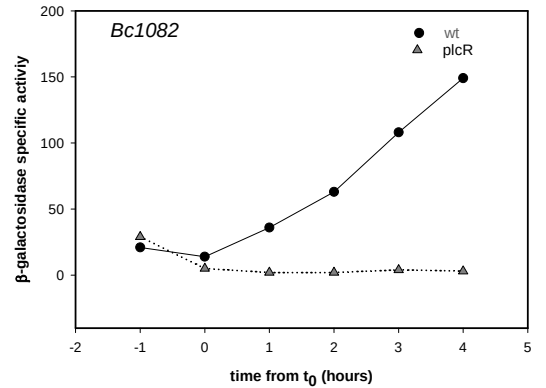
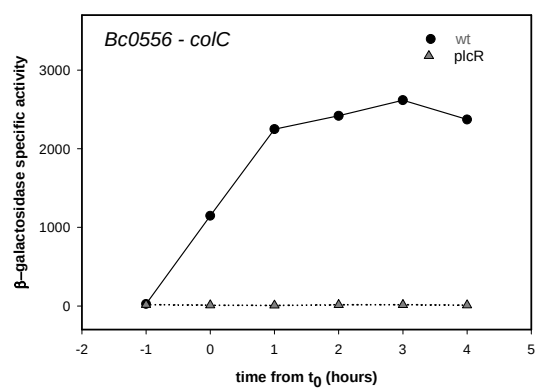
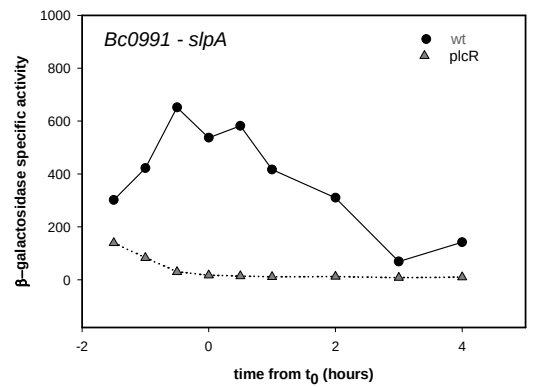
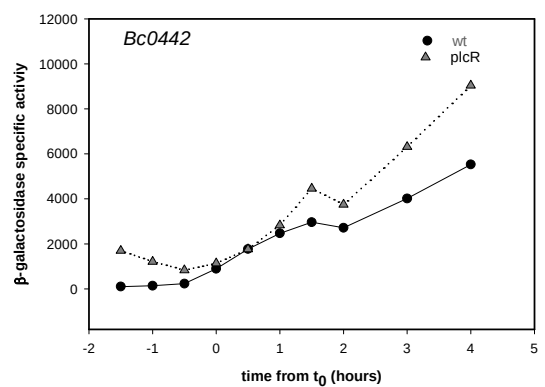
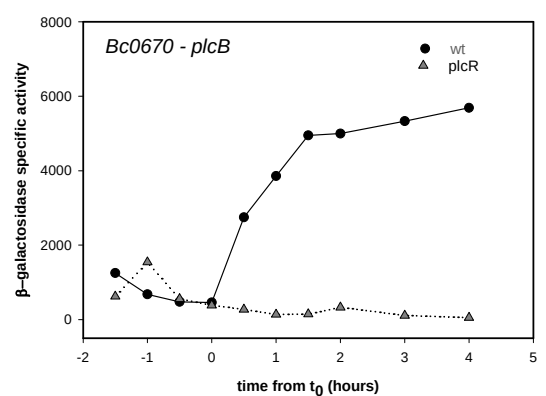
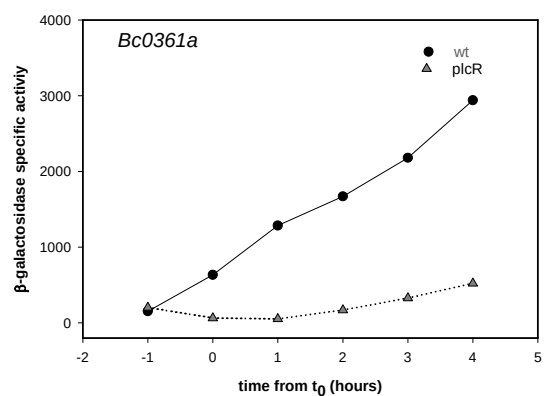
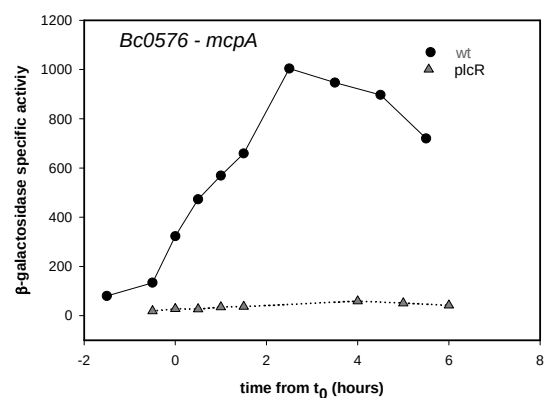
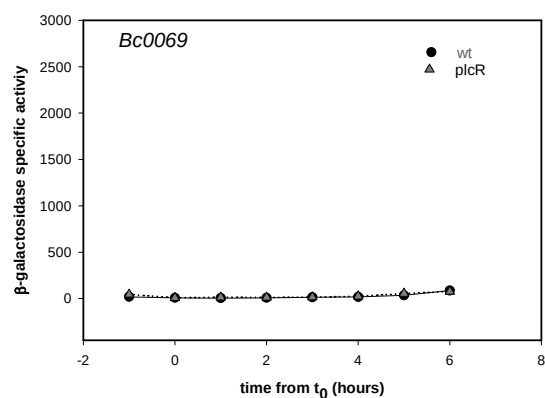
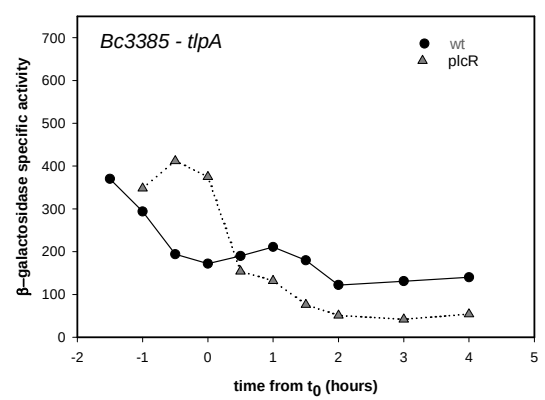
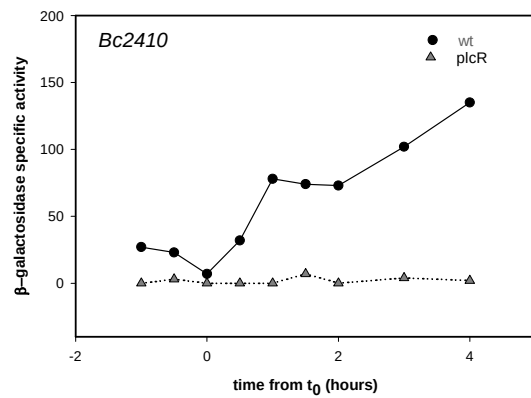
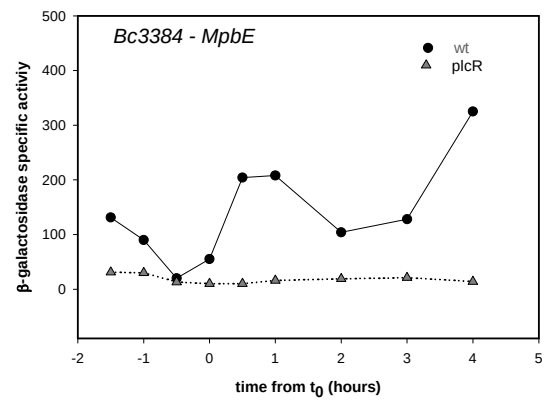
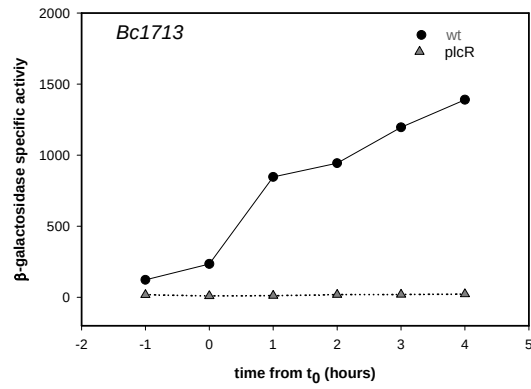
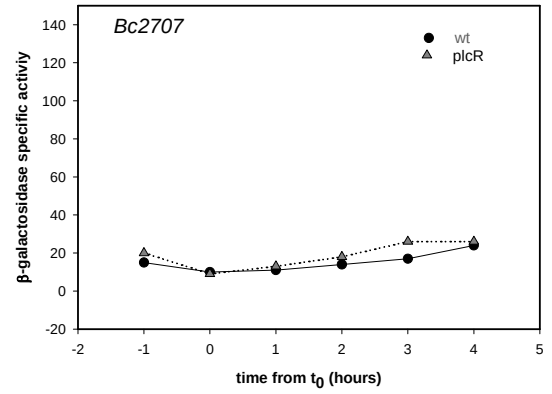
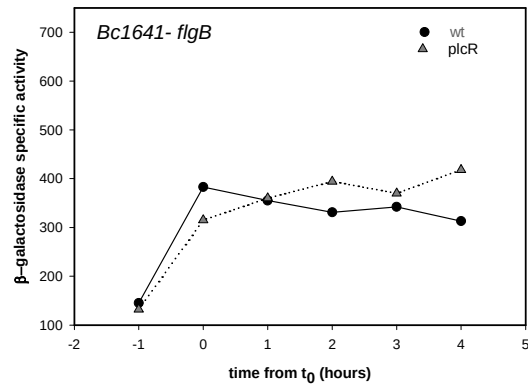
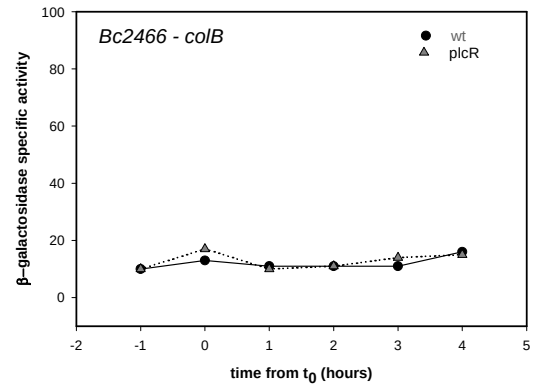
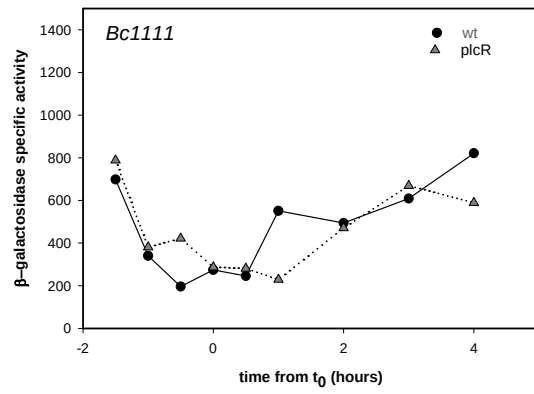
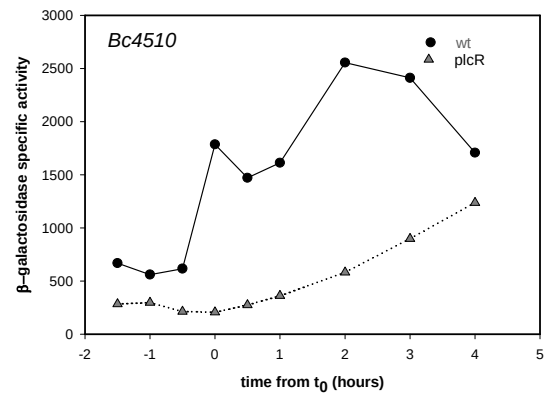
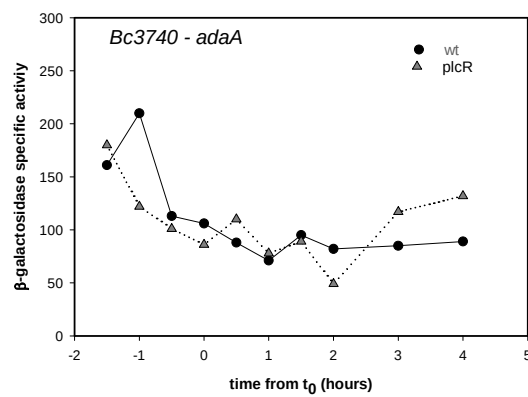
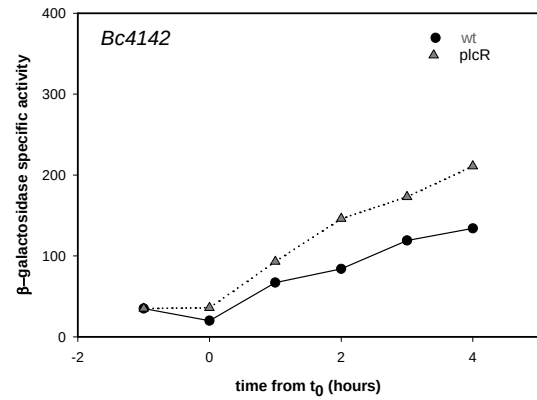
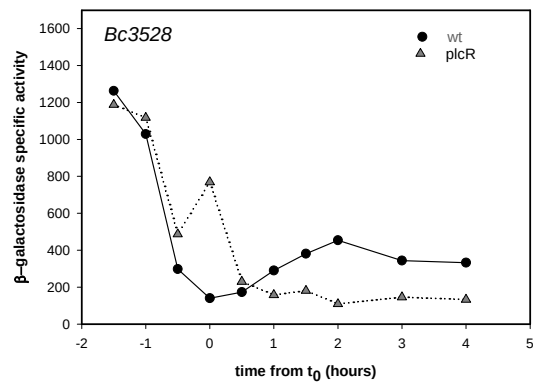
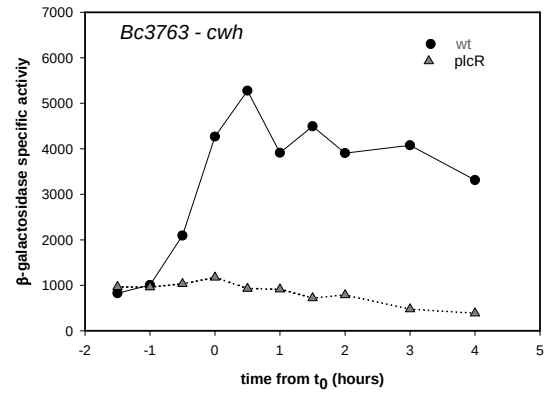
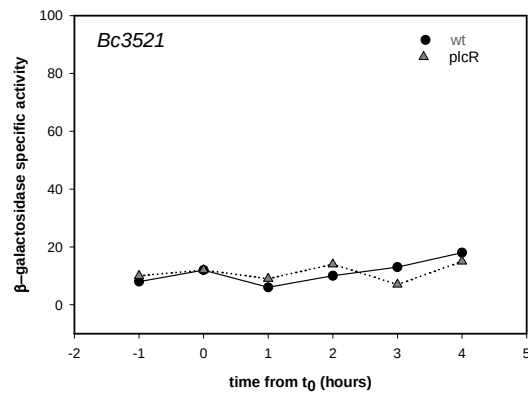
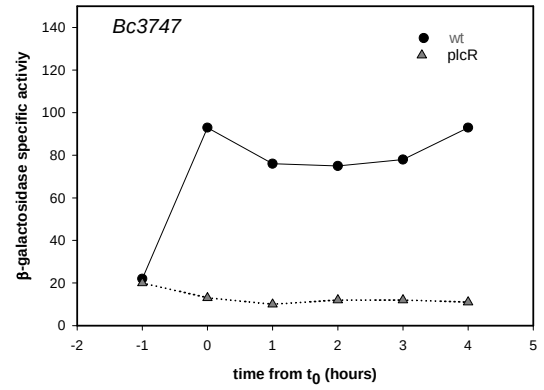
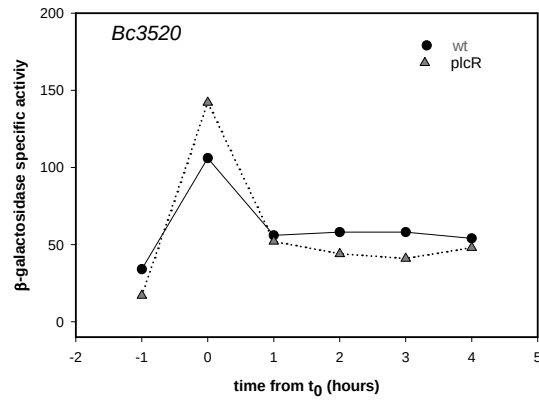








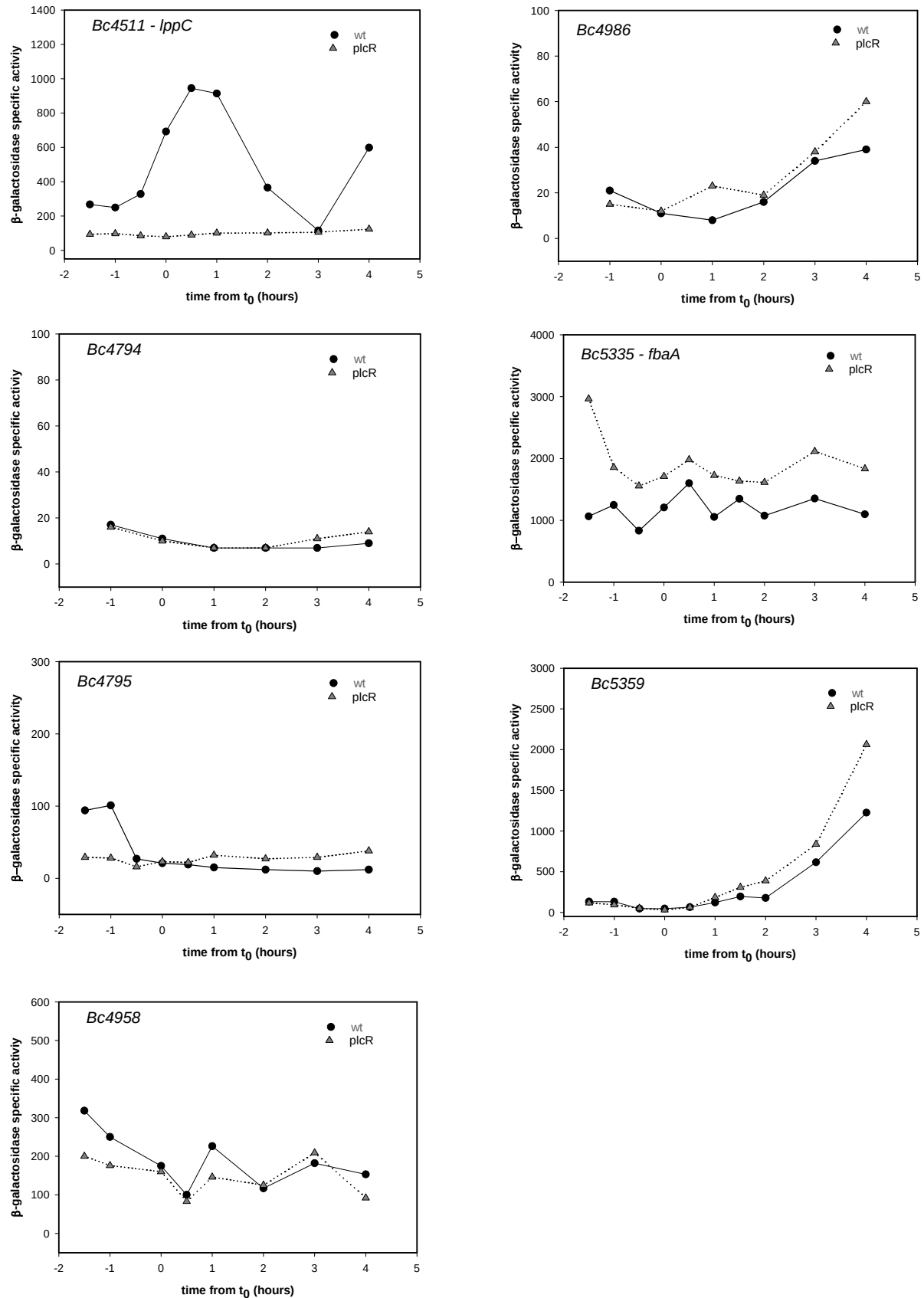


Figure S1: Results from *lacZ* fusions.

Transcriptional fusions were performed between genes promoter regions and *lacZ*. Beta-galactosidase activity is plotted as a function of time of for the ATCC14579 wild type and mutant strains.