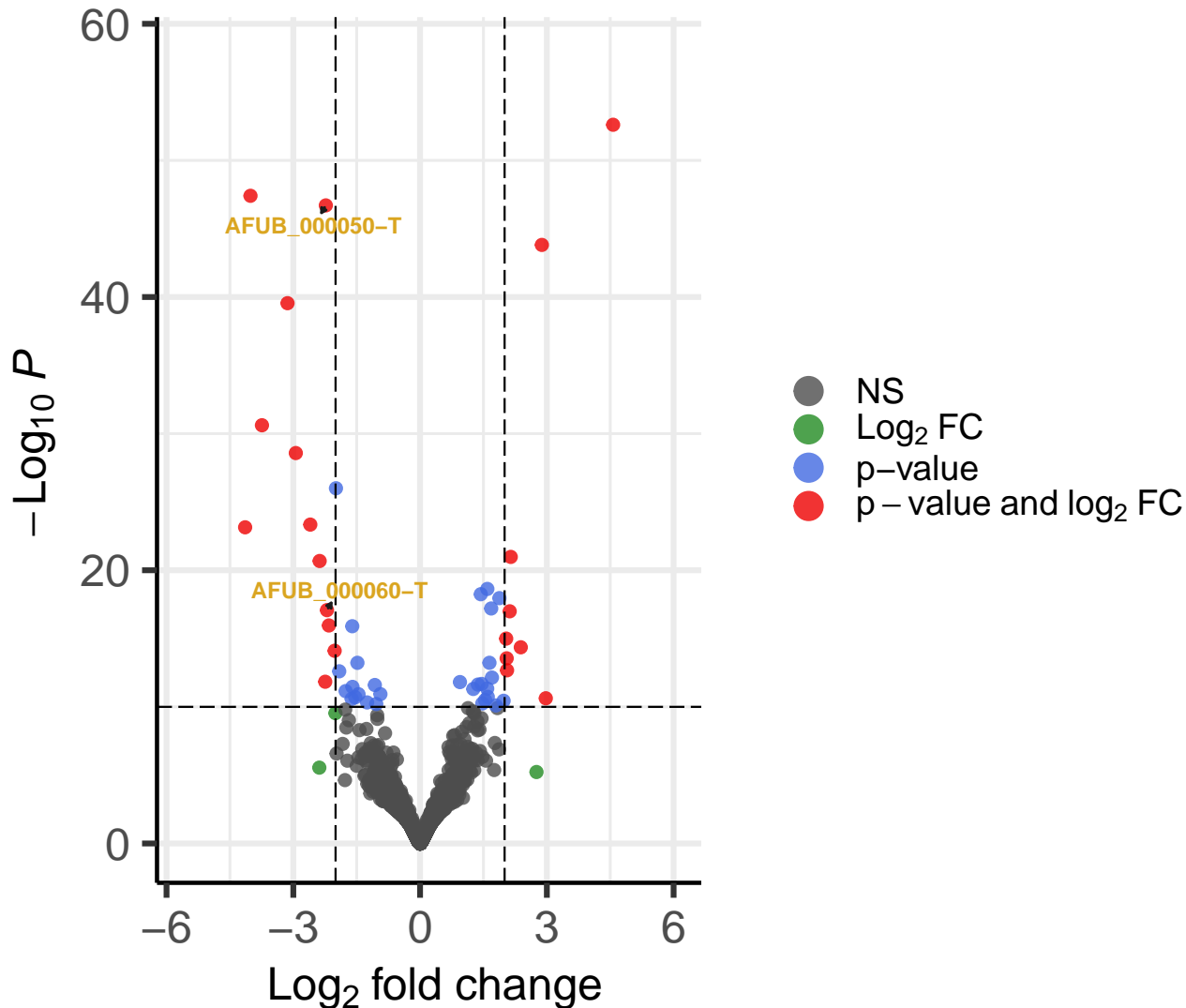






# Azole WT versus Delta

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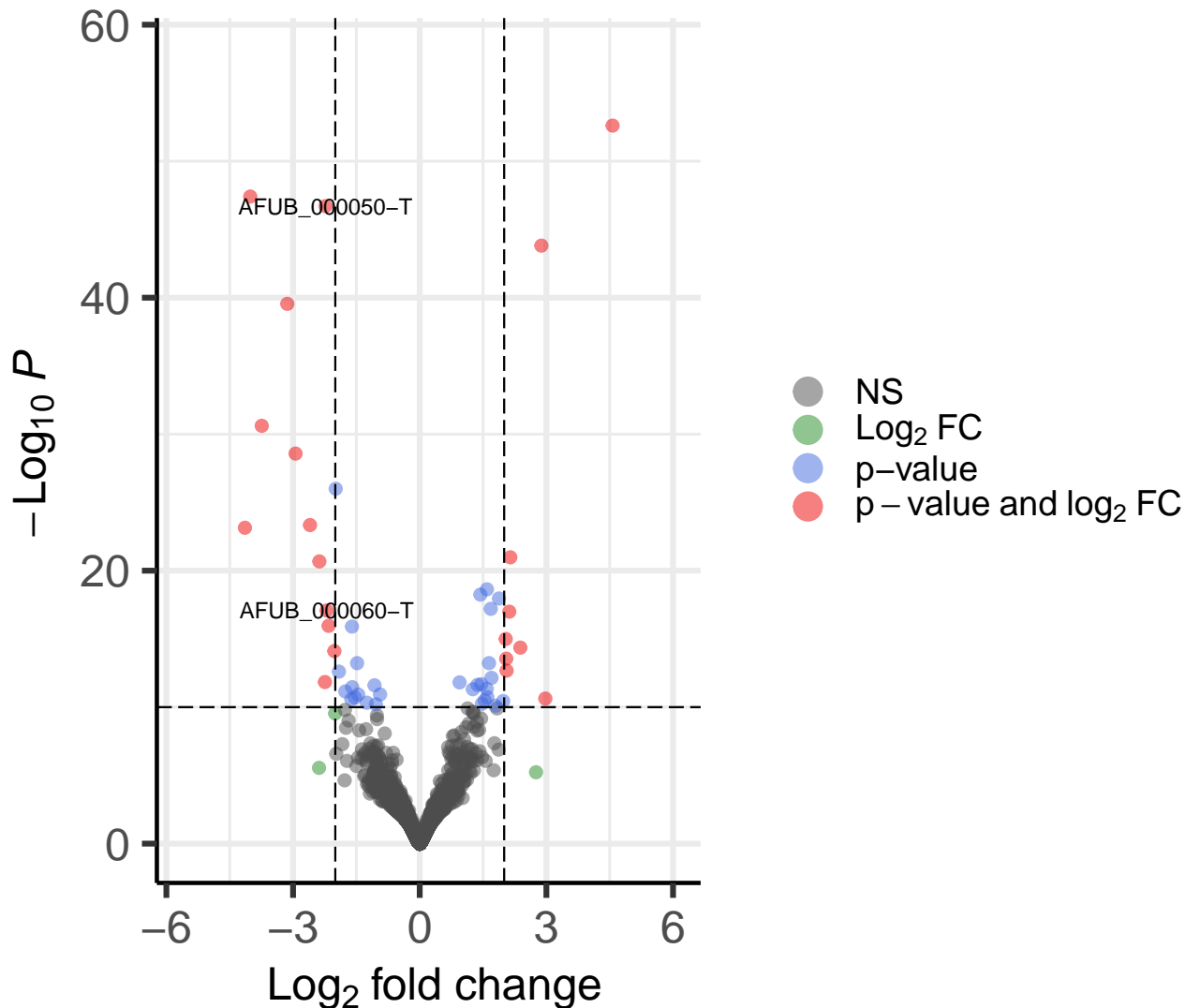


total = 10025 variables



# Azole WT versus Delta

*EnhancedVolcano*



total = 10025 variables

