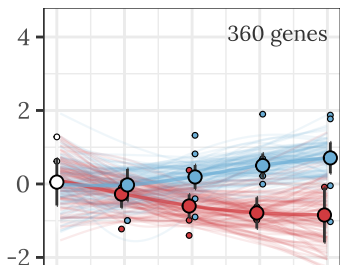


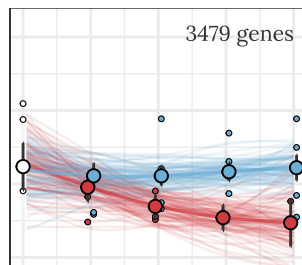
Cell type ○ Pre-grafting ● Queen ● Worker

Relative % methylation of genes in this module  
(eigengene, posterior estimates)

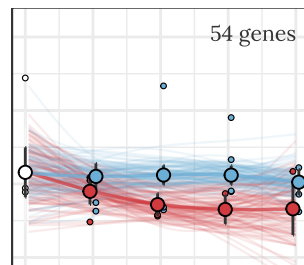
Module 0



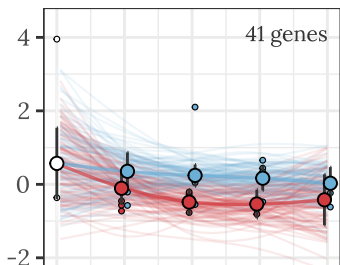
Module 1



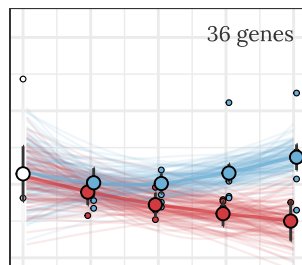
Module 2



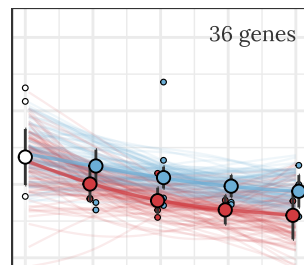
Module 3



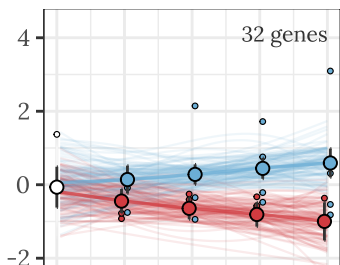
Module 4



Module 5



Module 6



Hours post-grafting into queen or worker cell