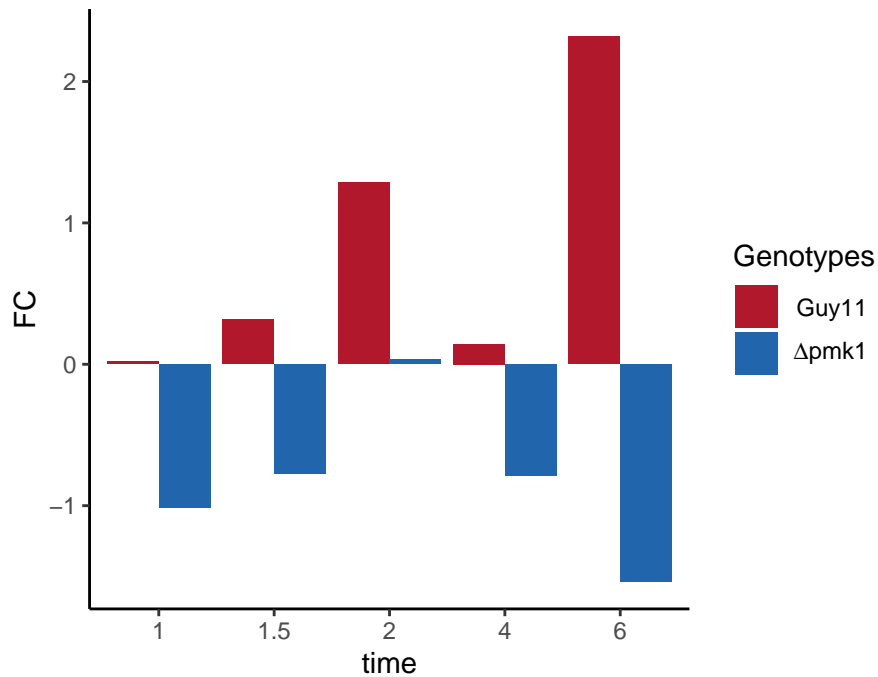
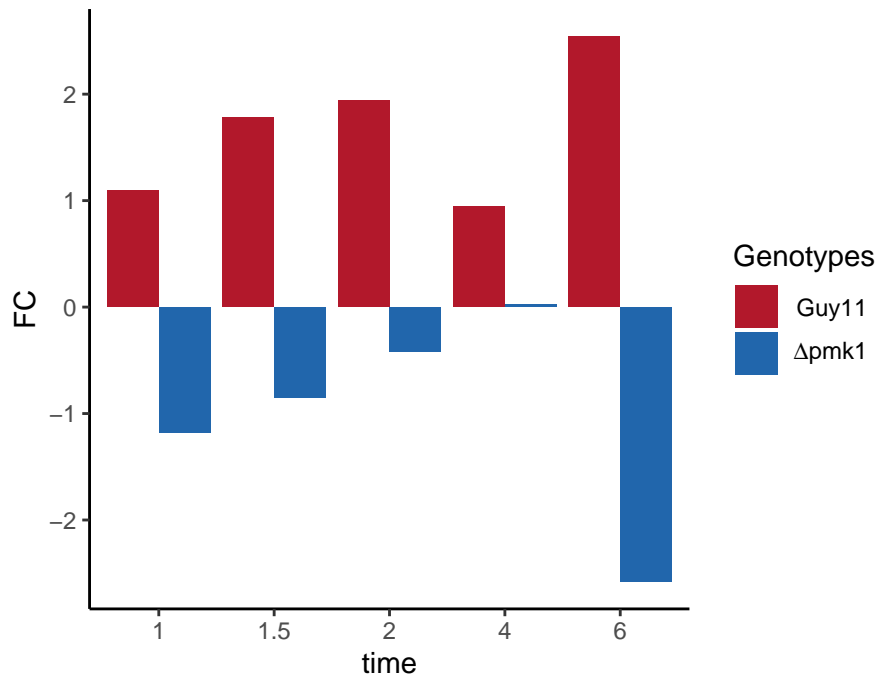


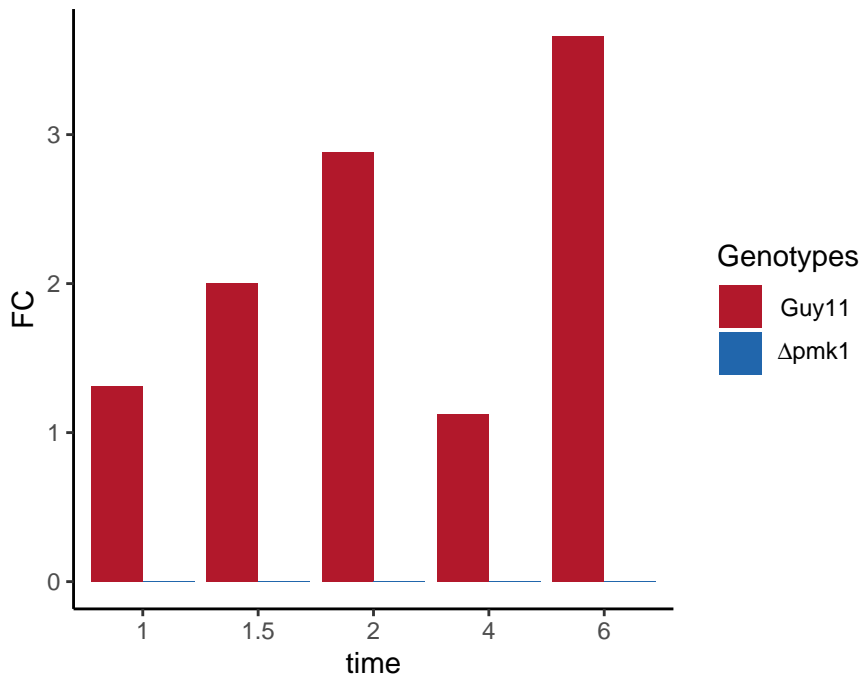
MGG_00345 RIM15 (S625 & S633)
S[+80]LILPGAVS[+80]PR



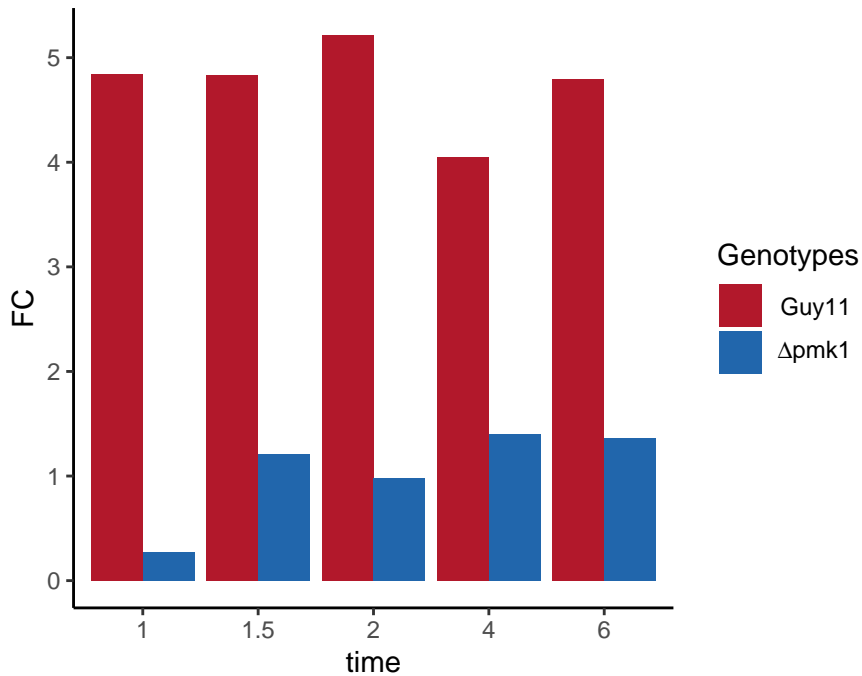
MGG_00454 ATG13 (S517 & S519)
GGPPIDIAGS[+80]AS[+80]PKPASR



MGG_00800 MST7 (S351 & S358)
DNRKS[+80]HLAPQLS[+80]PATADLLR

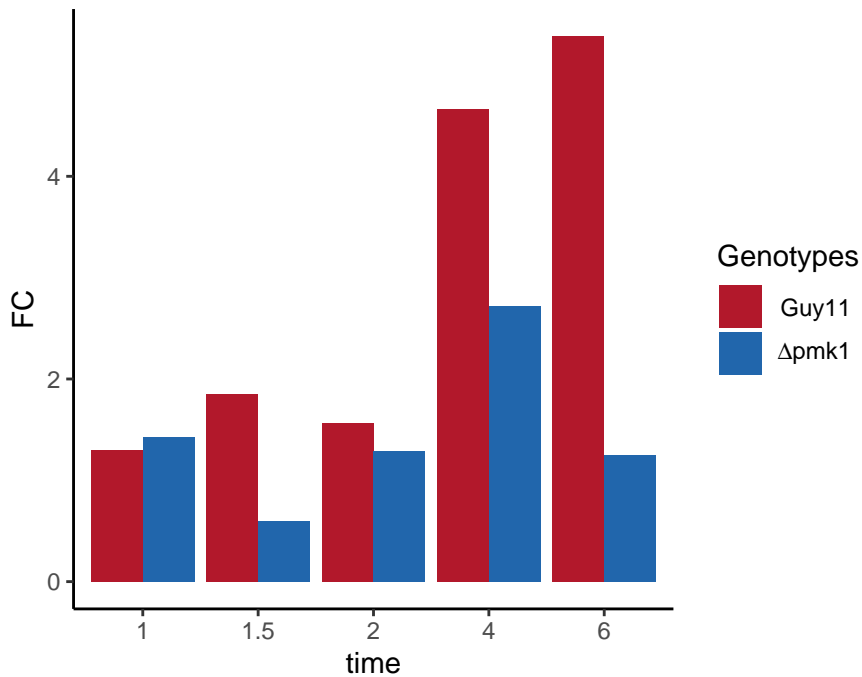


MGG_00800 MST7 (S370)
SSDS[+80]PTATYHGDDRPLETPTSAYR

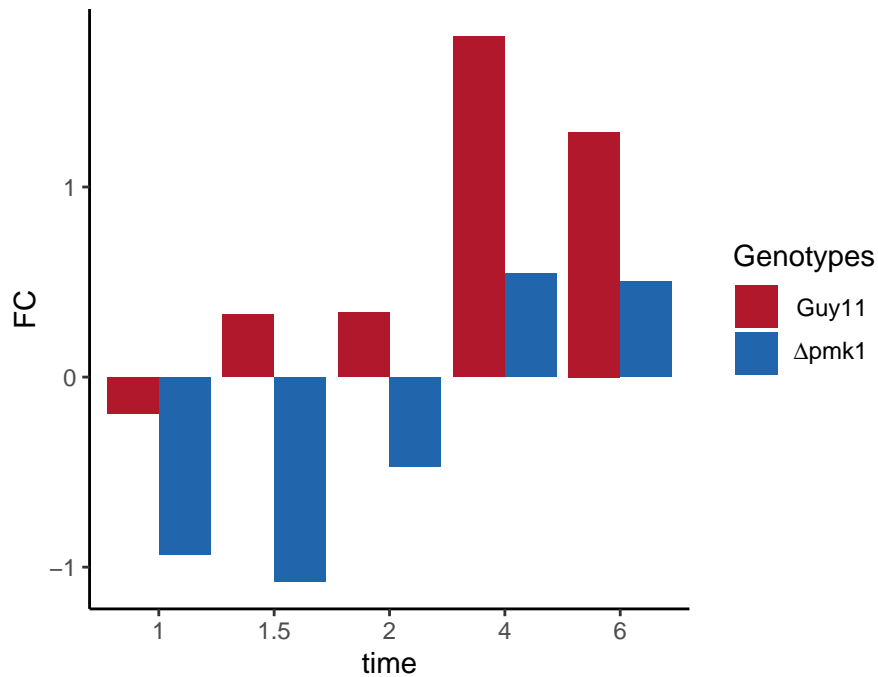


MGG_00803 SNF1 (S90)

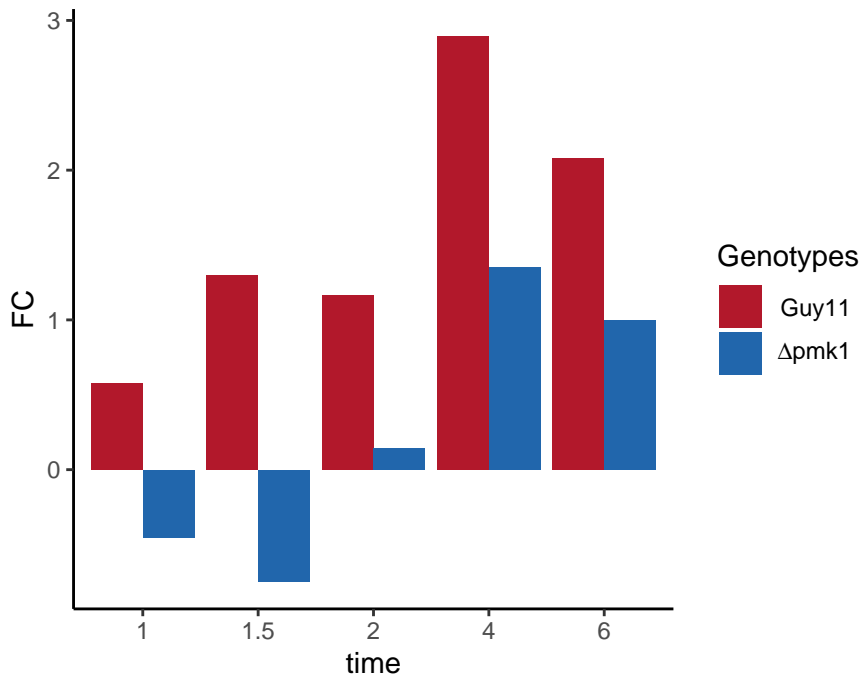
HSQHAQHQNQHQNQDQNQQS[+80]PPSAM[+16]PPTVHR



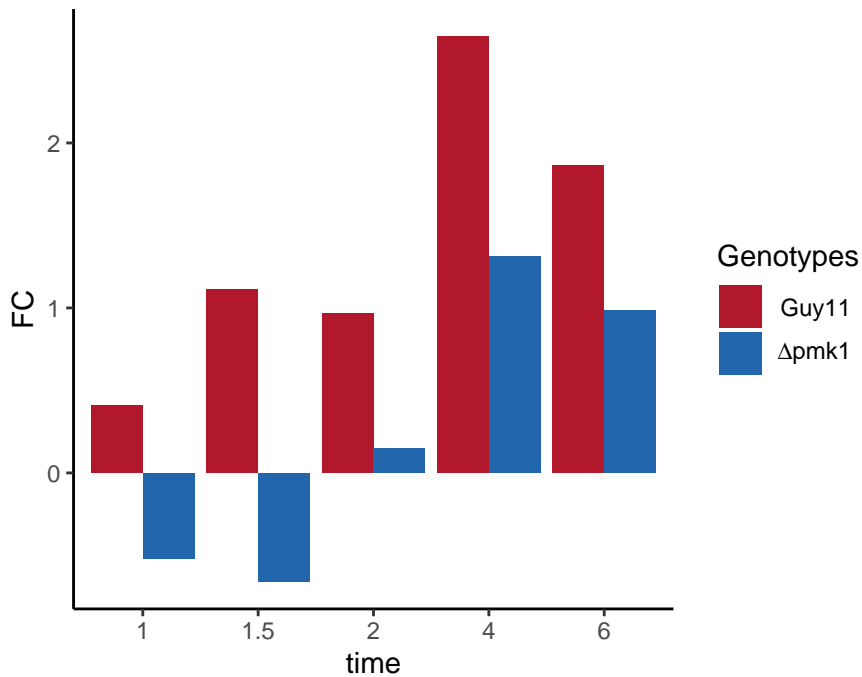
MGG_01279 KIN1 (S914)
HPSIDLQNVADNGAAS[+80]PAQTPSGAK



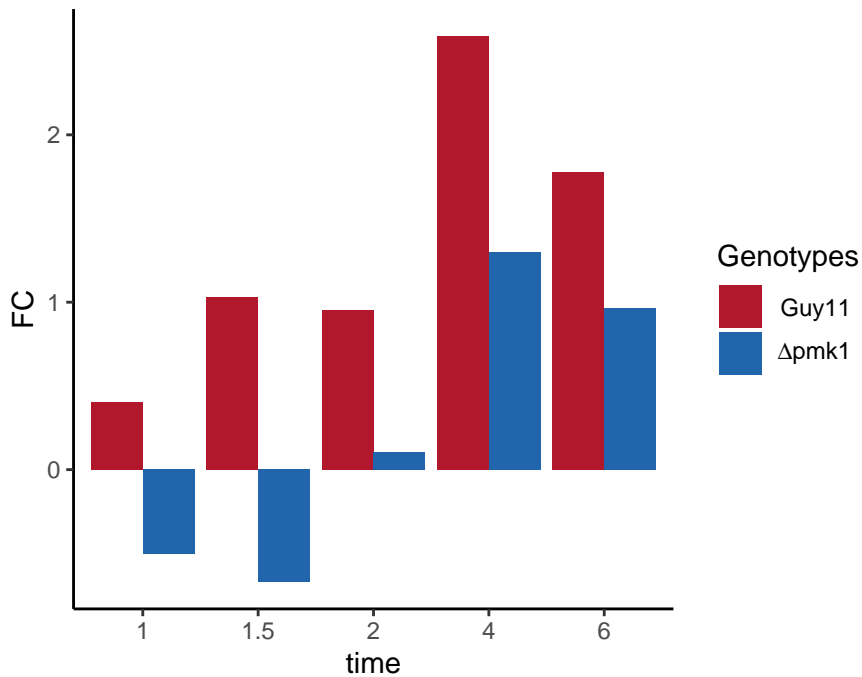
MGG_01279 KIN1 (S914)
HPSIDLQNVADNGAAS[+80]PAQTTPSGAKR



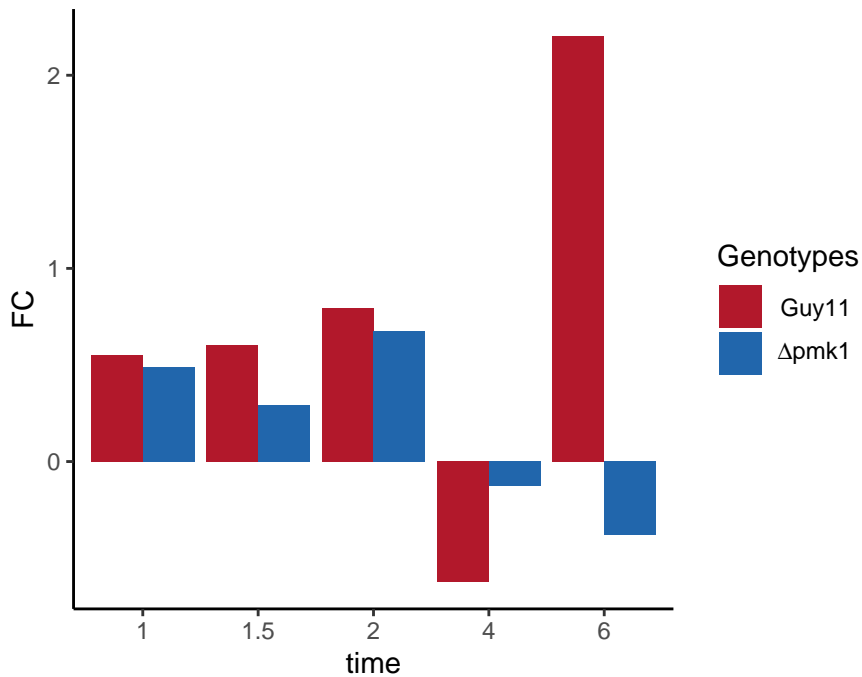
MGG_01279 KIN1 (S920)
HPSIDLQNVADNGAASPAQTPS[+80]GAKR



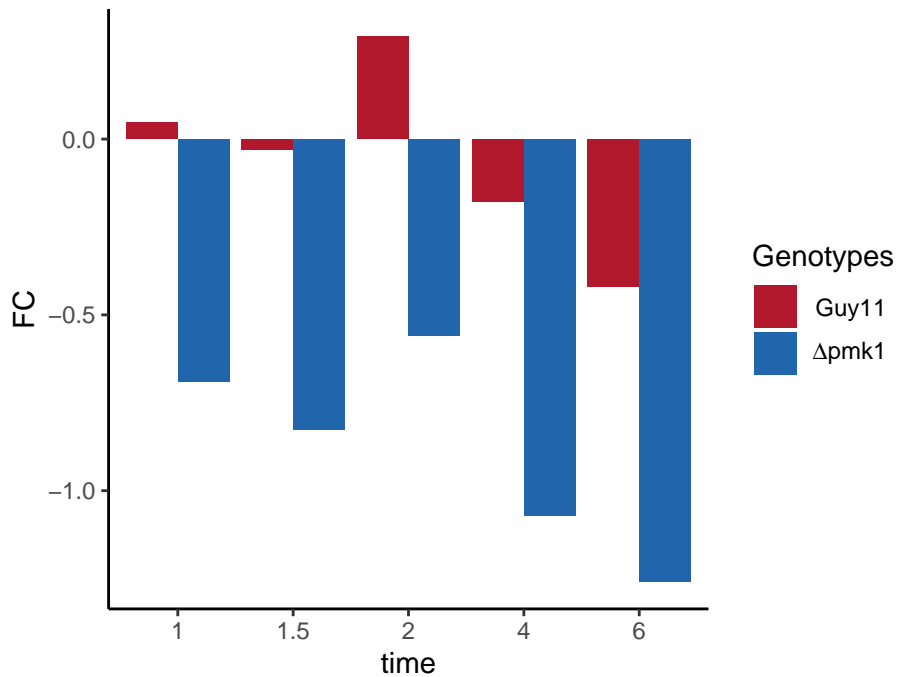
MGG_01279 KIN1 (T918)
HSPSIDLQNVADNGAASPAQT[+80]PSGAKR



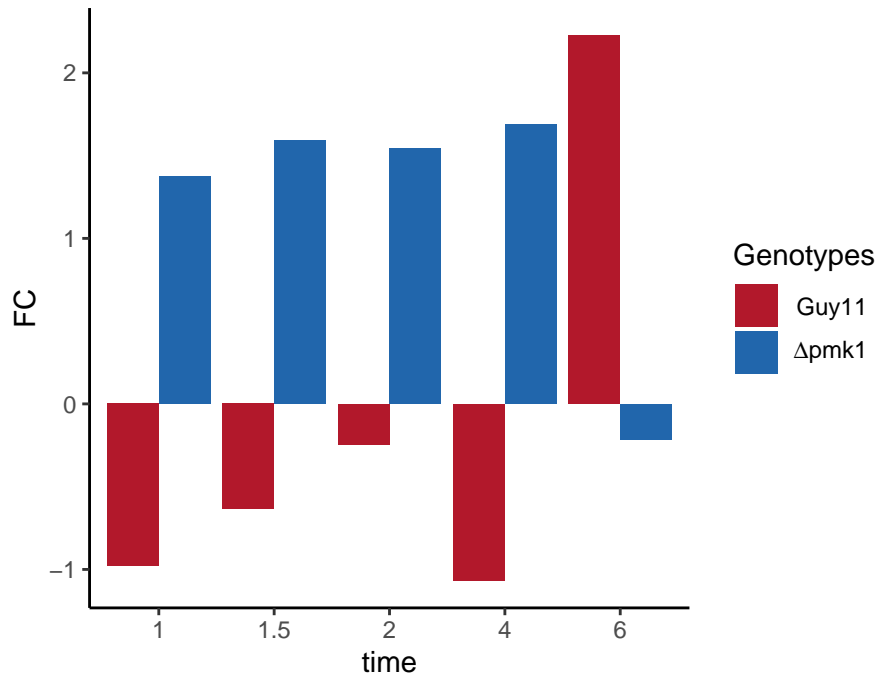
MGG_01311 (S216)
SQS[+80]PPPLTR



MGG_01376 PTP2 (S288)
NSEDVPINPPS[+80]PR

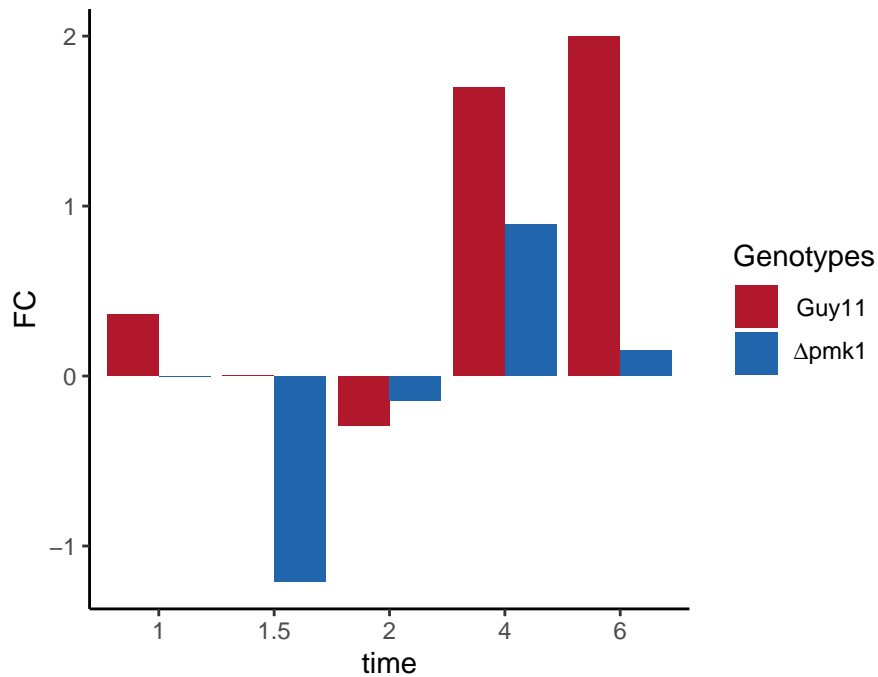


MGG_01836 FAR1 (S41)
TSGS[+80]PATANSNIPTNQPGHPSFRR

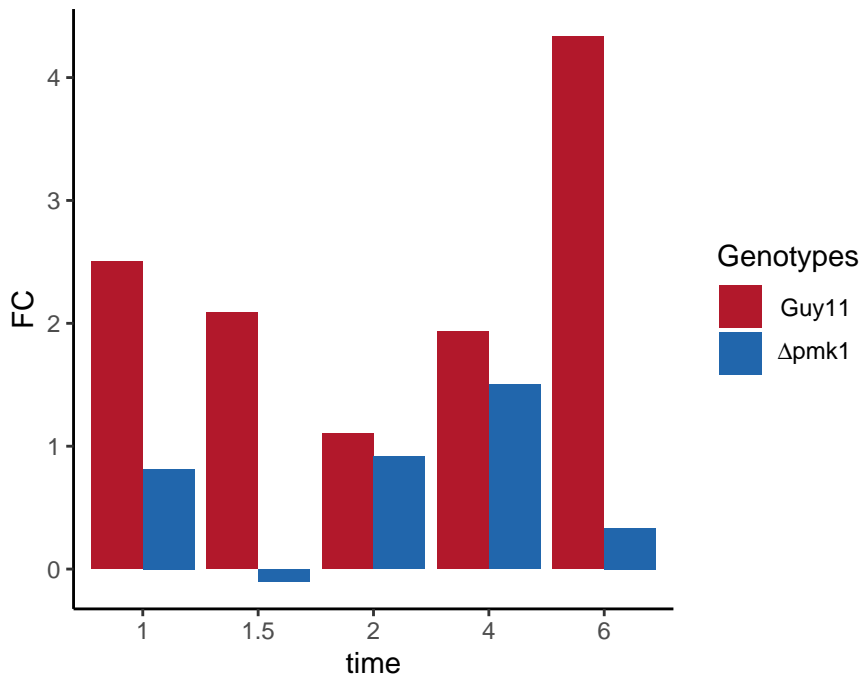


MGG_03139 ATG18 (S287)

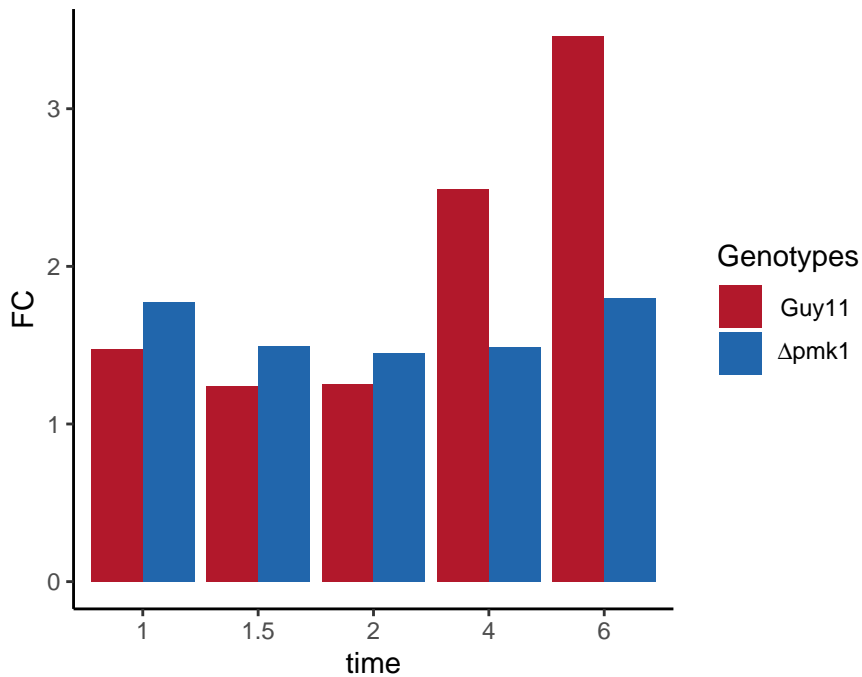
LGGPNGASGAAGAGSAGEVLAAS[+80]PGQDITGSPR



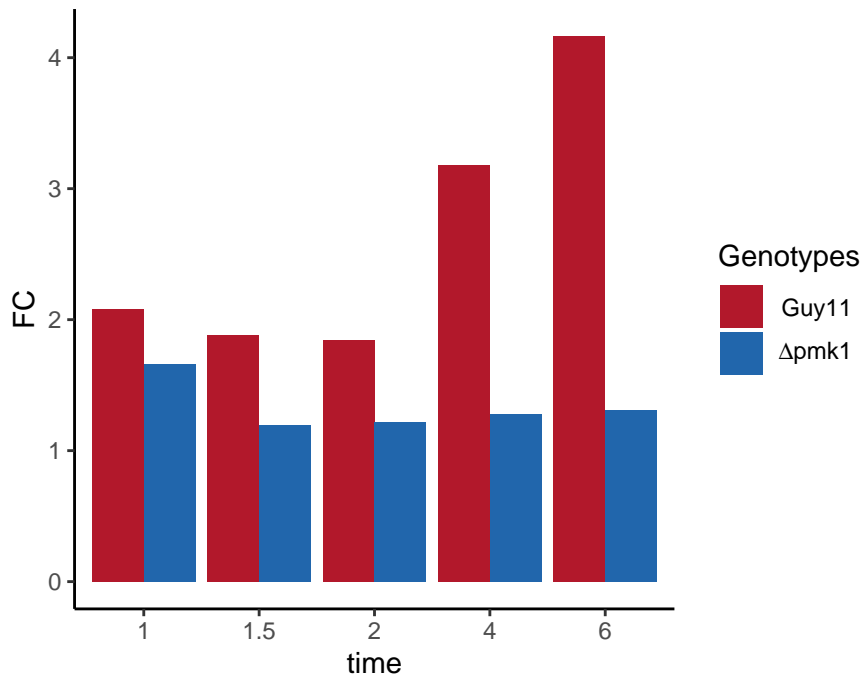
MGG_03196 RCM1 (S645)
SEQQRPTHDNRM[+16]PS[+80]PK



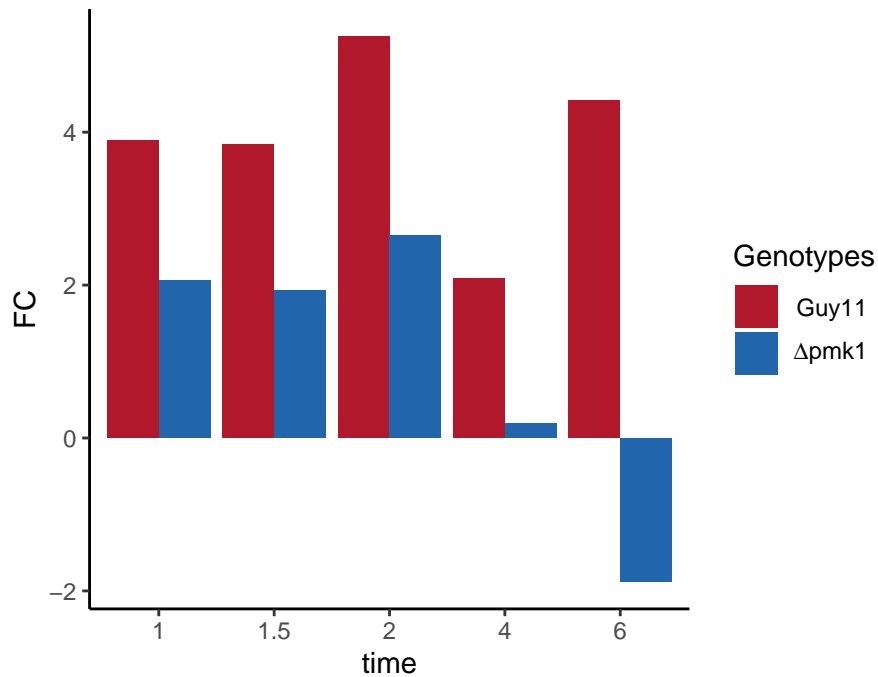
MGG_03218 (S193)
LHHNKDASGLSGVQTPIS[+80]PTGGNTR



MGG_03218 (T190)
LHHNKDASGLSGVQT[+80]PISPTGGNTR

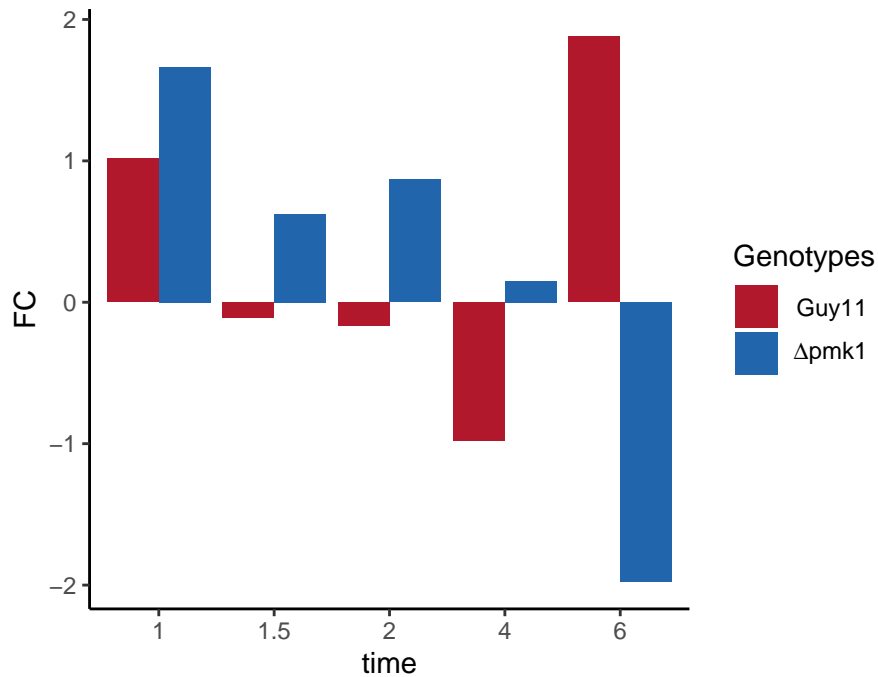


MGG_03558 (S688)
GLQVPQS[+80]PASVNSGVFQVGNR

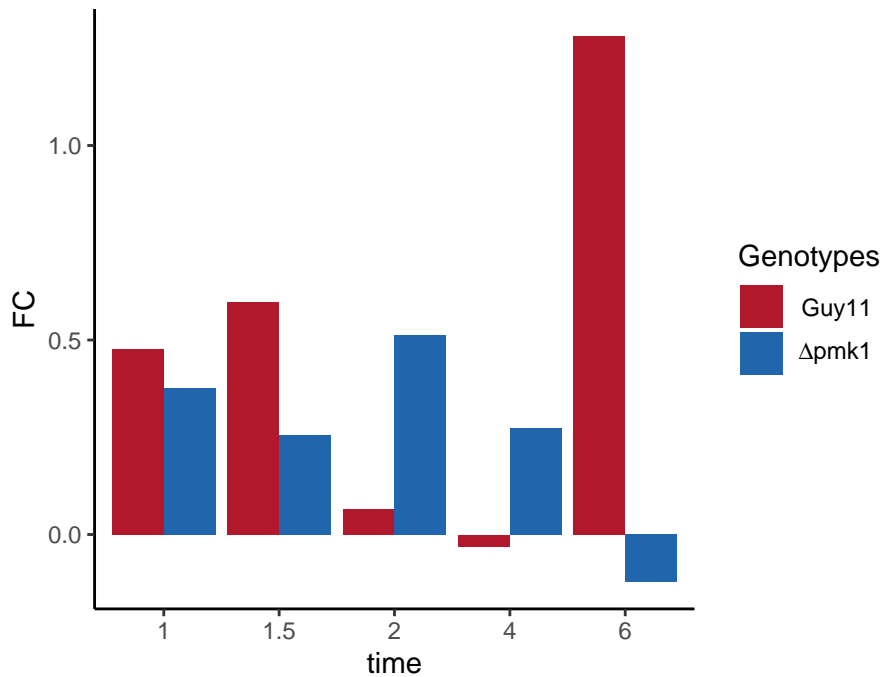


MGG_03558 (S875)

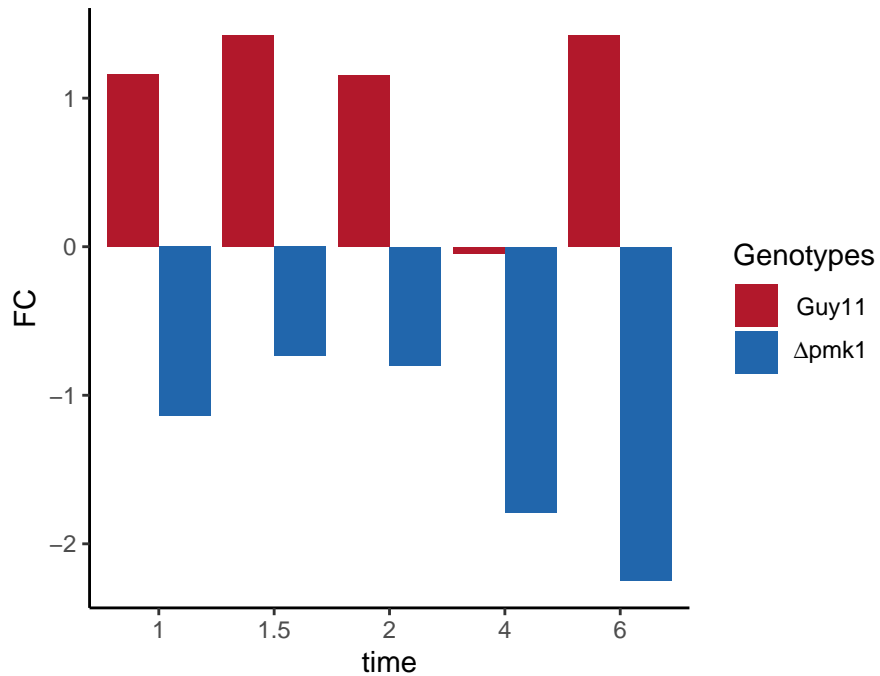
GRS[+80]PAPGVTATGNGHAGLGDDTNGRPVSGTM[+16]R



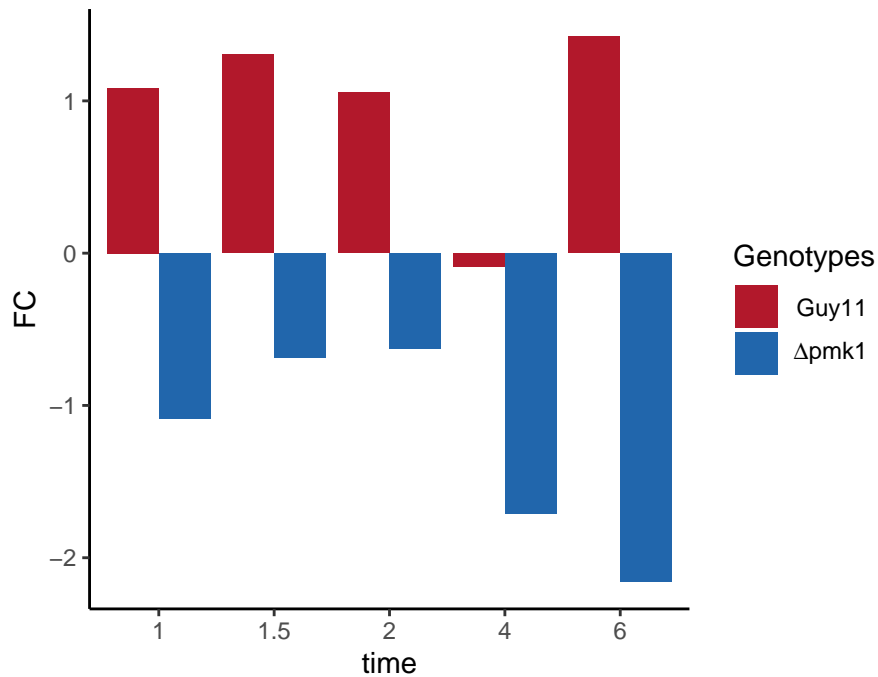
MGG_04708 SOM1 (S228)
QRPSS[+80]PSSADNAPSPSKR



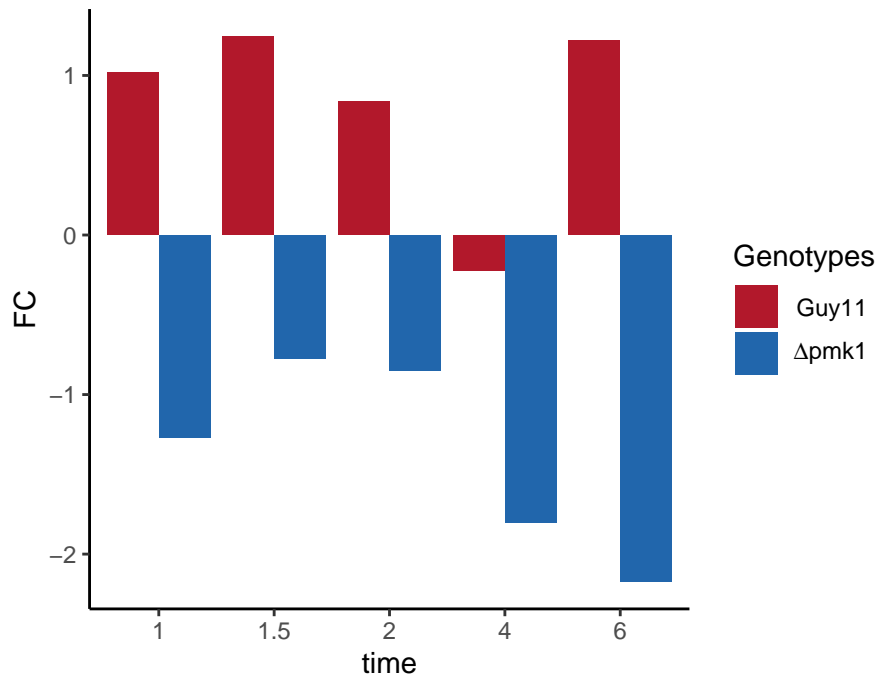
MGG_04708 SOM1 (S591 & S595)
NTTAS[+80]PQVS[+80]NAAPPTPSQANK



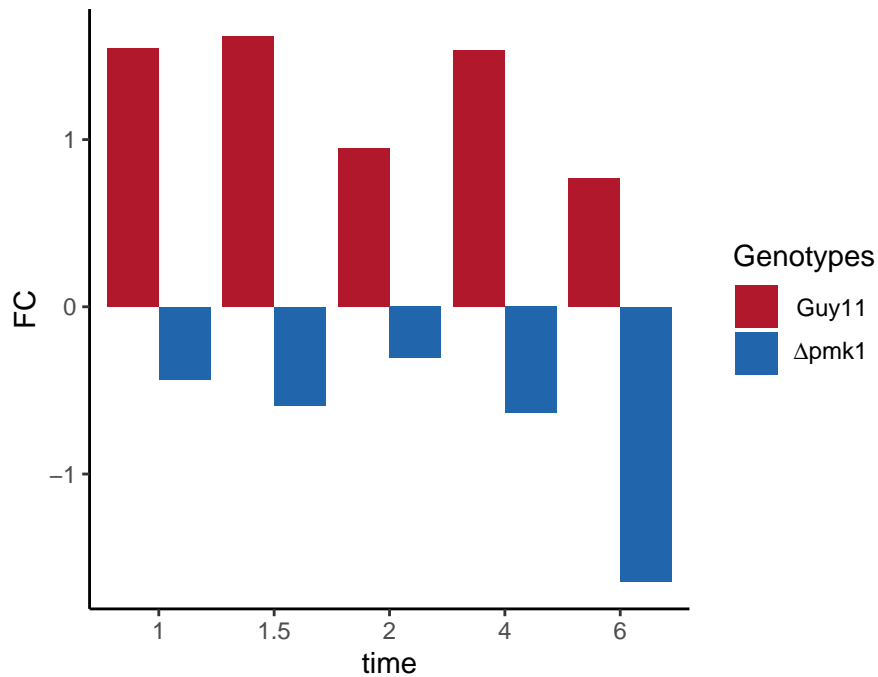
MGG_04708 SOM1 (T588 & S591)
NT[+80]TAS[+80]PQVSNAAPPTPSQANK



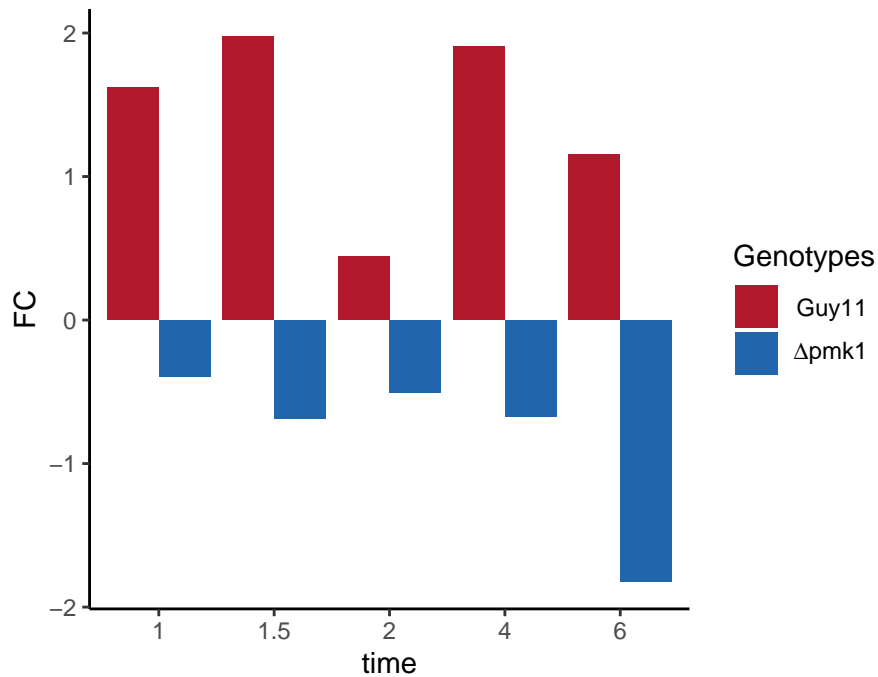
MGG_04708 SOM1 (T589 & S591)
NTT[+80]AS[+80]PQVSNAAPPTPSQANK



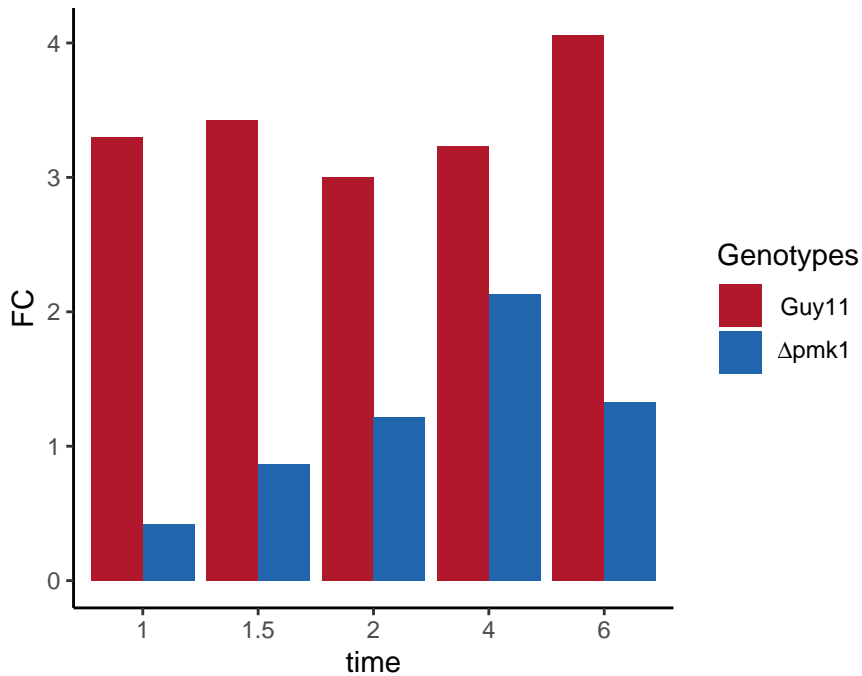
MGG_04790 CDS1 (S1174)
AYVGTKS[+80]P



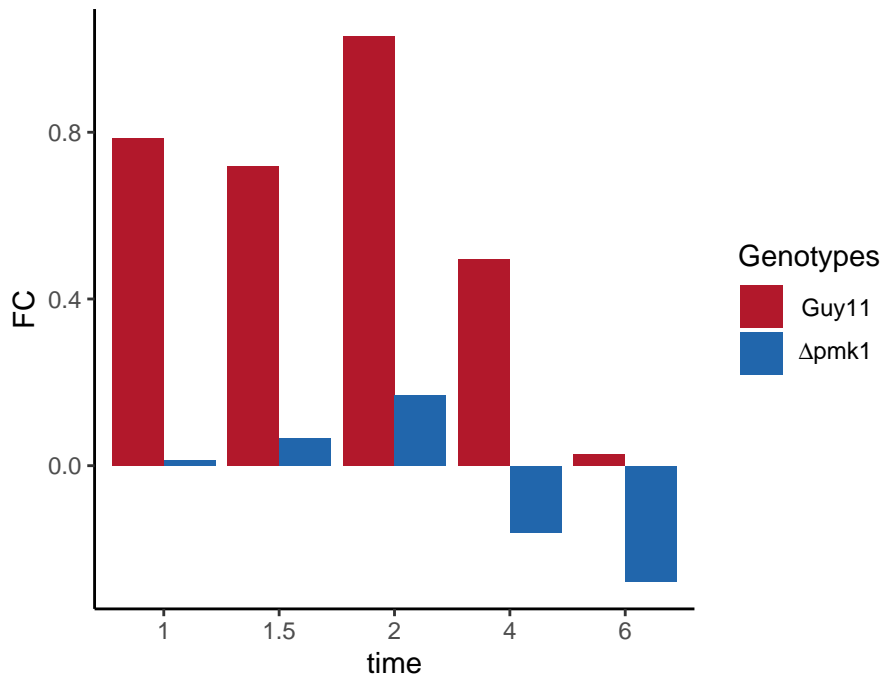
MGG_04790 CDS1 (S1174)
RAYVGTKS[+80]P



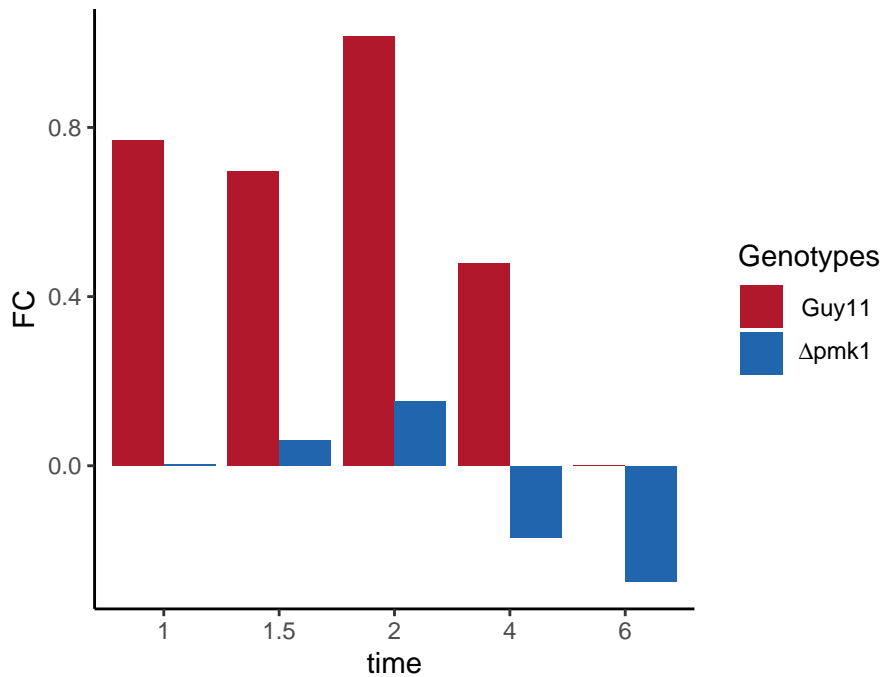
MGG_05199 MST50 (S260)
ATS[+80]PTHLQTAHDR



MGG_05220 (S267)
TSS[+80]PEPLEQSPDFPEGLPGQEPK

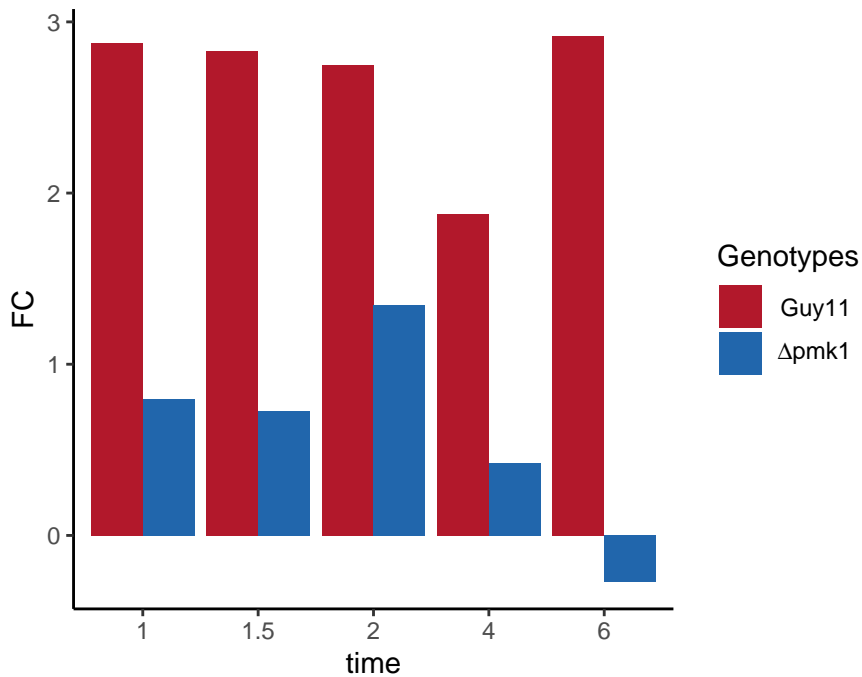


MGG_05220 (S274)
TSSPEPLEQS[+80]PDFPEGLPGQEPK

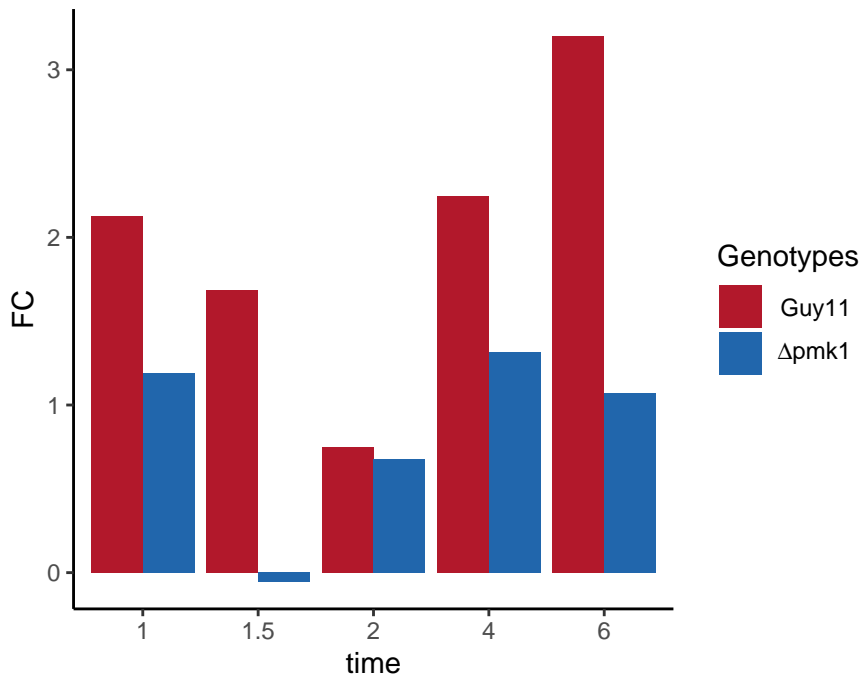


MGG_05220 (S320)

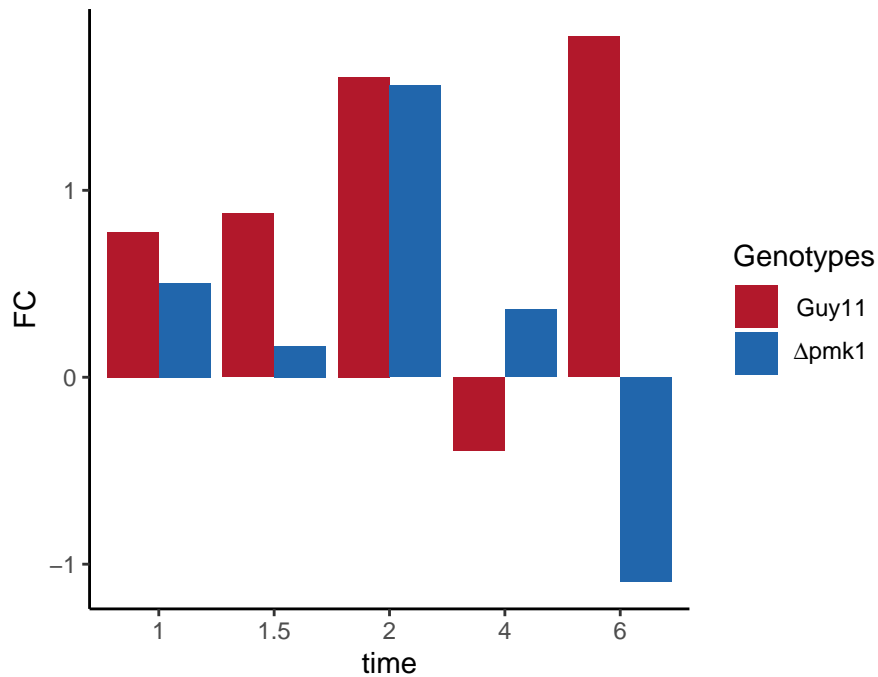
GTSDSAESSRTPTNESGASAQTAQTS[+80]PPISR



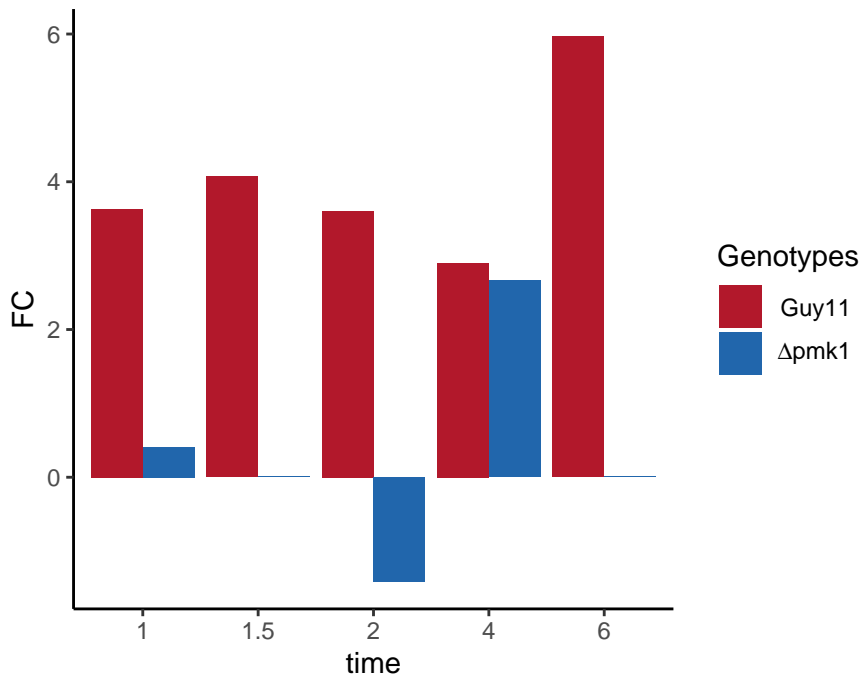
MGG_06258 FKH1 (S491)
RVS[+80]PSGSPQPR



MGG_06258 FKH1 (S641)
YADIGSTPM[+16]RPGALDFS[+80]PSR

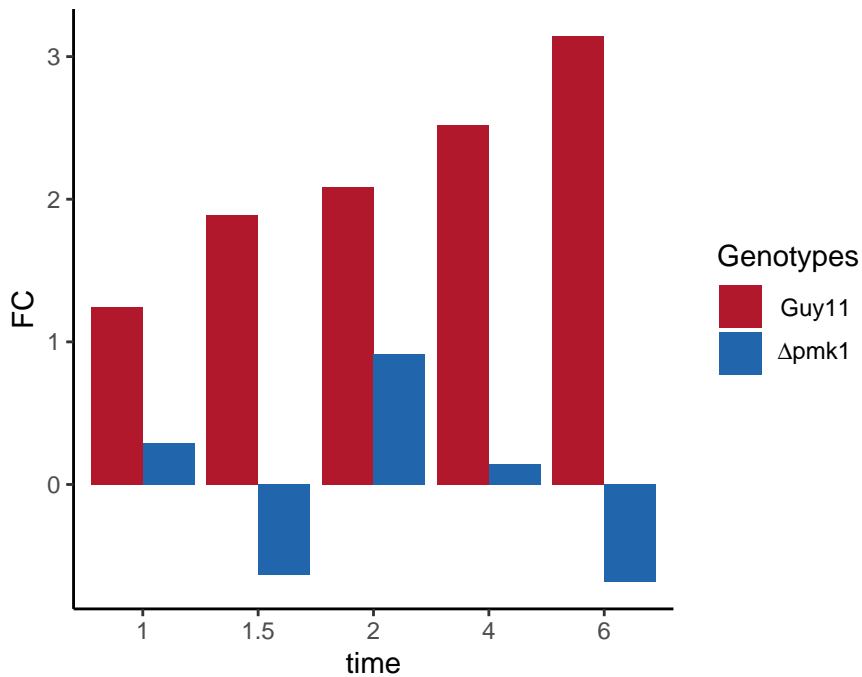


MGG_06334 VTS1 (S175)
NHPM[+16]SGVLS[+80]PANFDKDPM[+16]SNR

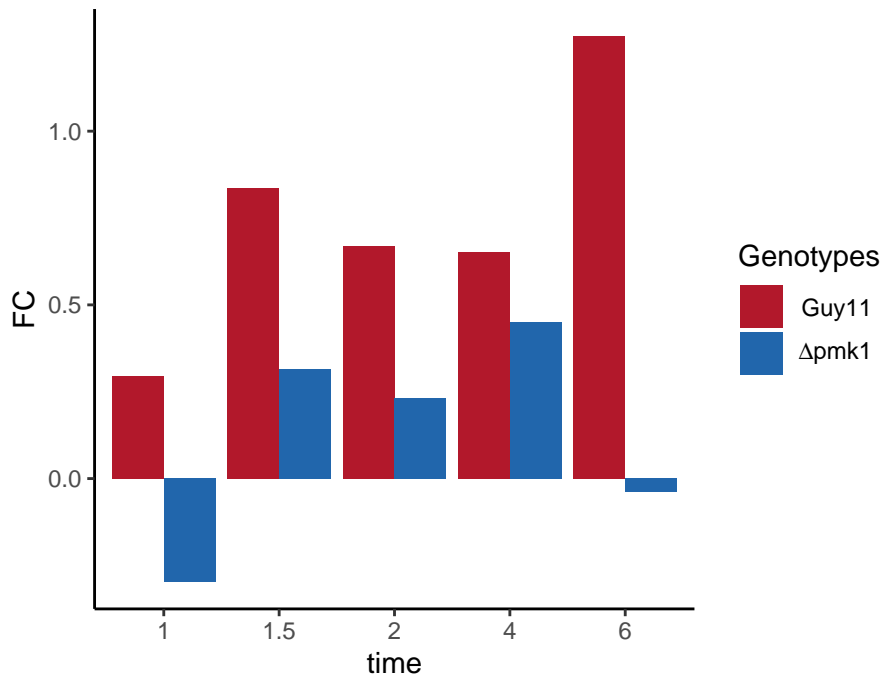


MGG_06334 VTS1 (S420)

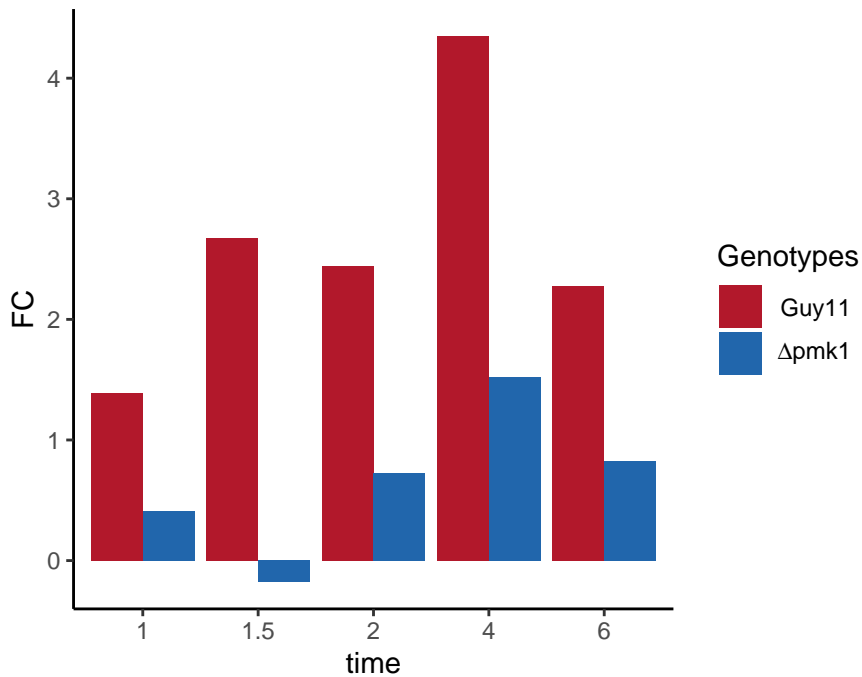
SNDNSGQNPLS[+80]PGM[+16]ISPNVVM[+16]VDEHGR



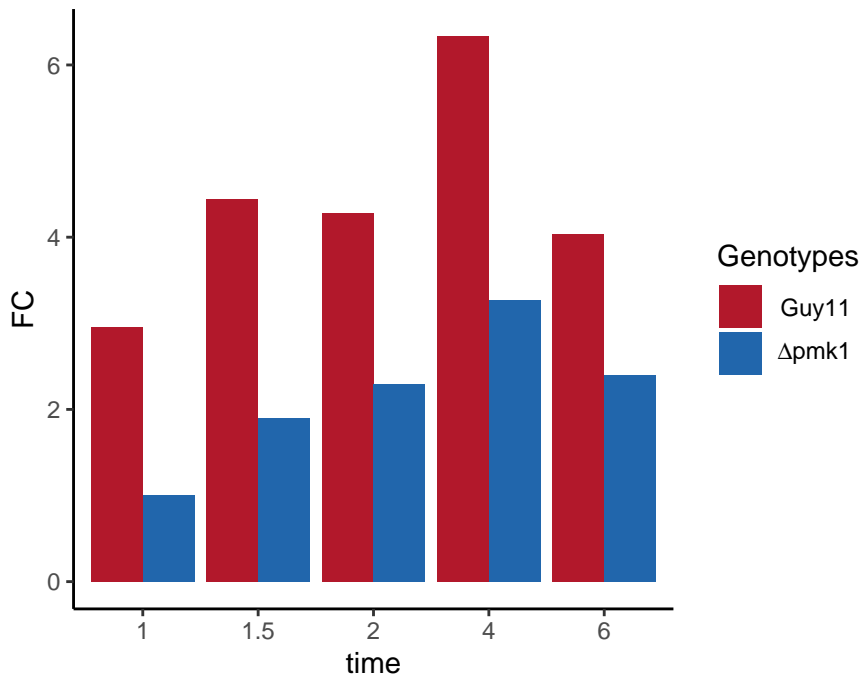
MGG_06393 ATG1 (S476)
SNRESSS[+80]PTSAALANDSAR



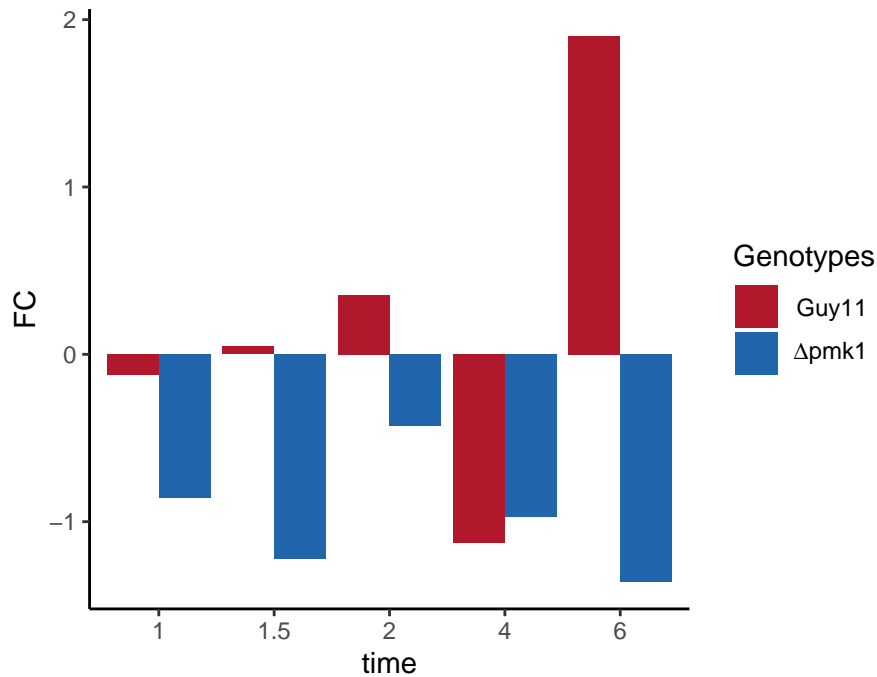
MGG_06403 (S272)
SAESVNQQS[+80]PSSPPDRPNLGGR



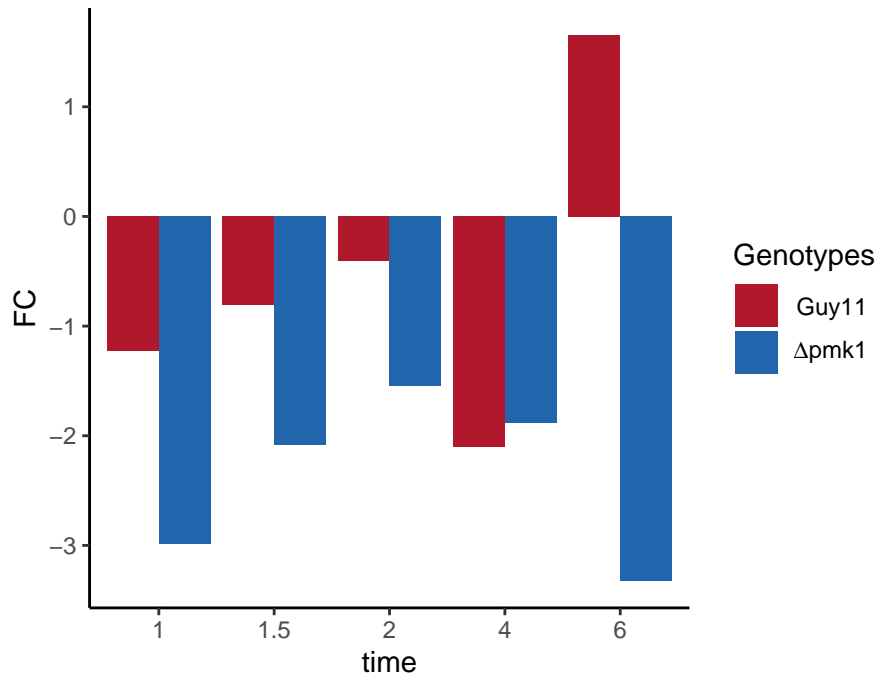
MGG_06403 (S275)
SAESVNQQSPSS[+80]PPDRPNLGGR



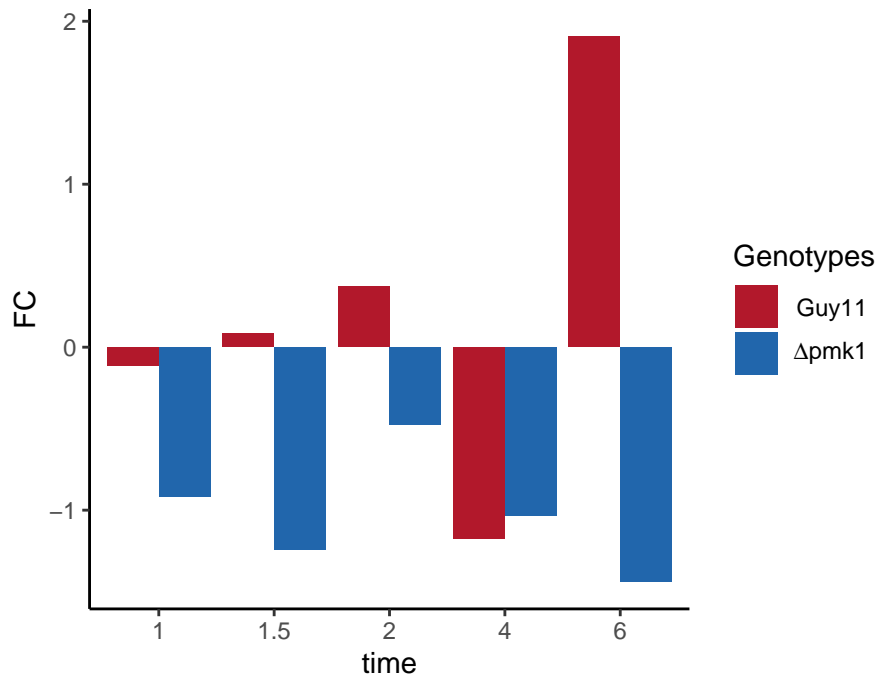
MGG_06726 SEP4p (S313 & S318)
QLLALKDPSAQGHS[+80]SRPIS[+80]PAAER



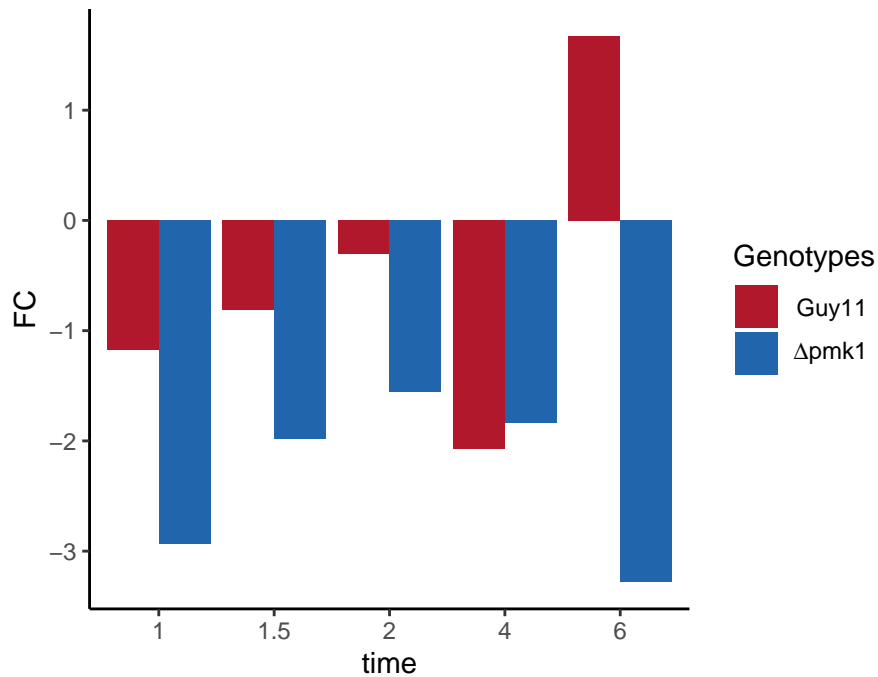
MGG_06726 SEP4p (S313 & S318)
TKQLLALKDPSAQGHS[+80]SRPIS[+80]PAAER



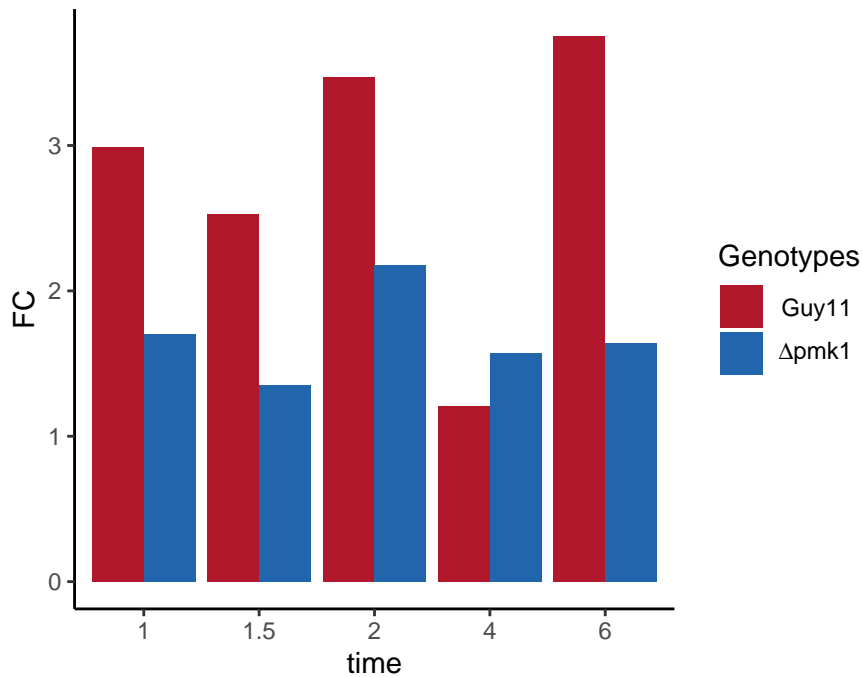
MGG_06726 SEP4p (S314 & S318)
QLLALKDPSAQGHSS[+80]RPIS[+80]PAAER



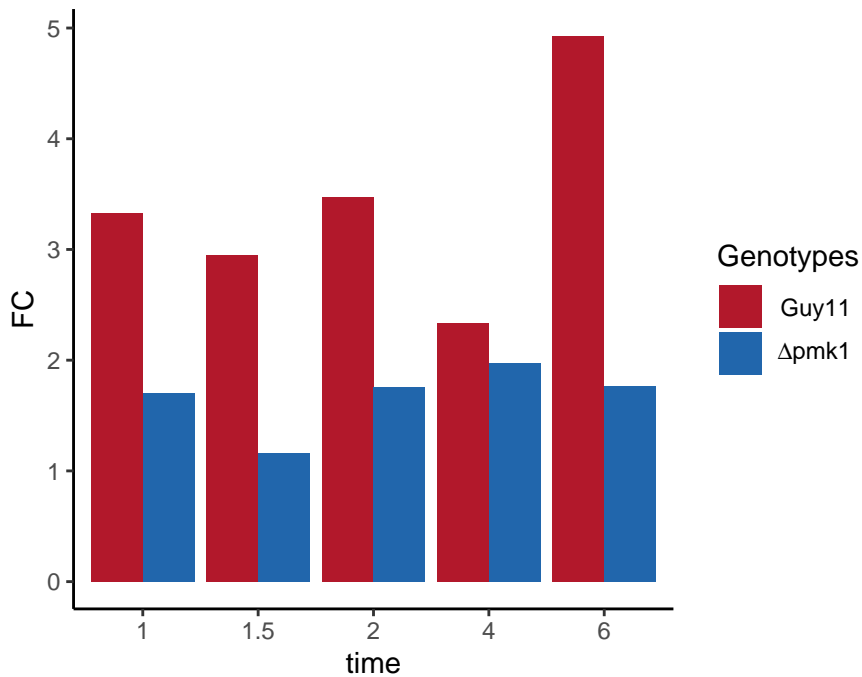
MGG_06726 SEP4p (S314 & S318)
TKQLLALKDPSAQGHSS[+80]RPIS[+80]PAAER



MGG_06726 SEP4p (S318)
DPSAQGHSSRPIS[+80]PAAER

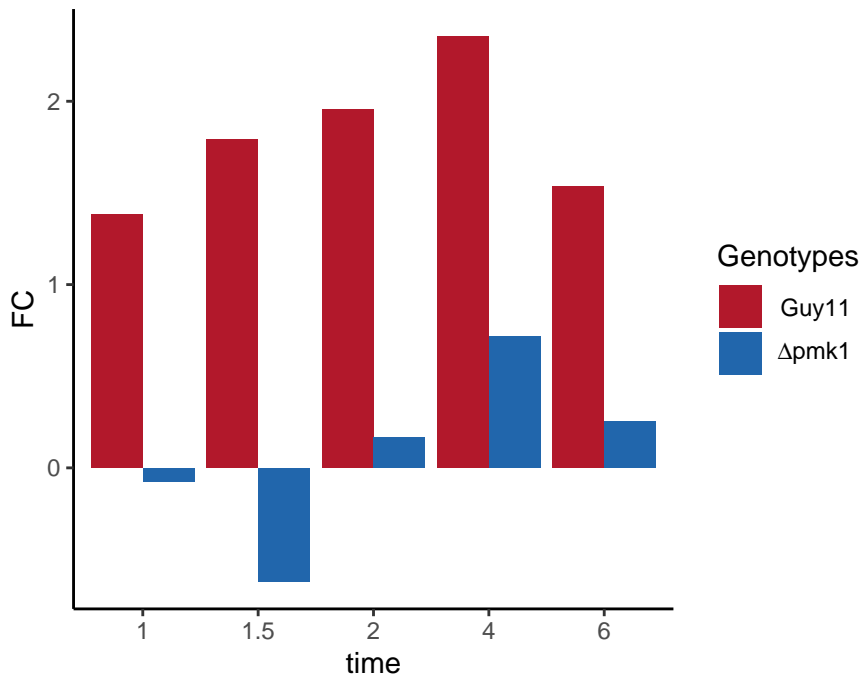


MGG_06726 SEP4p (S318)
QLLALKDPSAQGHSSRPIS[+80]PAAER

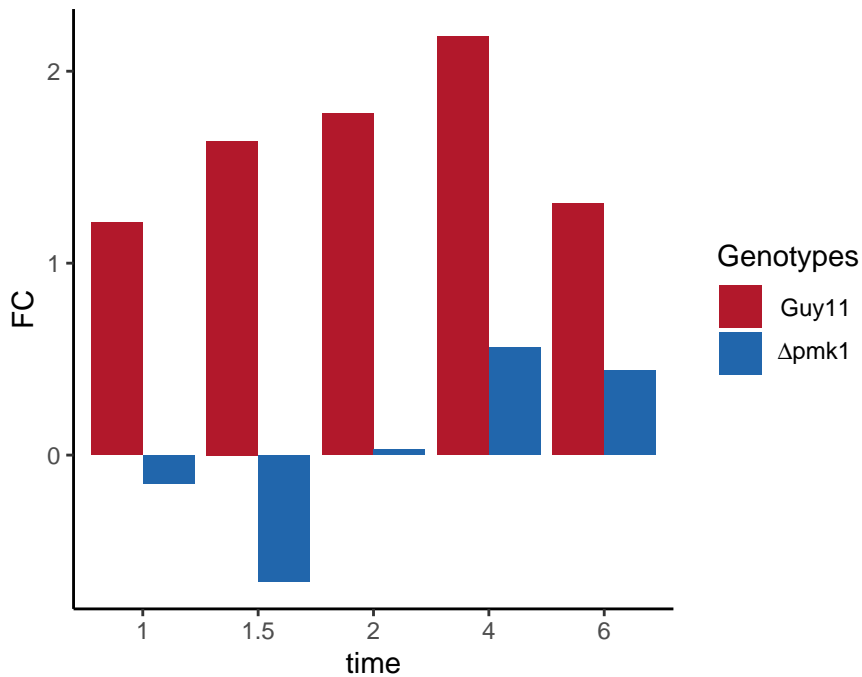


MGG_07667 ATG17 (S207 & S209)

HLLAAPGLPS[+80]PS[+80]NSVFSRPM[+16]ADLVR

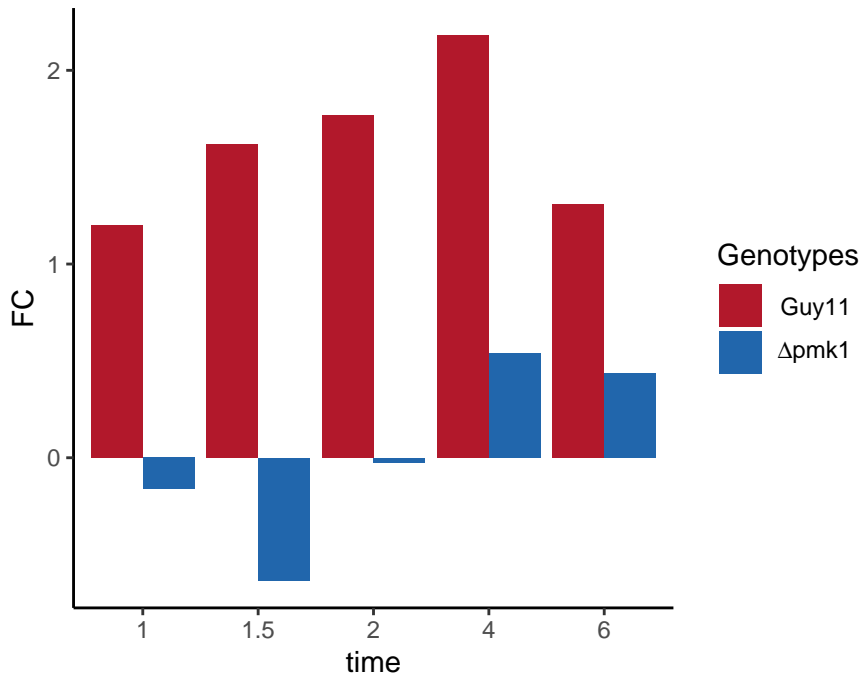


MGG_07667 ATG17 (S207 & S211)
HLLAAPGLPS[+80]PSNS[+80]VFSRPM[+16]ADLVR

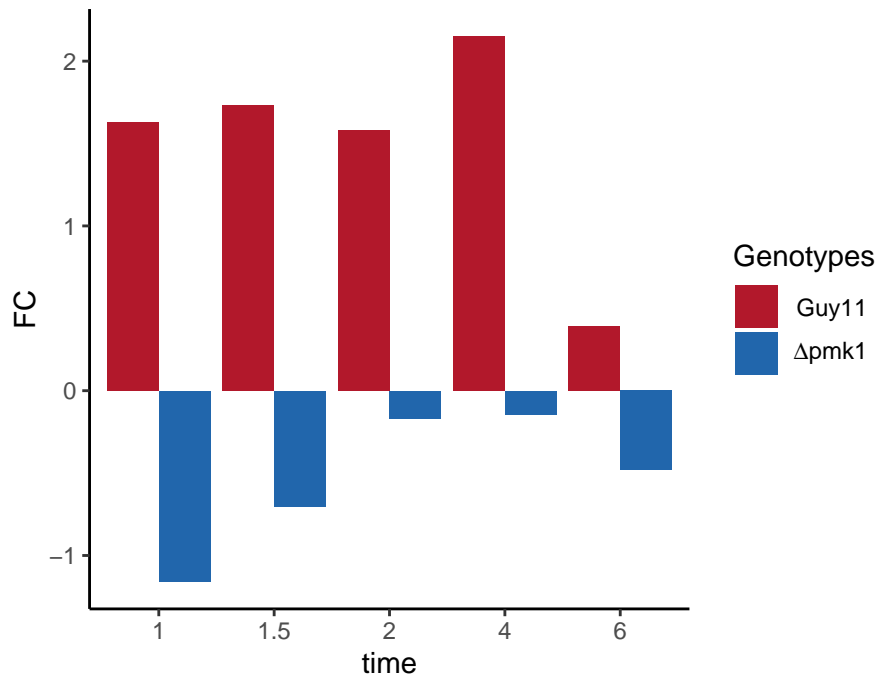


MGG_07667 ATG17 (S207 & S214)

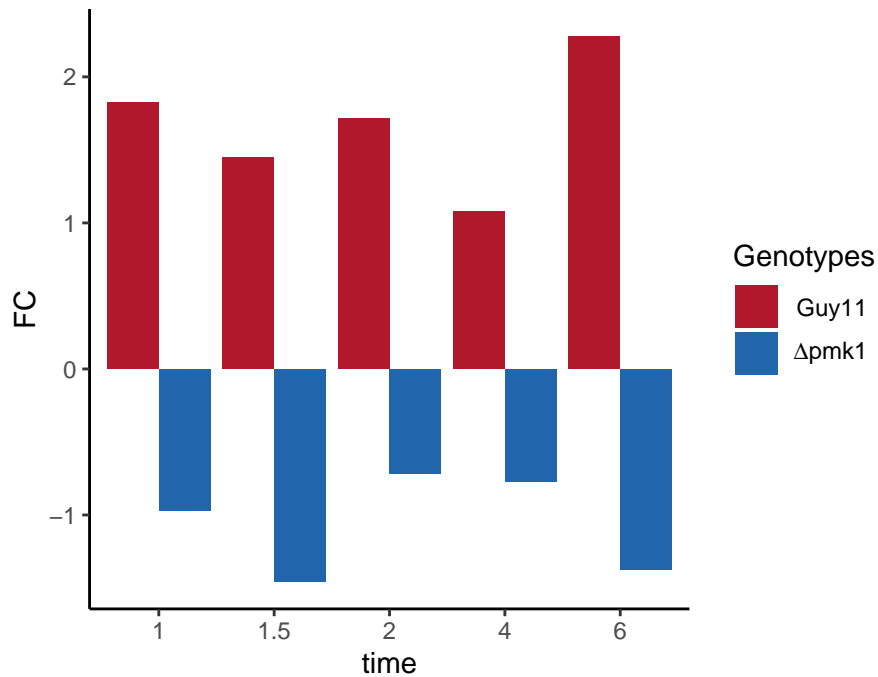
HLLAAPGLPS[+80]PSNSVFS[+80]RPM[+16]ADLVR



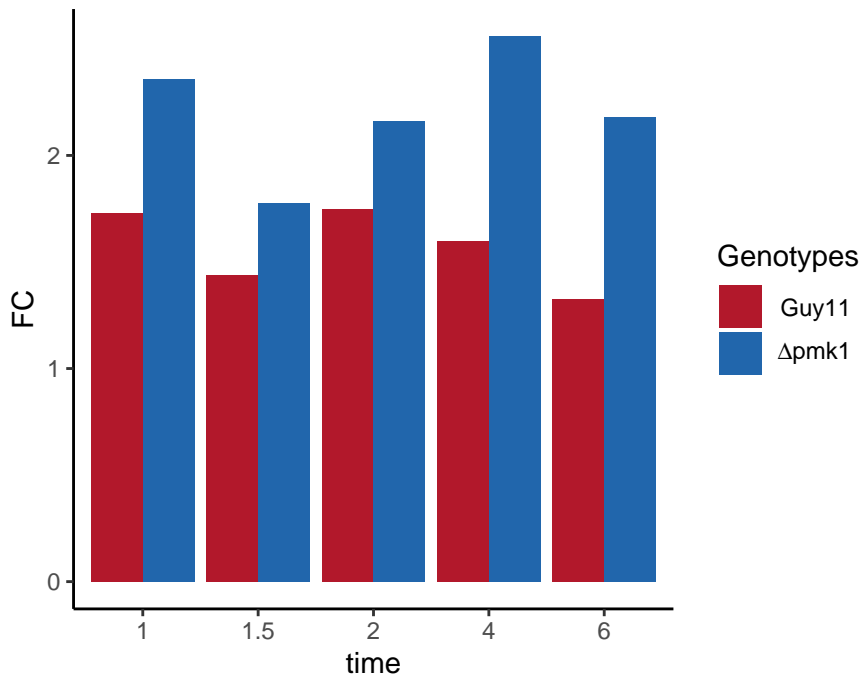
MGG_07714 (S258)
SGGGTPES[+80]PPSSPAIYPR



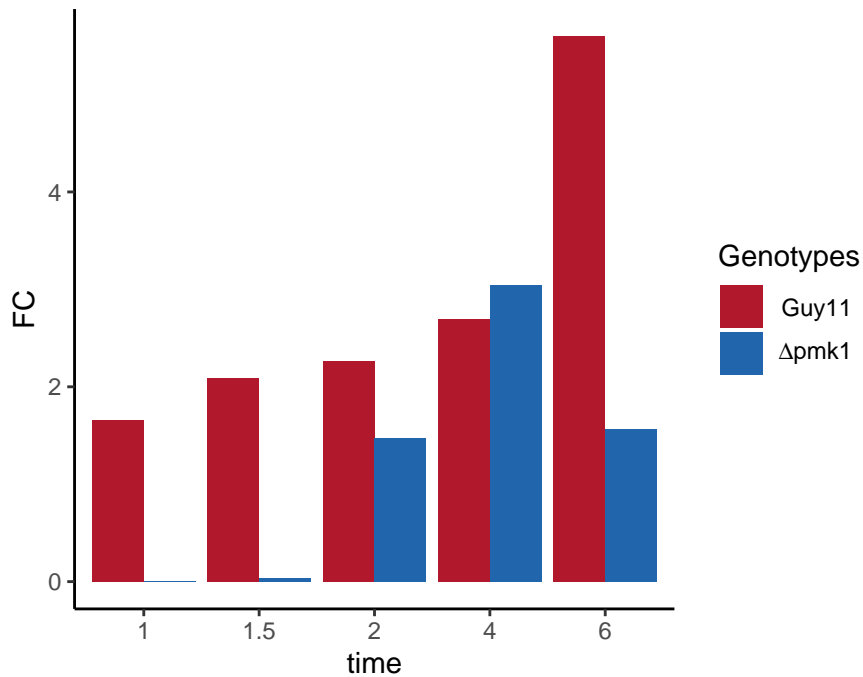
MGG_08061 (S399)
GVS[+80]PSGLEFLSLNPER



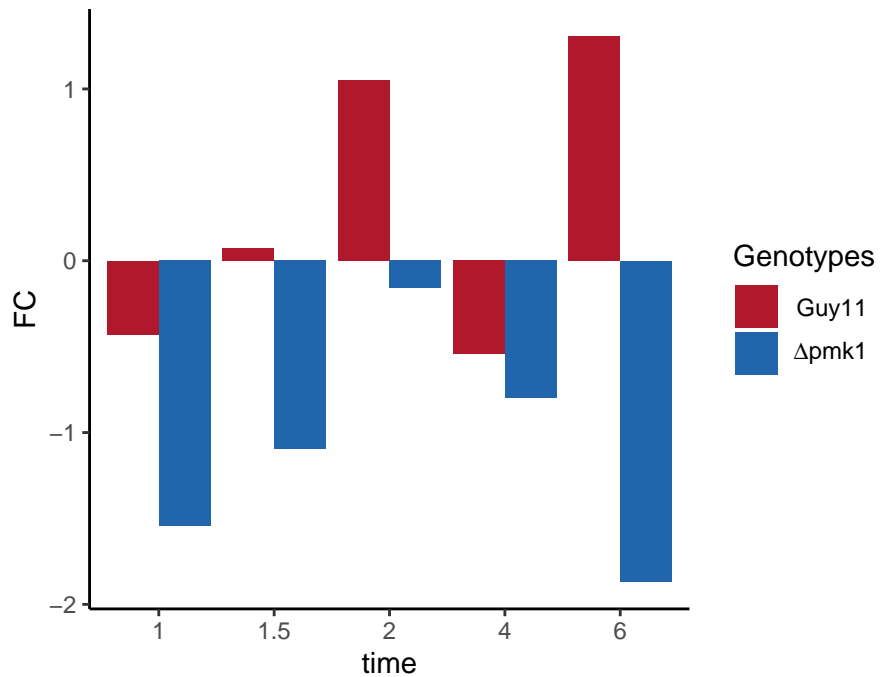
MGG_08061 (S399)
GVS[+80]PSGLEFLSLNPERR



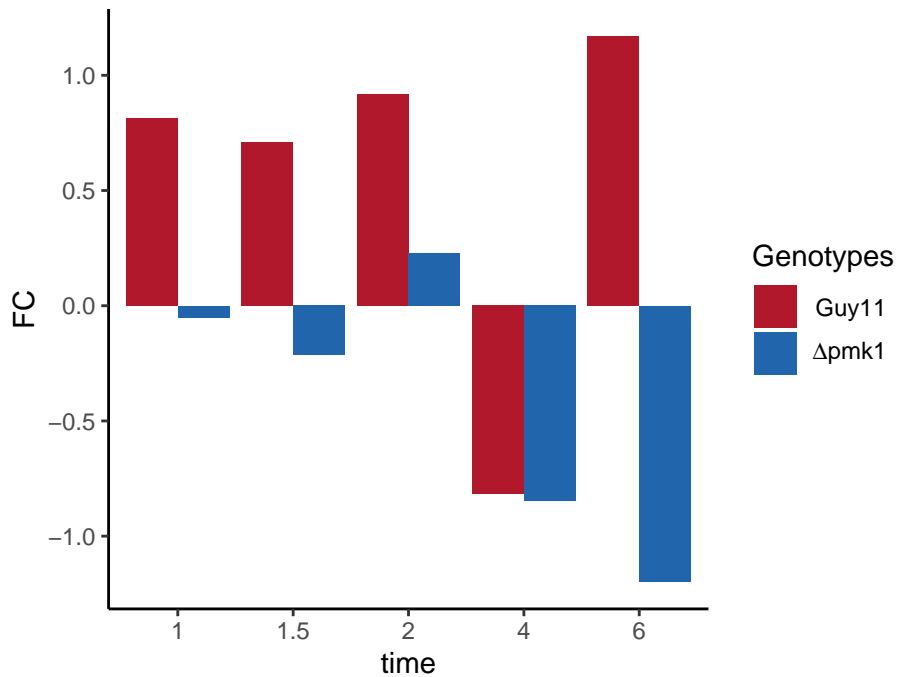
MGG_08097 YCK1 (S359)
LTVDRNLNAAQPPPPS[+80]PIKANQGK



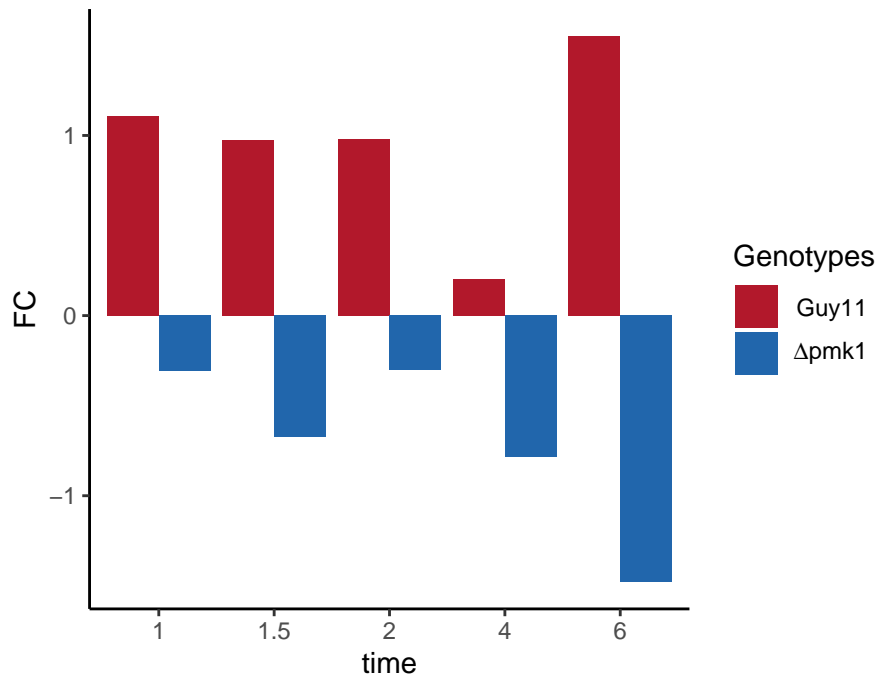
MGG_09293 (S125 & Y129)
ASGSGLIGESS[+80]PIRY[+80]ESSSPGR



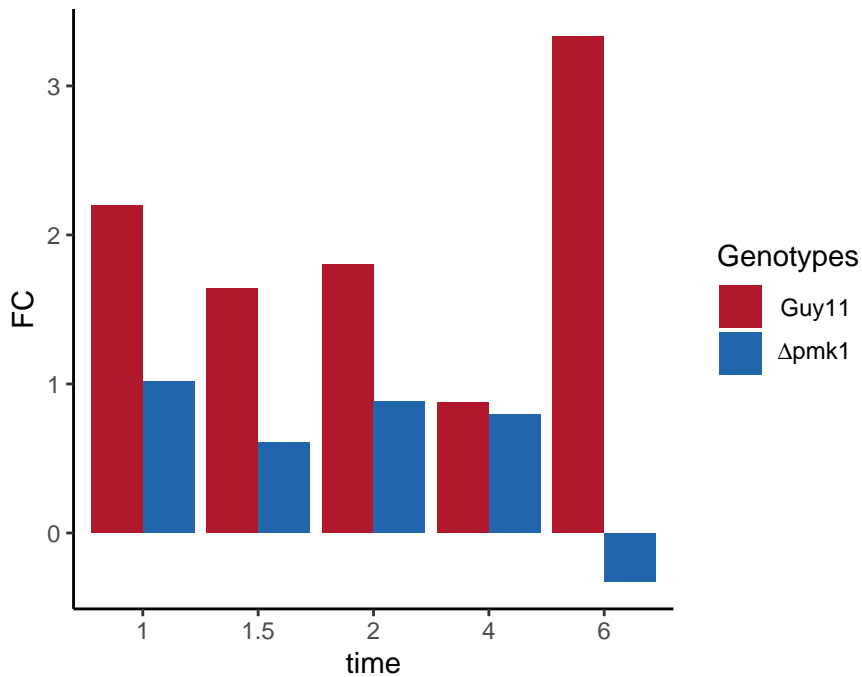
MGG_09293 (S133)
ASGSGLIGESSPIRYESSS[+80]PGR



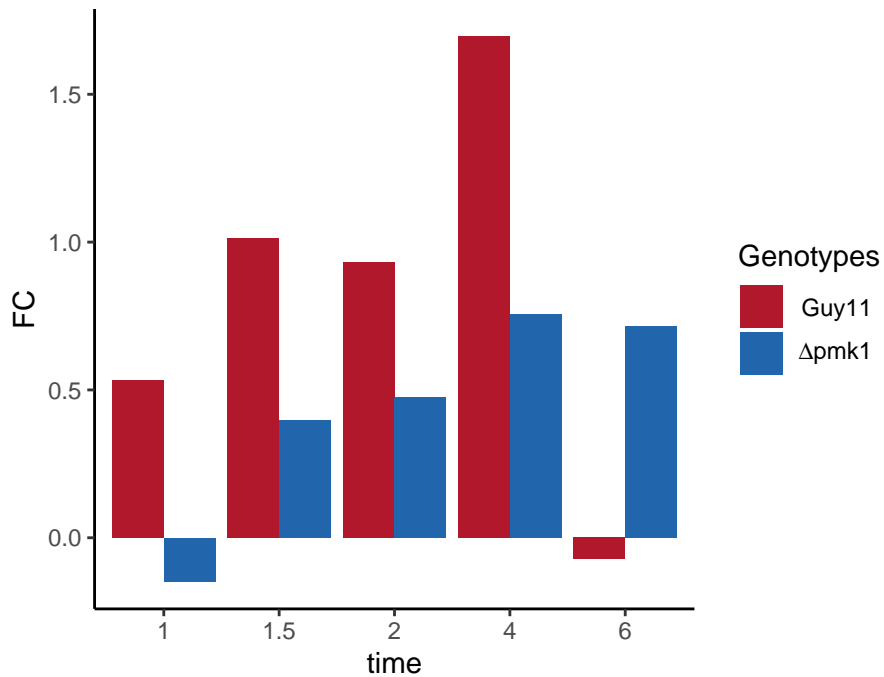
MGG_09554 (S295)
ASGSGGVDSVTAERS[+80]PR



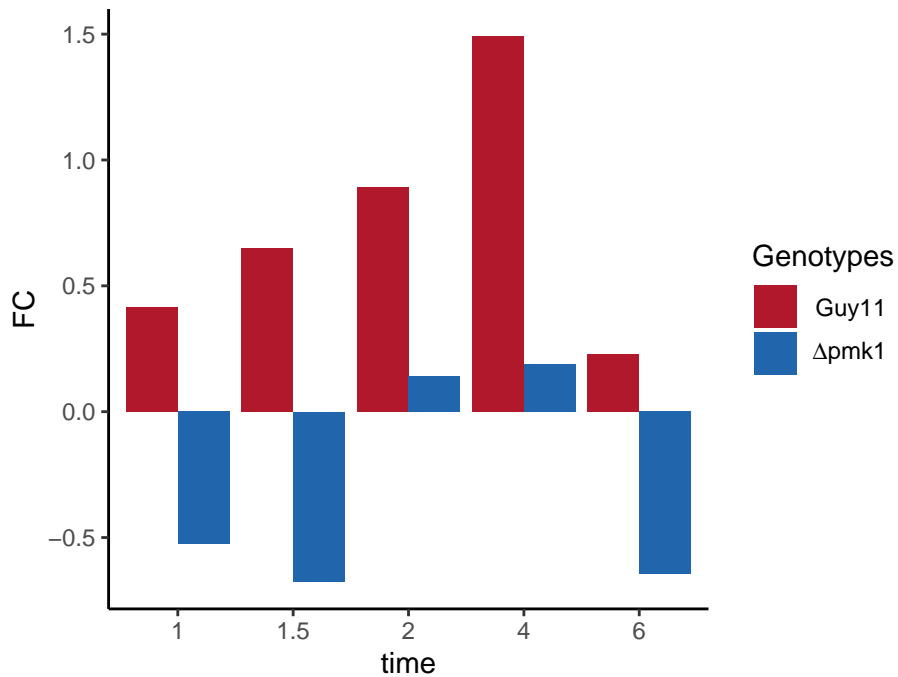
MGG_09554 (S295)
ASGSGGVDSVT AERS[+80]PRR



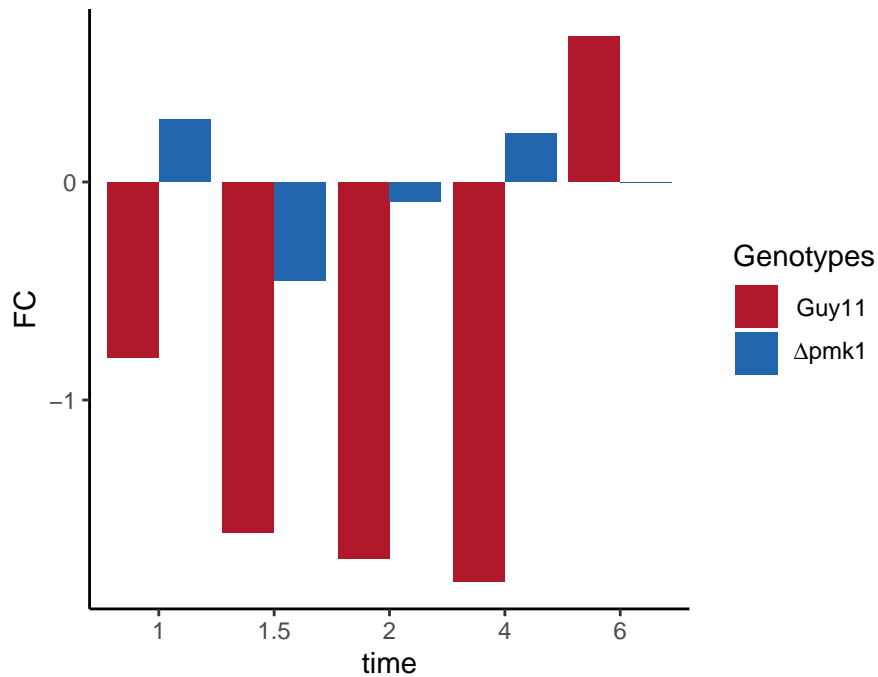
MGG_09697 (T798)
SRSYST[+80]PDIQGQGPPGM[+16]R



MGG_09697 (T798)
SYST[+80]PDIQQGQPPGM[+16]R

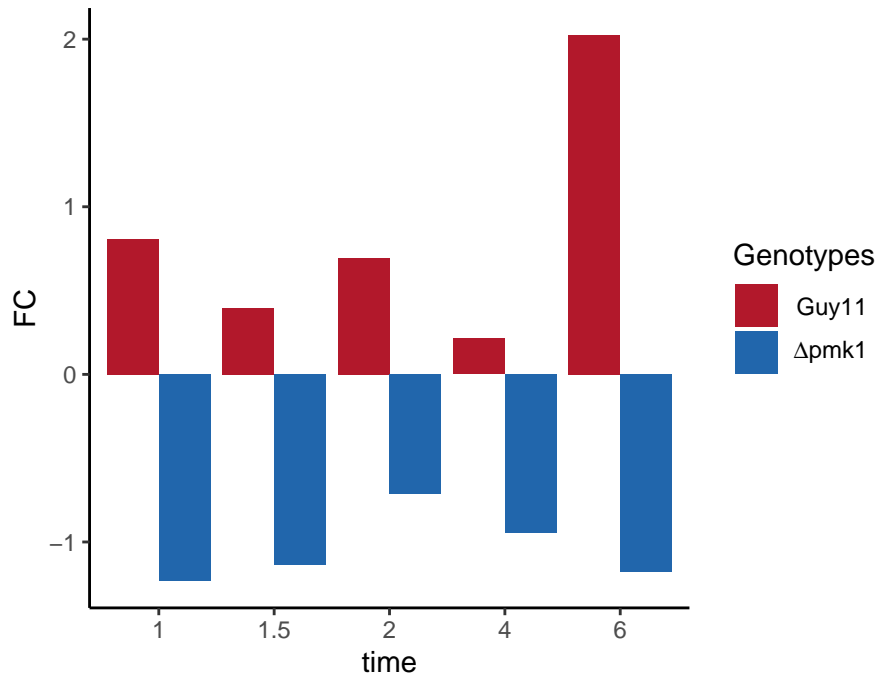


MGG_09697 (T899)
GGT[+80]PNPPGSGHTPAPLNLSADM[+16]GR

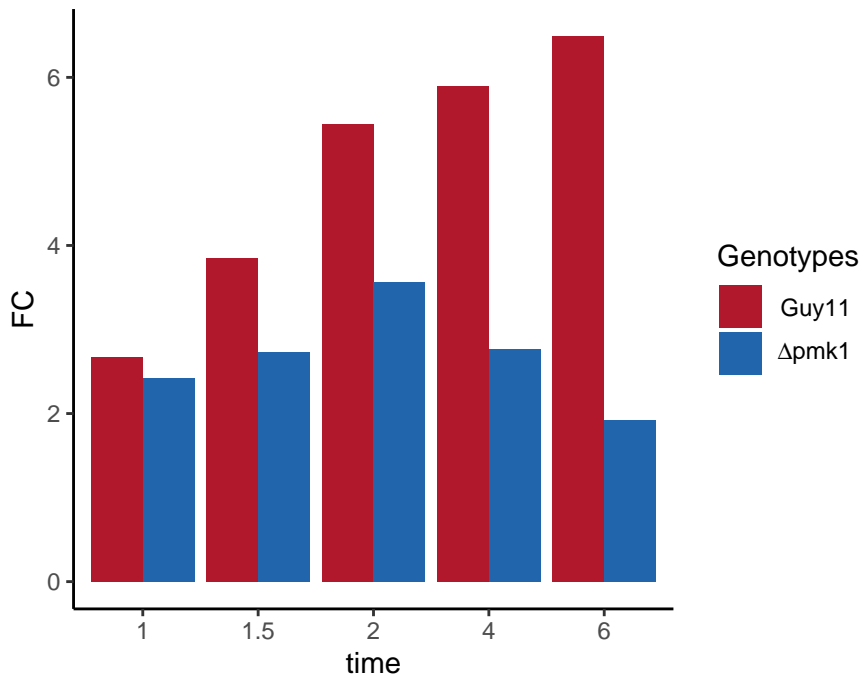


MGG_10538 (T512)

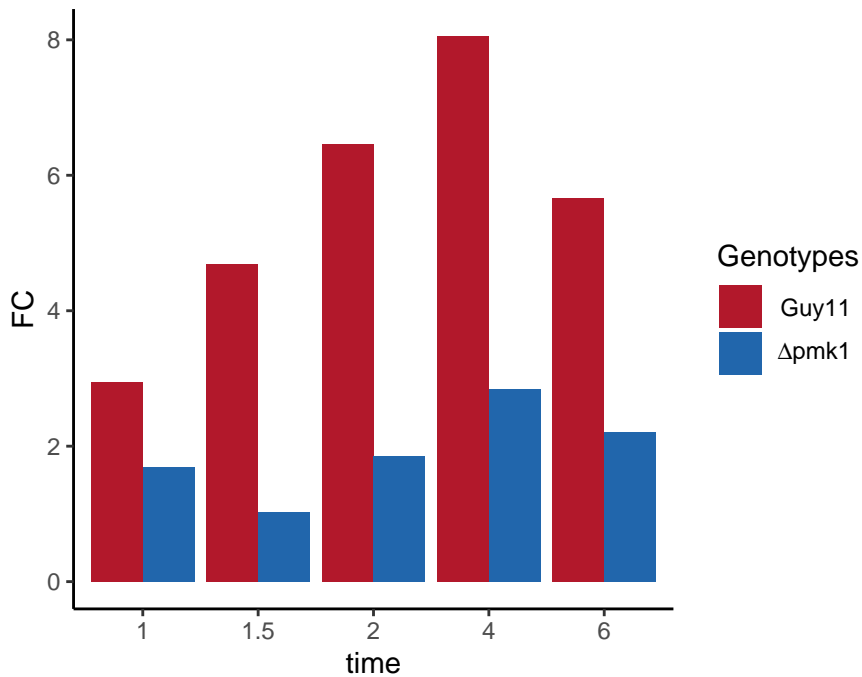
SATPGGRGT[+80]PDVGGSGGYGGGGSLTDAER



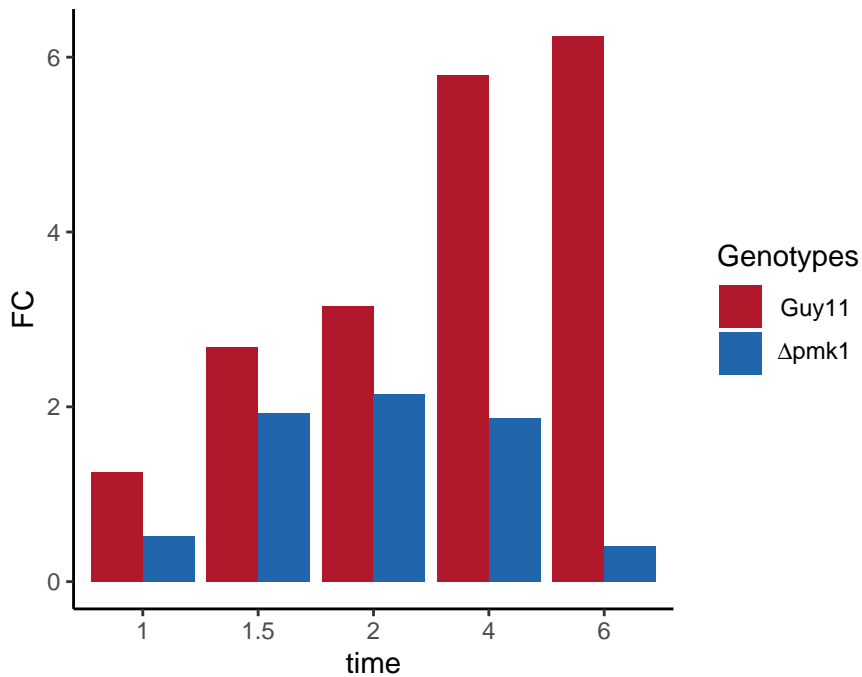
MGG_12865 HOX7 (S126)
RLS[+80]FGDDQEER



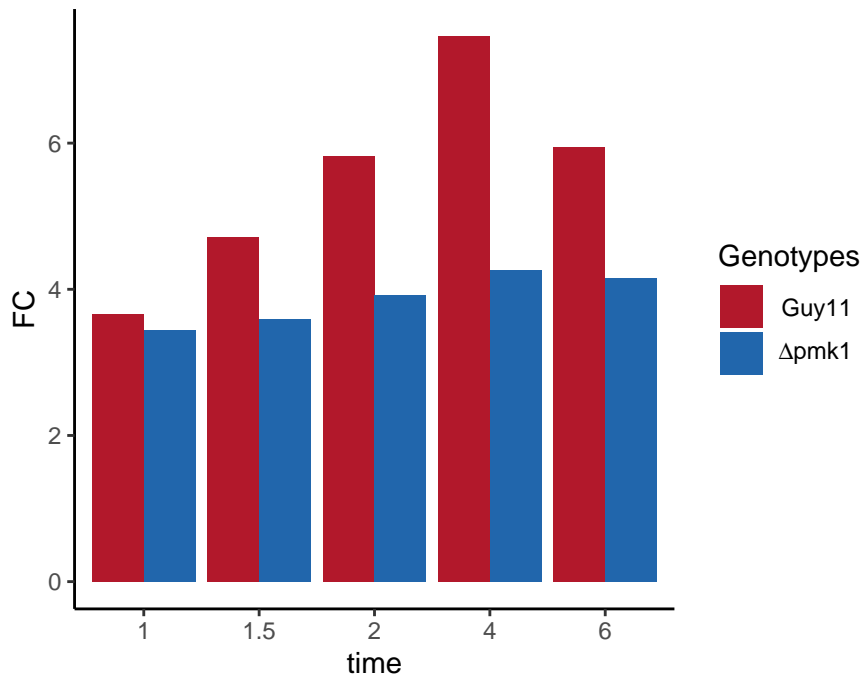
MGG_12865 HOX7 (S126)
RLS[+80]FGDDQEERVSQVPR



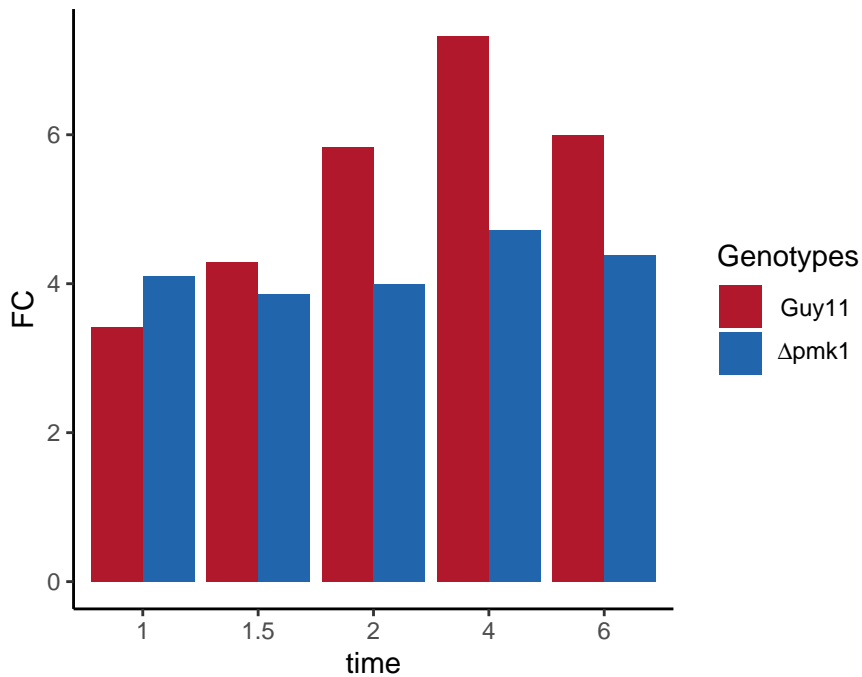
MGG_12865 HOX7 (S126)
RRLS[+80]FGDDQEER



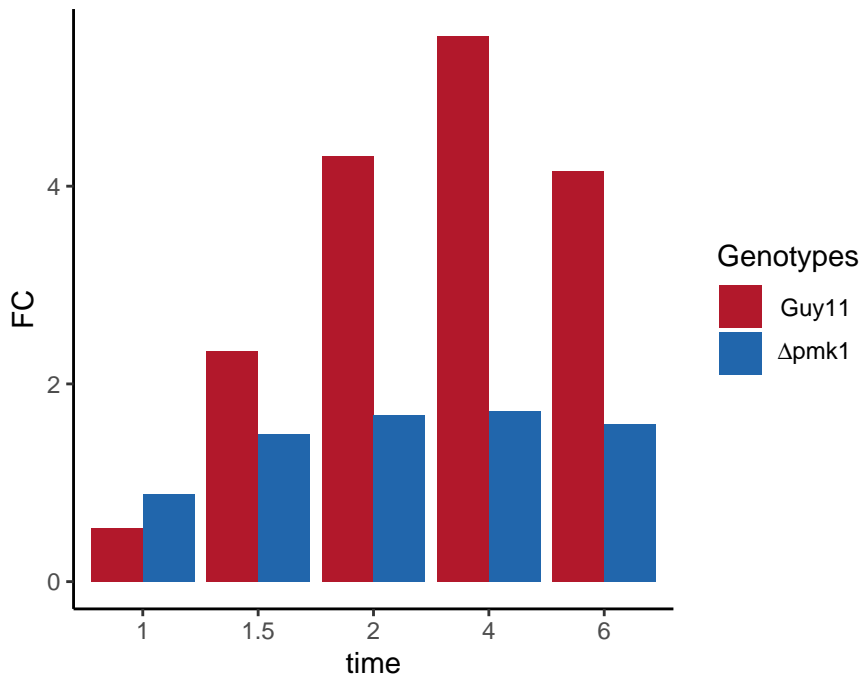
MGG_12865 HOX7 (S158)
DNRPLS[+80]PVTLSGGPR



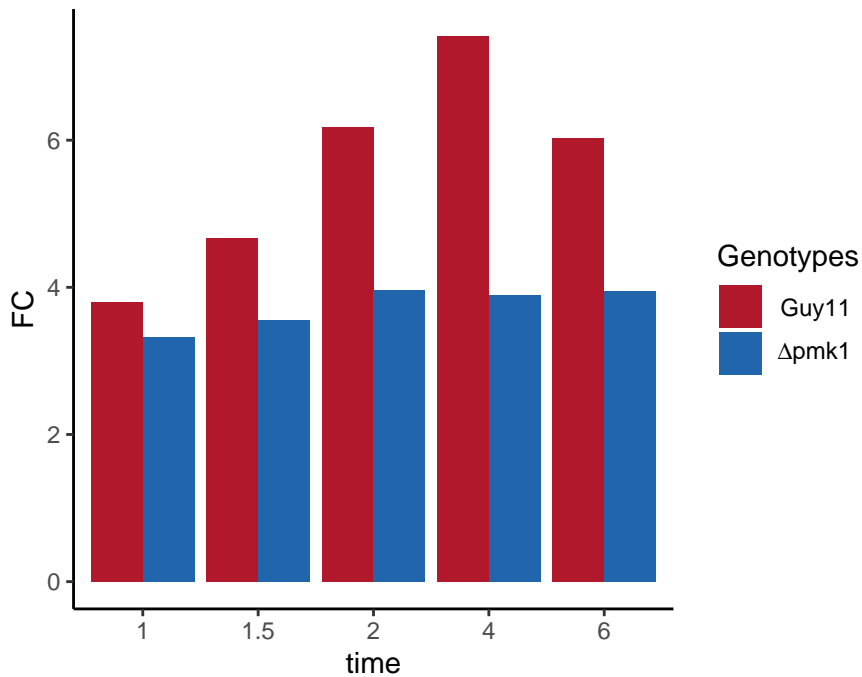
MGG_12865 HOX7 (S158)
RDNRPLS[+80]PVTLSGGPR



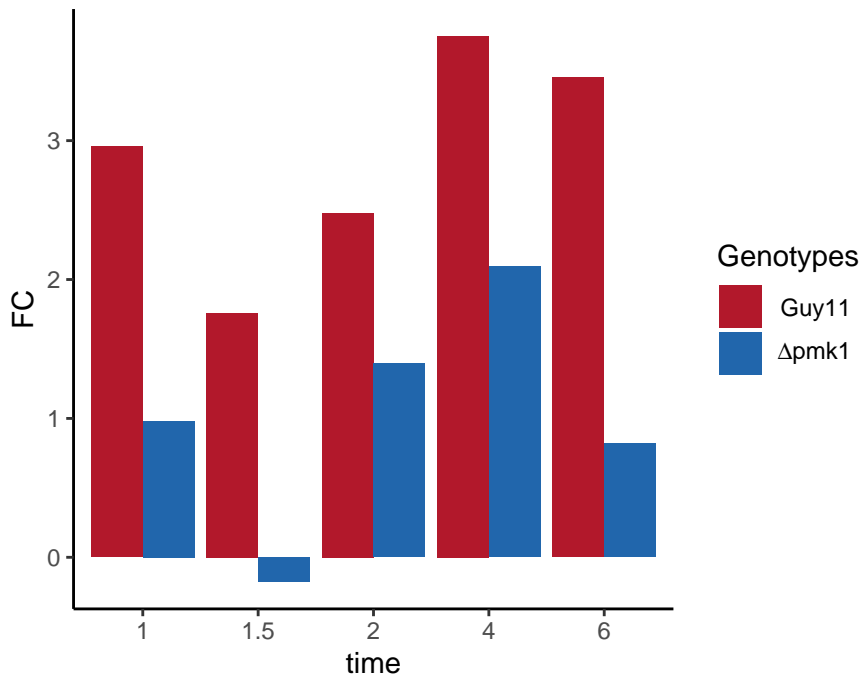
MGG_12865 HOX7 (S163)
RDNRPLSPVTL[+80]GGPR



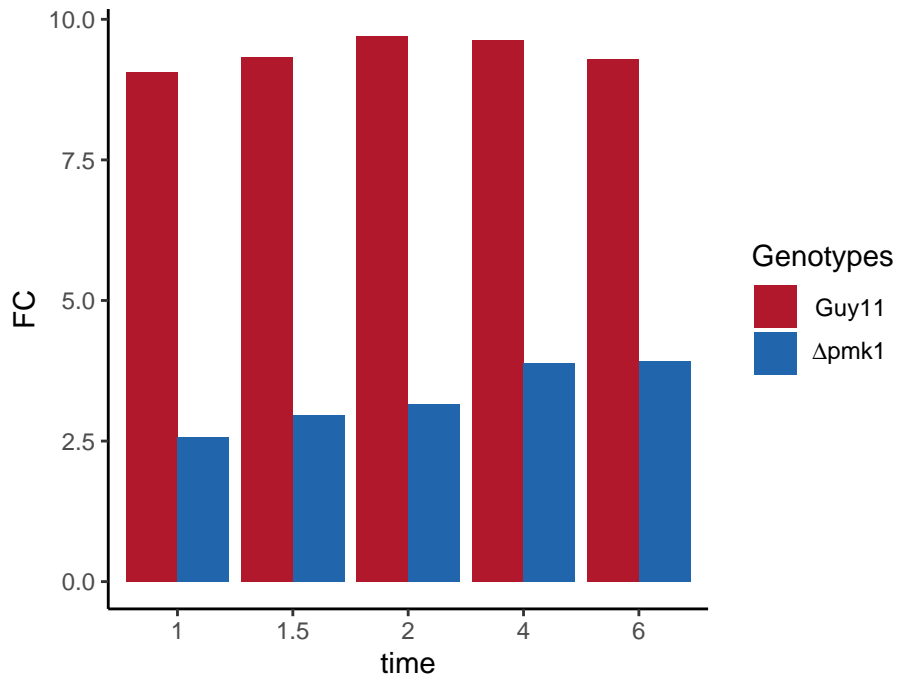
MGG_12865 HOX7 (T161)
DNRPLSPVT[+80]LSGGPR



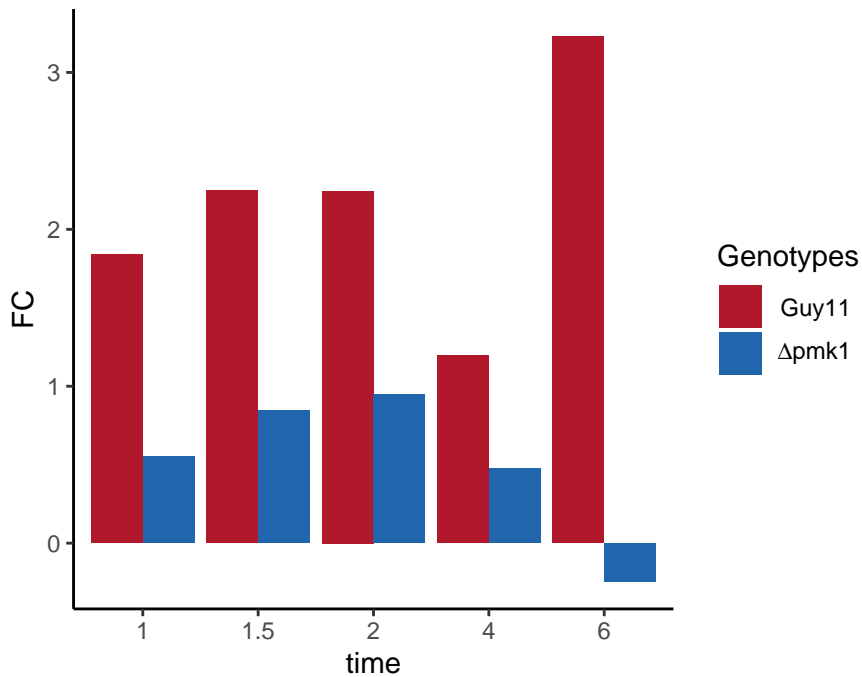
MGG_14847 MST11 (S557)
LQHGYRDSVASSM[+16]LDTLQEEGS[+80]PIEPNRK



MGG_00800* MST7 (S318)
SSDS[+80]PTATYHGDDRPLETPTSGEIPILGAIVSPR

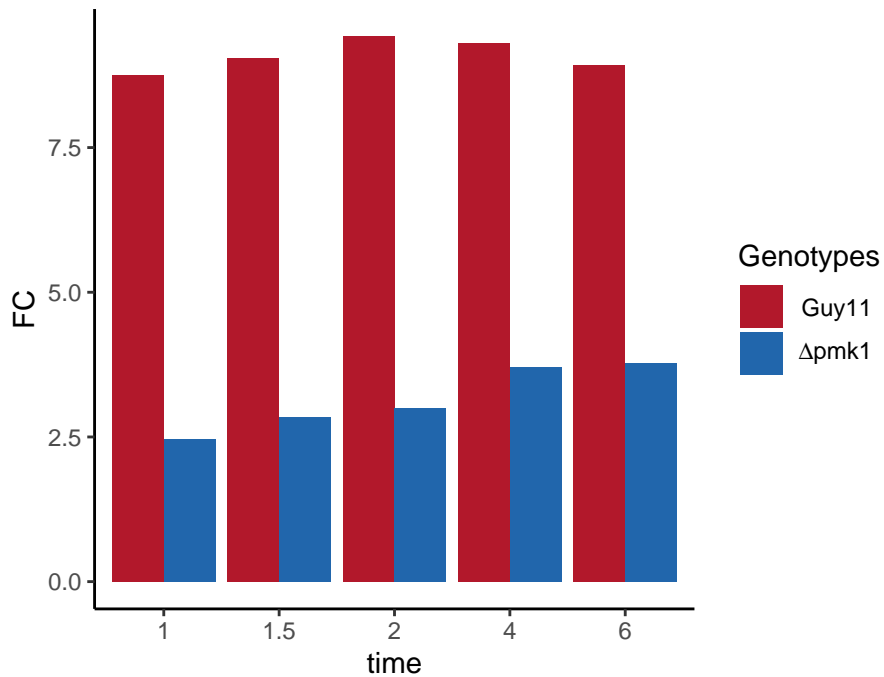


MGG_00800* MST7 (S358)
DQYGSISSRS[+80]PTR



MGG_00800* MST7 (T332)

SSDSPTATYHGDDRPLET[+80]PTSGEIPILGAIVSPR



MGG_09898* MAC1 (S75)
ITASPLRDDTQVS[+80]PTNTFGTR

