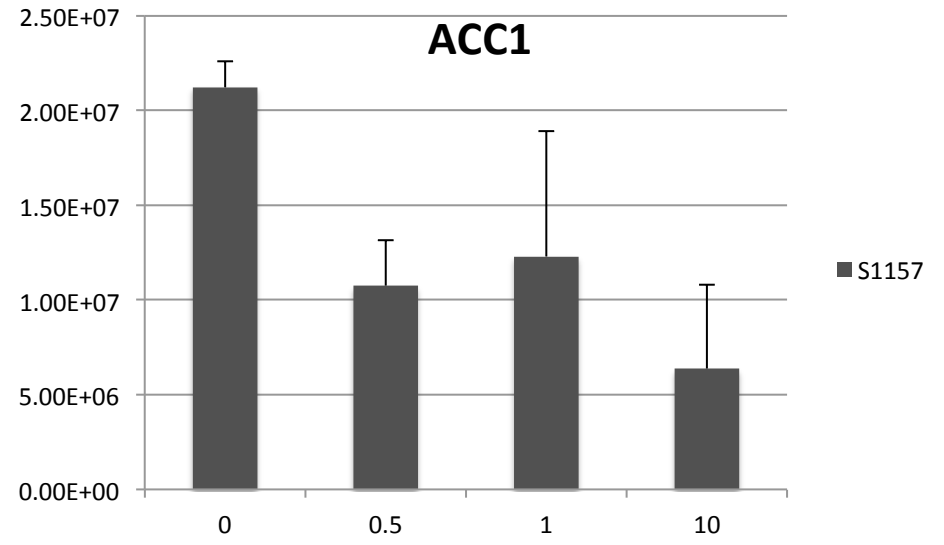
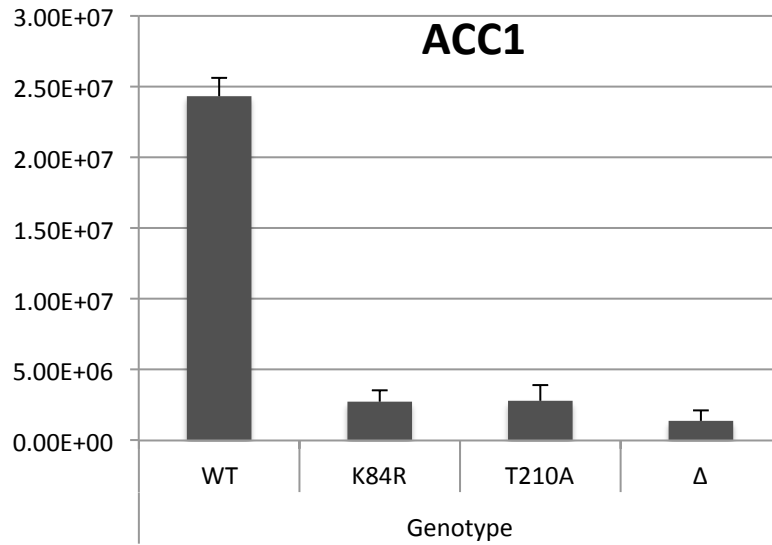


Supplementary File 1

- First panel: Phosphopeptide abundance after two hours of glucose depletion in *SNF1* mutant strain CKY18(*snf1Δ*) carrying WT *SNF1* (WT), a catalytically inactive mutant (*snf1-K84R*), an activation-defective mutant (*snf1-T210A*), or no *SNF1* (Δ) on low copy number *CEN URA3* plasmids. The *SNF1* genes were expressed from their own promoter.
- Second panel: Phosphopeptide abundance in strains TYY923, TYY924, and TYY925 (*SNF1^{as}* (I132G)) after 2.5 hours of glucose depletion and treatment with the indicated concentrations of 2NM-PP1 or DMSO (vehicle) beginning at two hours.
- Third panel: Kinetics of phosphorylation and dephosphorylation of Snf1 target phosphopeptides in strains TYY923, TYY924, and TYY925 (*SNF1^{as}* (I132G)) during glucose depletion and after treatment with 10 μ M 2NM-PP1 at 40 min. At t=0 the cells were shifted to low (0.05%) glucose.
- The numbers on the ordinate represent the abundance measured as ion counts of the phosphopeptides. The values on the graphs represent the means and standard deviations of three biological replicates.
- Black solid and dashed lines and bars represent putative *SNF1*-dependent phosphopeptides based on data in experiments 1 and 2. Black dotted lines represent phosphopeptides likely to be Snf1-dependent based on inhibition by 2NM-PP1 in experiment 3. Colored lines indicate Snf1-independent phosphopeptides.

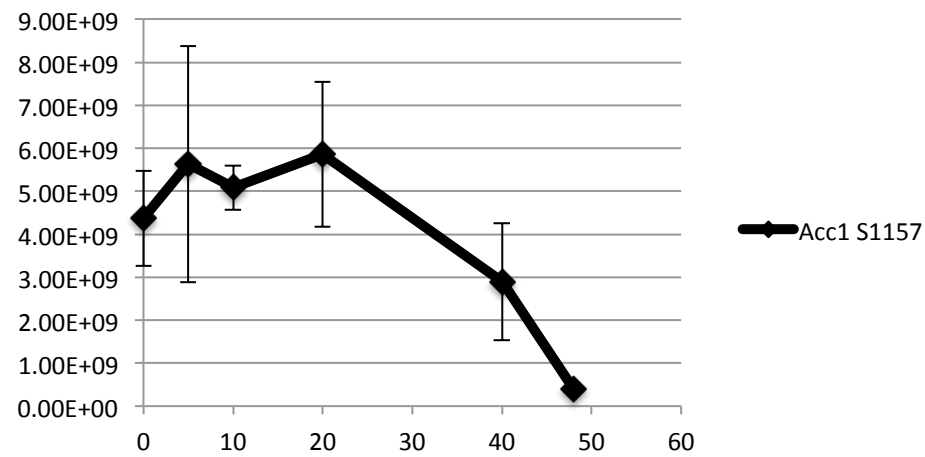
ACC1



SNF1 genotype

2 NM-PP1 (μM)

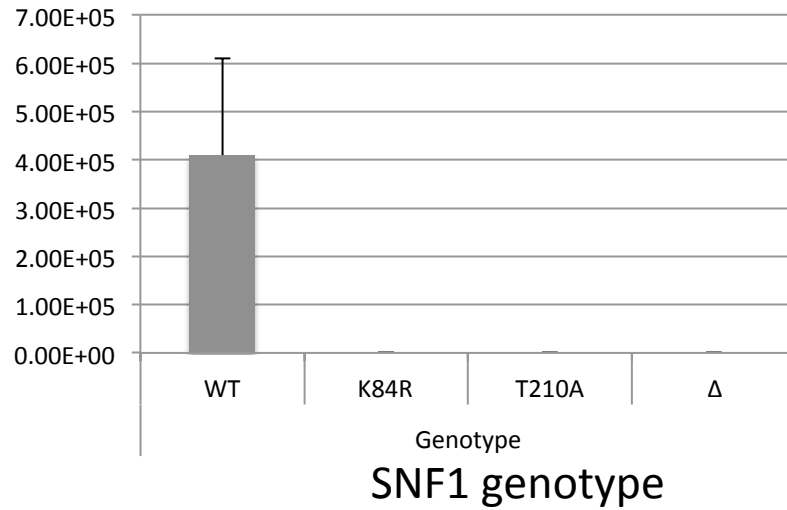
ACC1



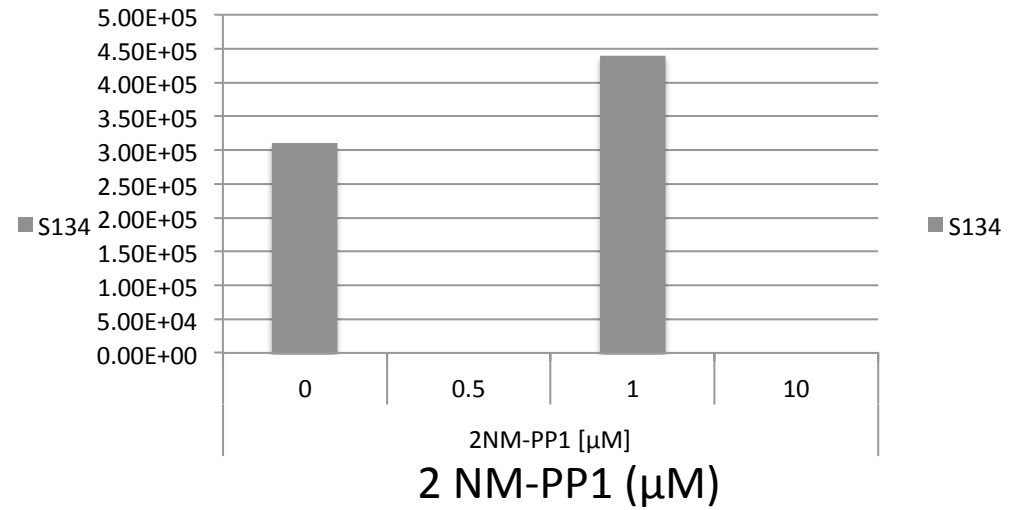
Min of derepression (48 = + 2 NM-PP1 (10 μM))

ADA3/NGG1

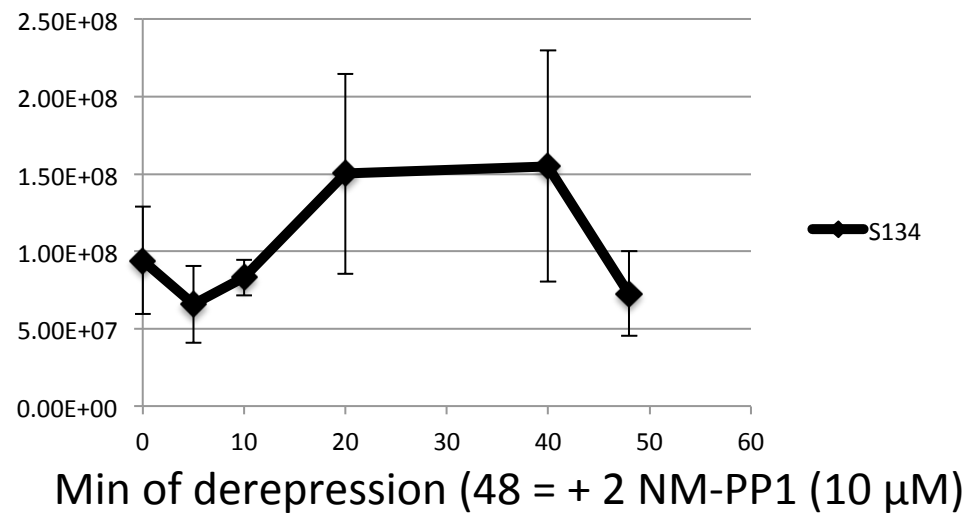
ADA3/NGG1



ADA3/NGG1

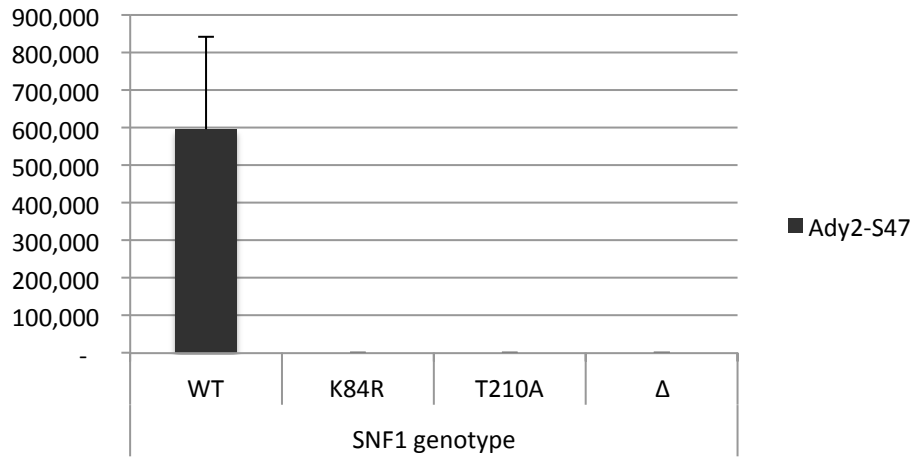


ADA3/NGG1

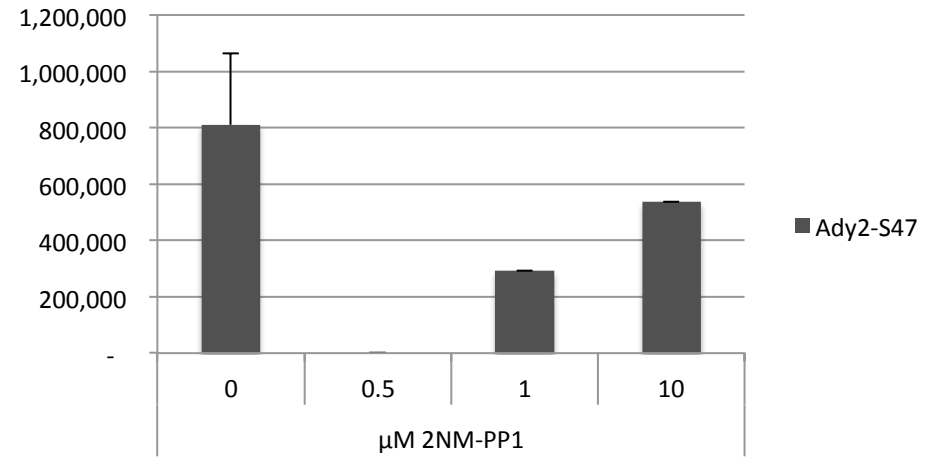


ADY2

Ady2-S47



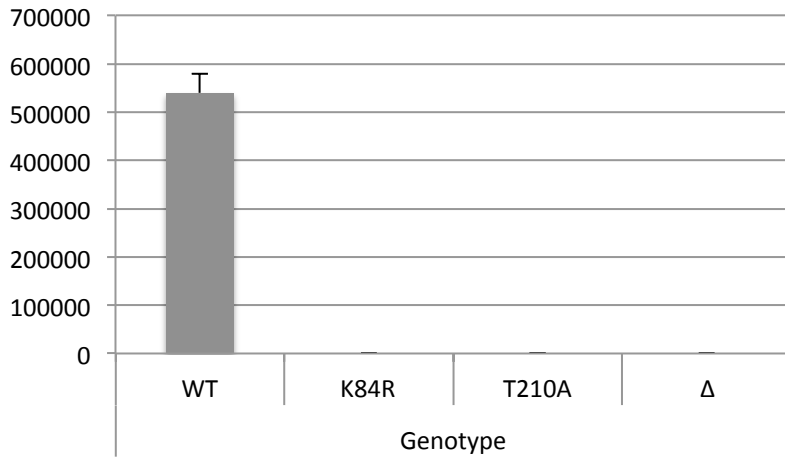
Ady2-S47



No time course data for ADY2

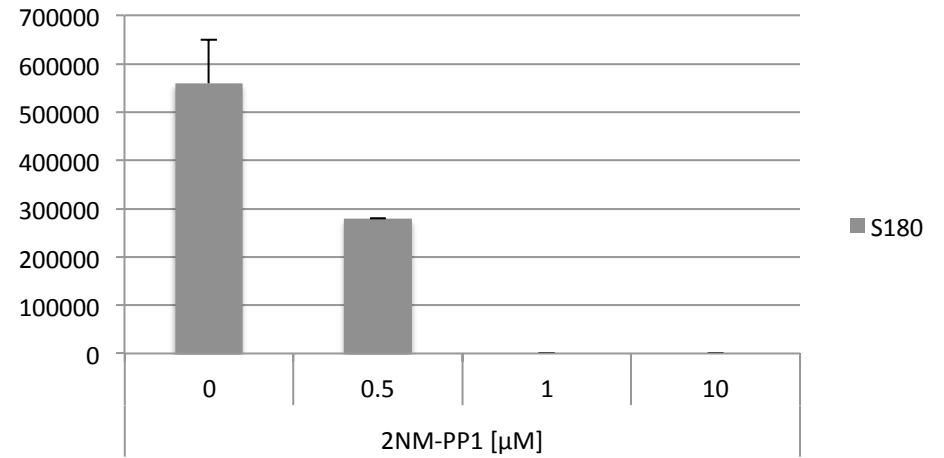
AGE2

AGE2



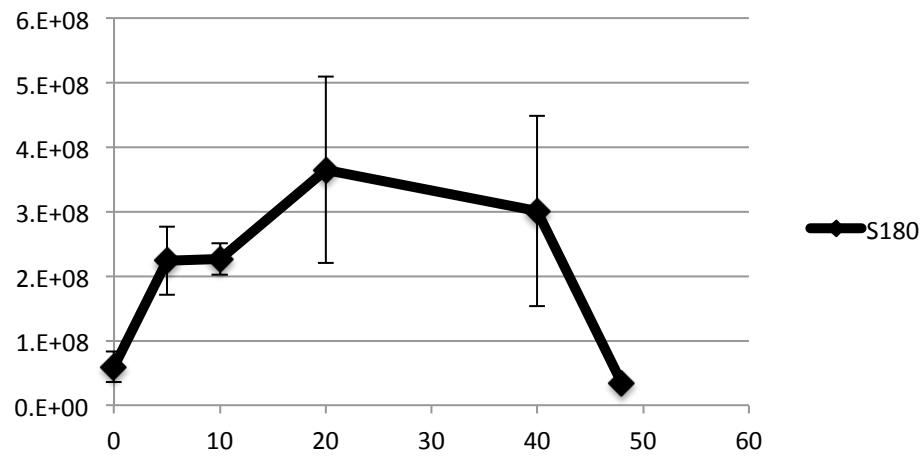
SNF1 genotype

AGE2



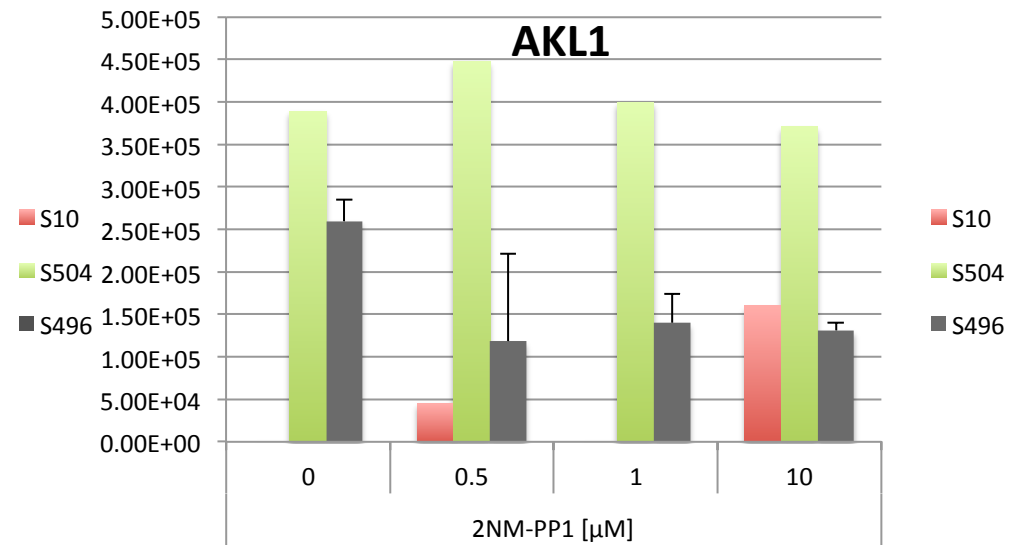
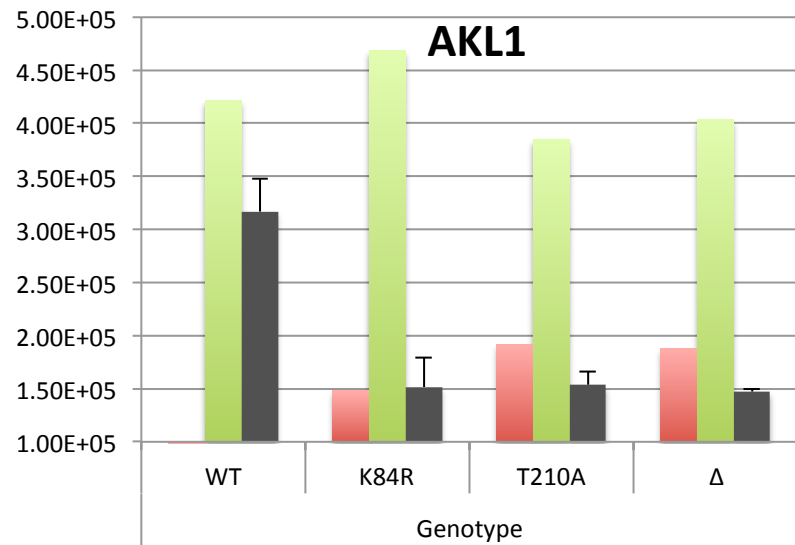
2 NM-PP1 (μM)

AGE2

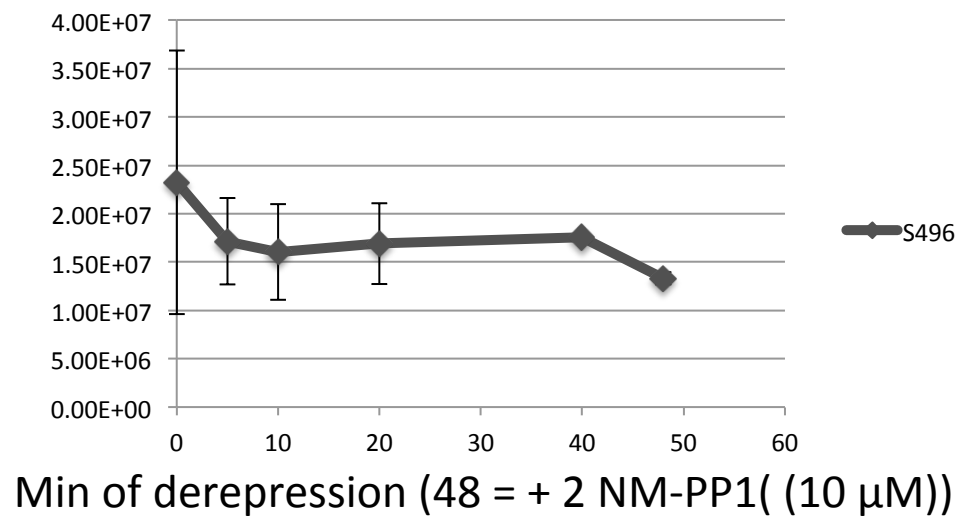


Min of derepression (48 = + 2 NM-PP1 (10 μM))

AKL1

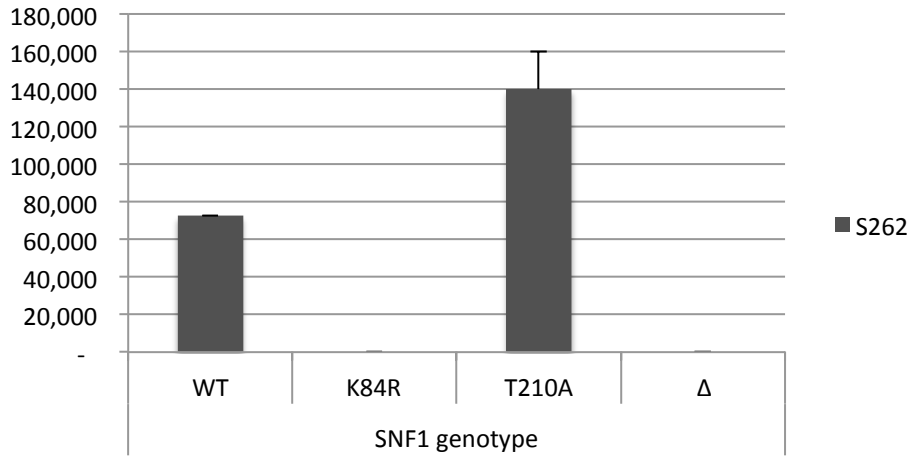


AKL1

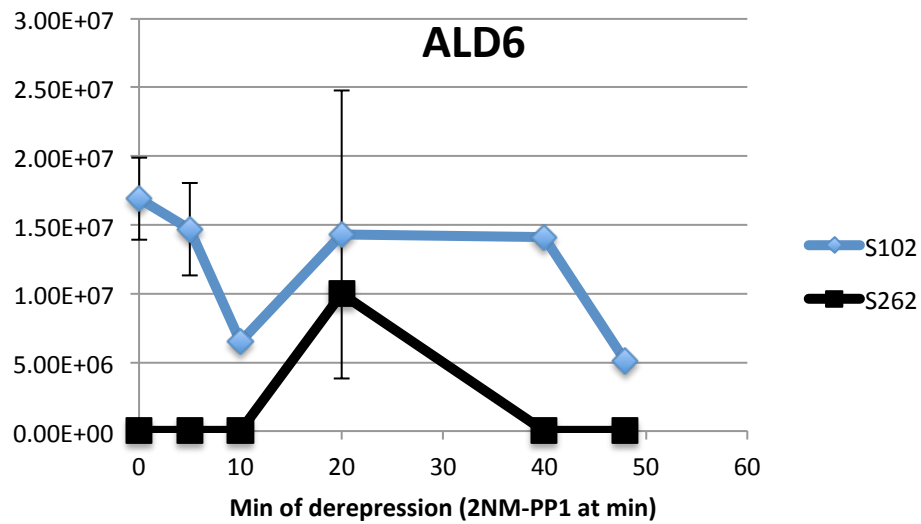
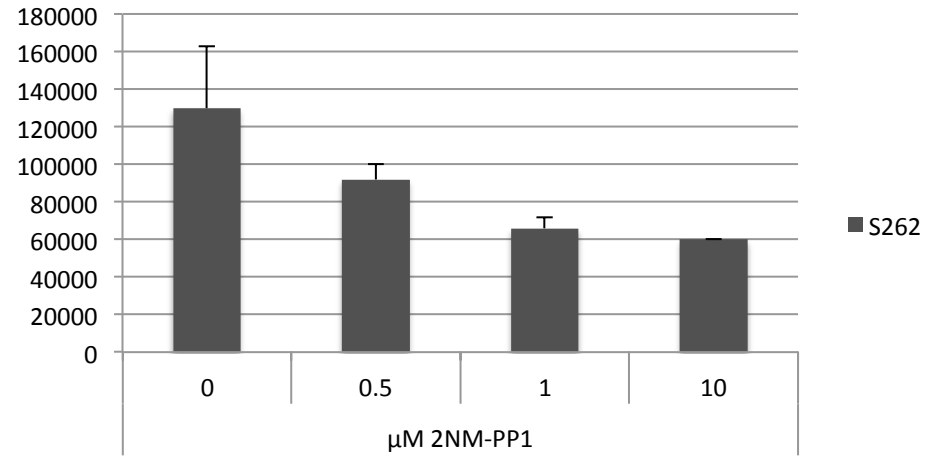


ALD6

ALD6

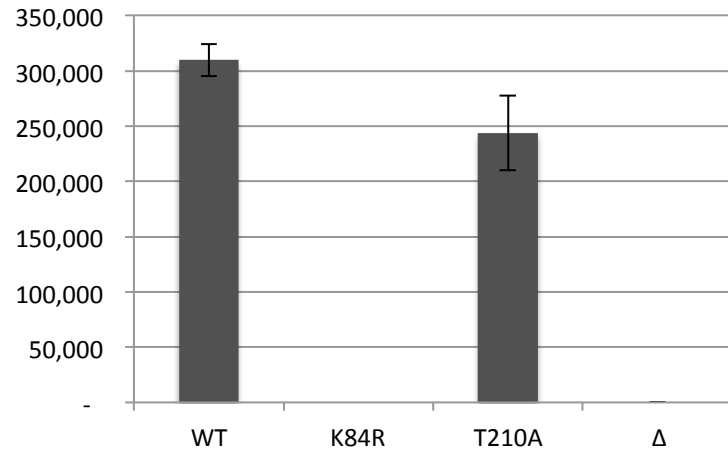


ALD6



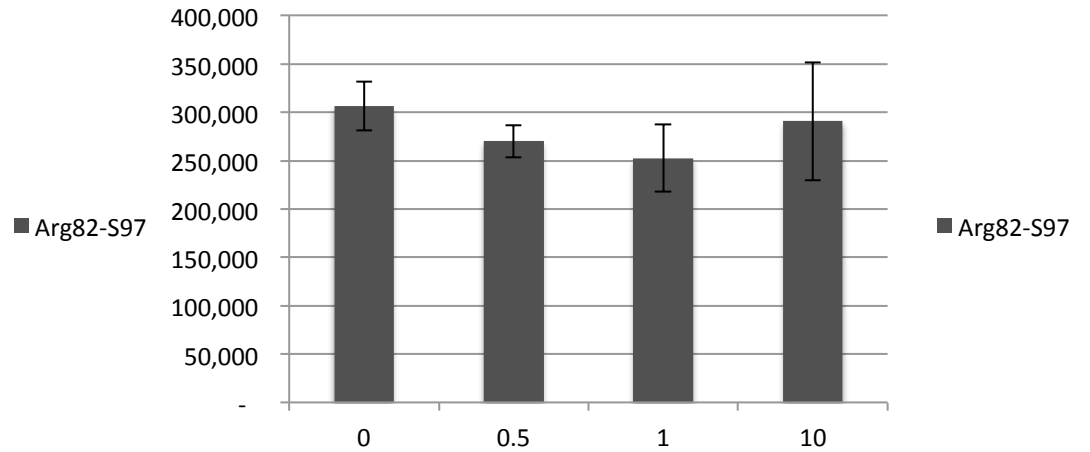
ARG82

ARG82



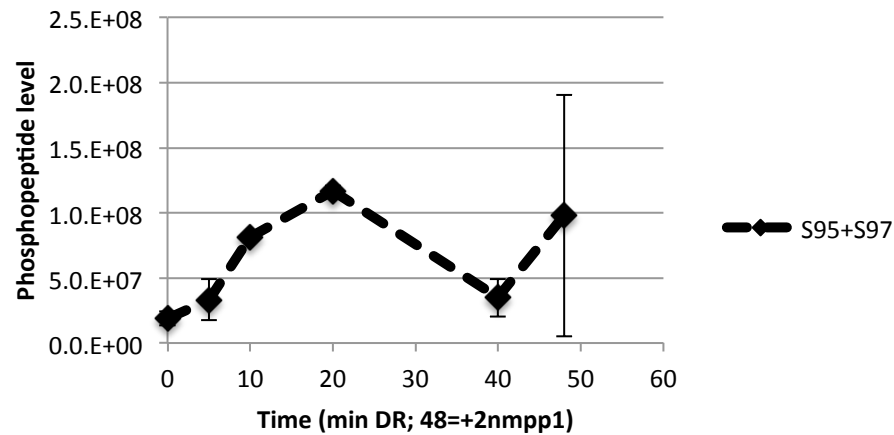
SNF1 genotype

ARG82



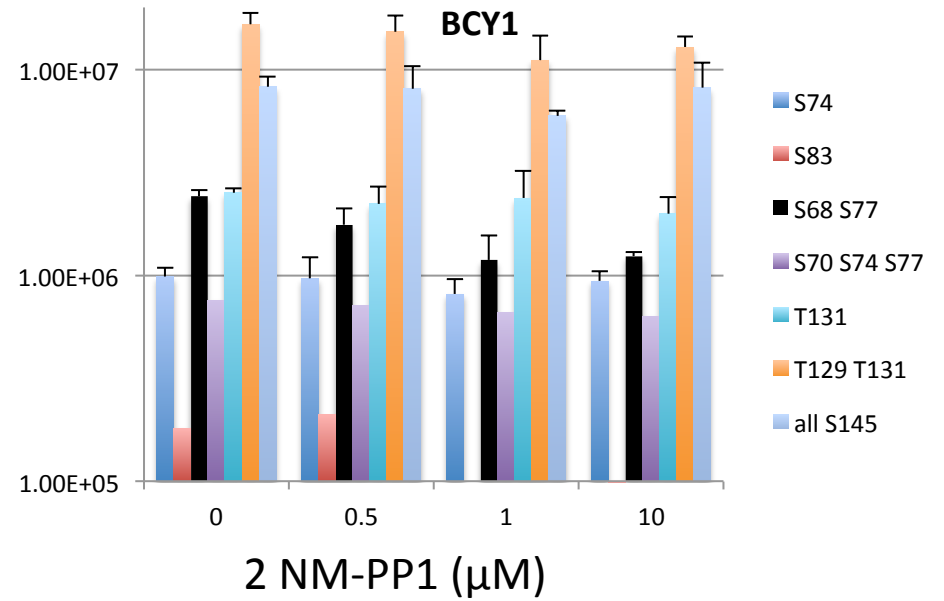
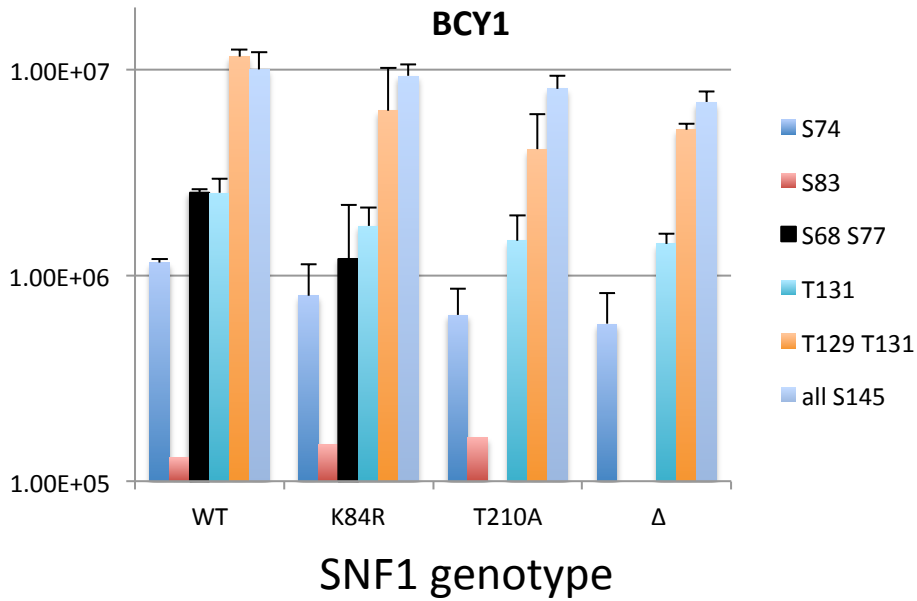
2 NM-PP1 (μM)

ARG82

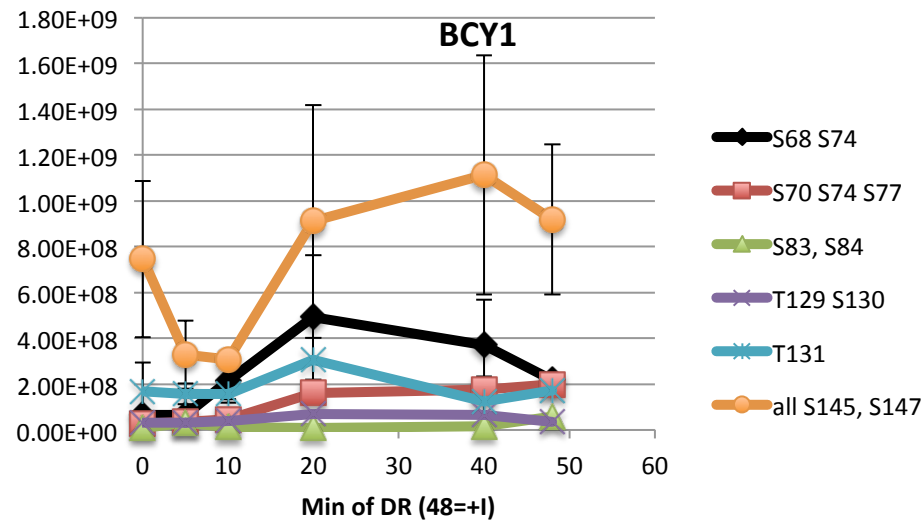


Min of derepression (48 = + 2 NM-PP1 (10 μM))

BCY1



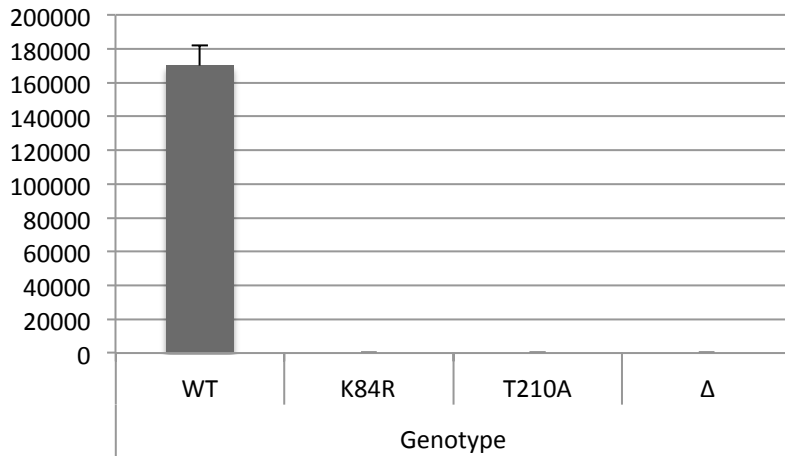
NB: Only one K84R sample (#3) detected pS68 pS77.



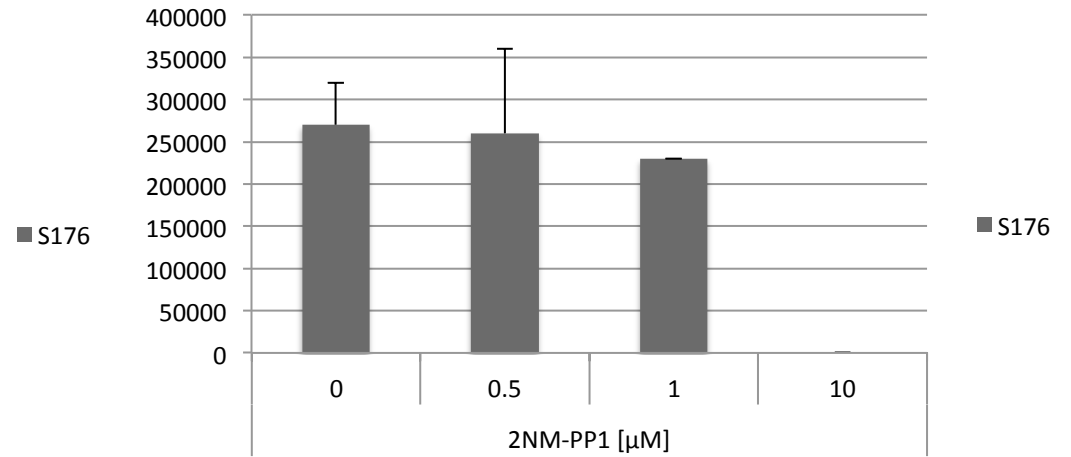
Min of derepression (48 = + 2 NM-PP1 (10 μM))

BNA1

BNA1

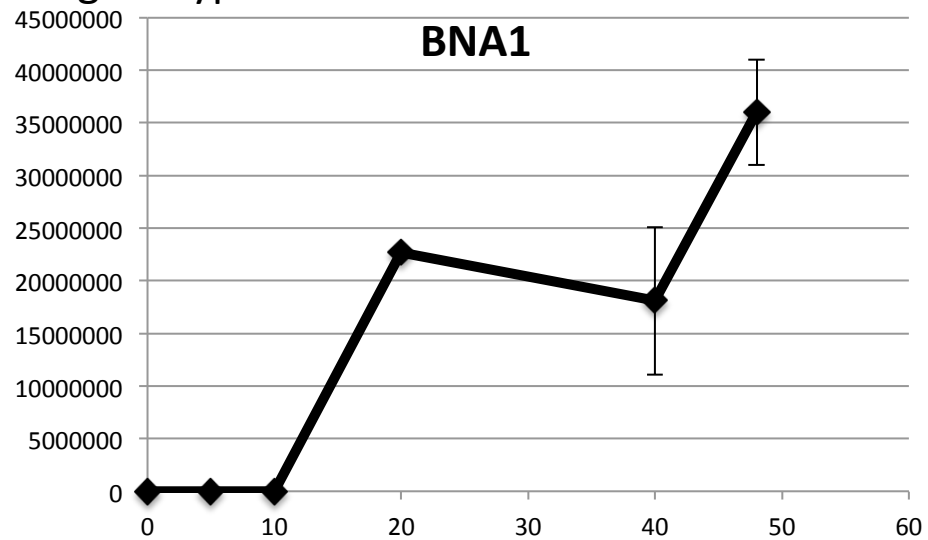


BNA1



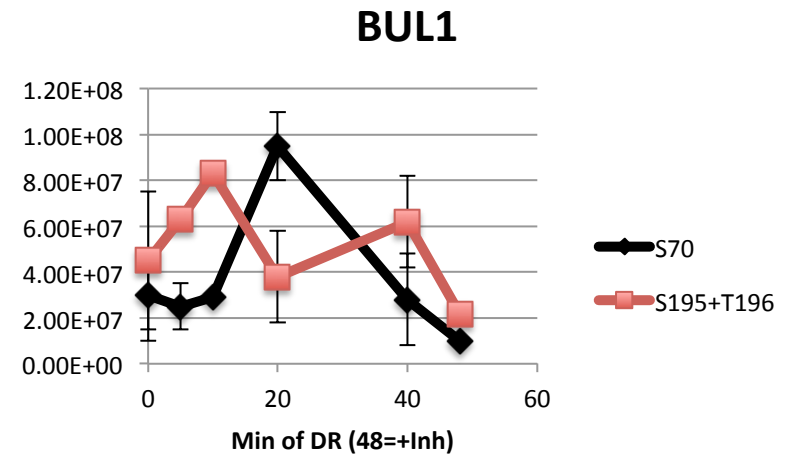
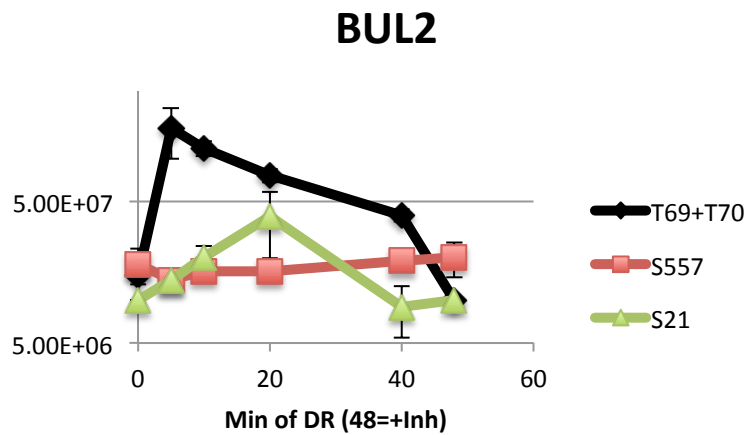
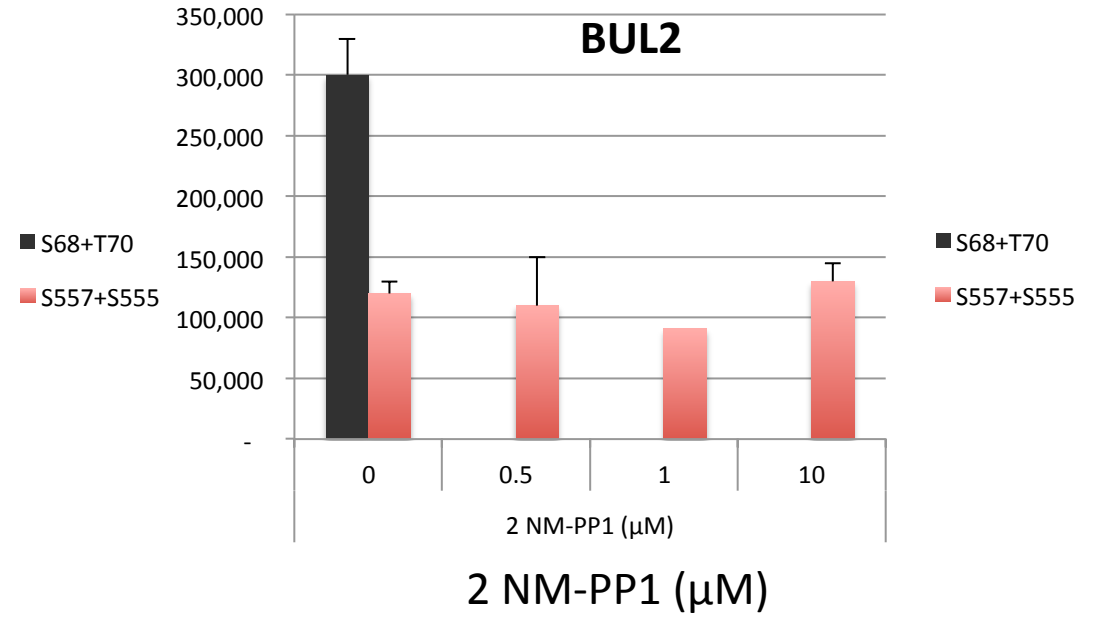
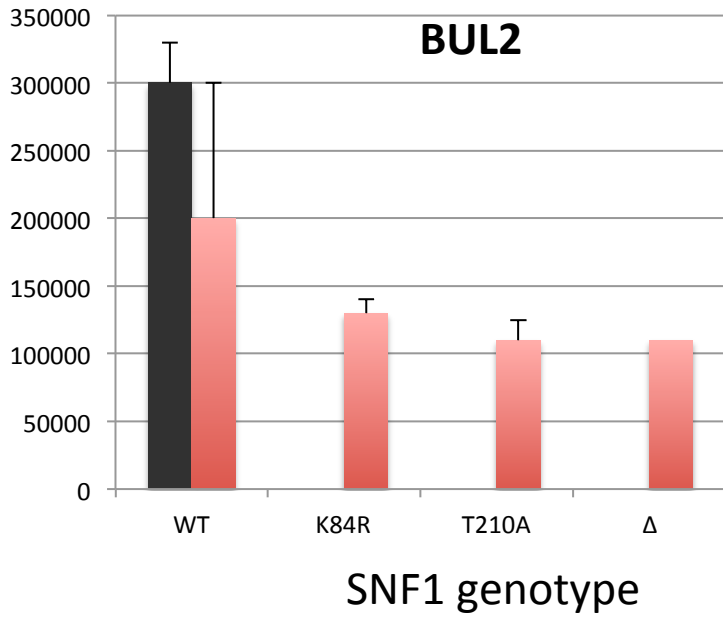
SNF1 genotype

2 NM-PP1 (μM)



Min of derepression (48 = + 2 NM-PP1 (10 μM))

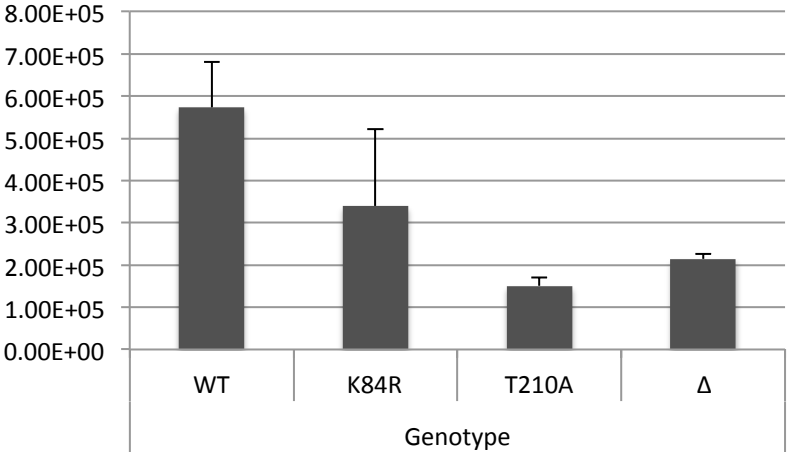
BUL2



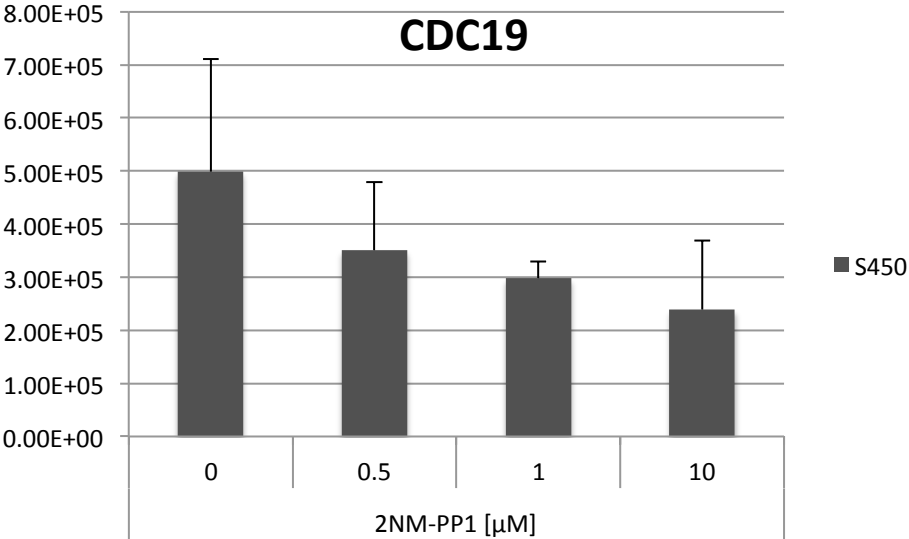
Min of derepression (48 = + 2 NM-PP1 (10 μM))

CDC19

CDC19

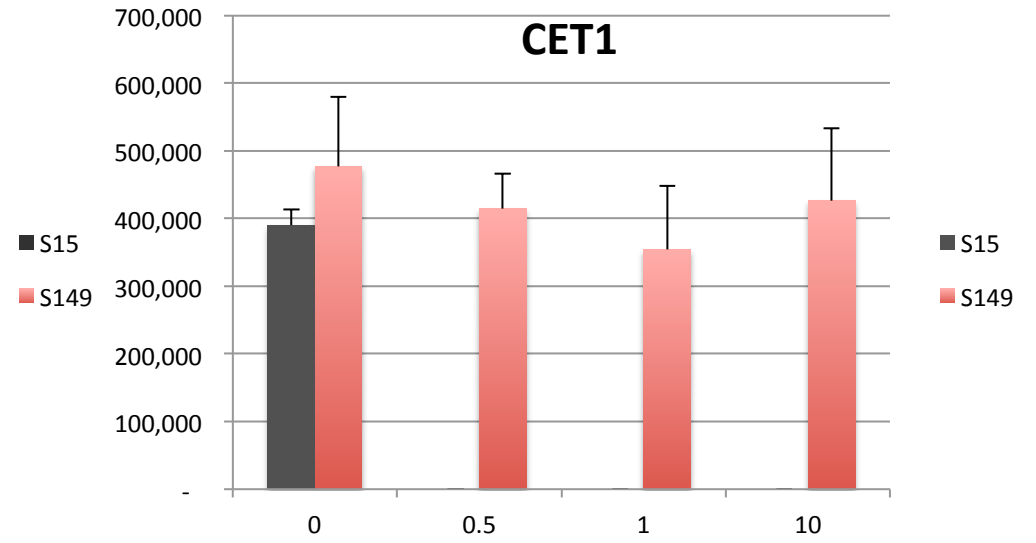
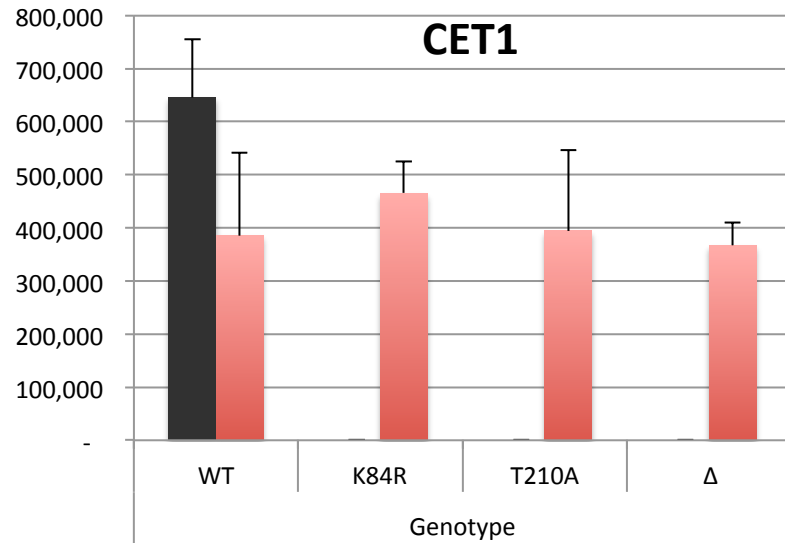


CDC19



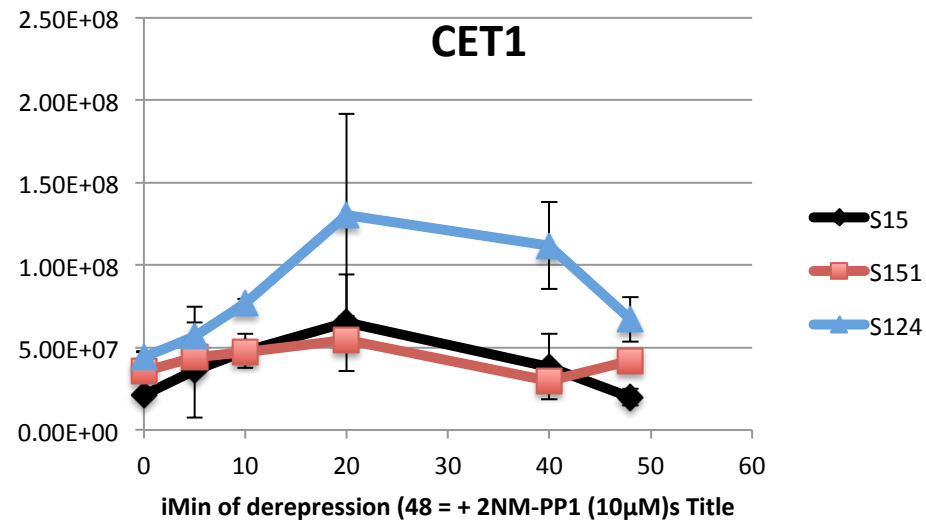
No time course data for Cdc19-S450.

CET1



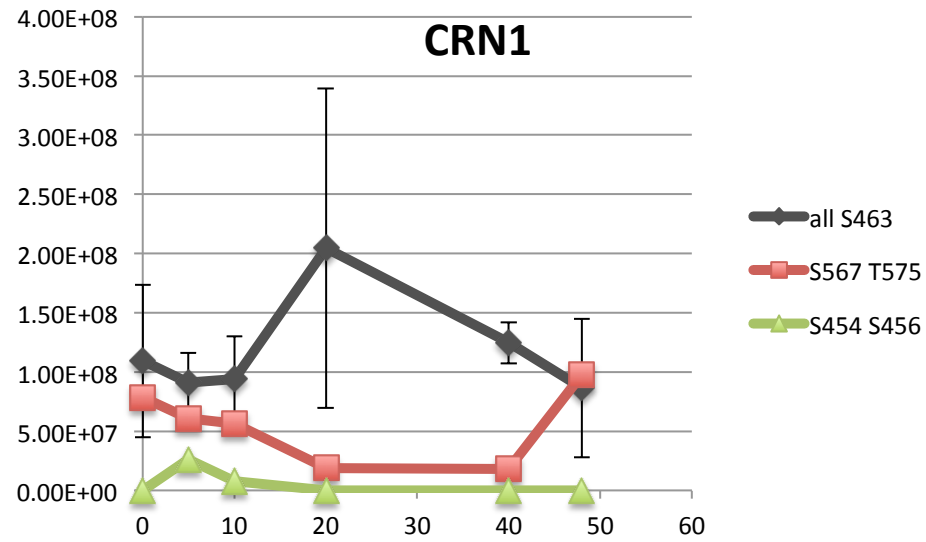
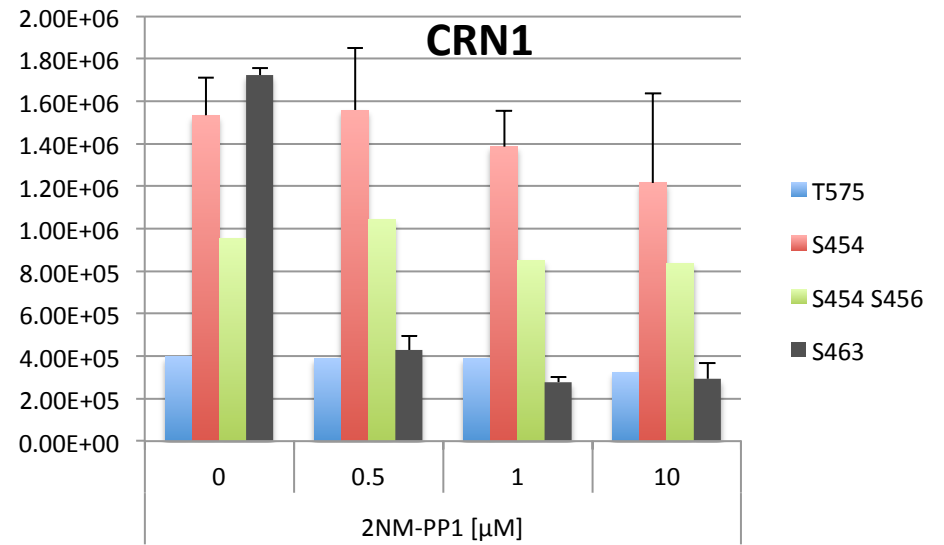
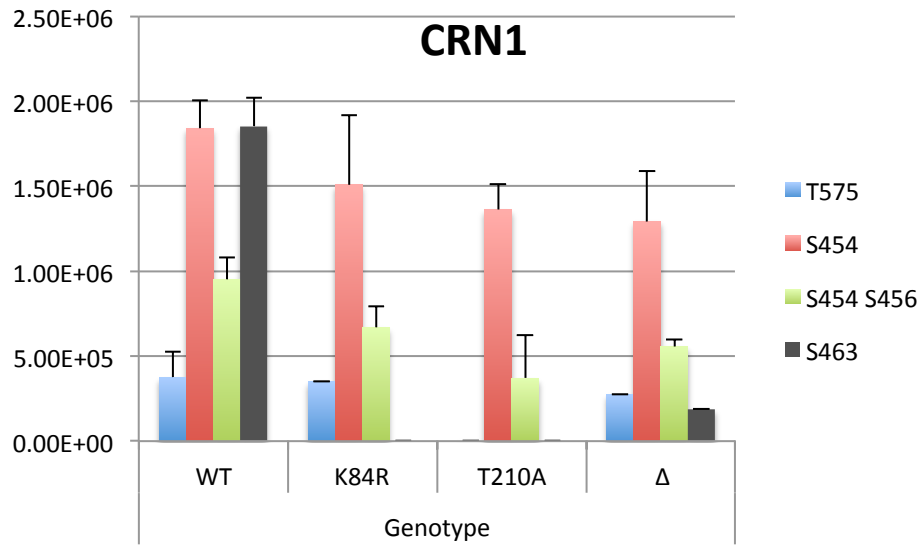
SNF1 genotype

2 NM-PP1 (μM)



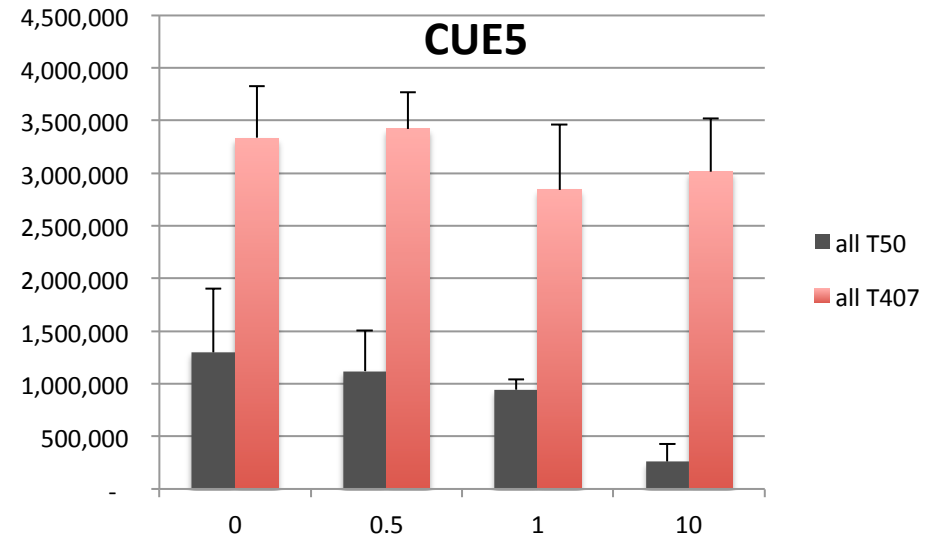
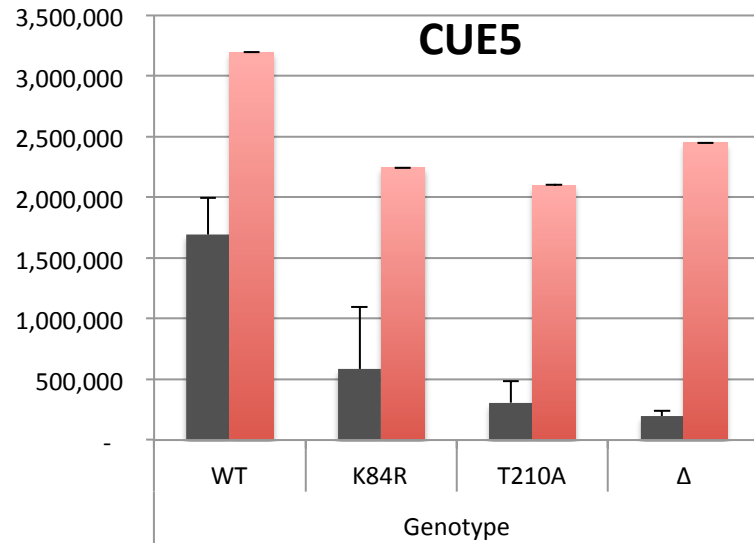
Min of derepression (48 = + 2 NM-PP1 (10 μM))

CRN1

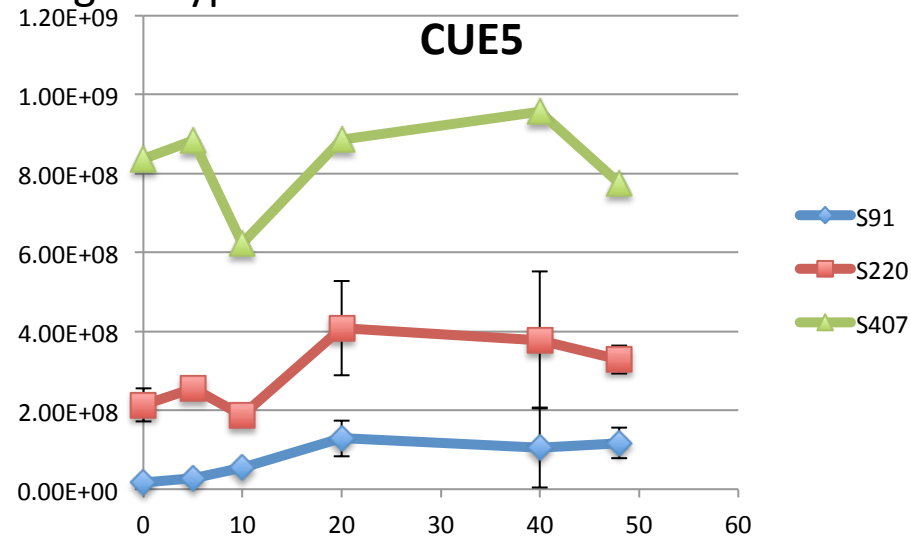


Min of derepression (48 = + 2 NM-PP1 (10 μM))

CUE5



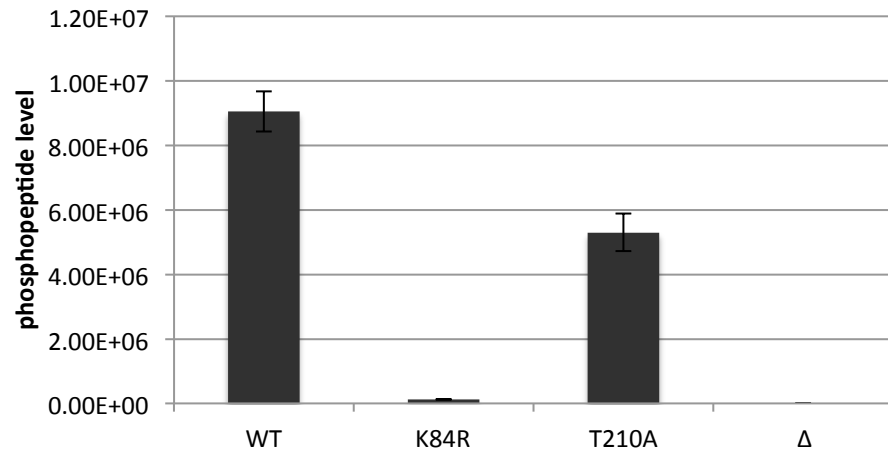
SNF1 genotype



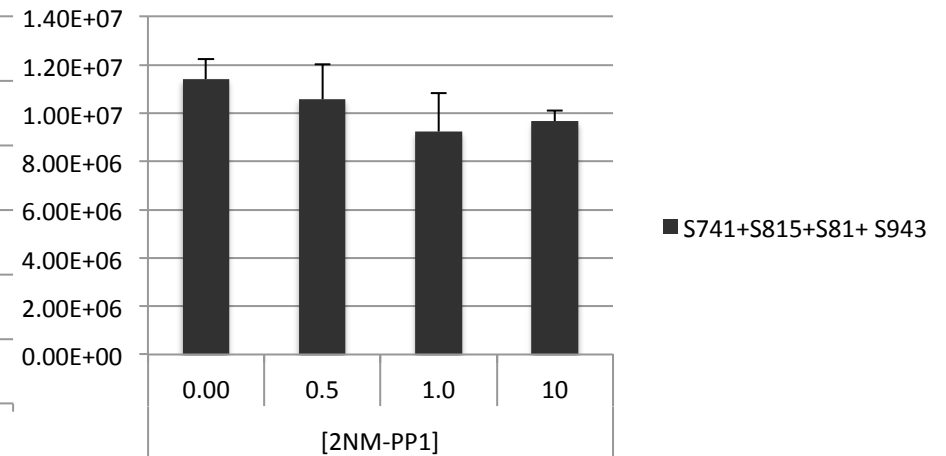
Min of derepression (48 = + 2 NM-PP1 (10 μM))

CYC8

CYC8-all phosphopeptides (average)



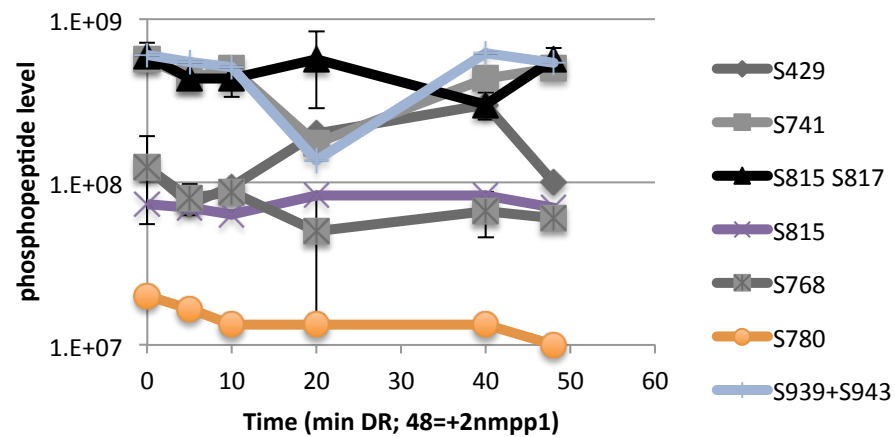
CYC8-all phosphopeptides (average)



SNF1 genotype

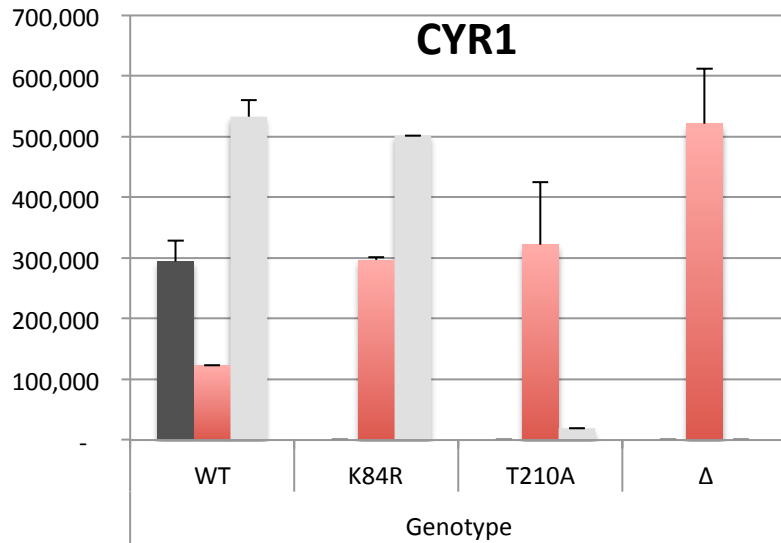
2 NM-PP1 (μM)

CYC8

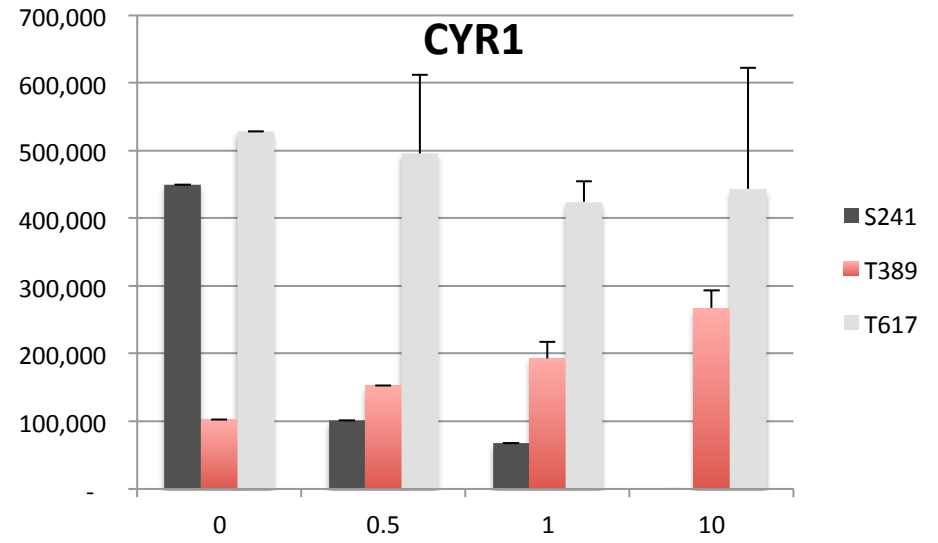


Min of derepression (48 = + 2 NM-PP1 (10 μM))

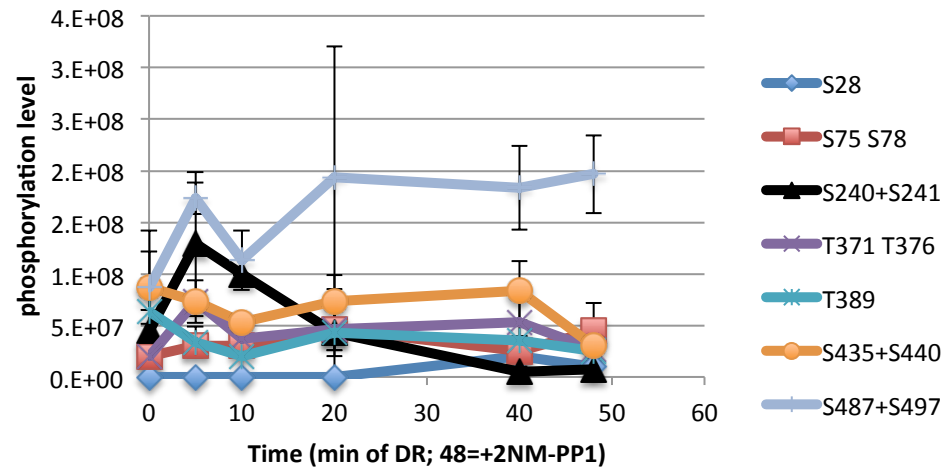
CYR1



SNF1 genotype

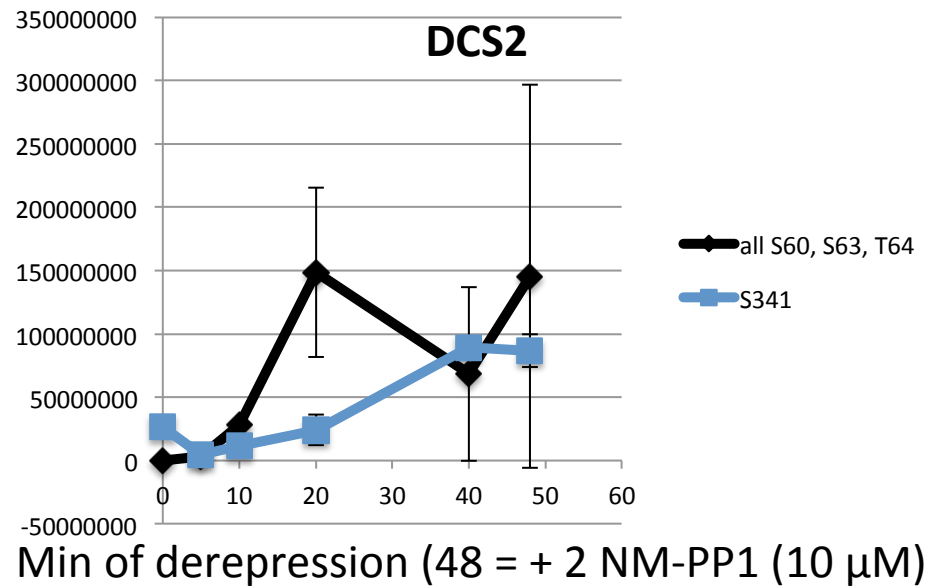
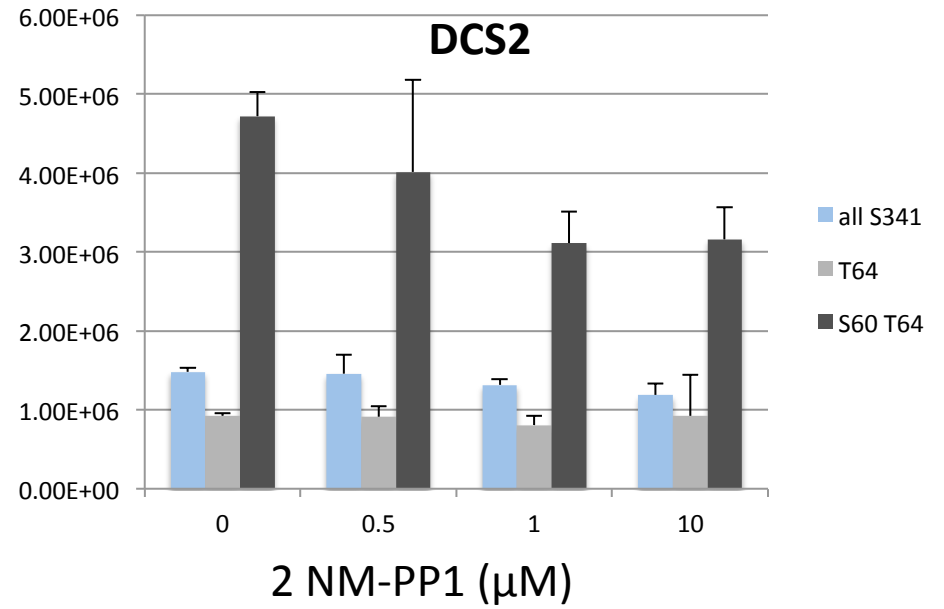
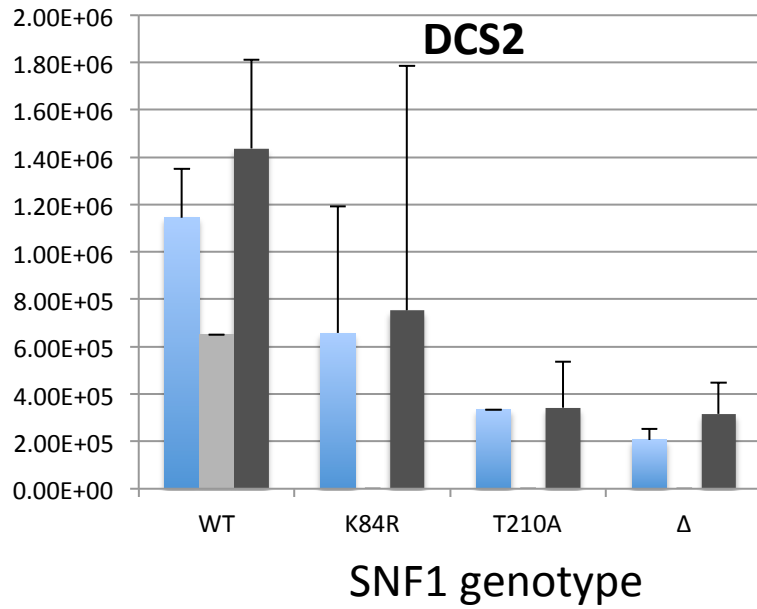


2 NM-PP1 (μM)



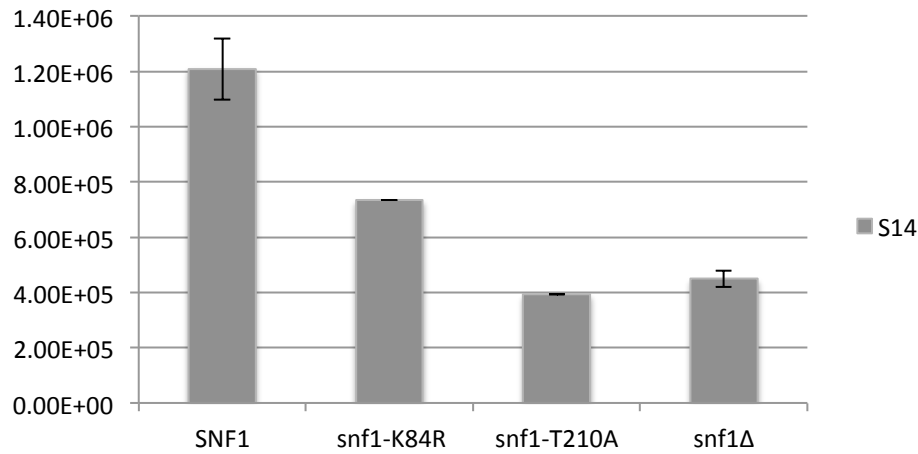
Min of derepression (48 = + 2 NM-PP1 (10 μM))

DCS2



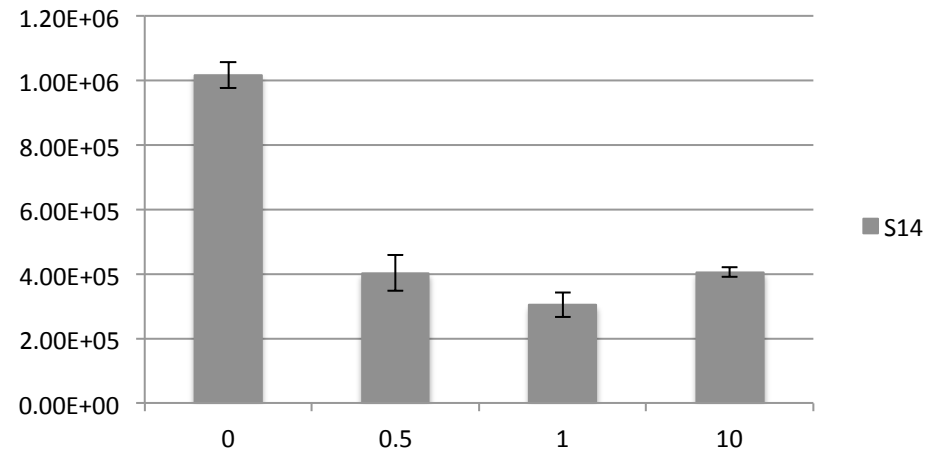
DHH1

DHH1



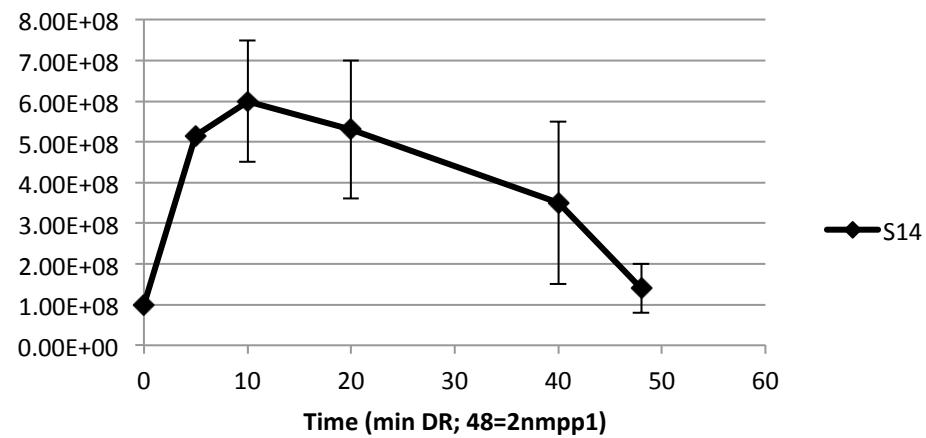
SNF1 genotype

DHH1



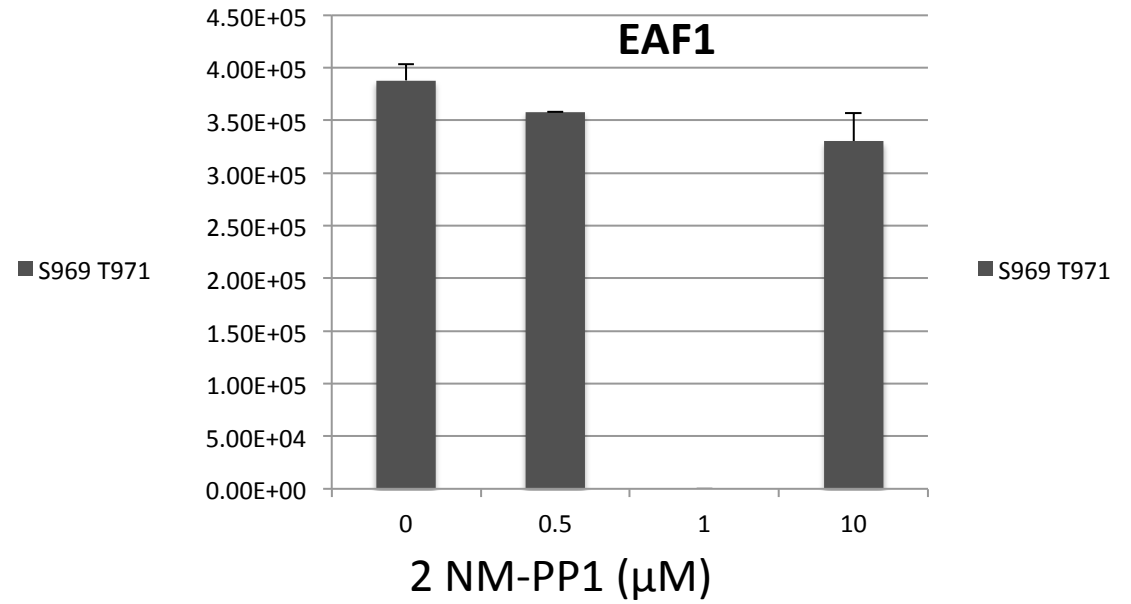
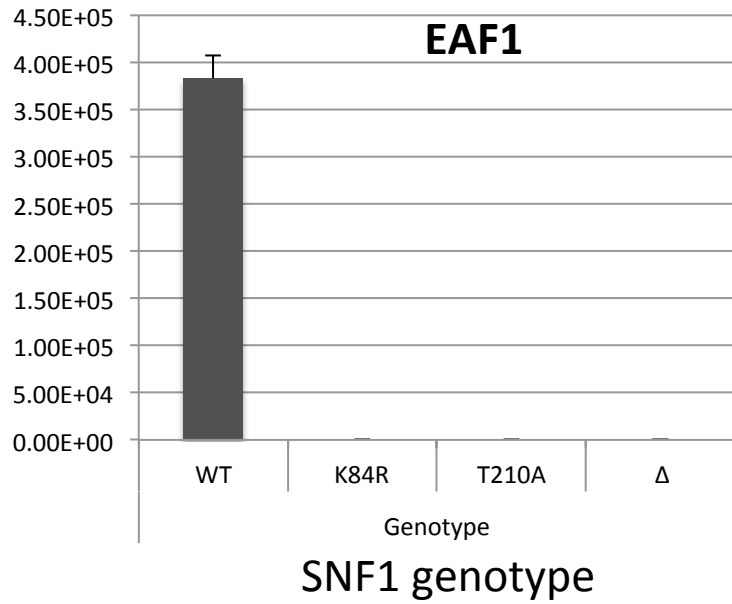
2 NM-PP1 (nM)

DHH1



Min of derepression (48 = + 2 NM-PP1 (10 μM))

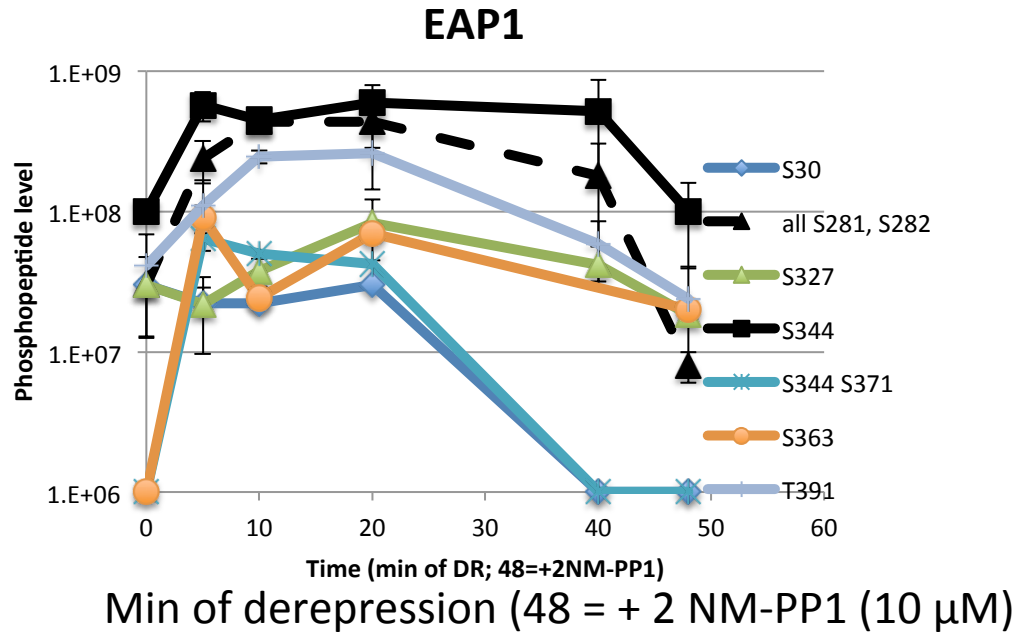
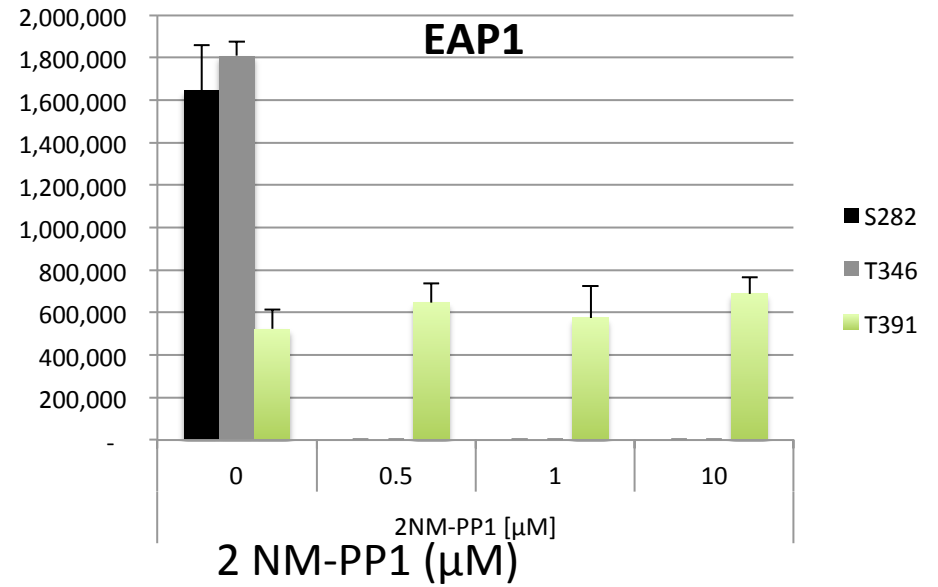
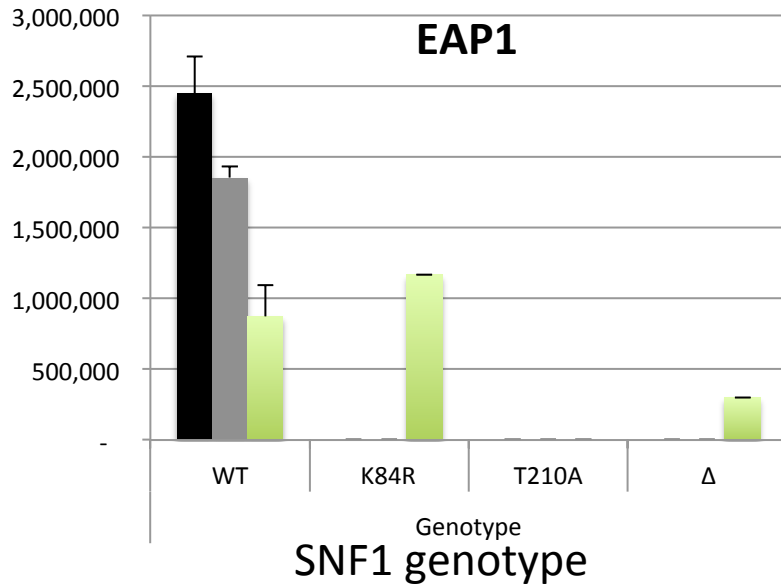
EAF1



No data

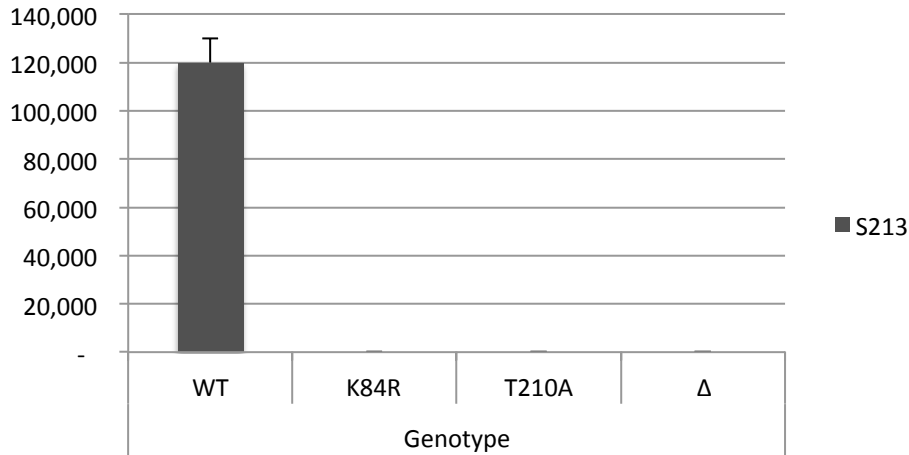
Min of derepression (48 = + 2 NM-PP1 (10 μM))

EAP1

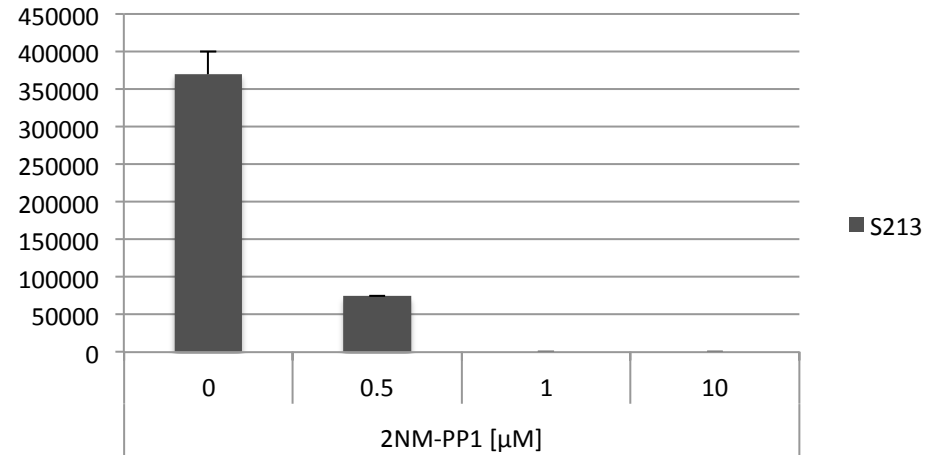


ECM32

ECM32

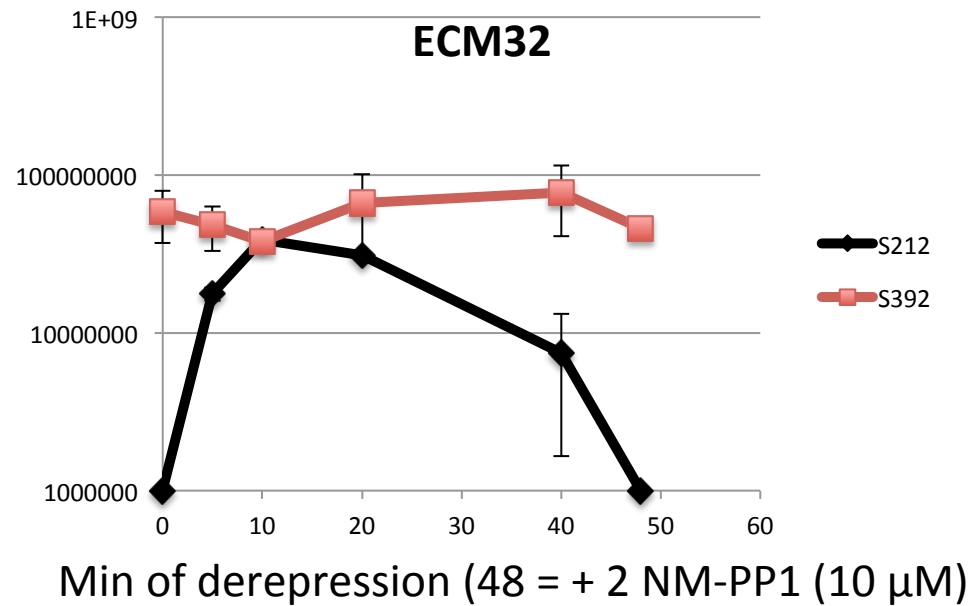


ECM32

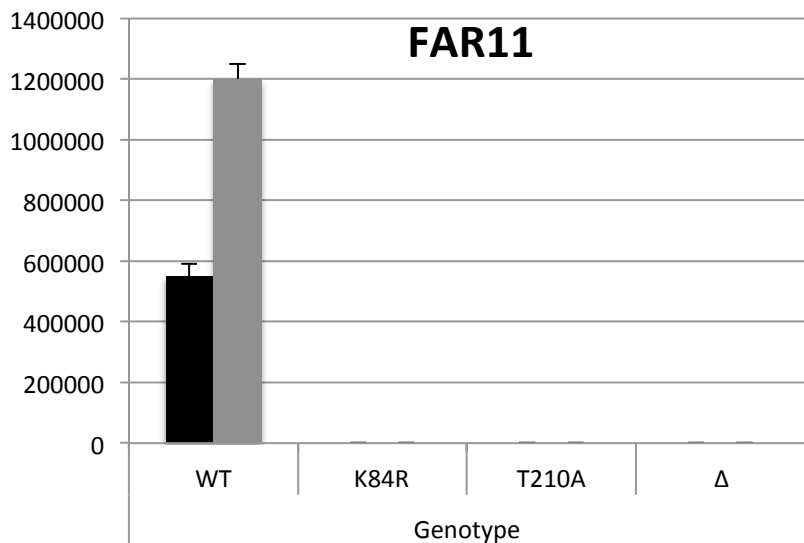


SNF1 genotype

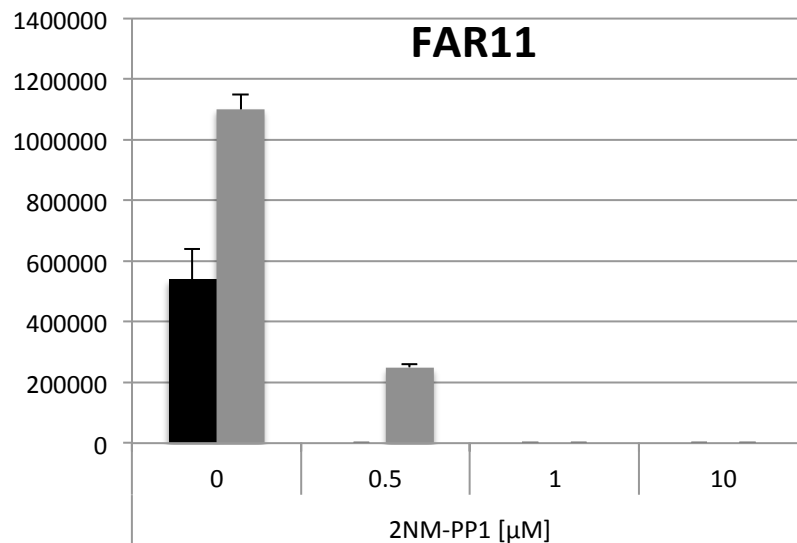
2 NM-PP1 (μM)



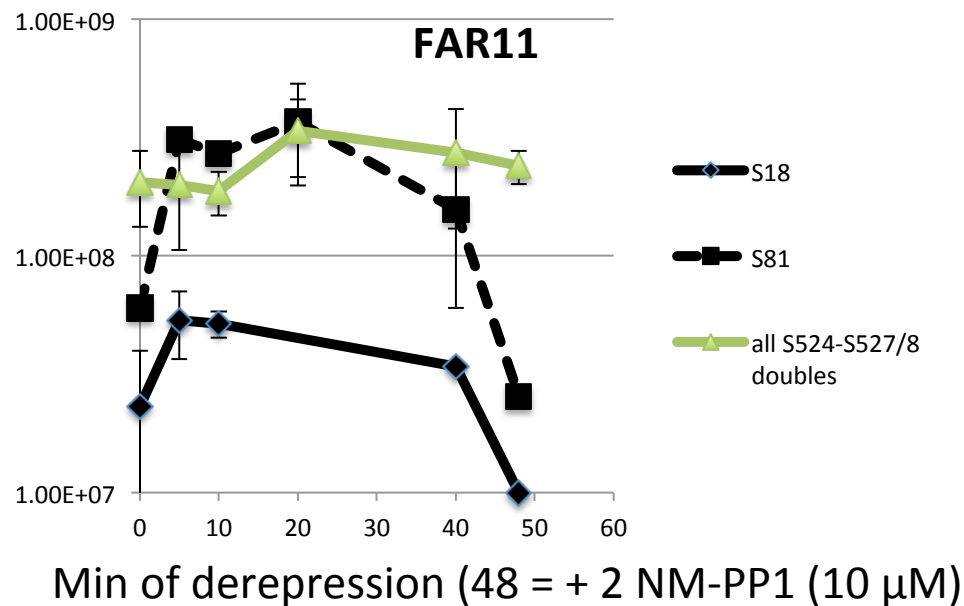
FAR11



SNF1 genotype

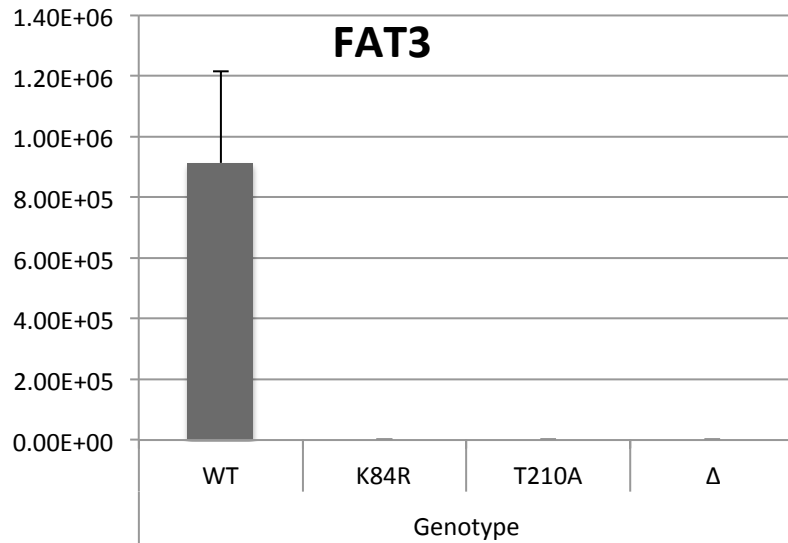


2 NM-PP1 (μM)

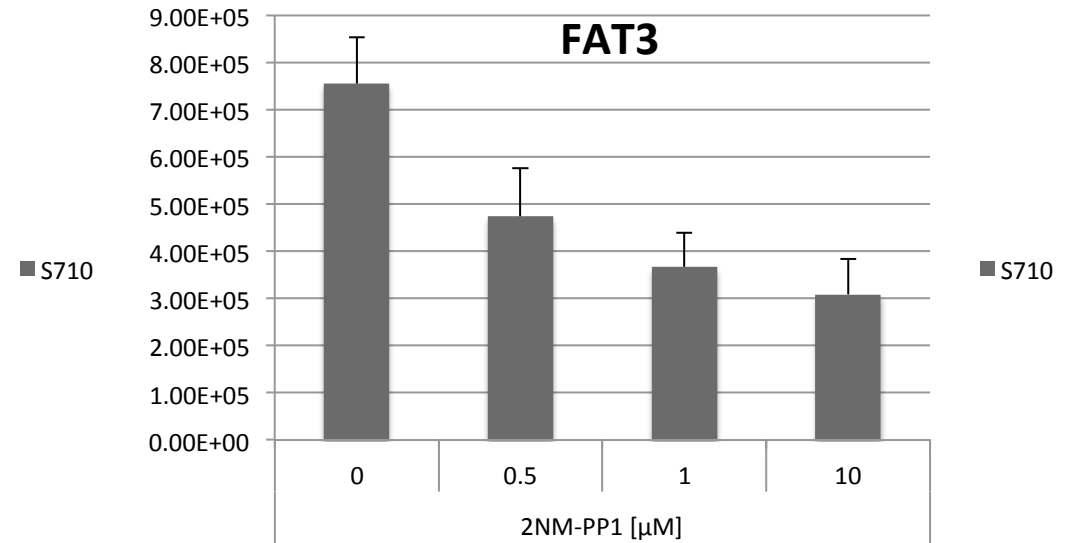


Min of derepression (48 = + 2 NM-PP1 (10 μM))

FAT3



SNF1 genotype



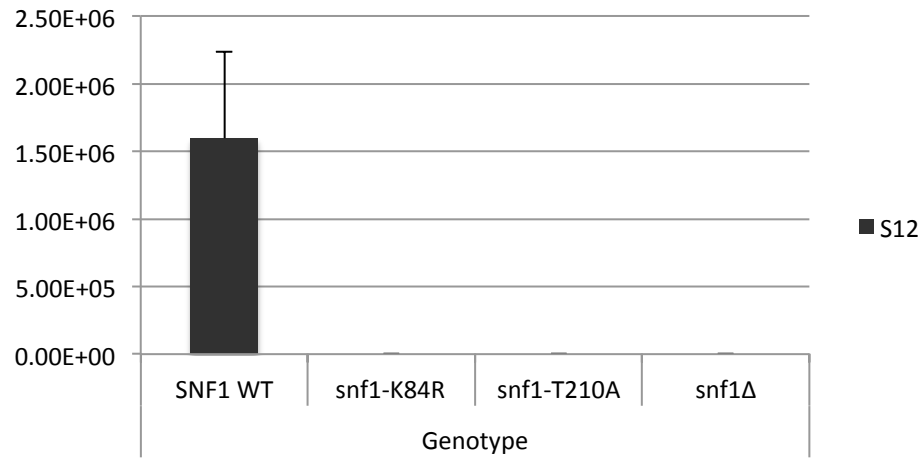
2 NM-PP1 (μ M)

No data

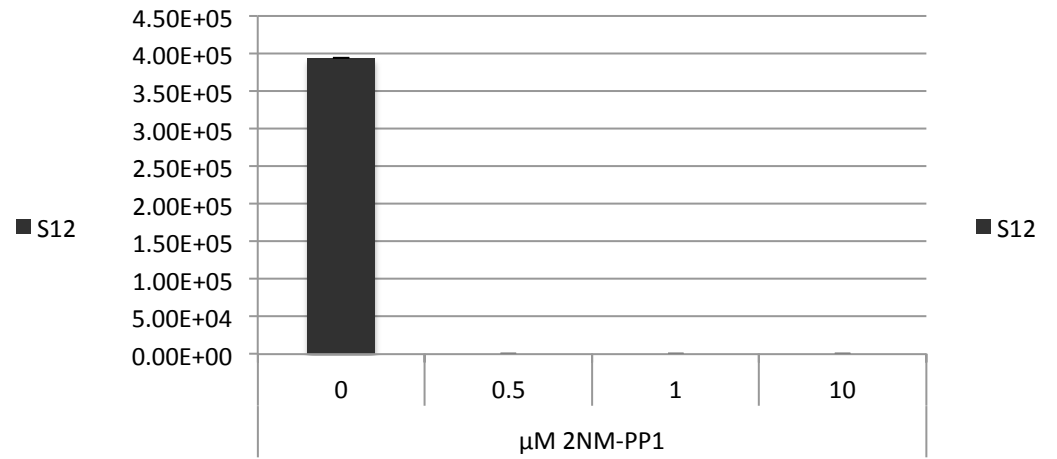
Min of derepression (48 = + 2 NM-PP1 (10 μ M))

FBP1

FBP1



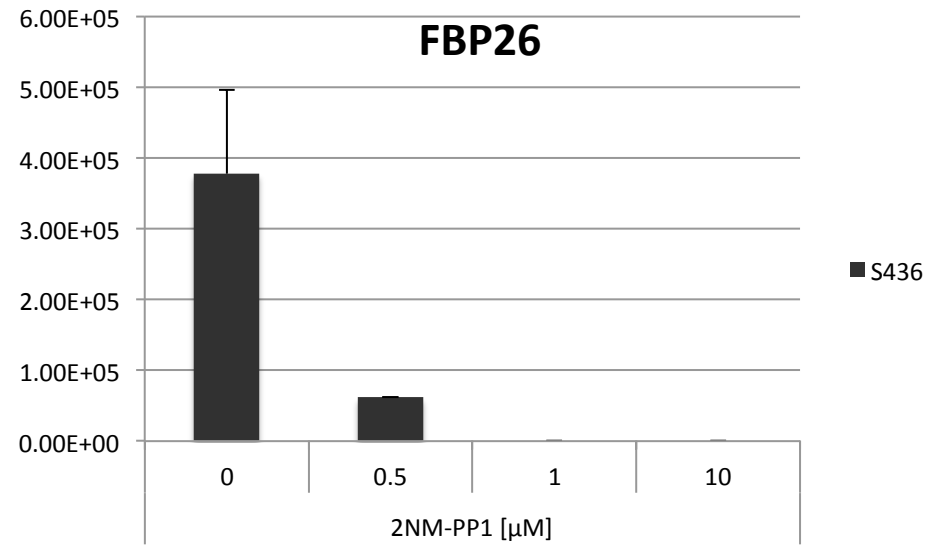
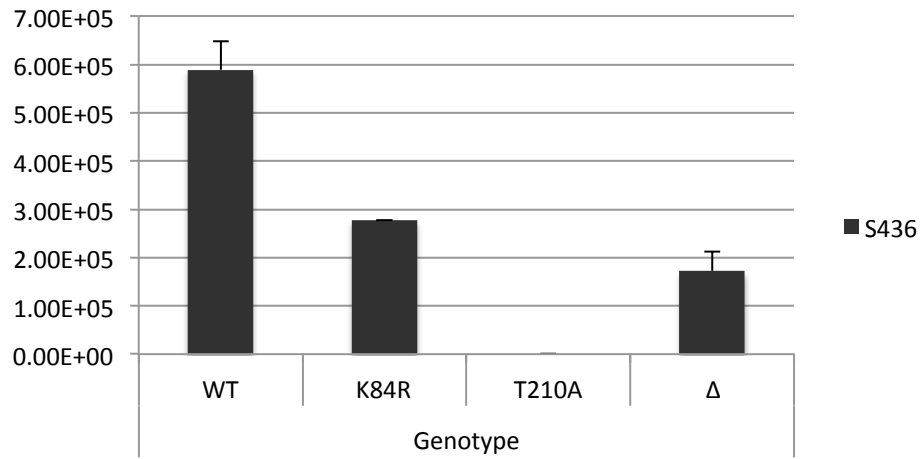
FBP1



No data for FBP1 in time course experiment

FBP26

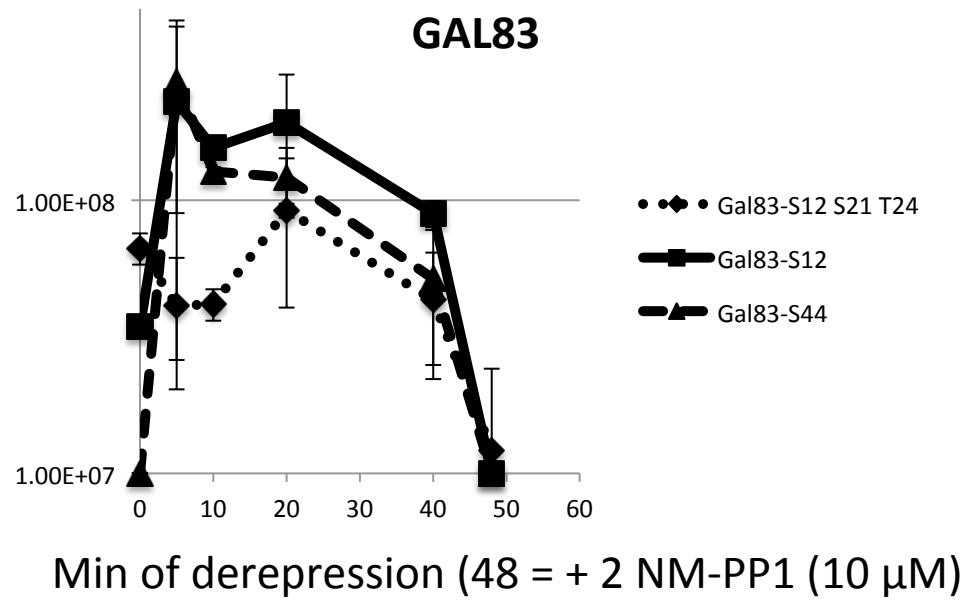
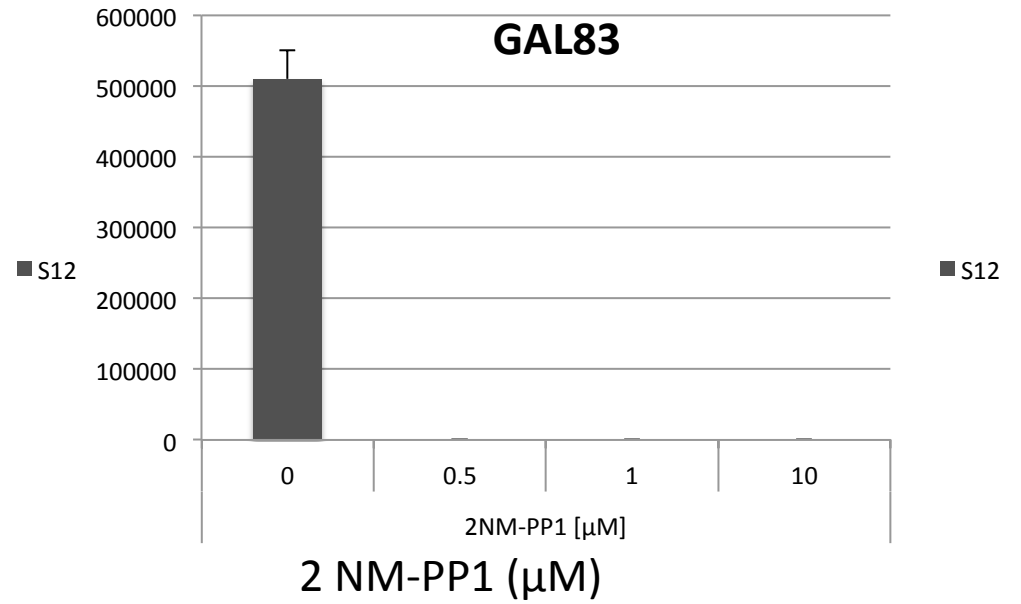
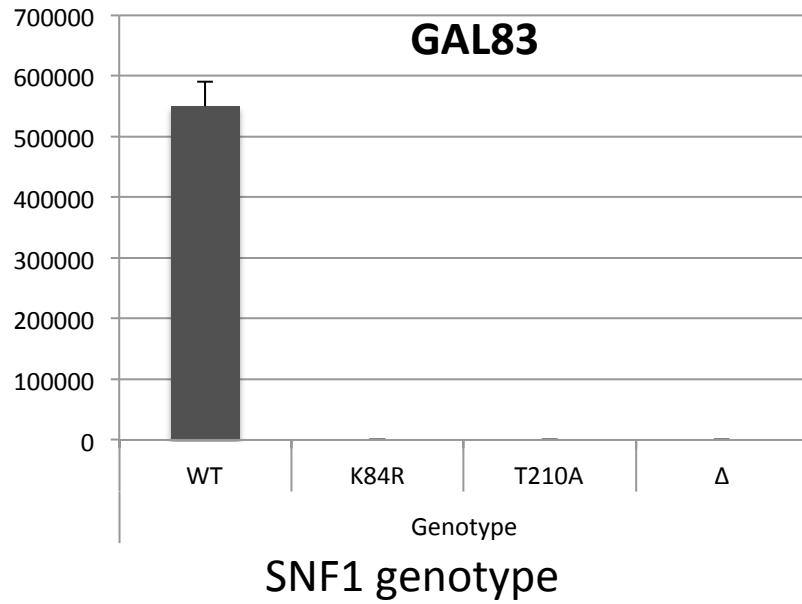
FBP26



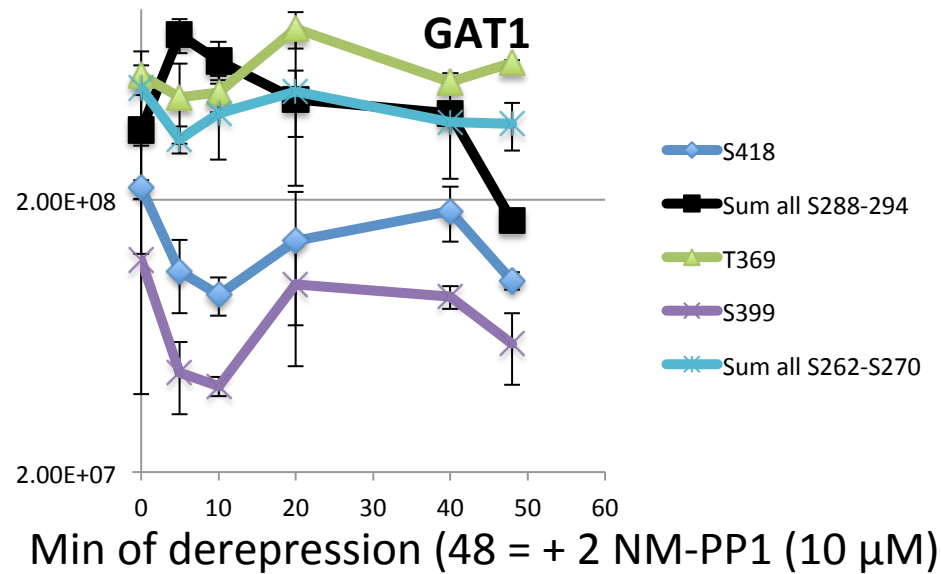
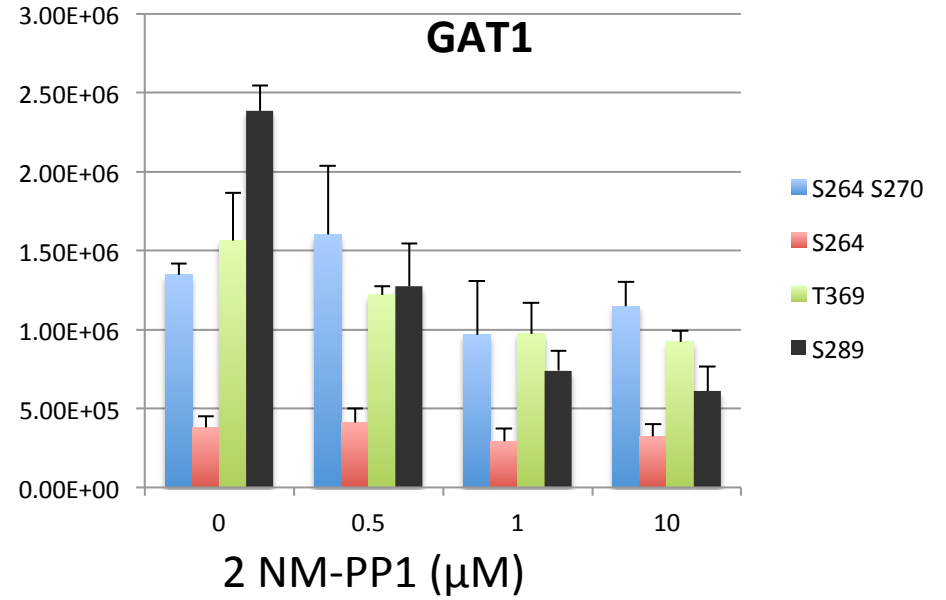
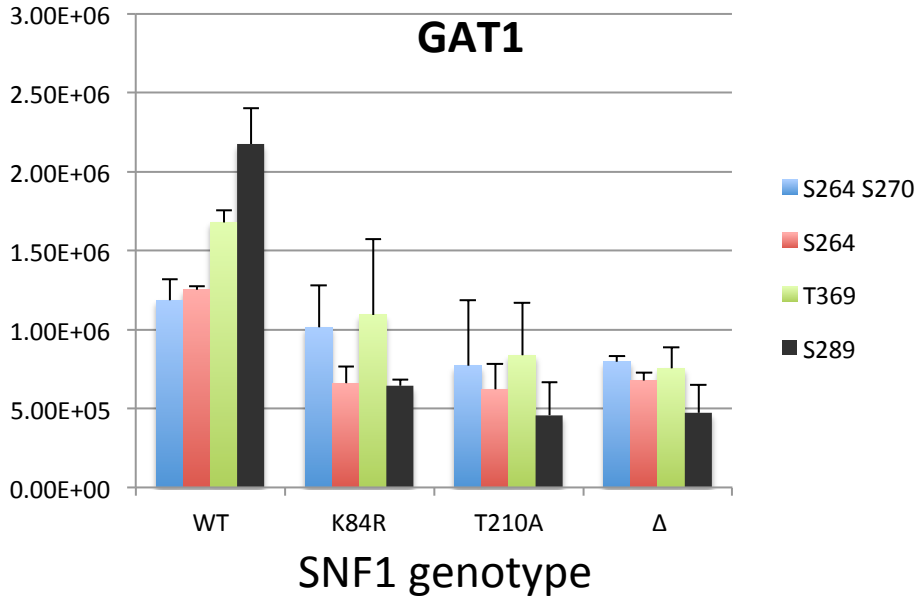
No data

Min of derepression (48 = + 2 NM-PP1 (10 μM))

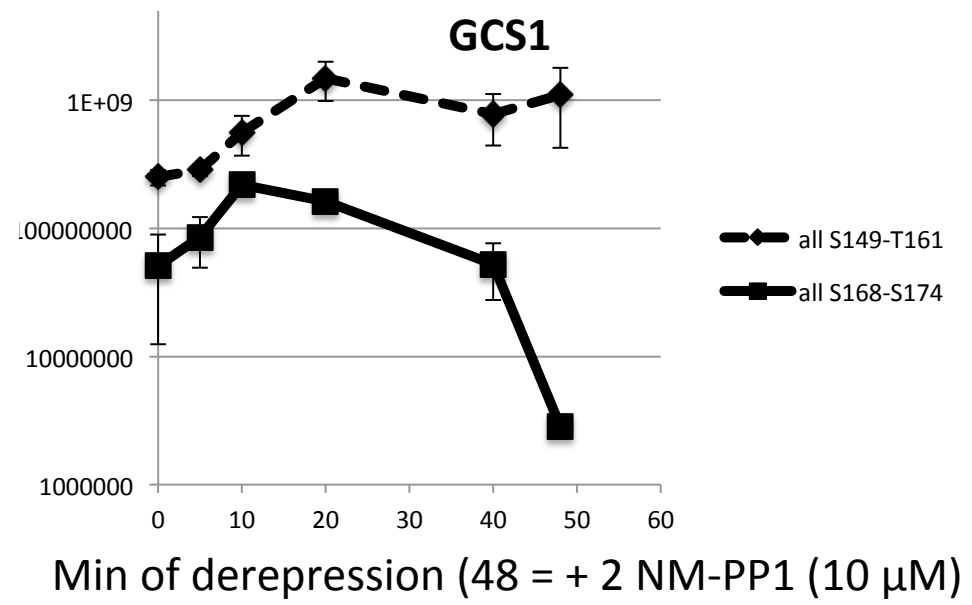
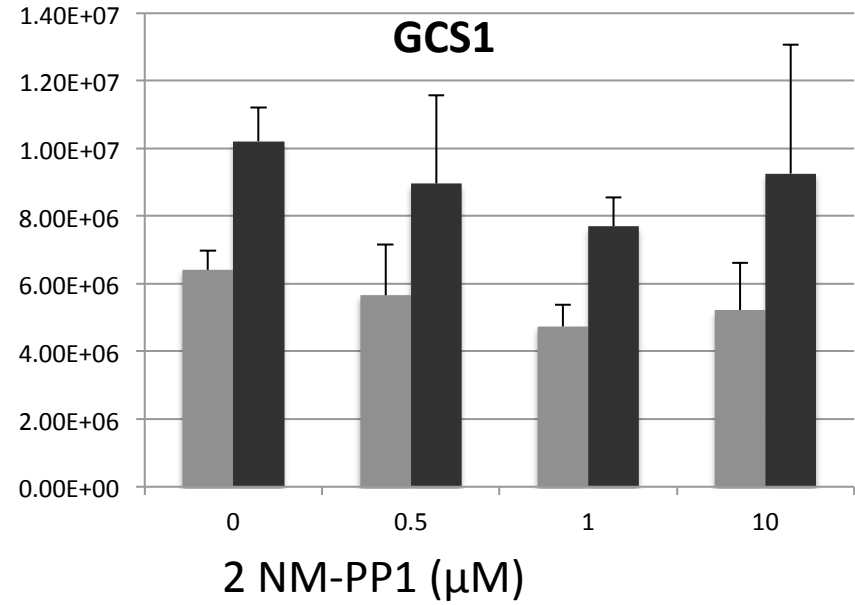
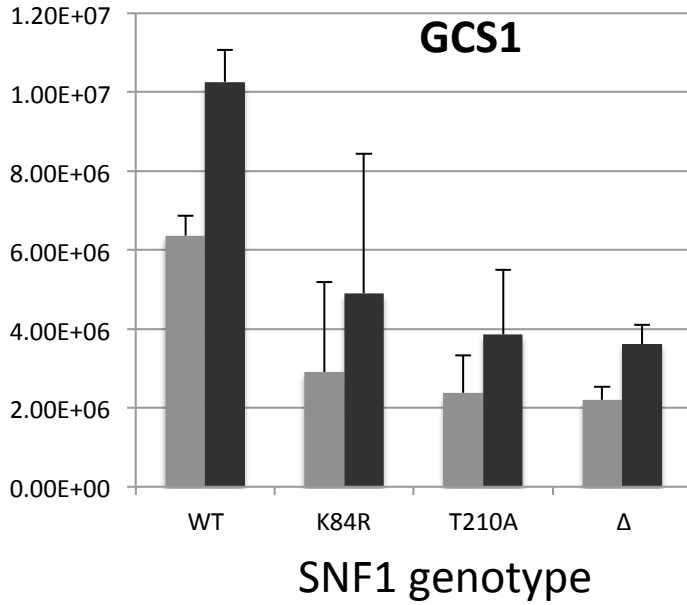
GAL83



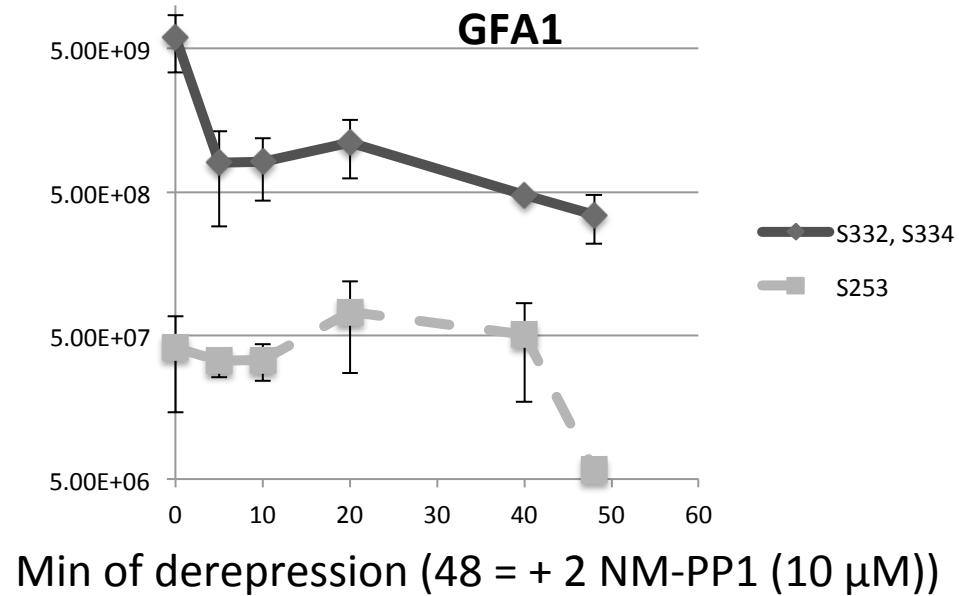
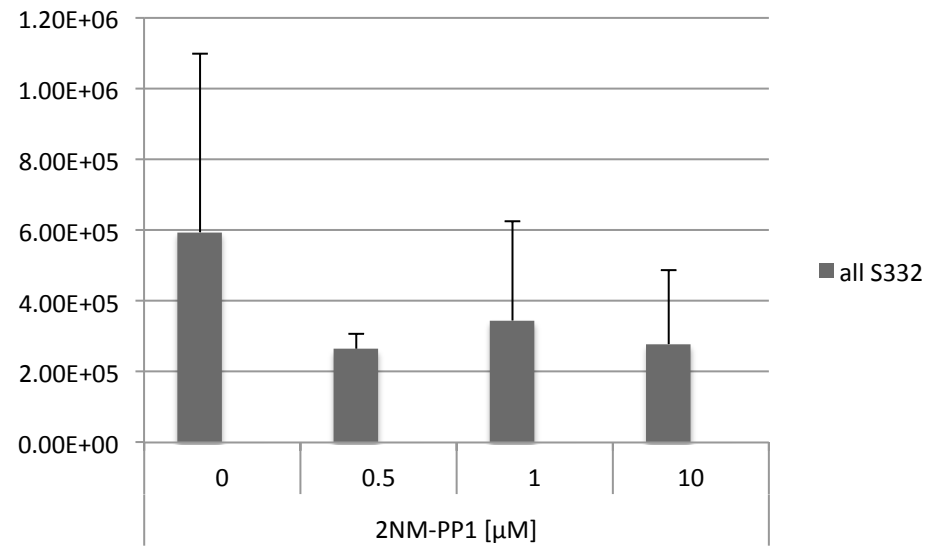
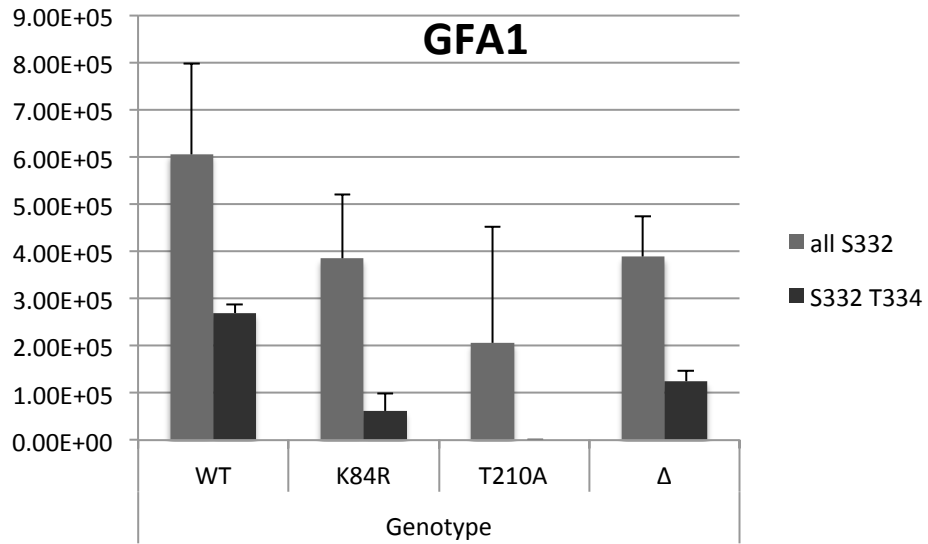
GAT1



GCS1

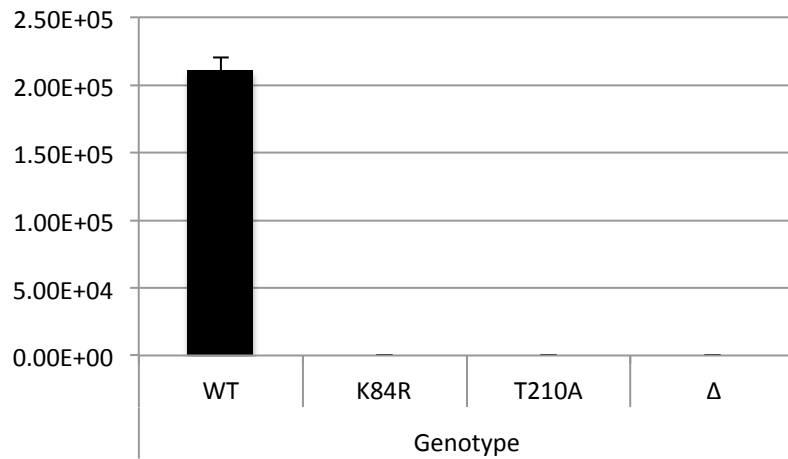


GFA1

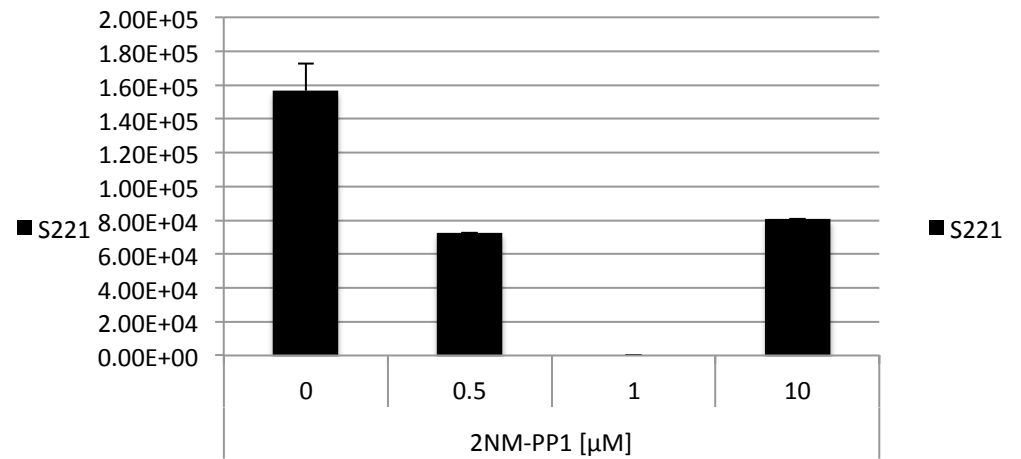


GIP2

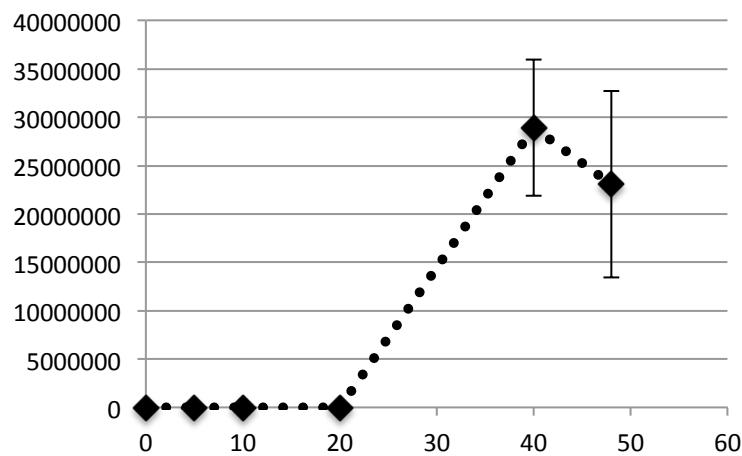
GIP2



GIP2



GIP2

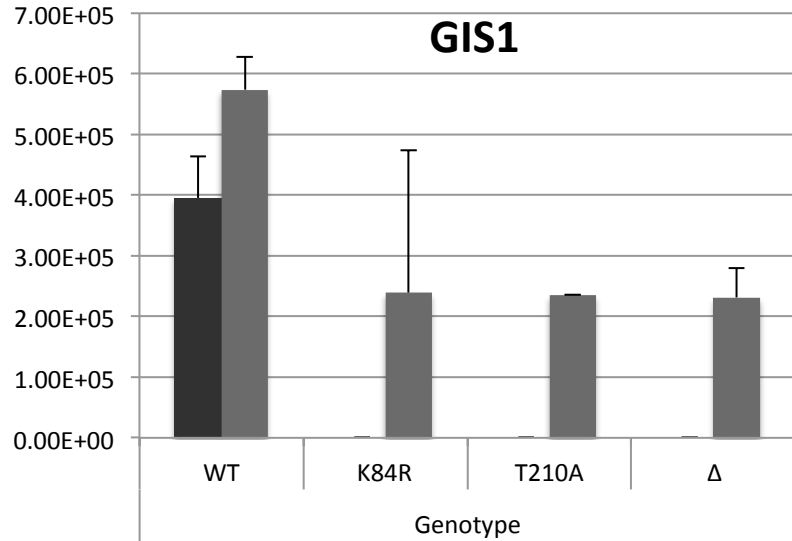


NB: different phosphosite
Than in experiments 1 and 2.

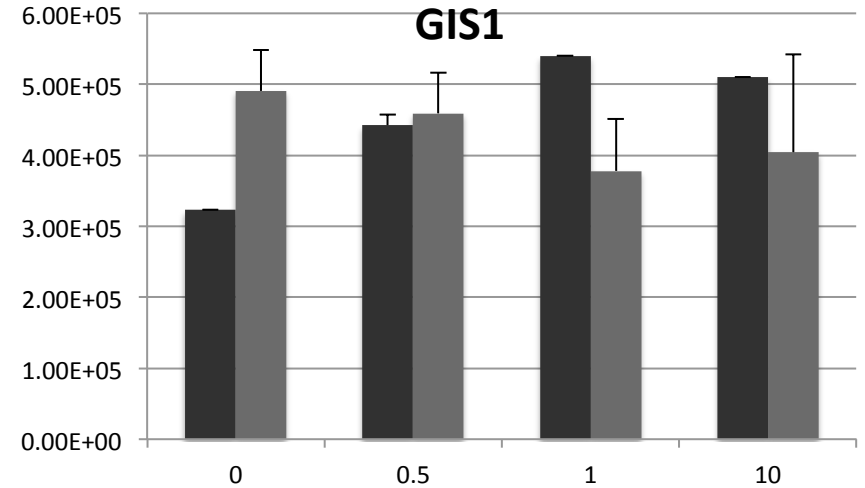
••• S137

Min of derepression (48 = + 2 NM-PP1 (10 μM))

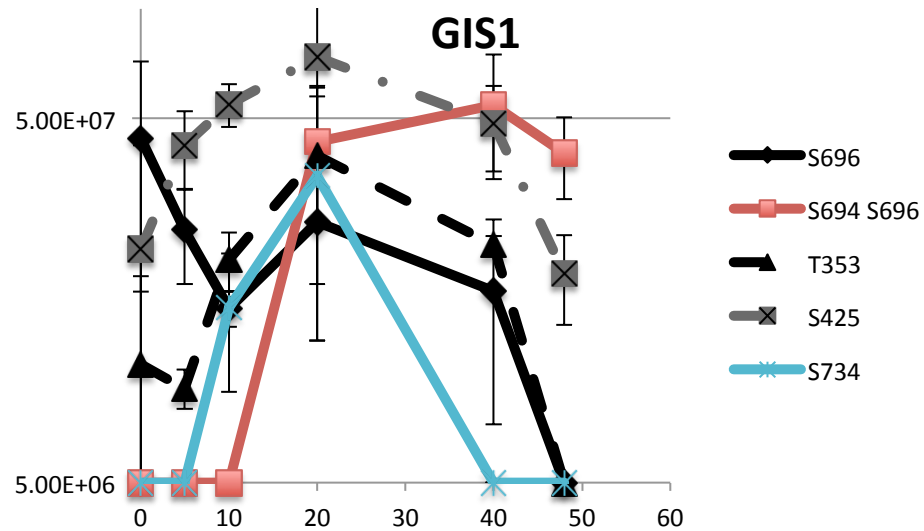
GIS1



SNF1 genotype

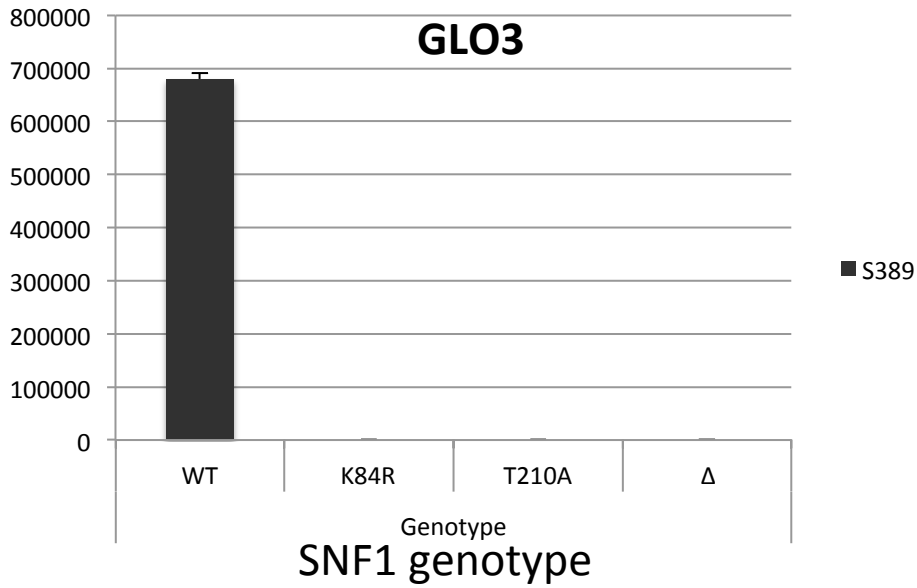


2 NM-PP1 (μM)



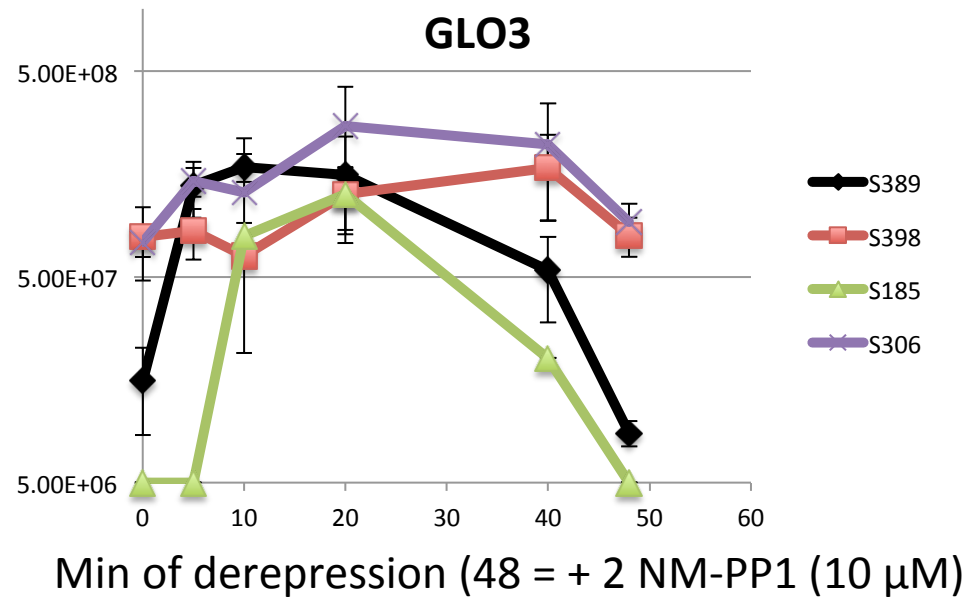
Min of derepression (48 = + 2 NM-PP1 (10 μM))

GLO3

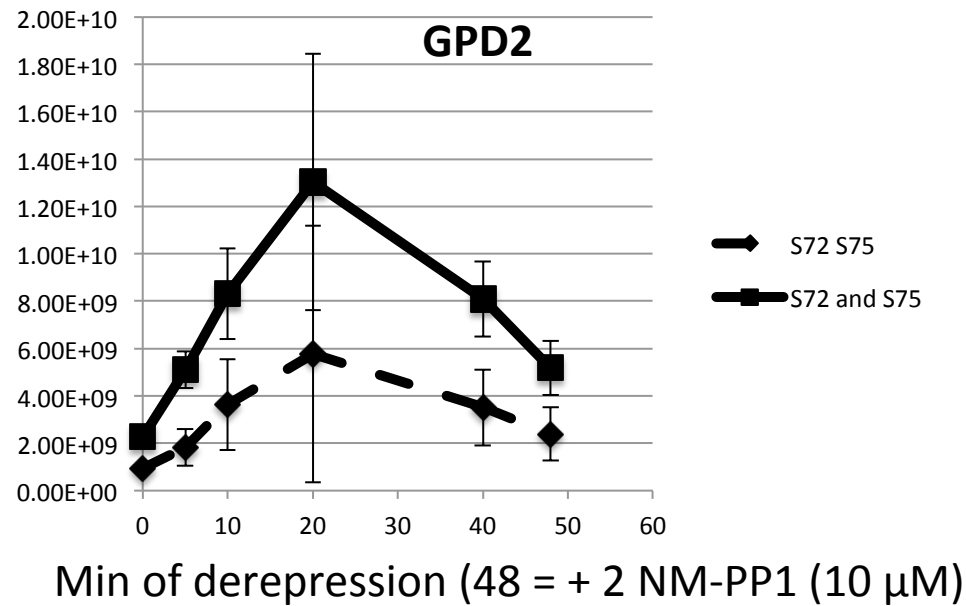
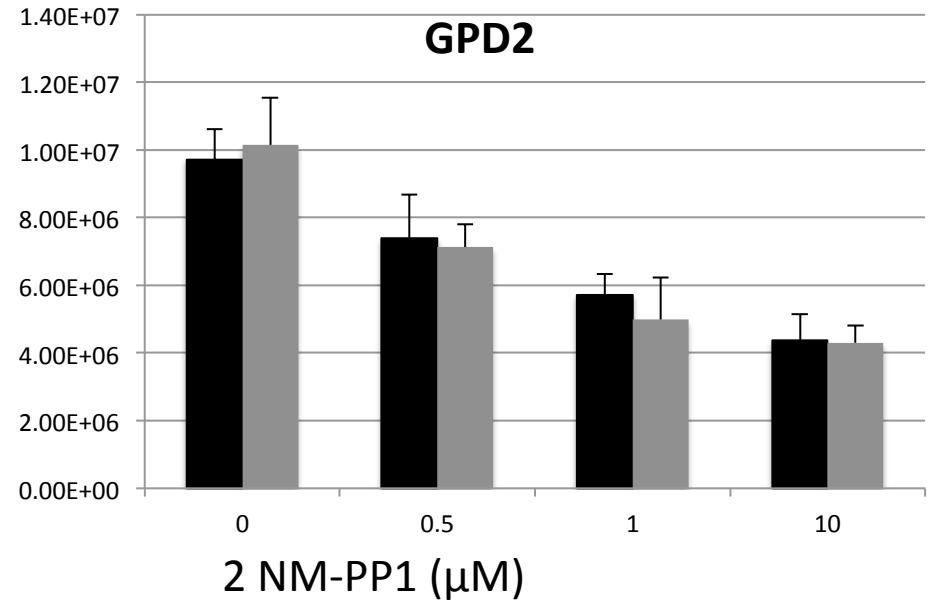
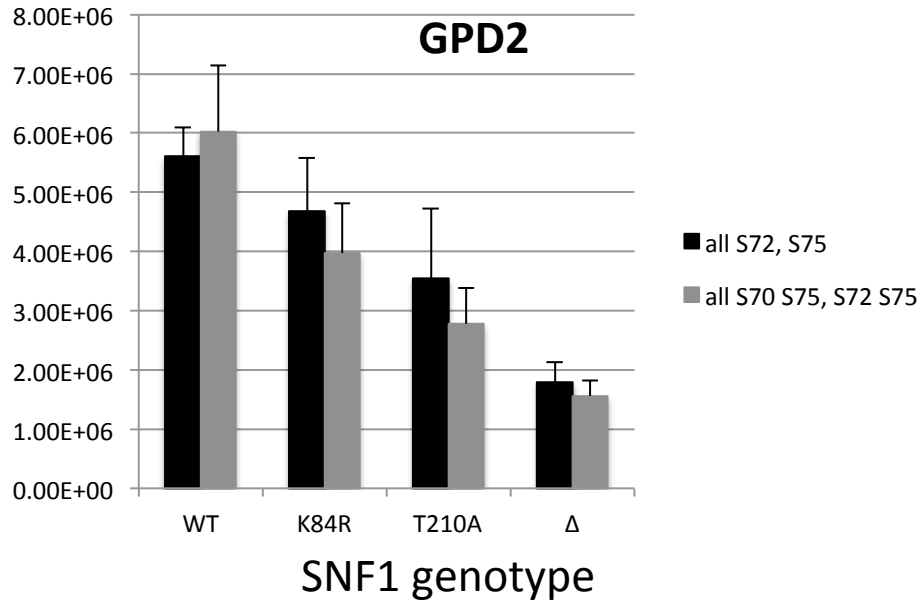


No data

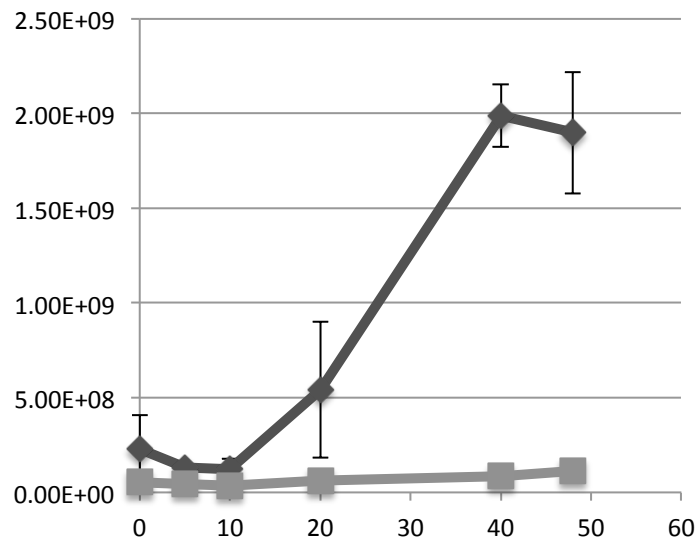
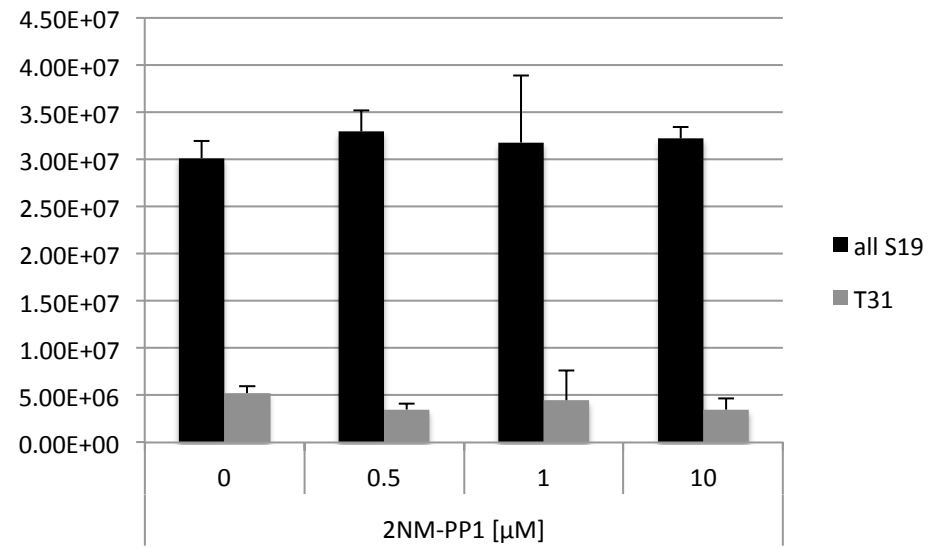
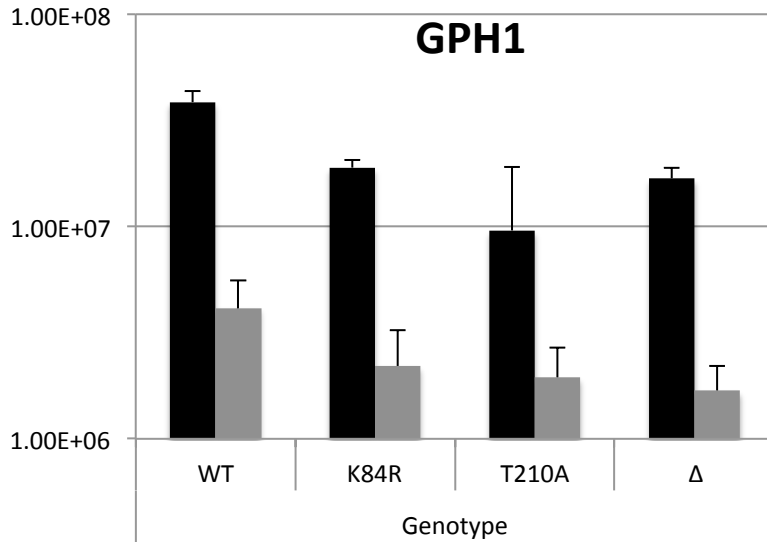
2 NM-PP1 (μM)



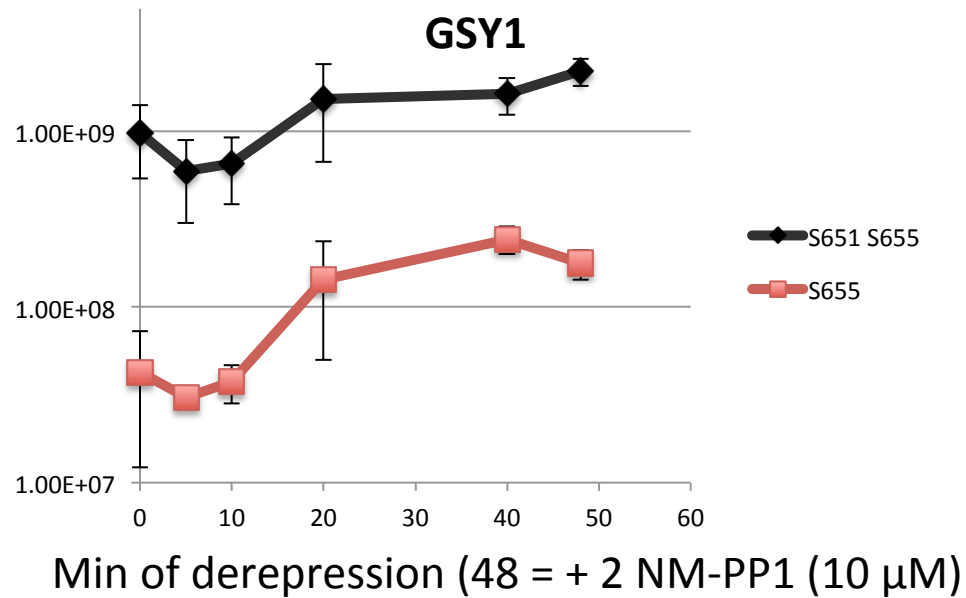
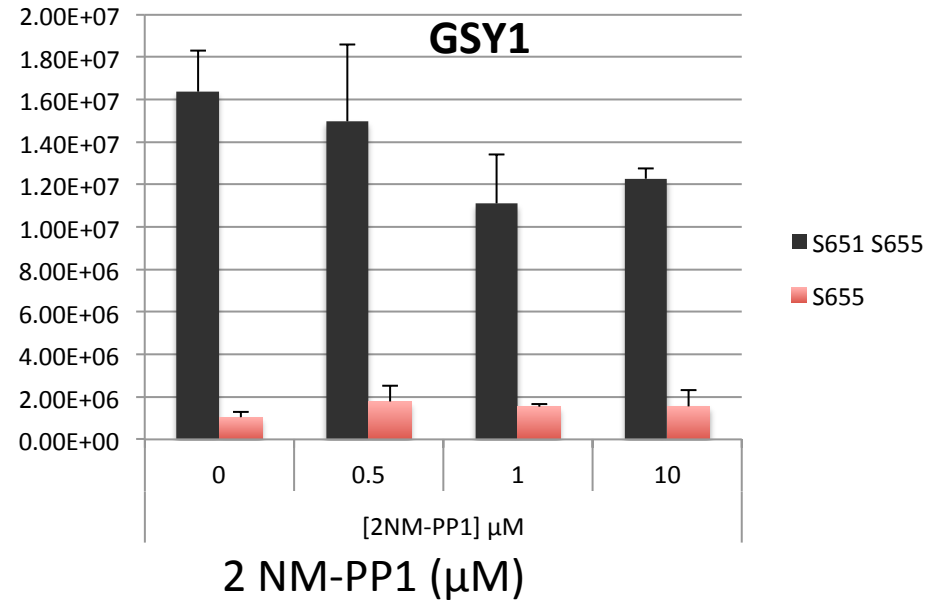
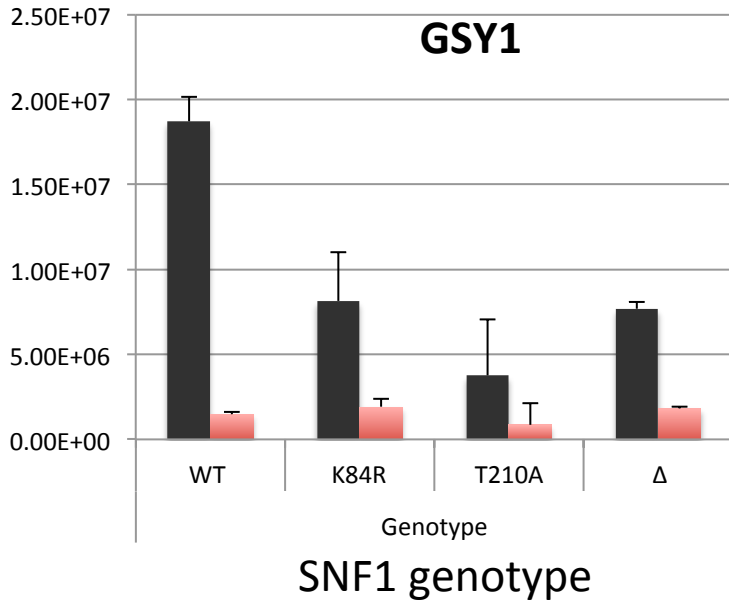
GPD2



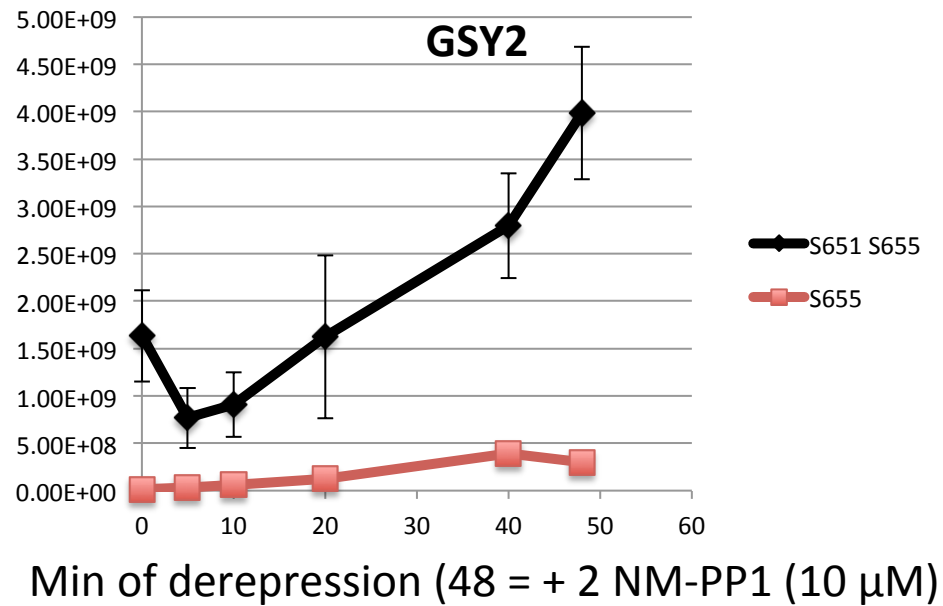
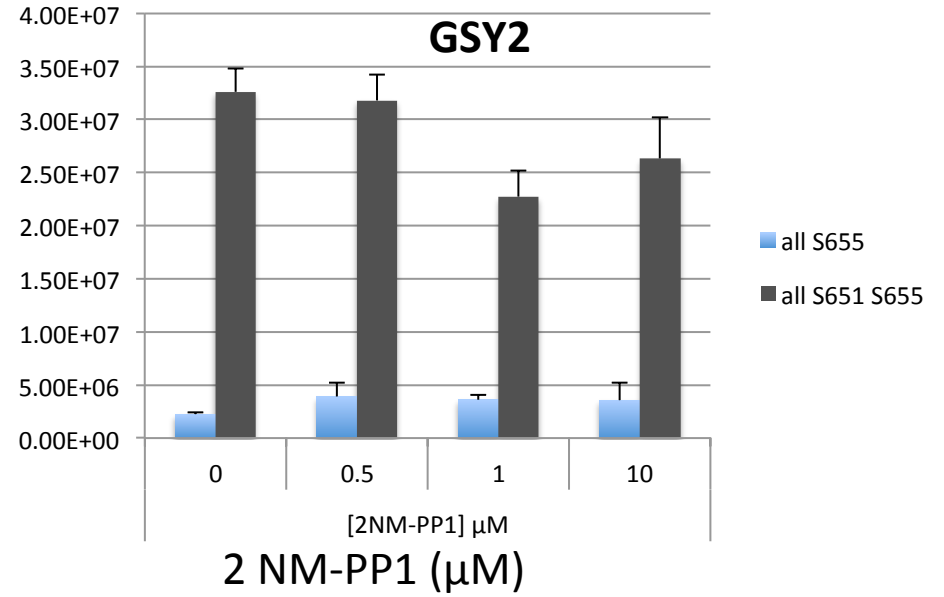
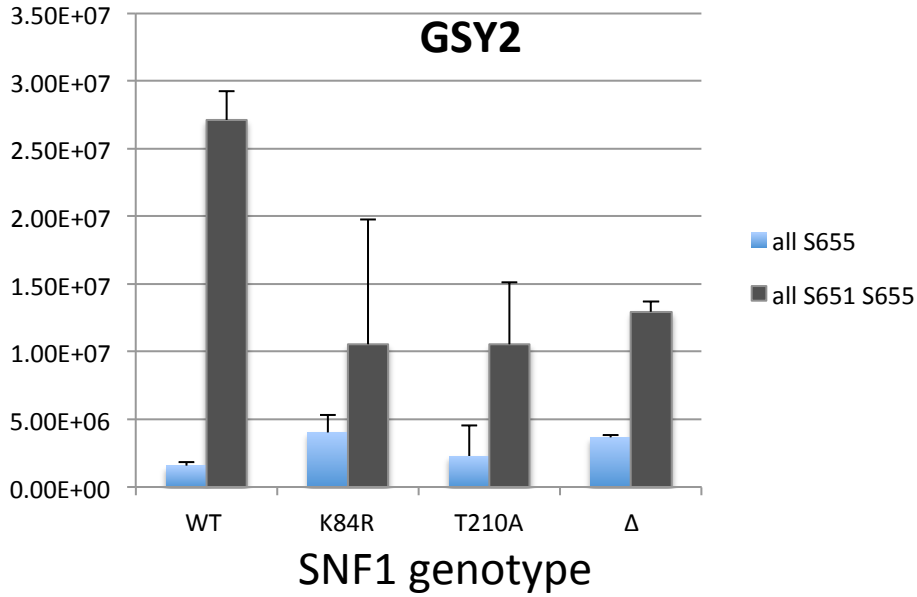
GPH1



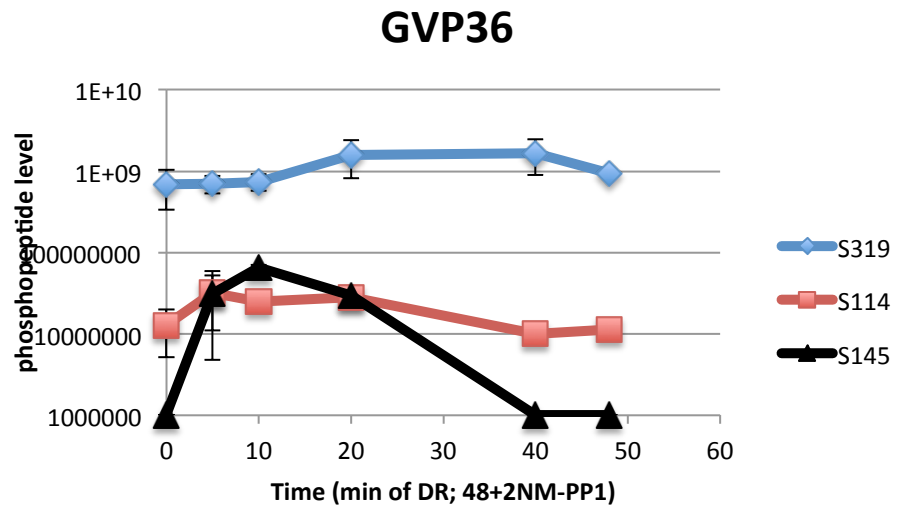
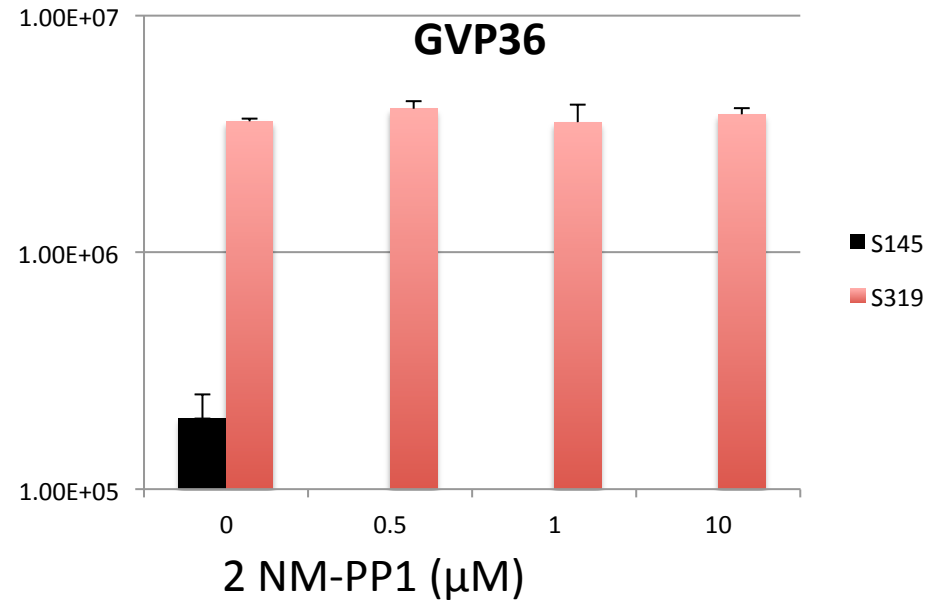
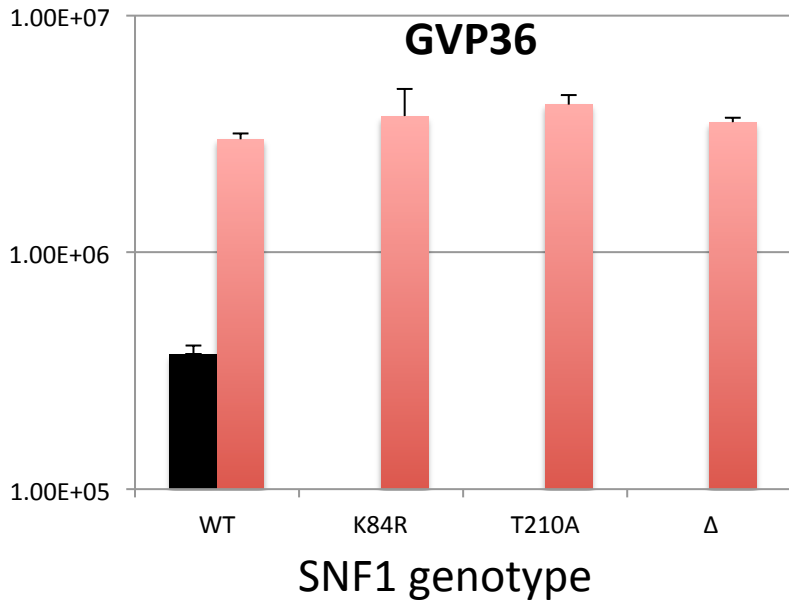
GSY1



GSY2

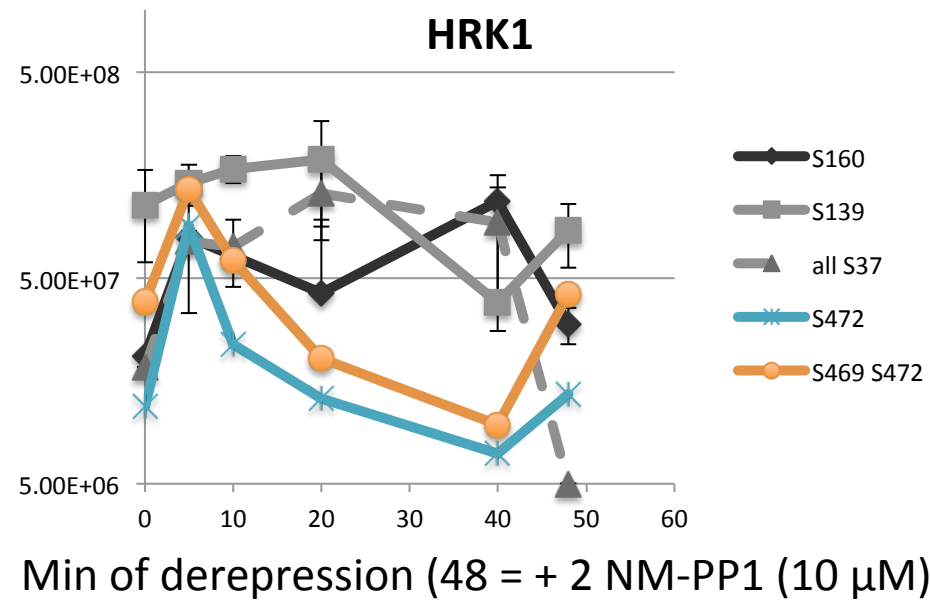
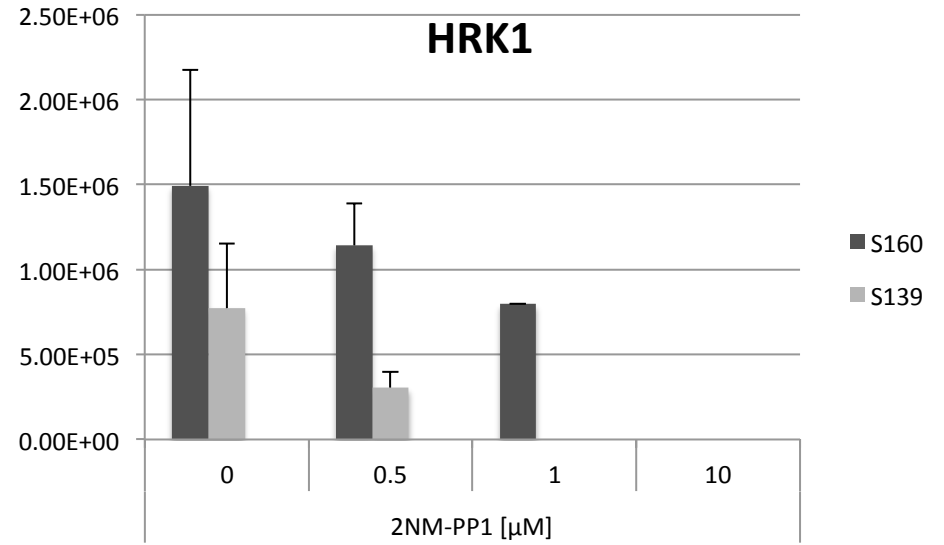
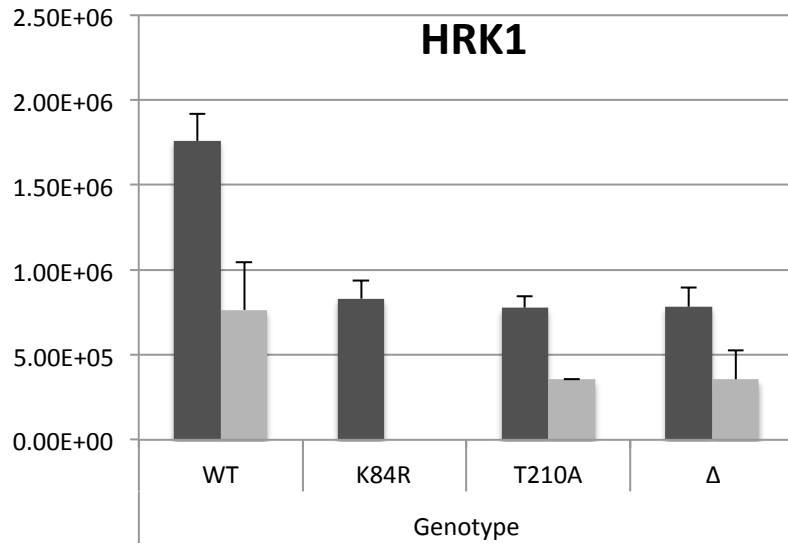


GVP36



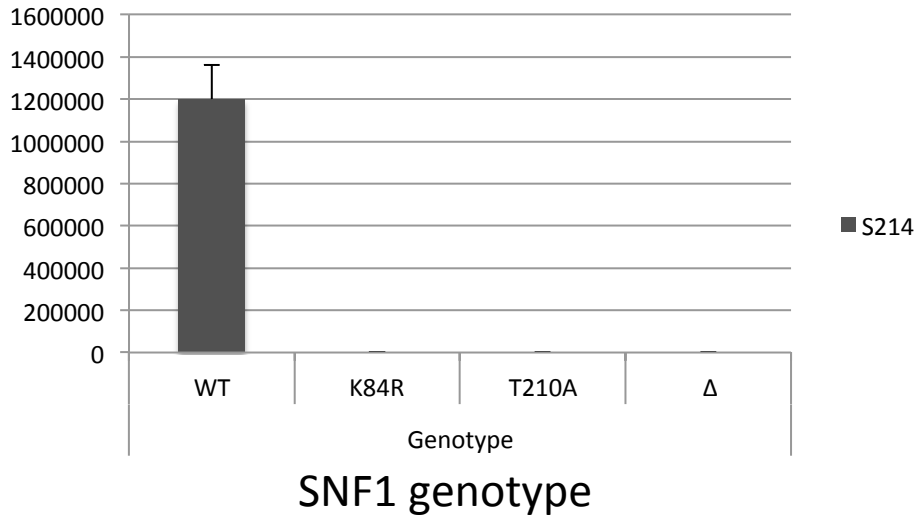
Min of derepression (48 = + 2 NM-PP1 (10 μM))

HRK1

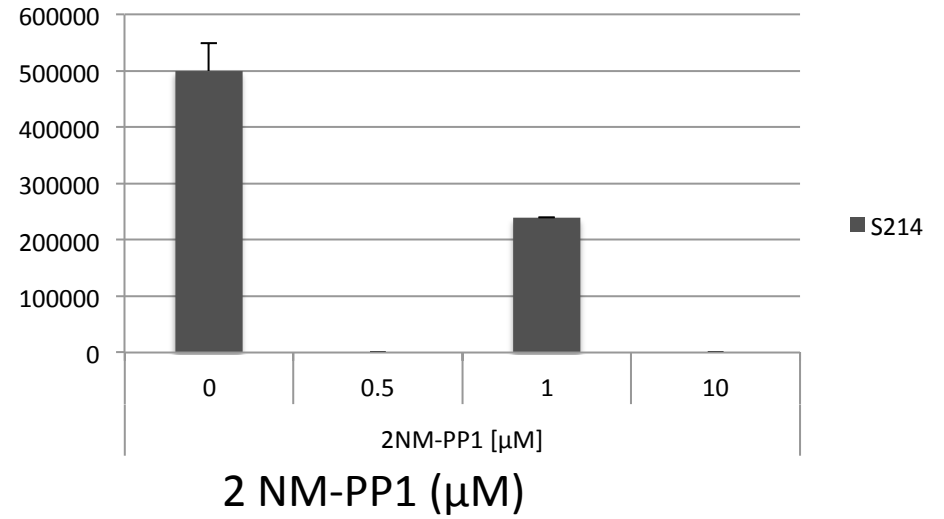


HSP42

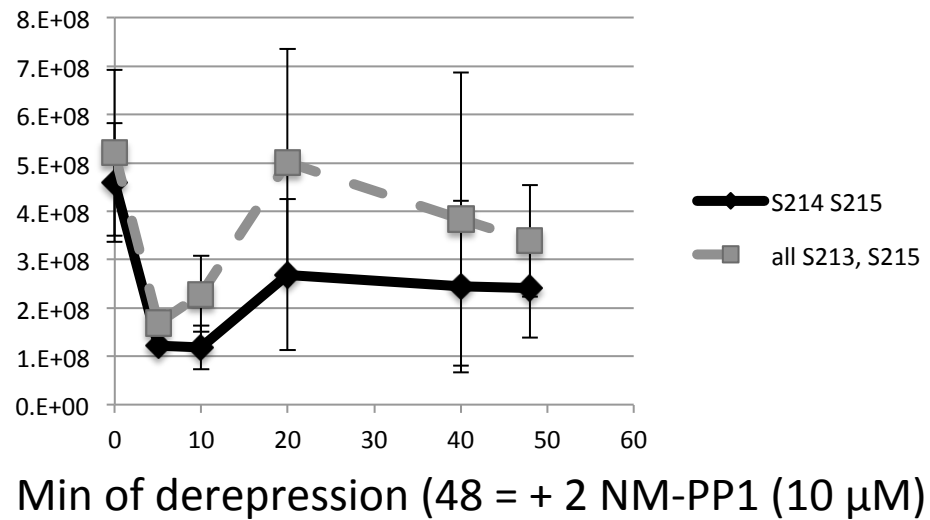
HSP42



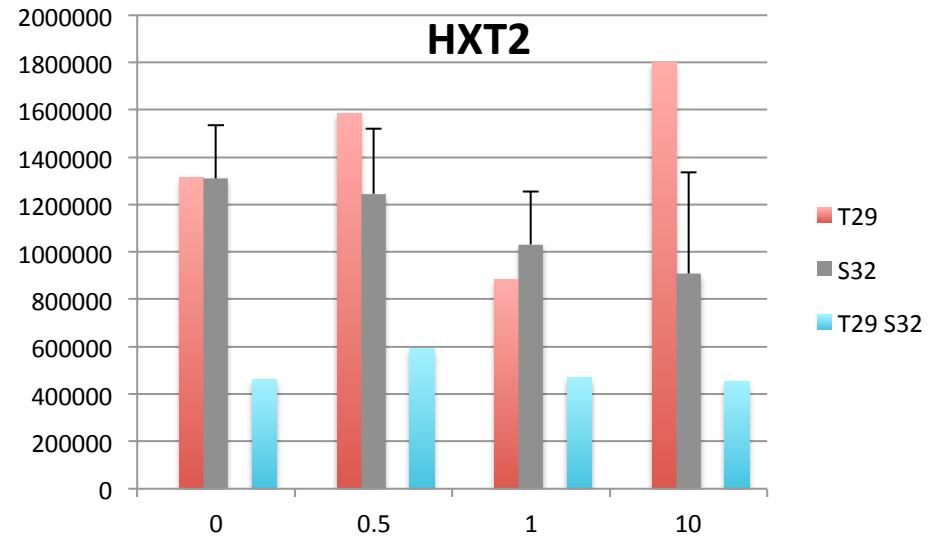
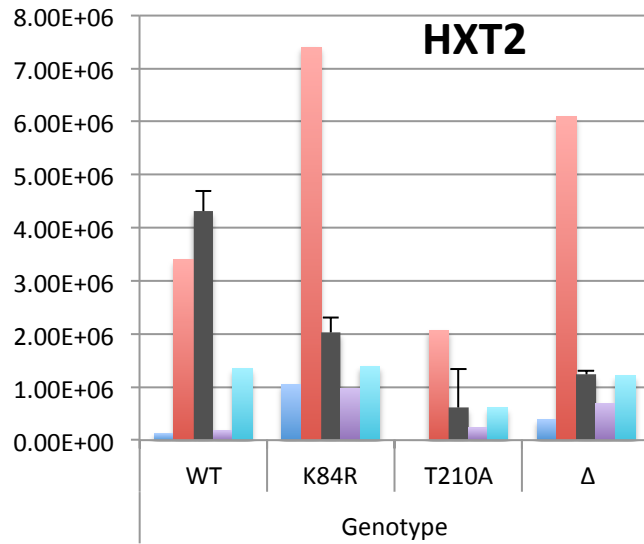
HSP42



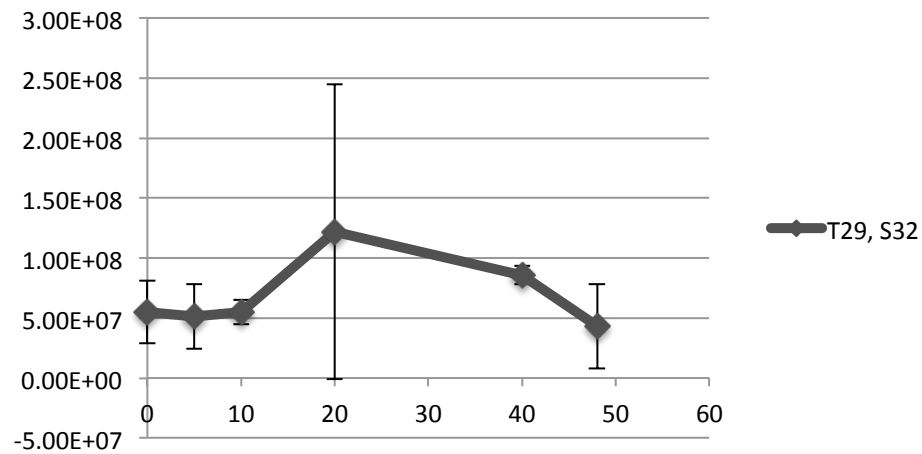
HSP42



HXT2

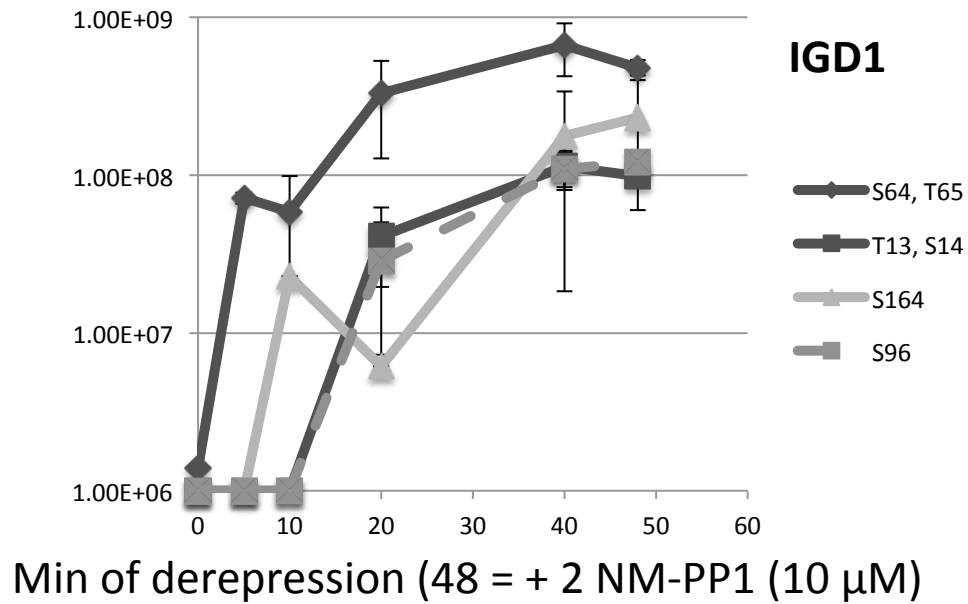
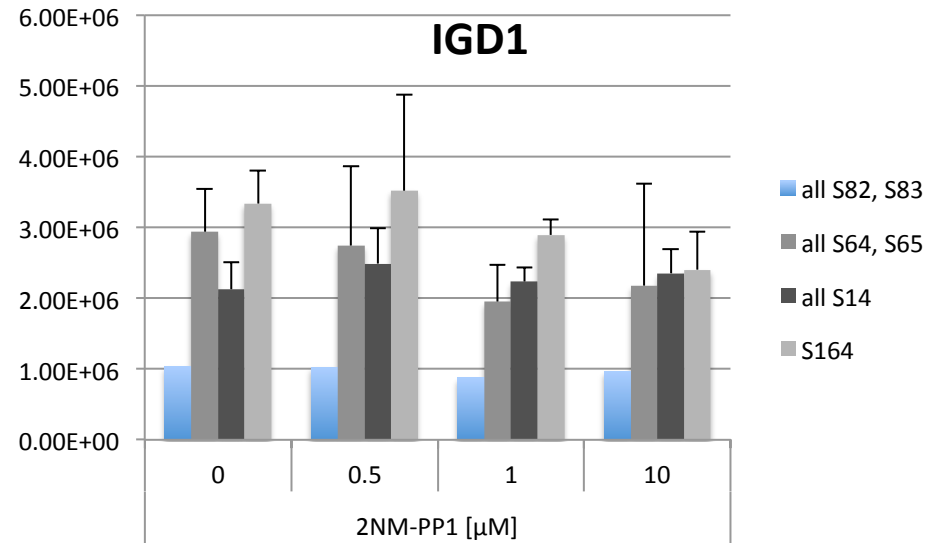
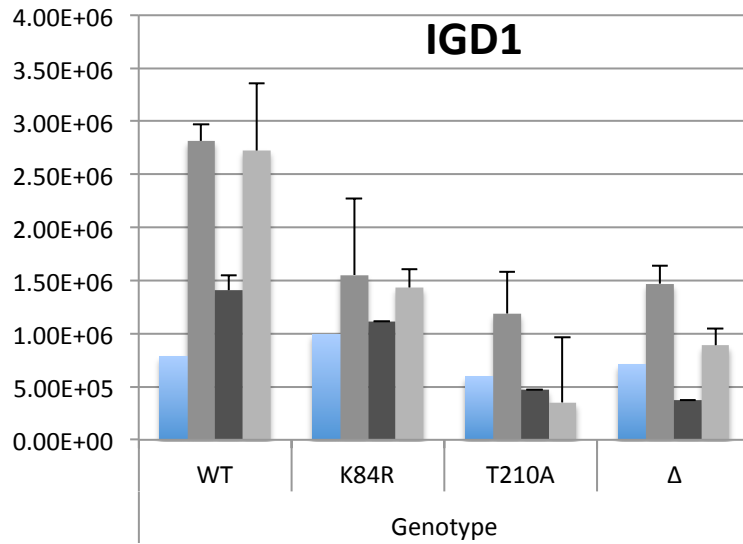


HXT2

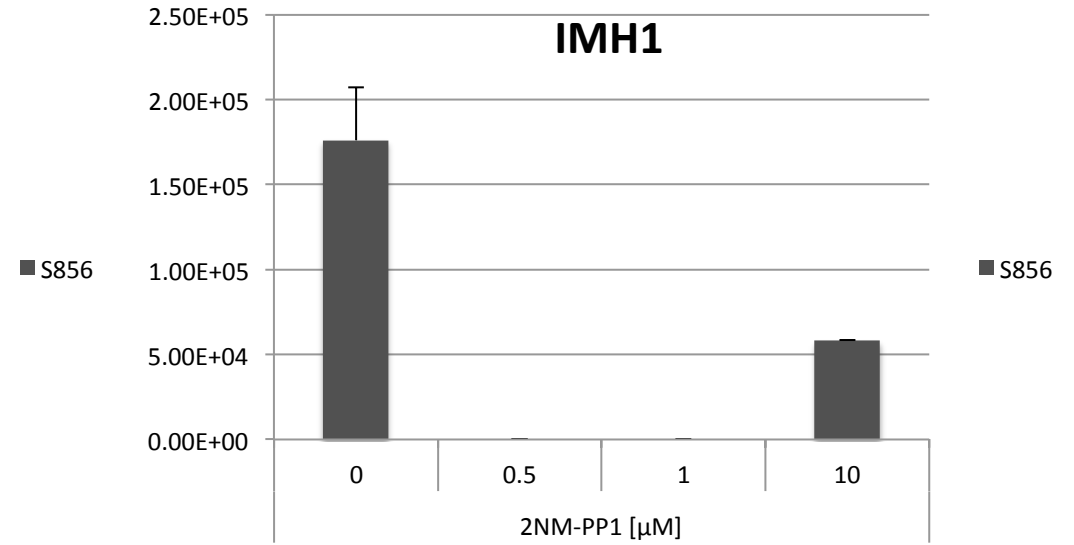
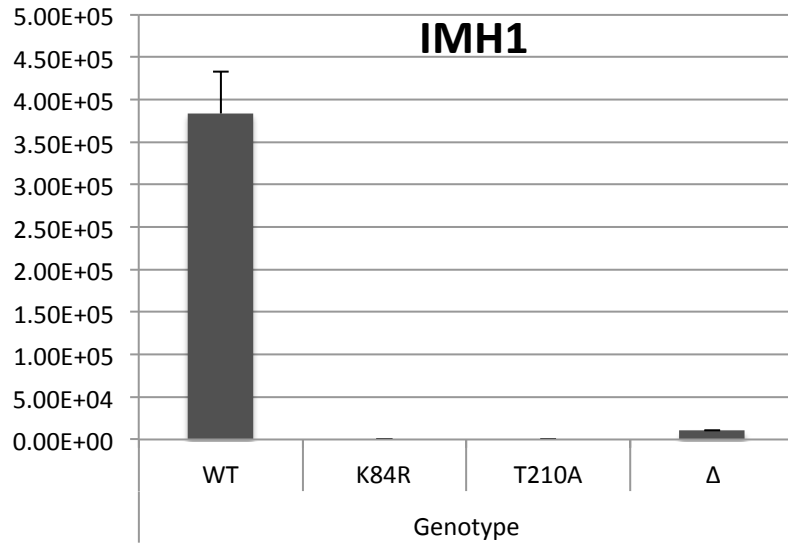


Min of derepression (48 = + 2 NM-PP1 (10 μM))

IGD1



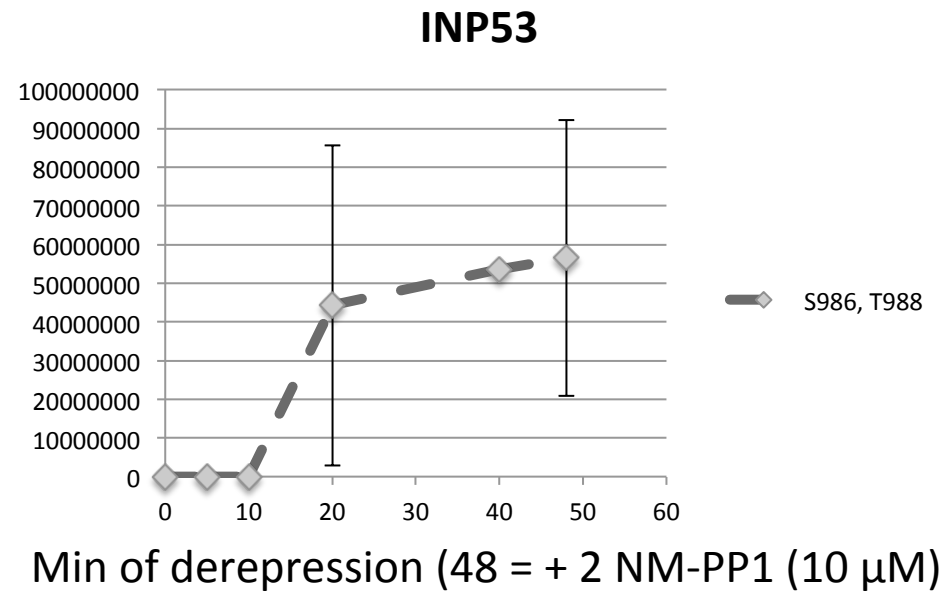
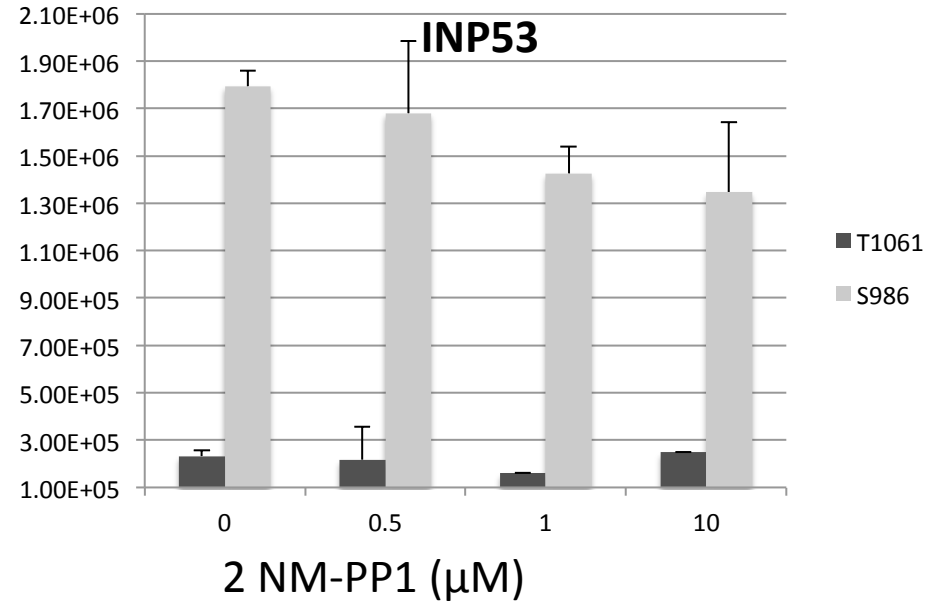
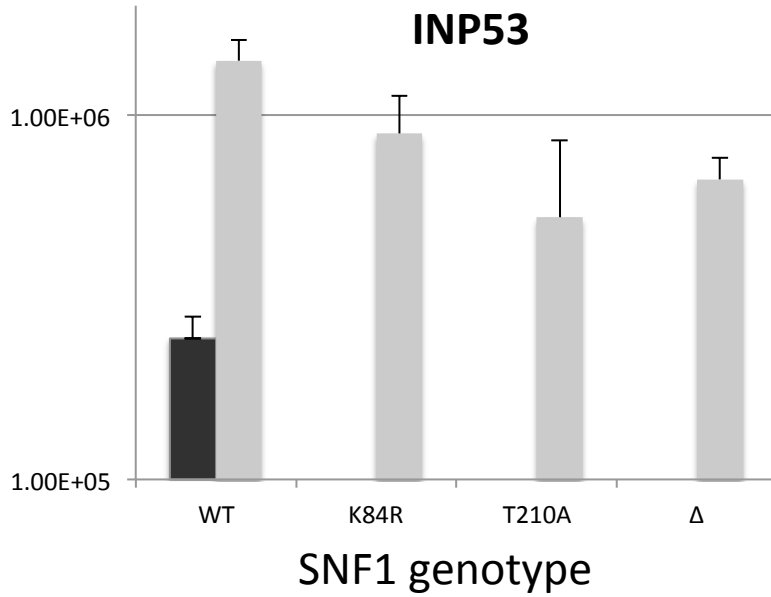
IMH1



No data

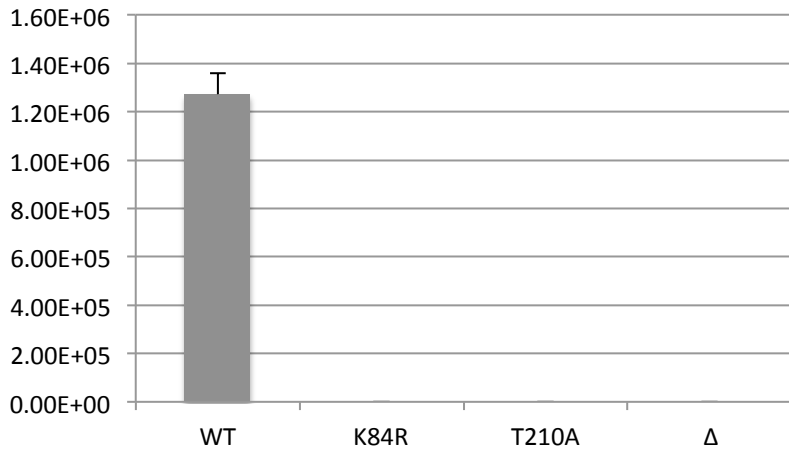
Min of derepression (48 = + 2 NM-PP1 (10 μM))

INP53



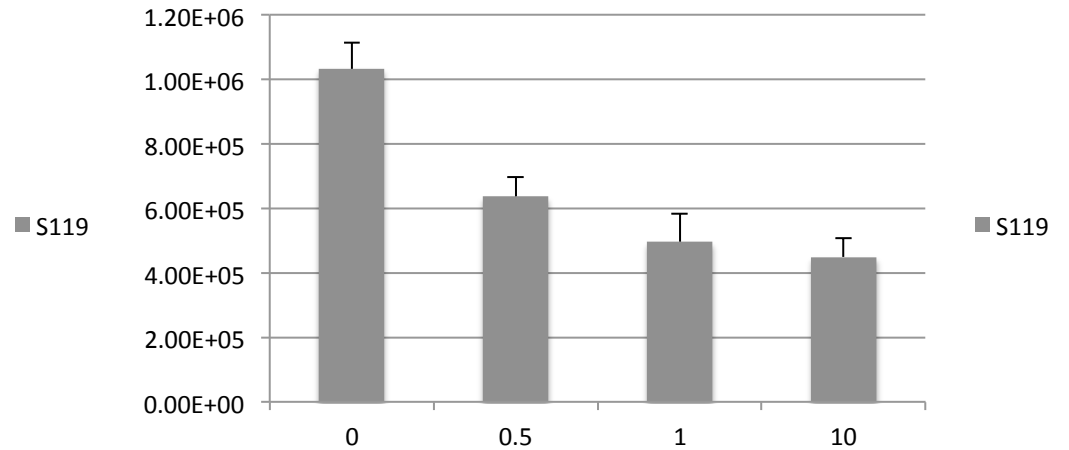
ISF1

ISF1

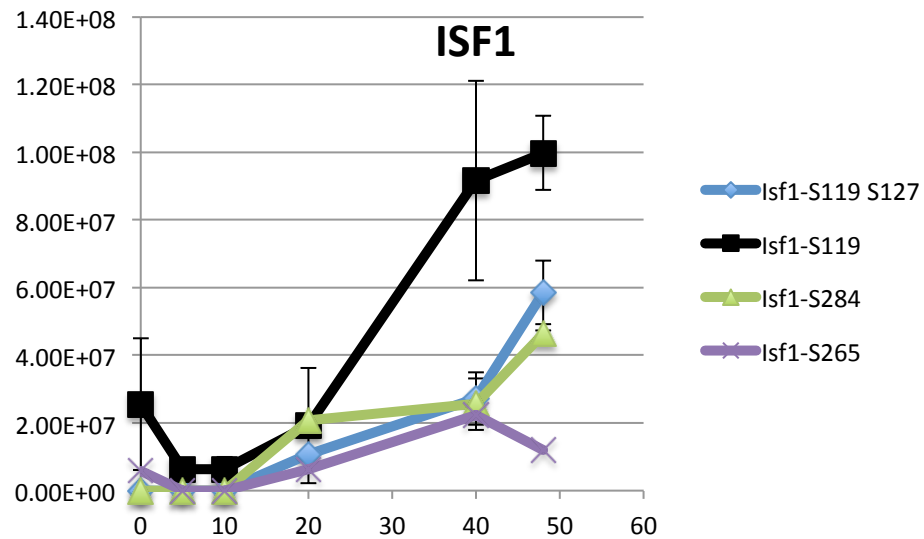


SNF1 genotype

ISF1



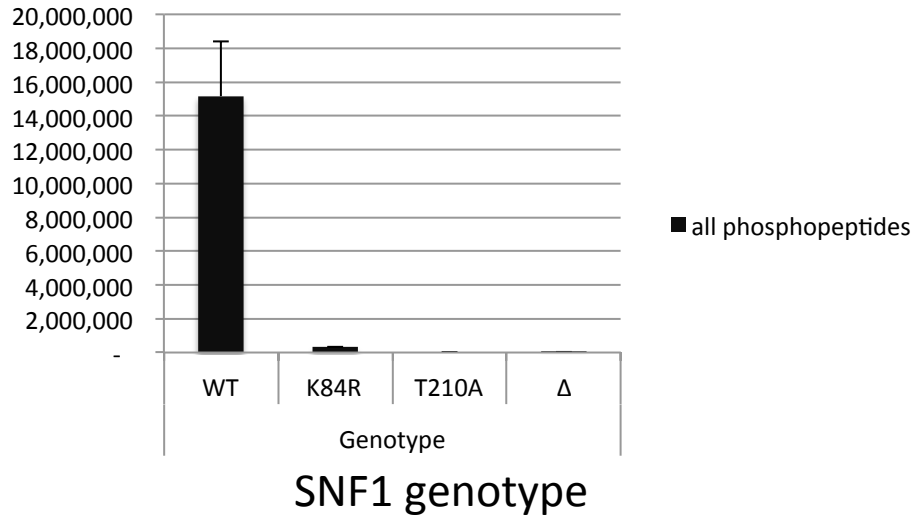
2 NM-PP1 (μM)



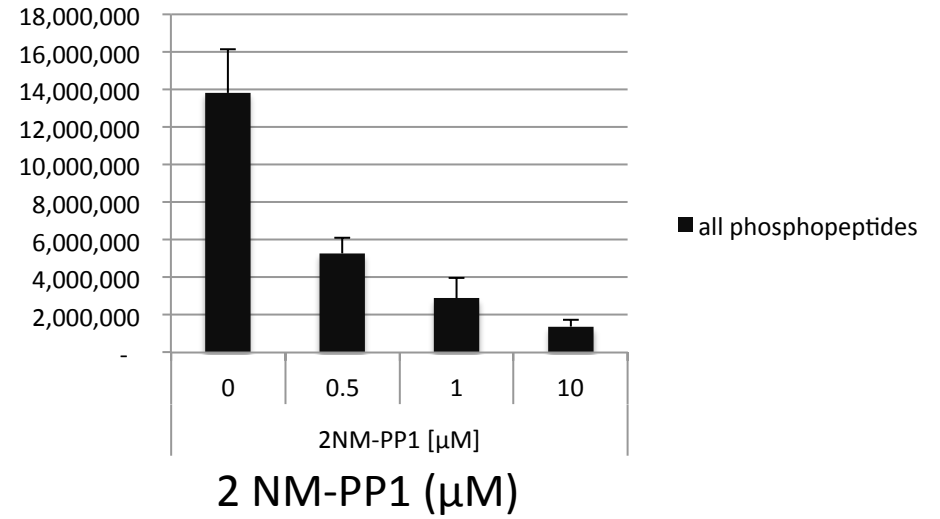
Min of derepression (48 = + 2 NM-PP1 (10 μM))

JEN1

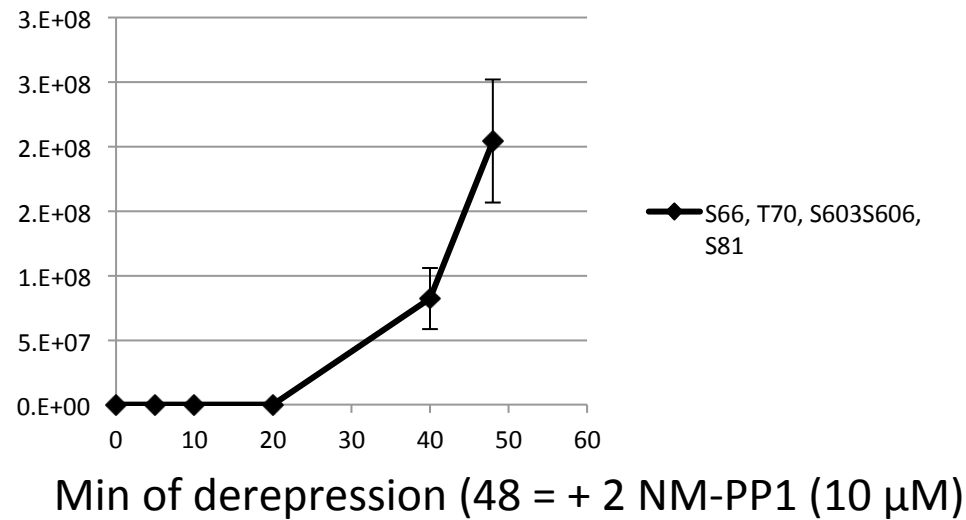
JEN1



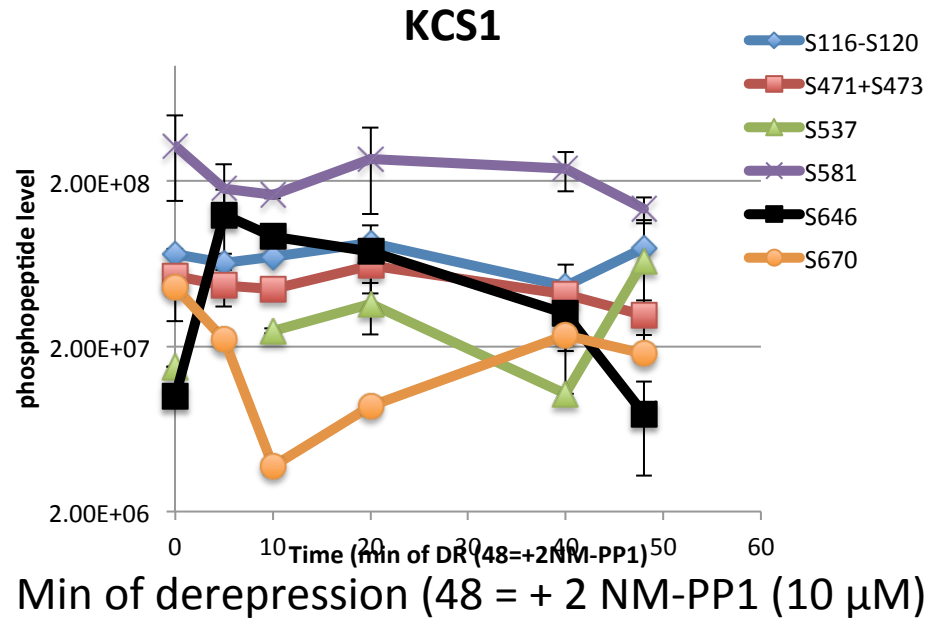
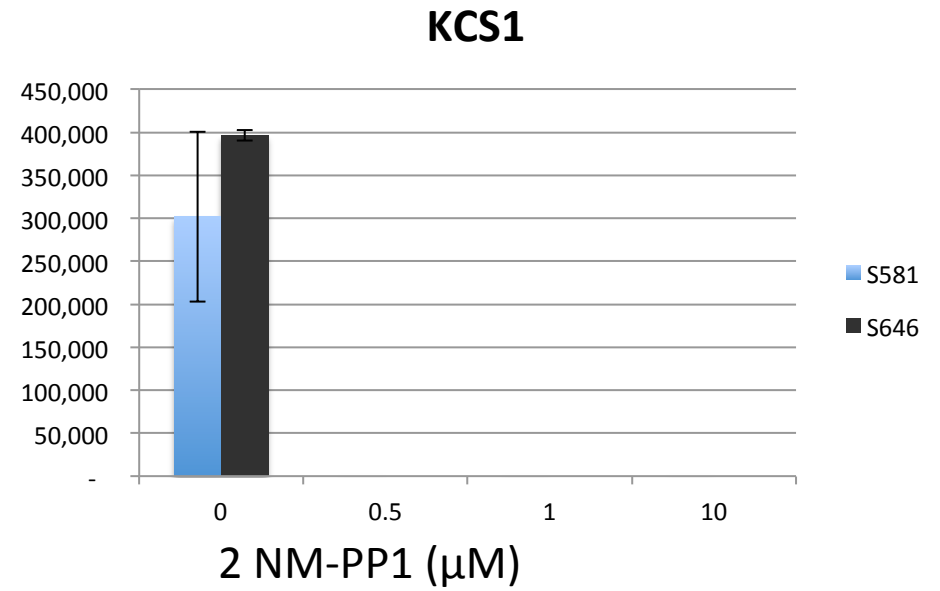
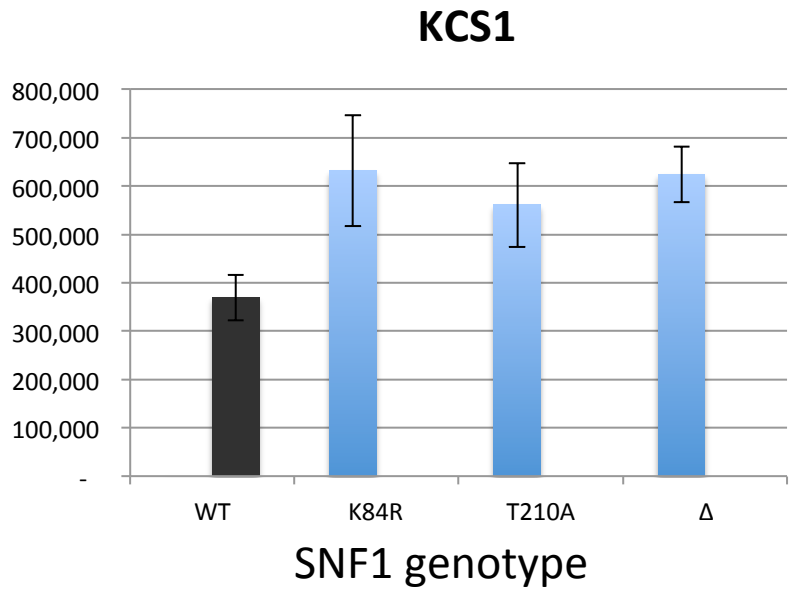
JEN1



JEN1

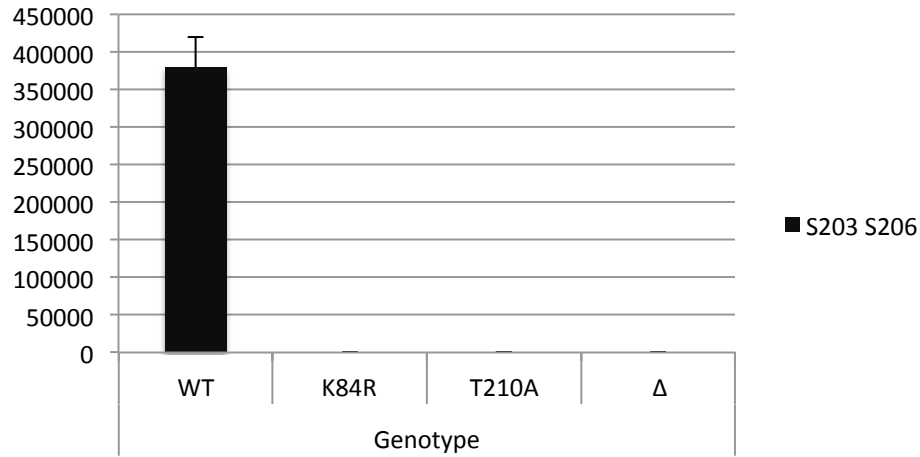


KCS1



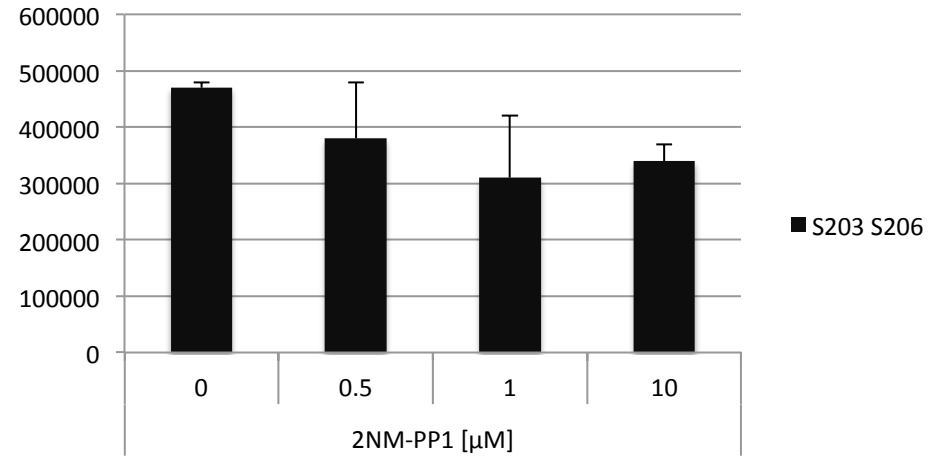
KIN82

KIN82

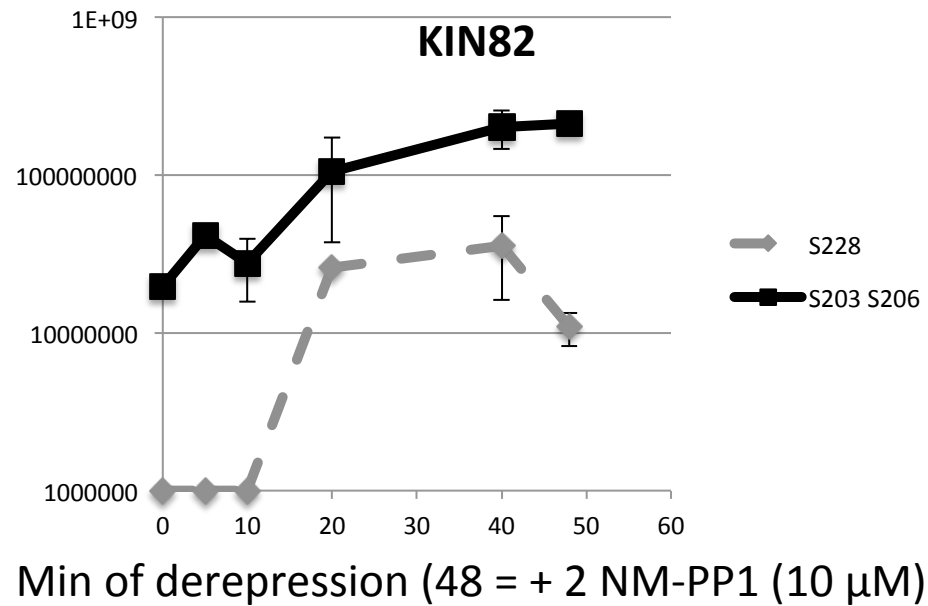


SNF1 genotype

KIN82



2 NM-PP1 (μM)



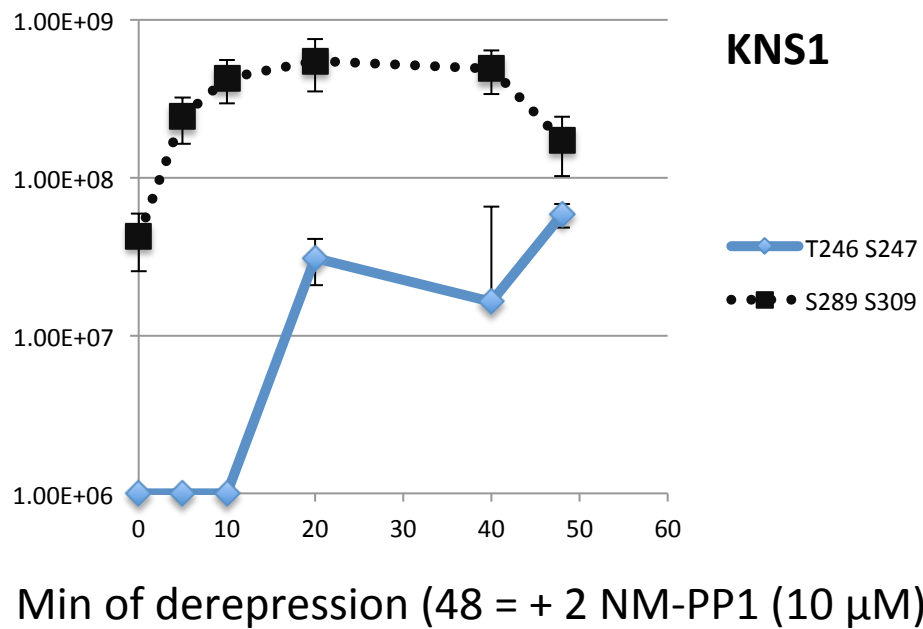
KNS1

No data

SNF1 genotype

No data

2 NM-PP1 (μM)



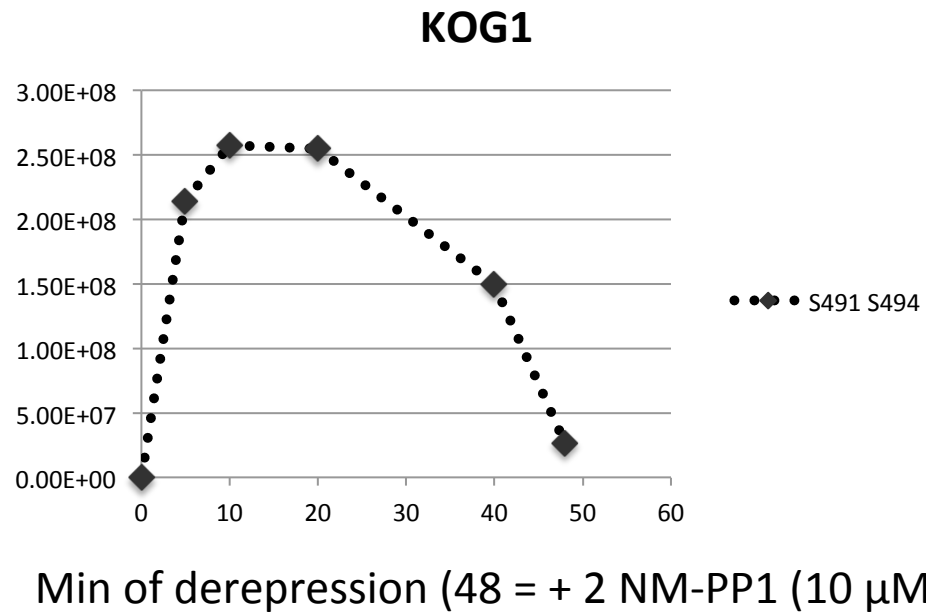
KOG1

No data

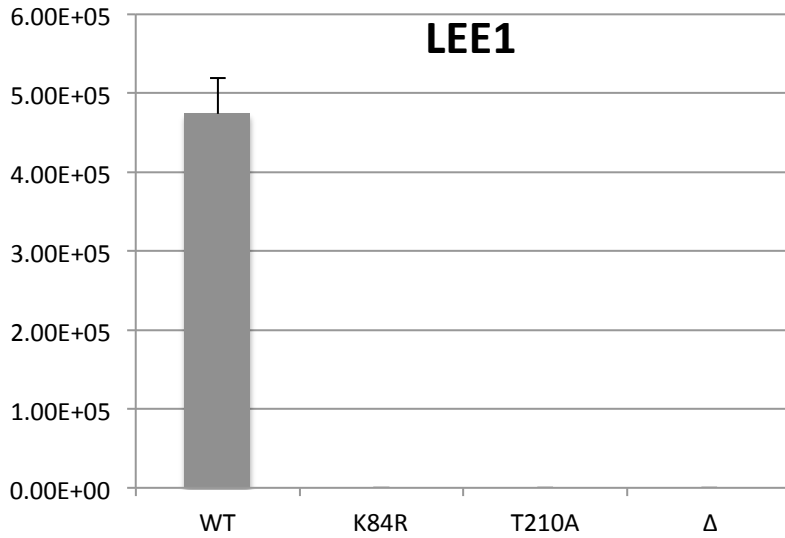
SNF1 genotype

No data

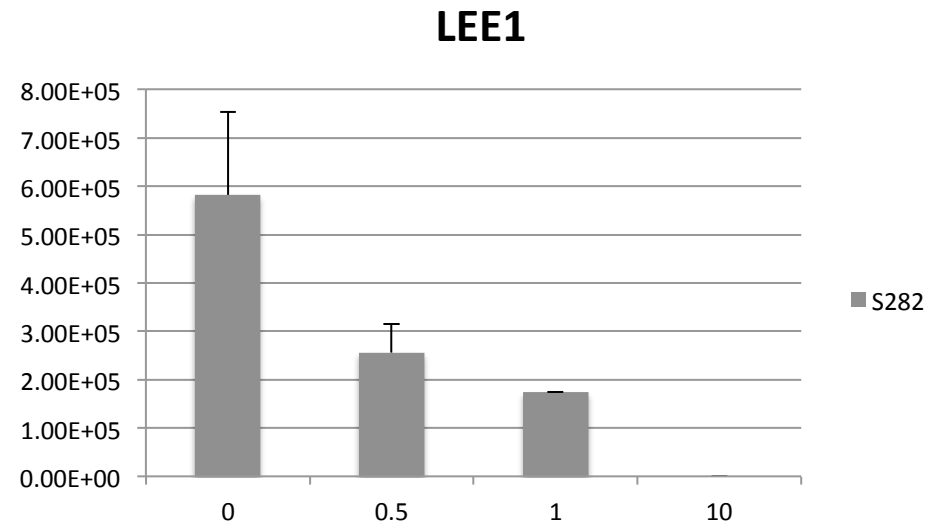
2 NM-PP1 (μM)



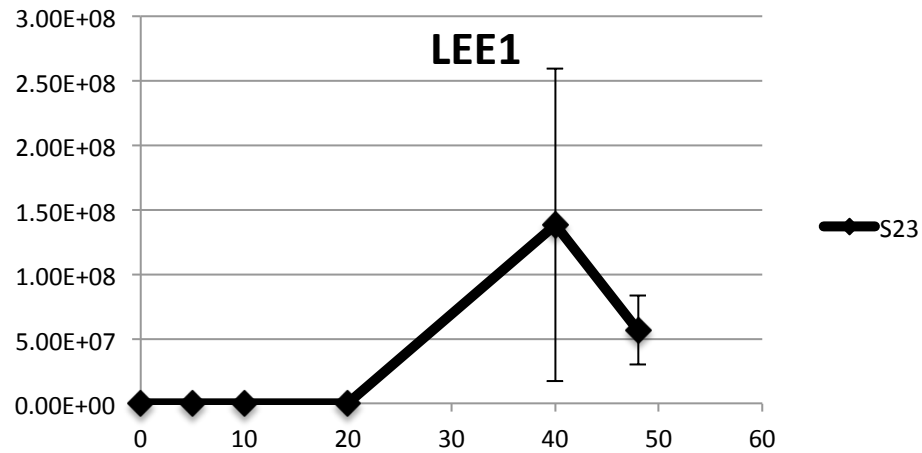
LEE1



SNF1 genotype

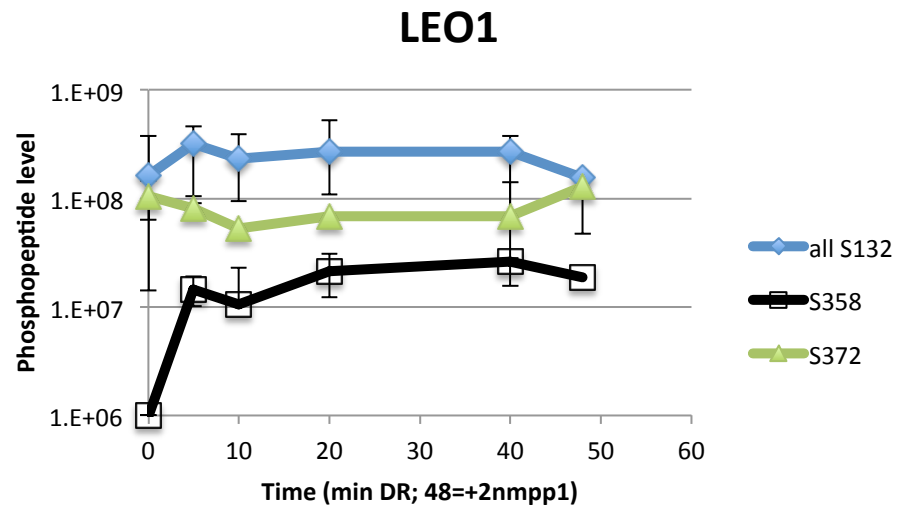
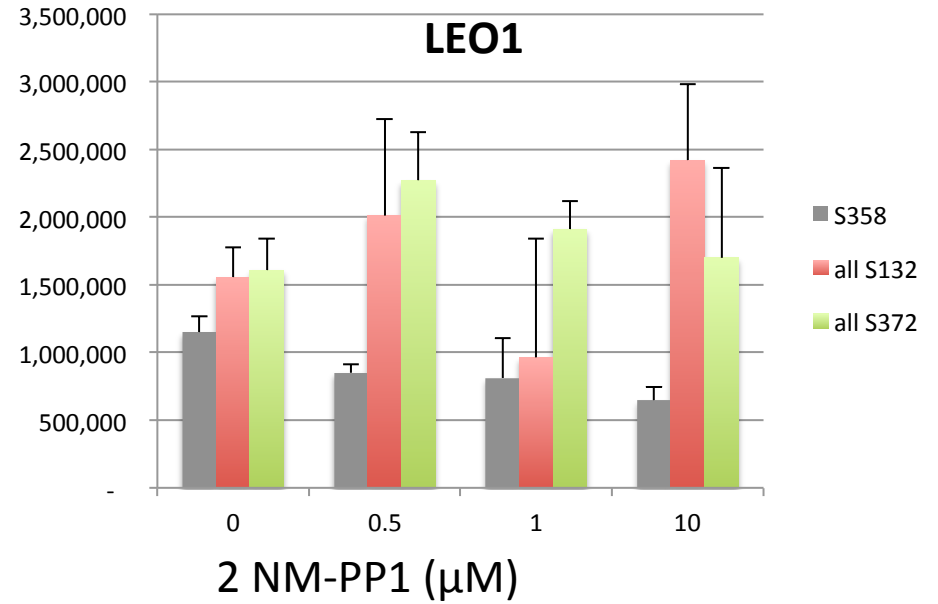
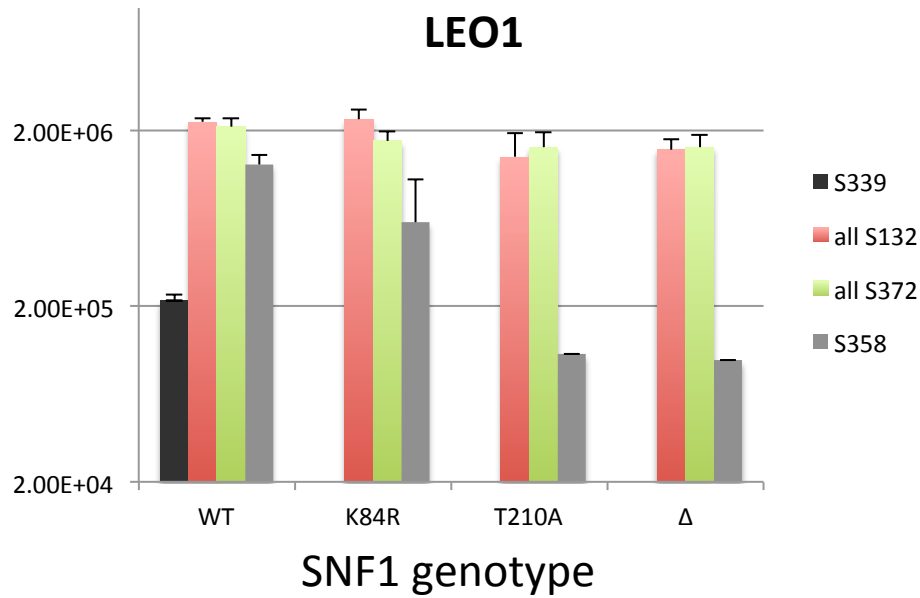


2 NM-PP1 (μM)



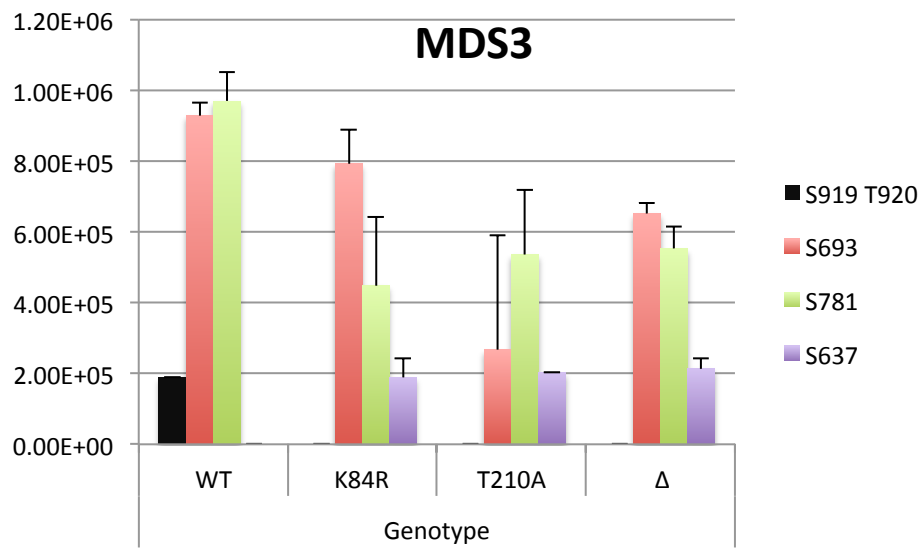
Min of derepression (48 = + 2 NM-PP1 (10 μM))

LEO1

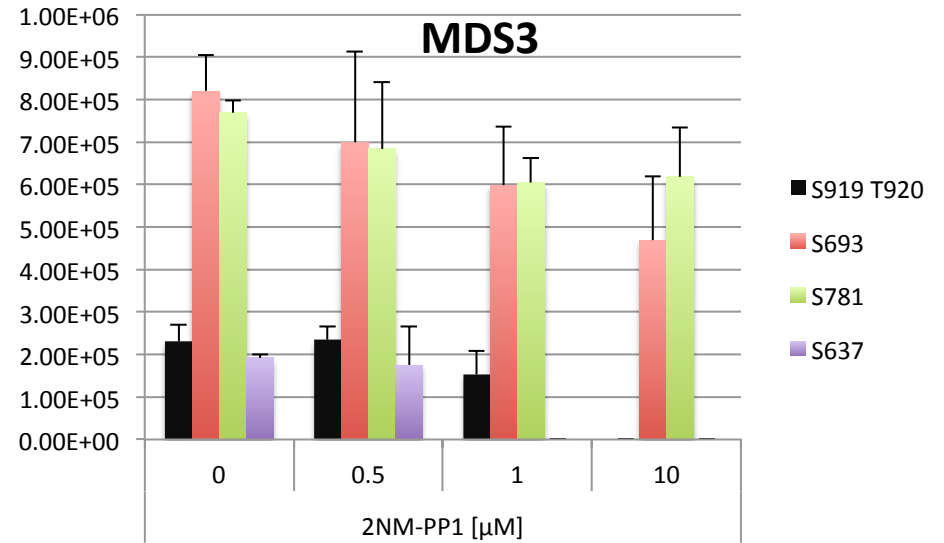


Min of derepression (48 = + 2 NM-PP1 (10 μM))

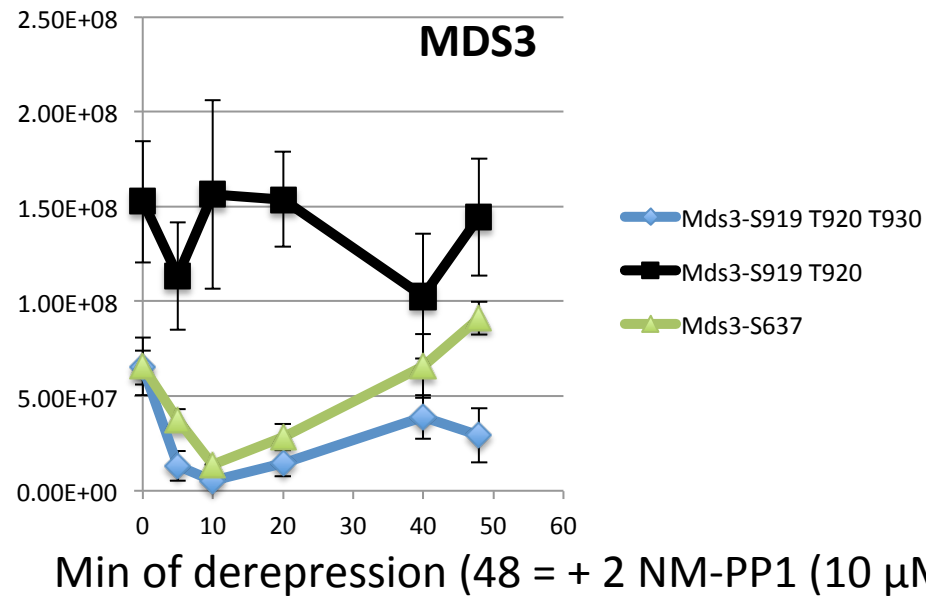
MDS3



SNF1 genotype



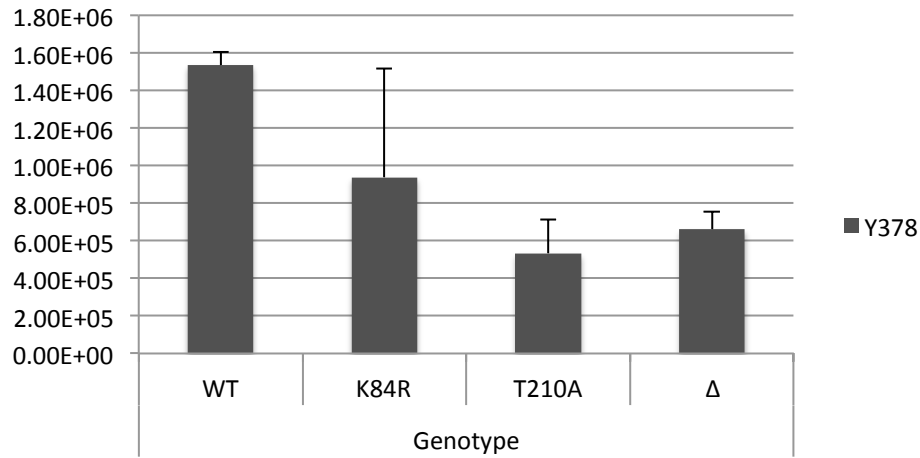
2 NM-PP1 (μM)



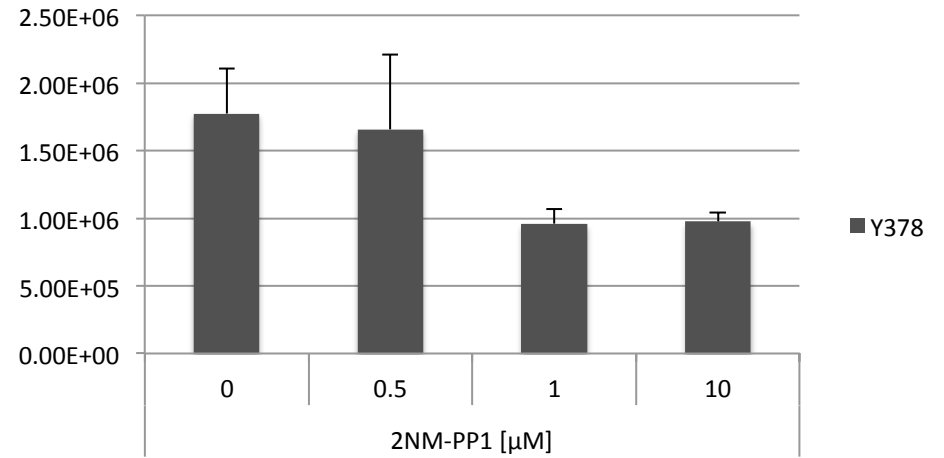
Min of derepression (48 = + 2 NM-PP1 (10 μM))

MES1

MES1

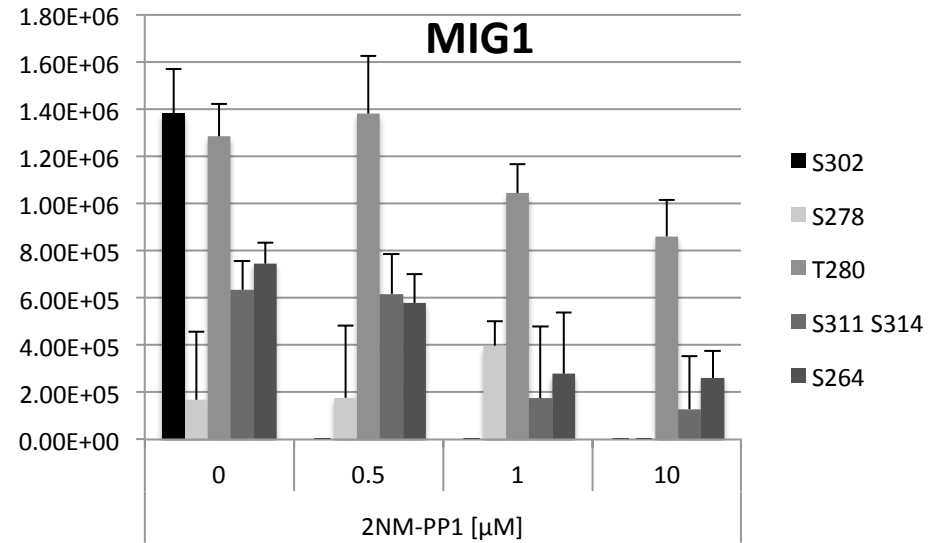
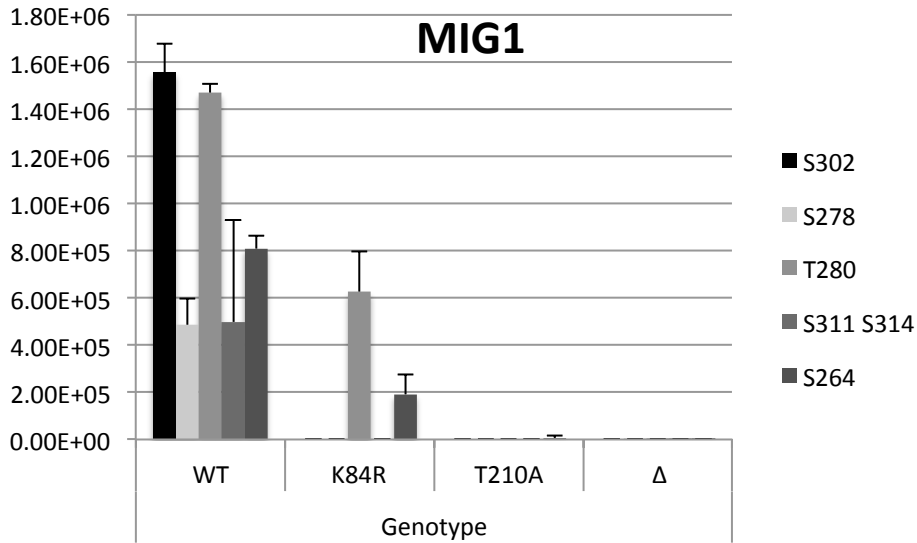


MES1



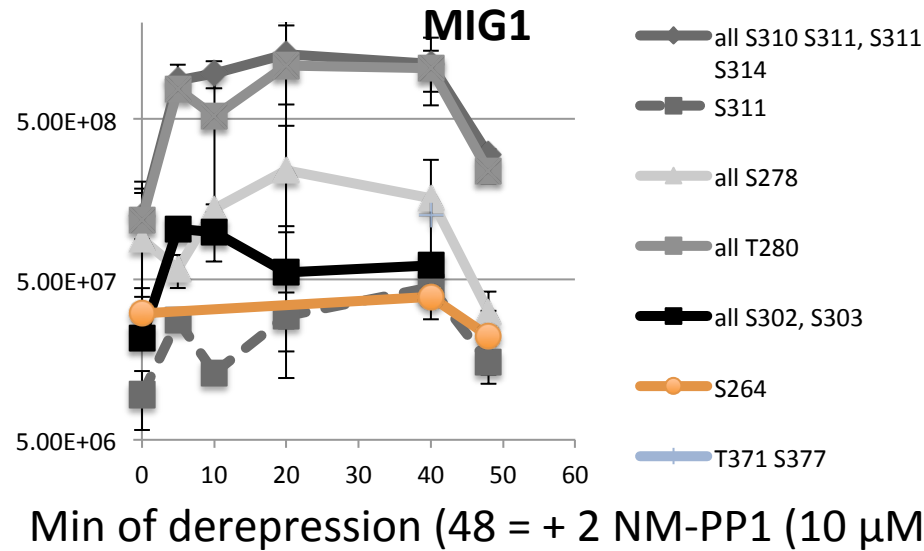
No data from time course experiment

MIG1



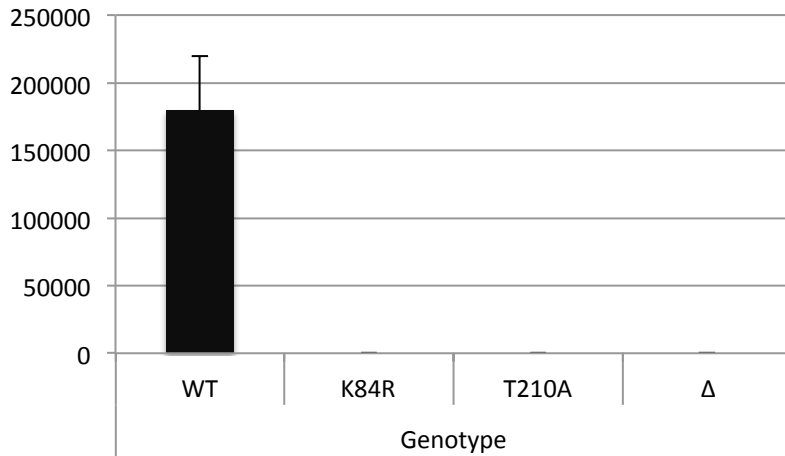
SNF1 genotype

2 NM-PP1 (μM)



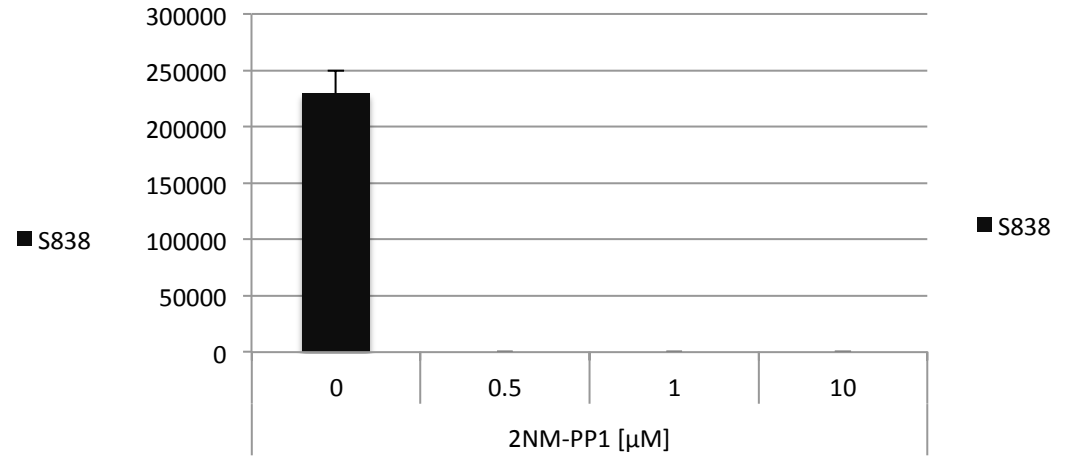
MPT5

MPT5

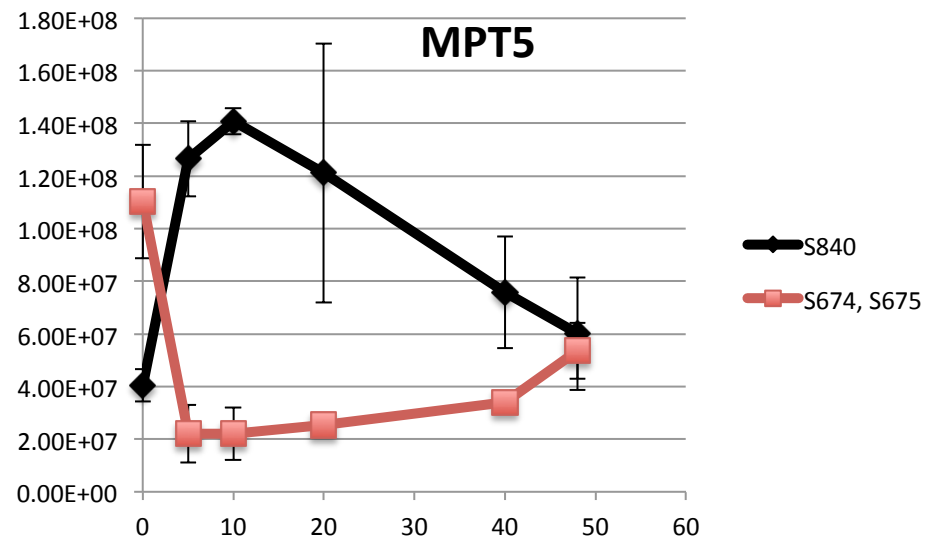


SNF1 genotype

MPT5

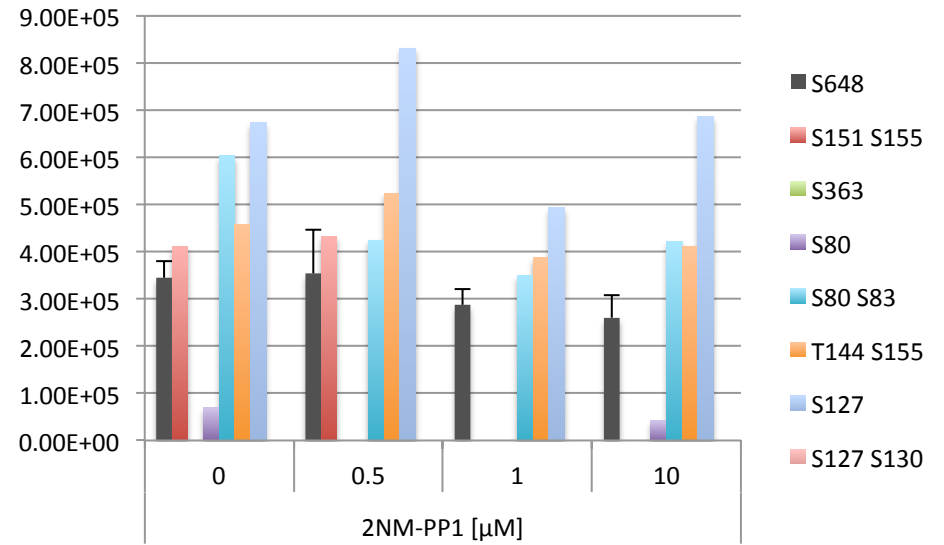
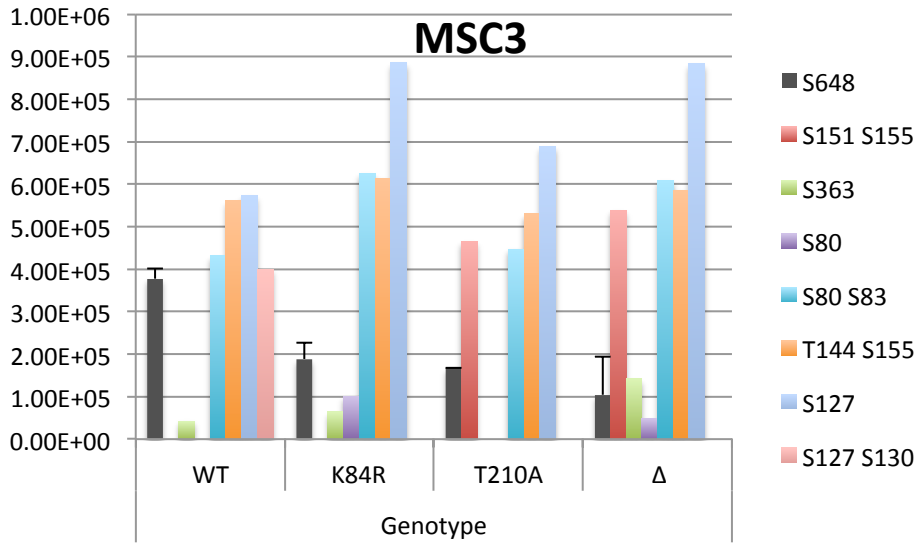


2 NM-PP1 (μM)

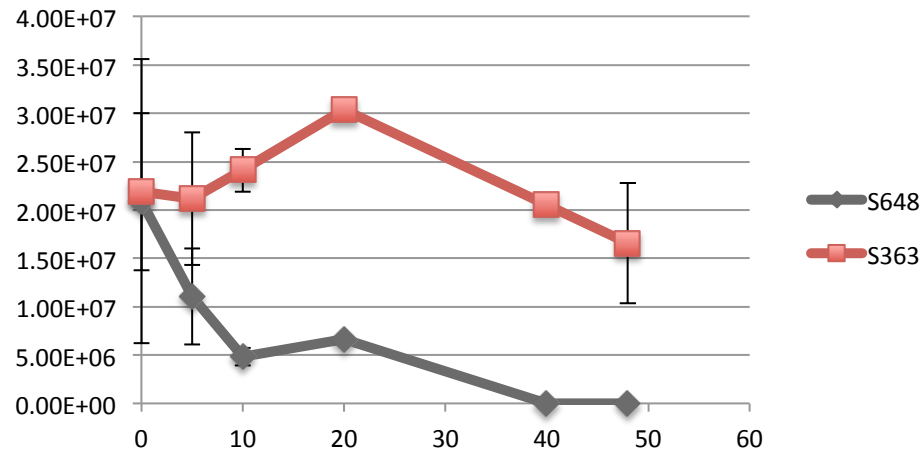


Min of derepression (48 = + 2 NM-PP1 (10 μM))

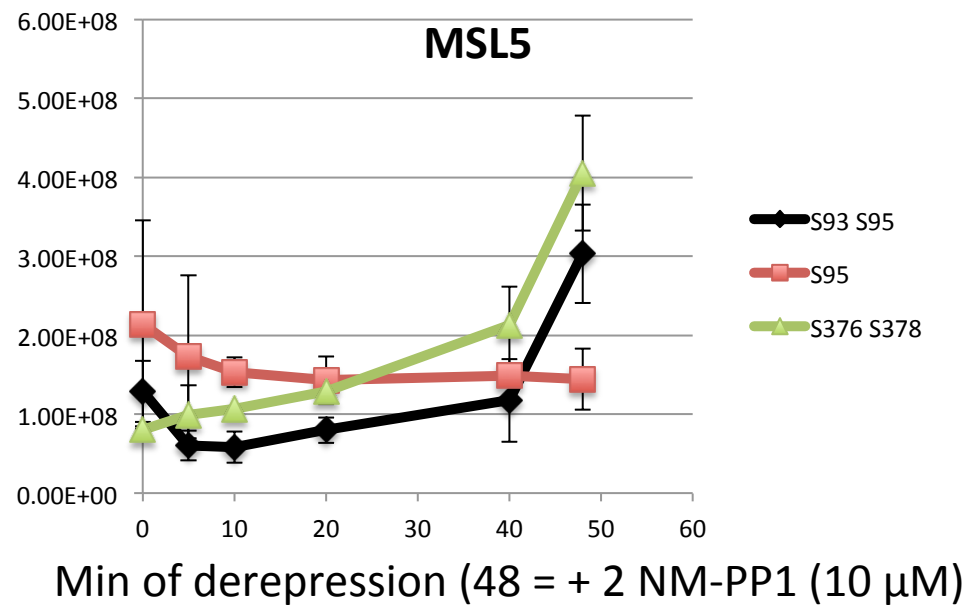
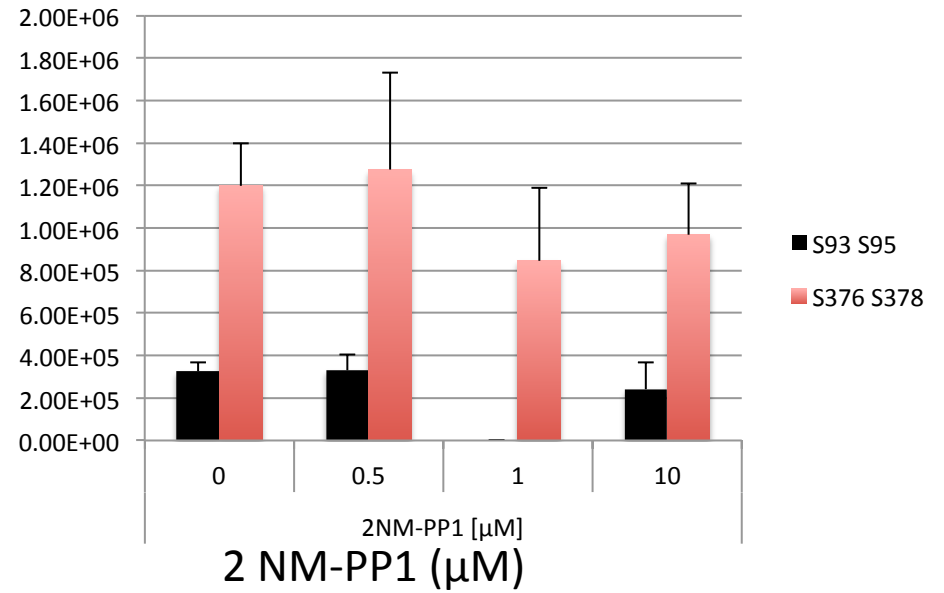
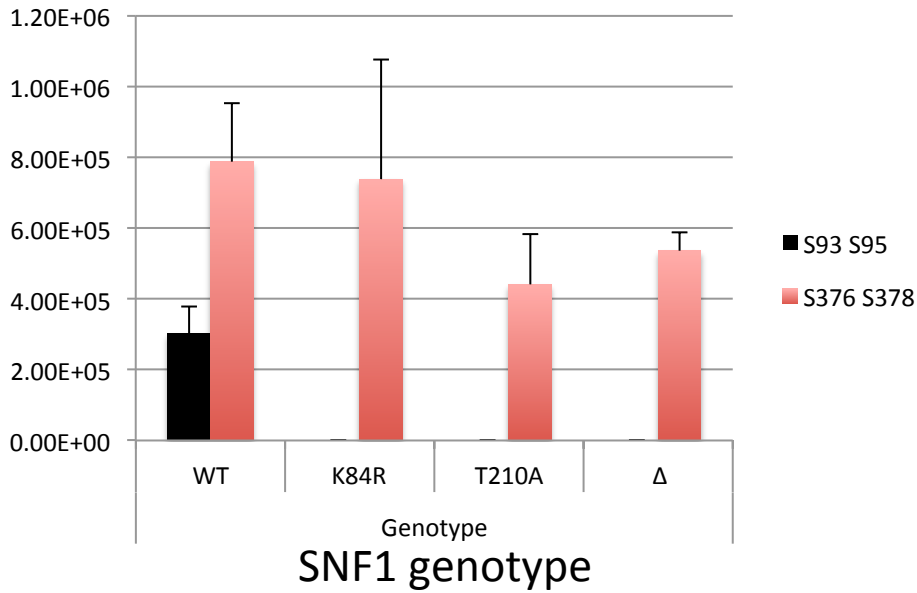
MSC3



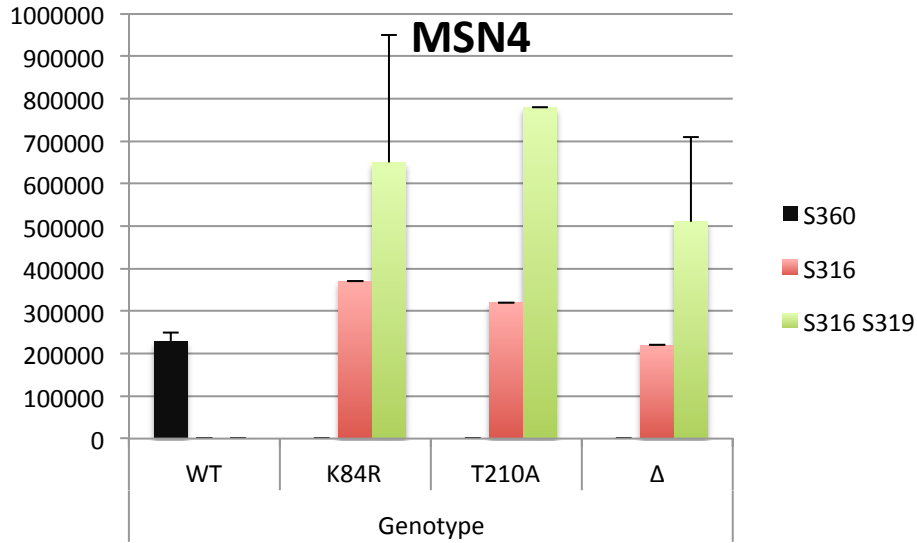
MSC3



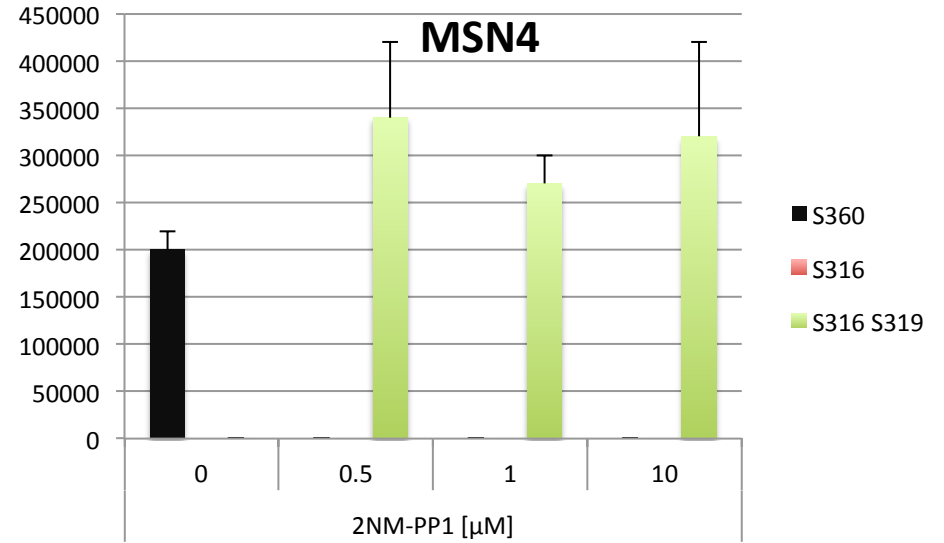
MSL5



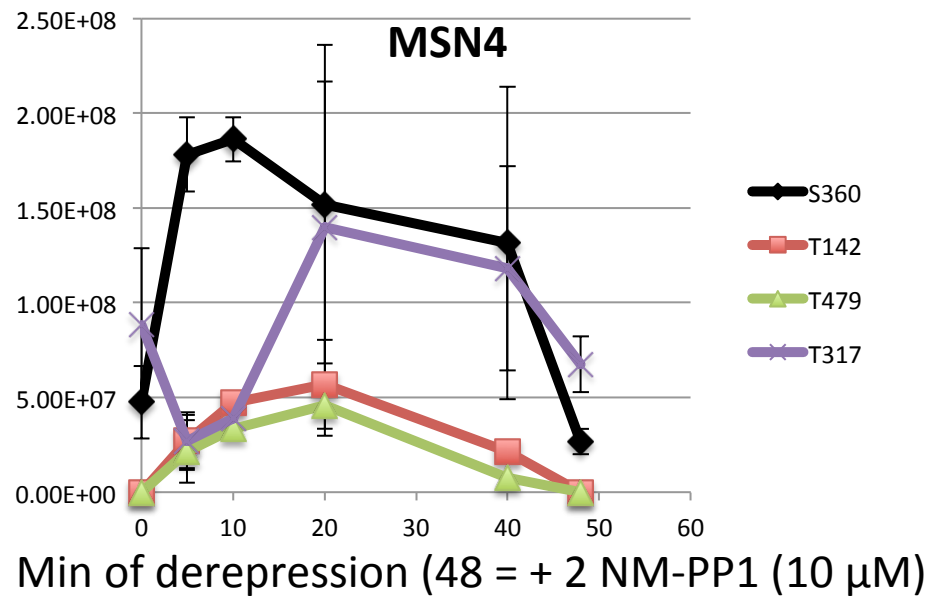
MSN4



SNF1 genotype

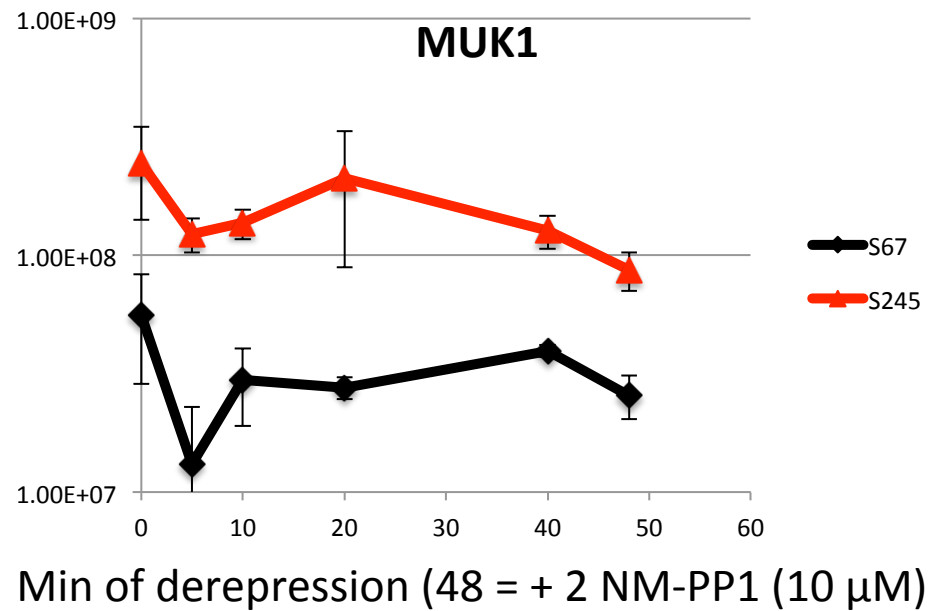
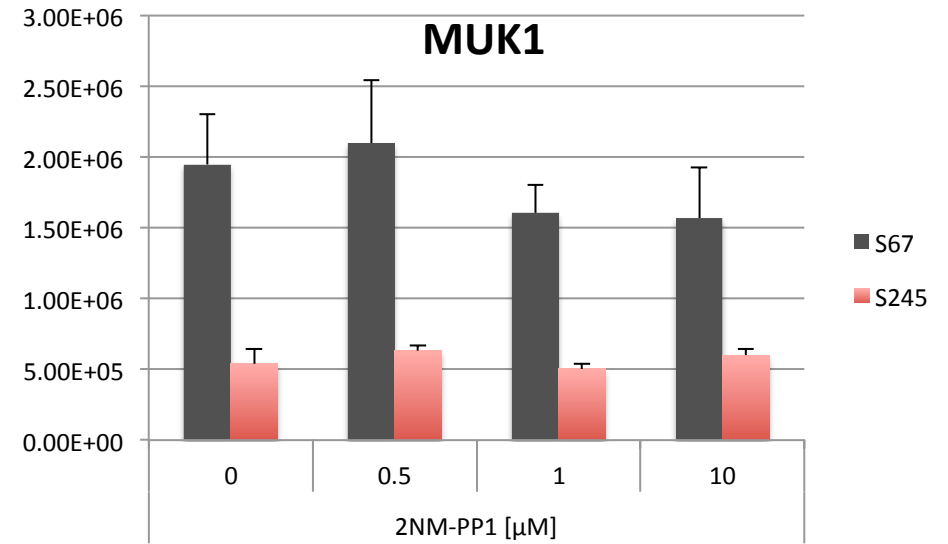
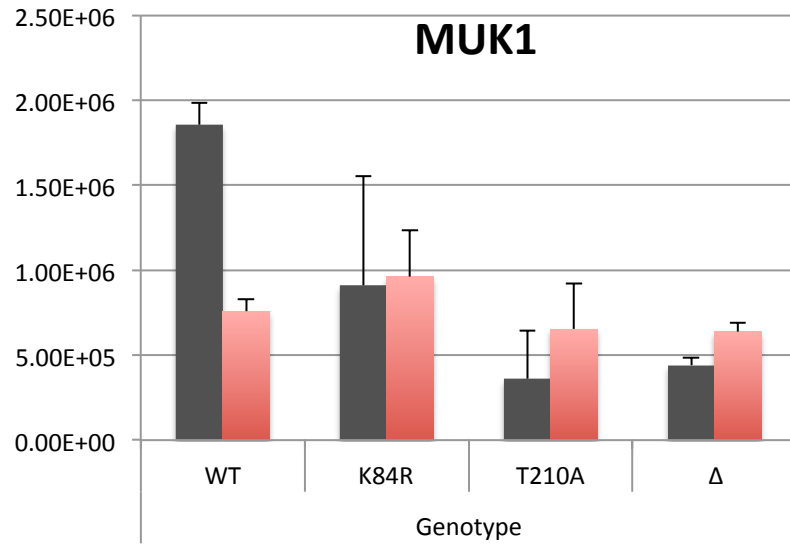


2 NM-PP1 (μM)

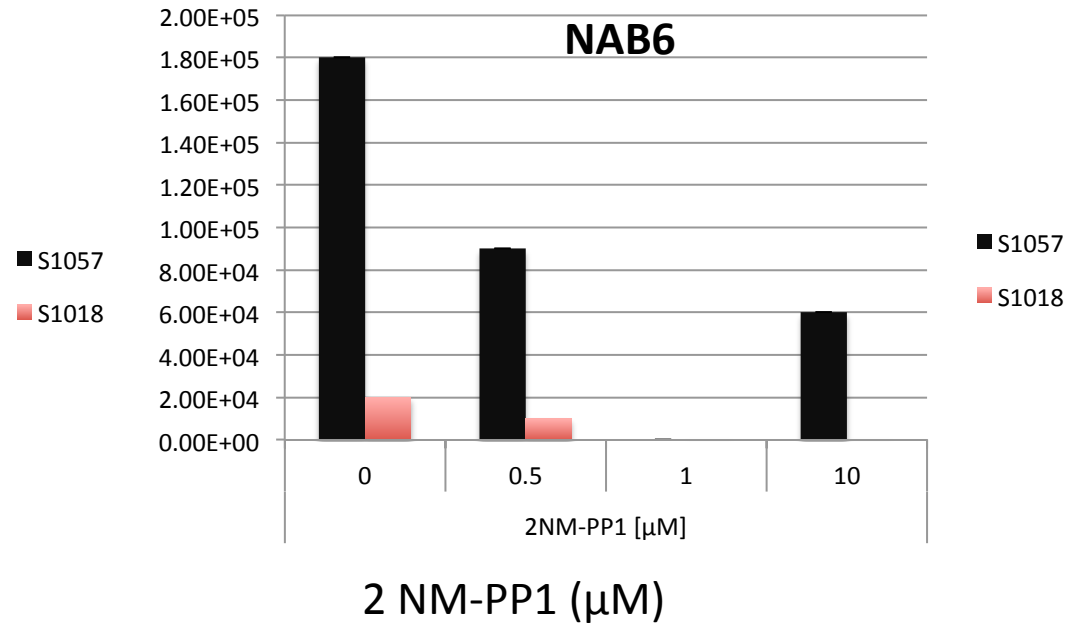
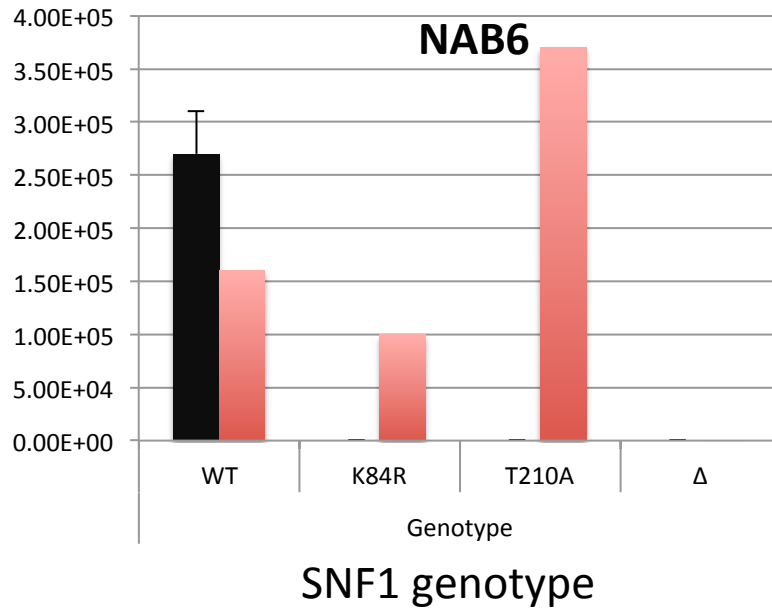


Min of derepression (48 = + 2 NM-PP1 (10 μM))

MUK1



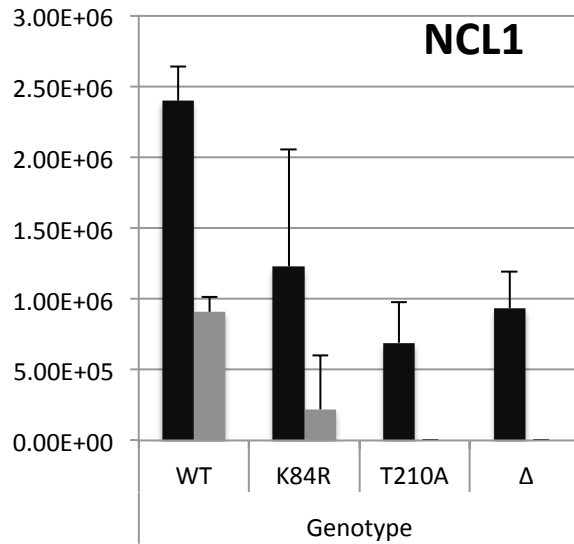
NAB6



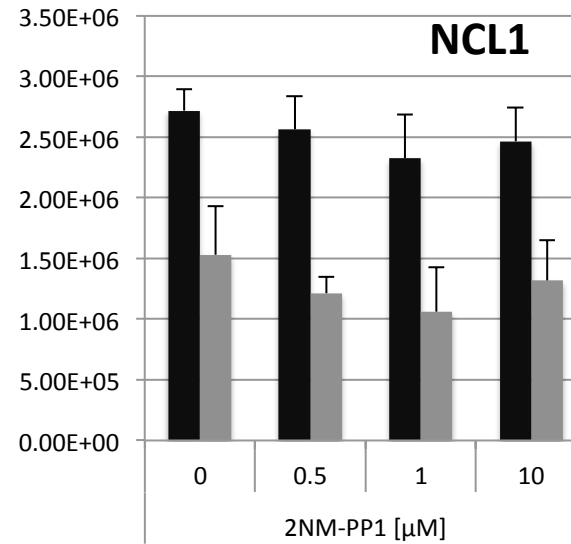
no data for S1057

Min of derepression (48 = + 2 NM-PP1 (10 μM))

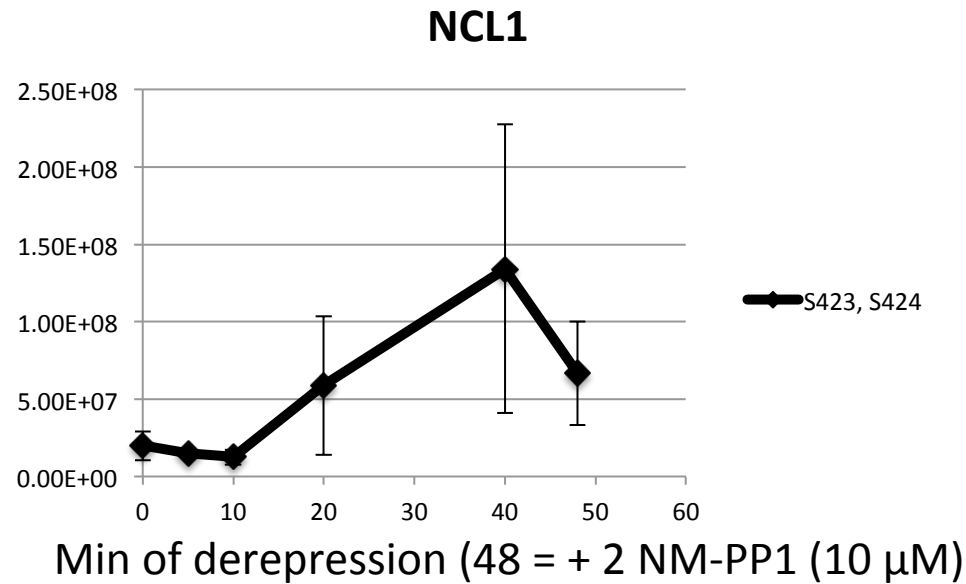
NCL1



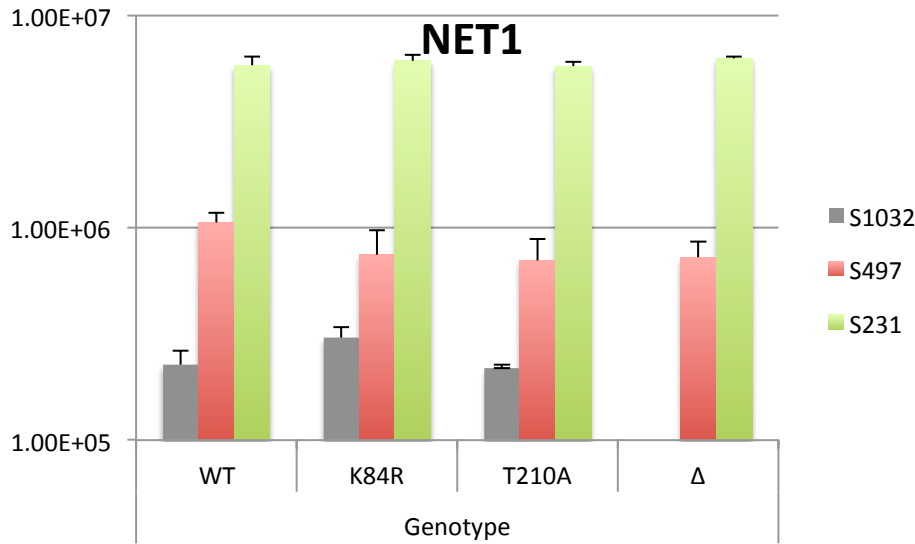
SNF1 genotype



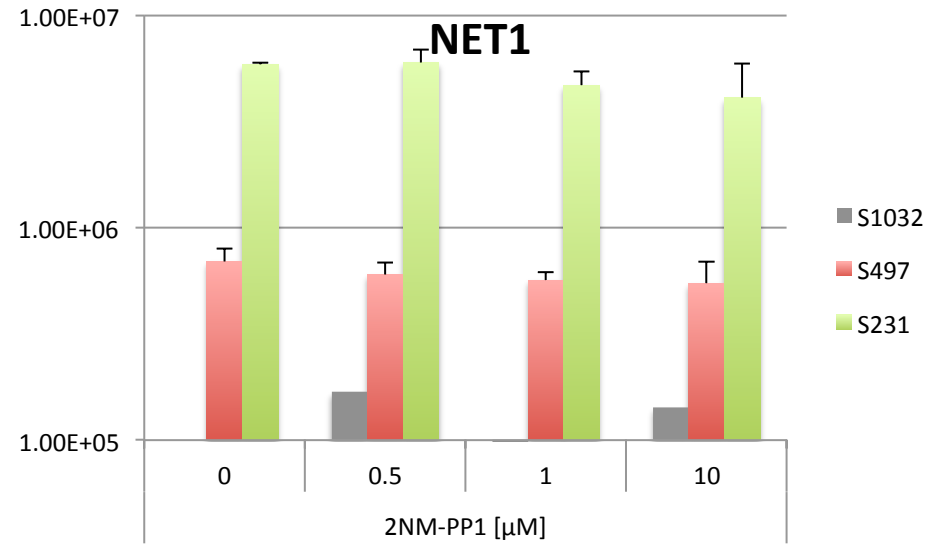
2 NM-PP1 (μM)



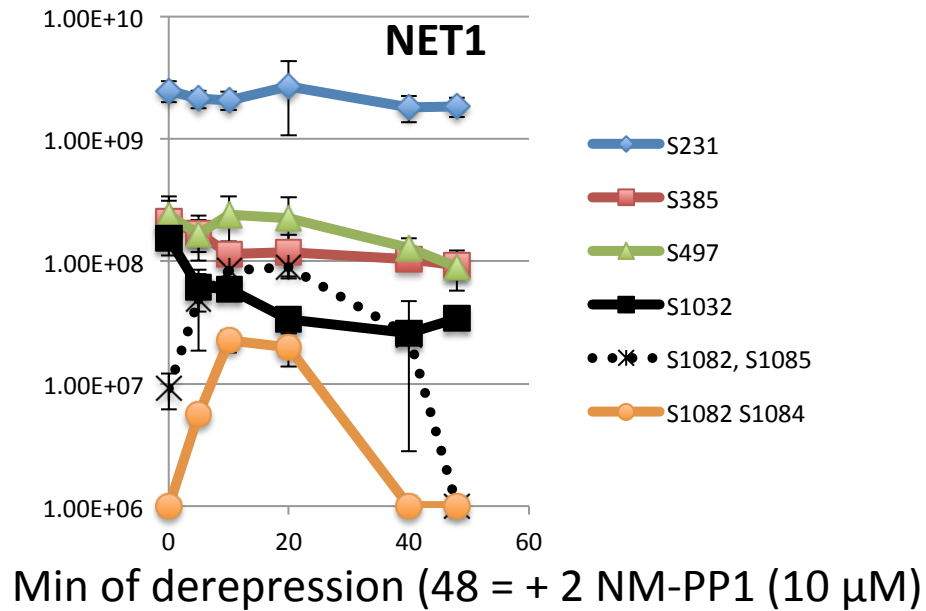
NET1



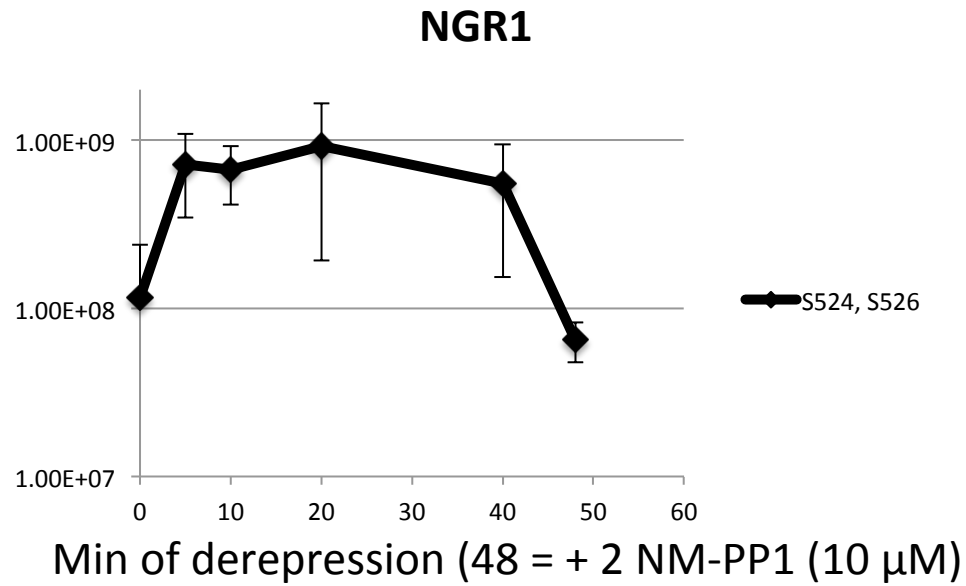
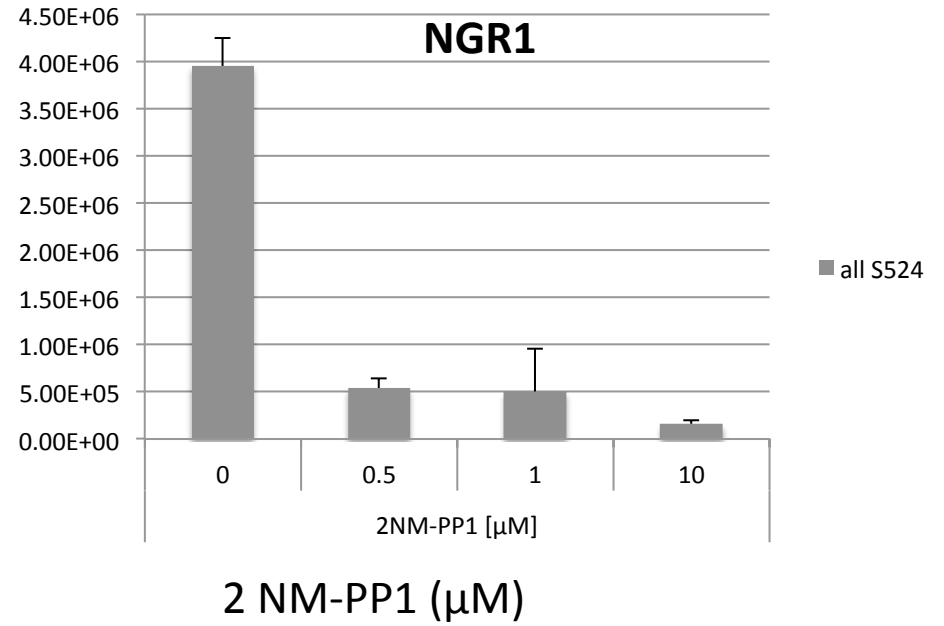
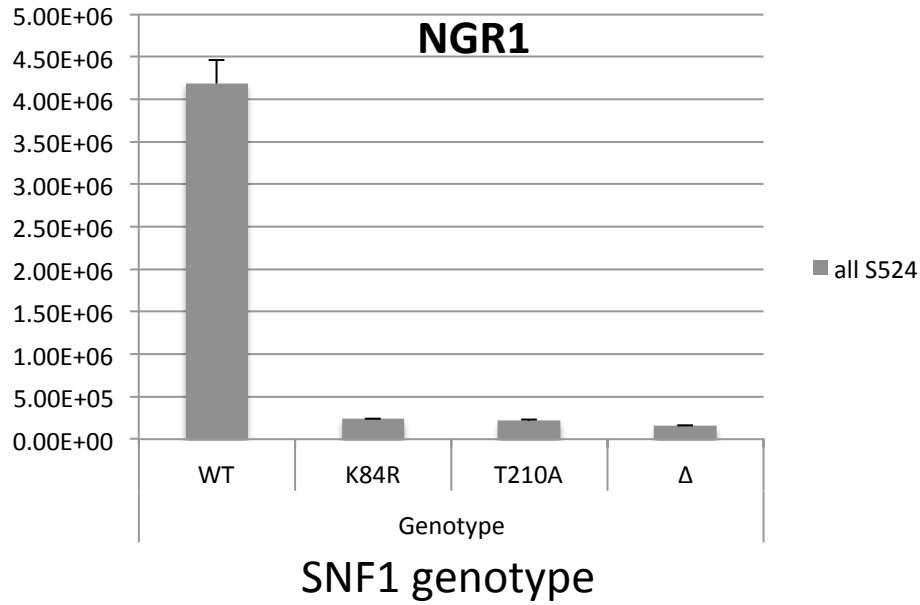
SNF1 genotype



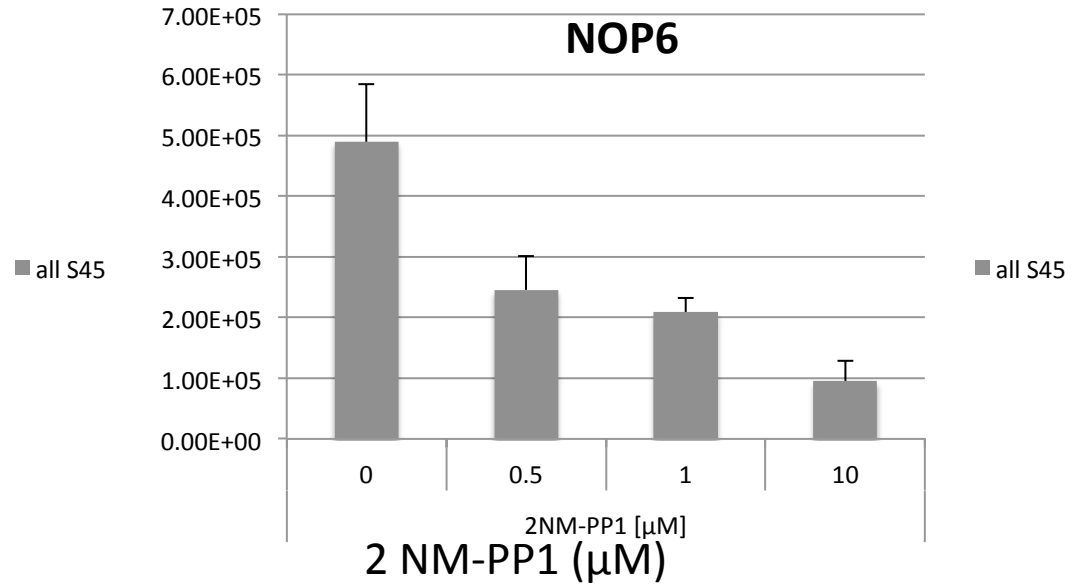
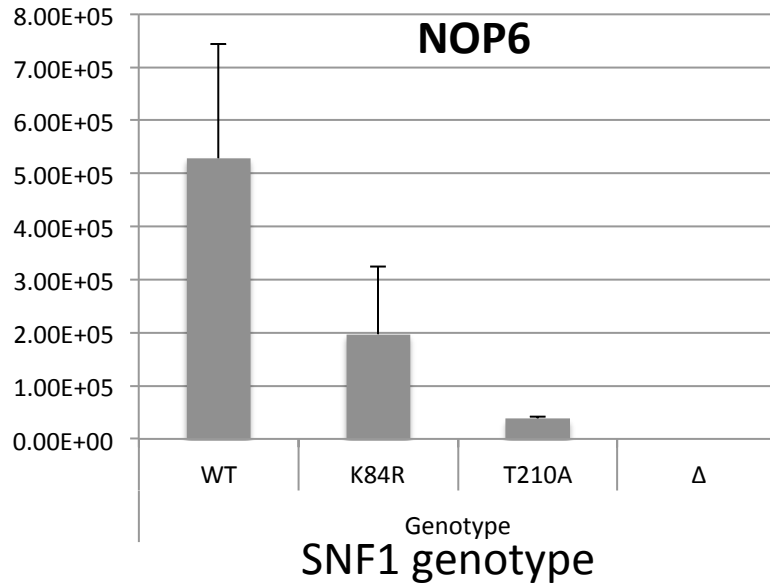
2 NM-PP1 (μM)



NGR1



NOP6

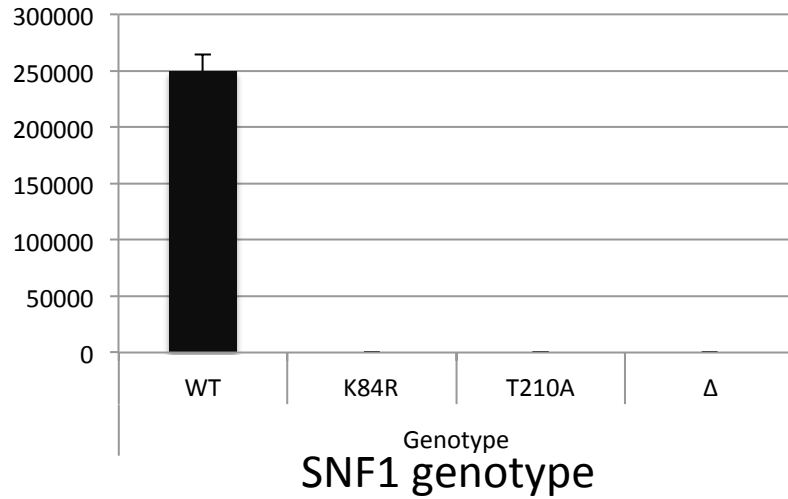


No data

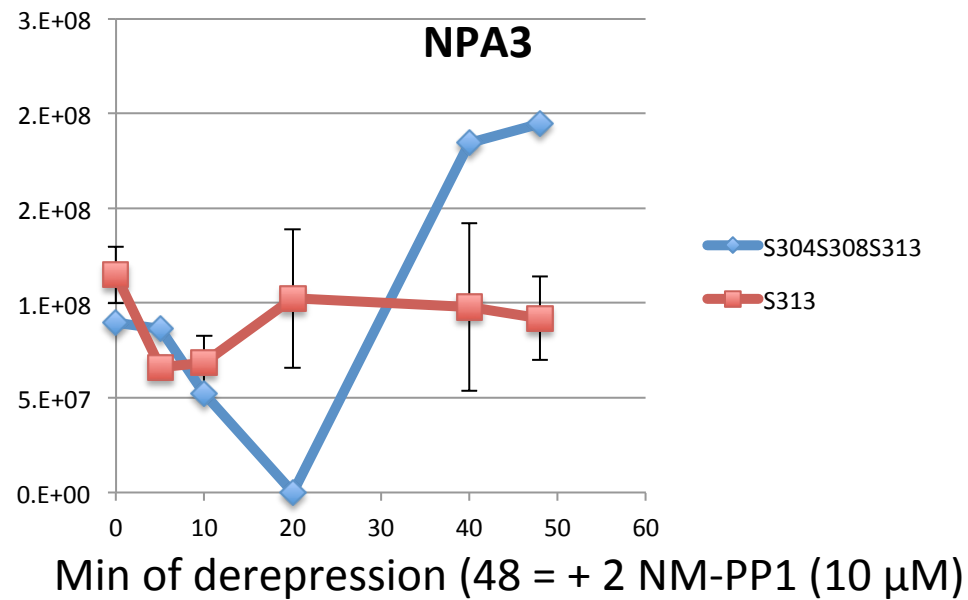
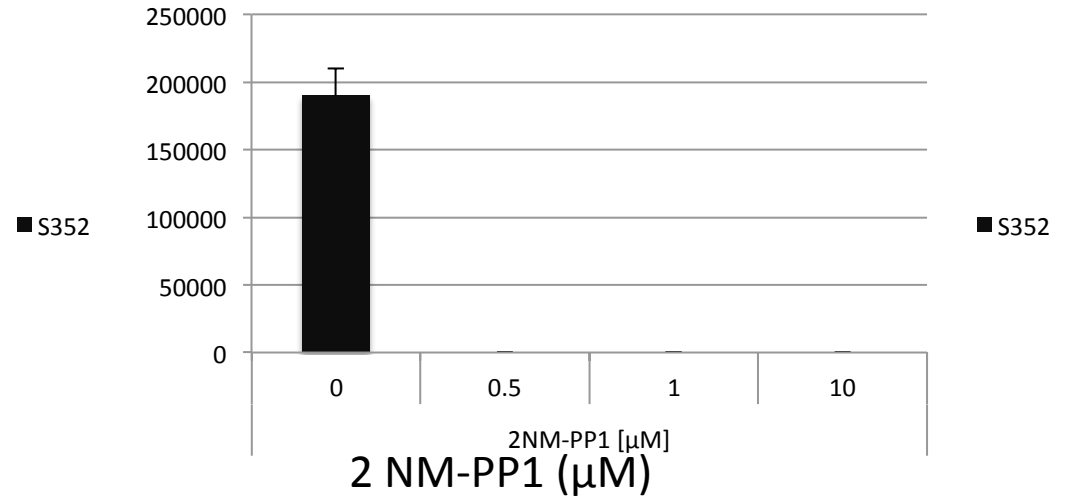
Min of derepression (48 = + 2 NM-PP1 (10 μM))

NPA3

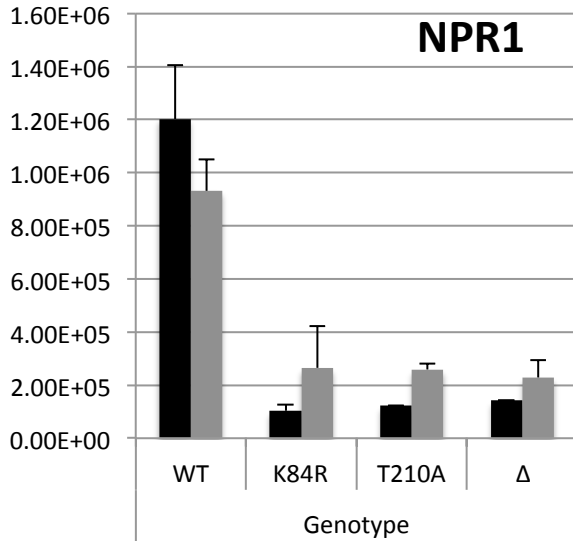
NPA3



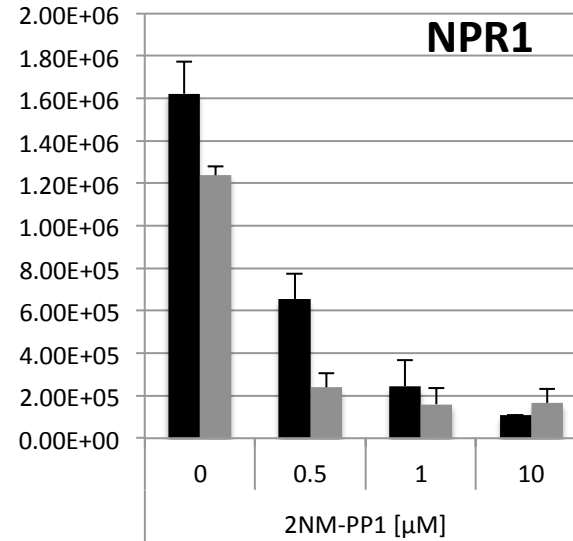
NPA3



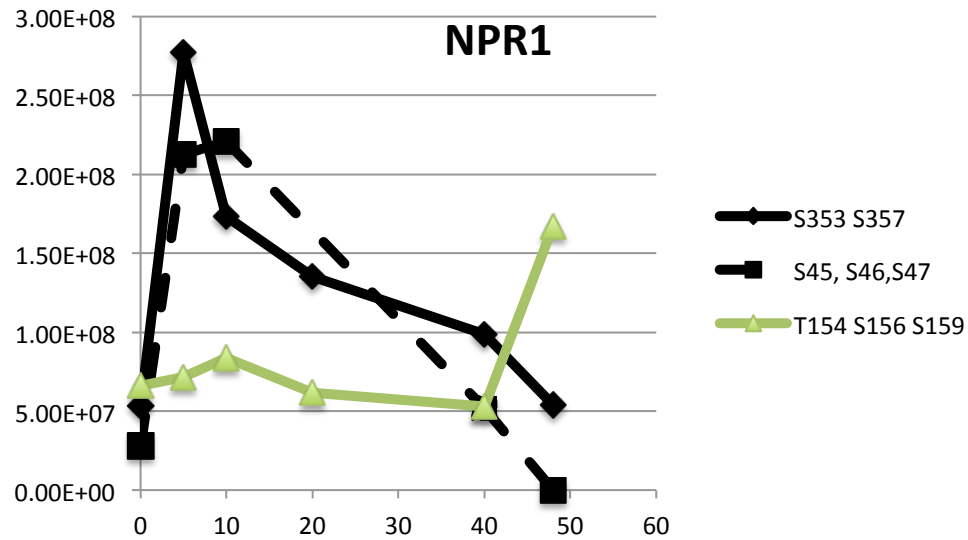
NPR1



SNF1 genotype



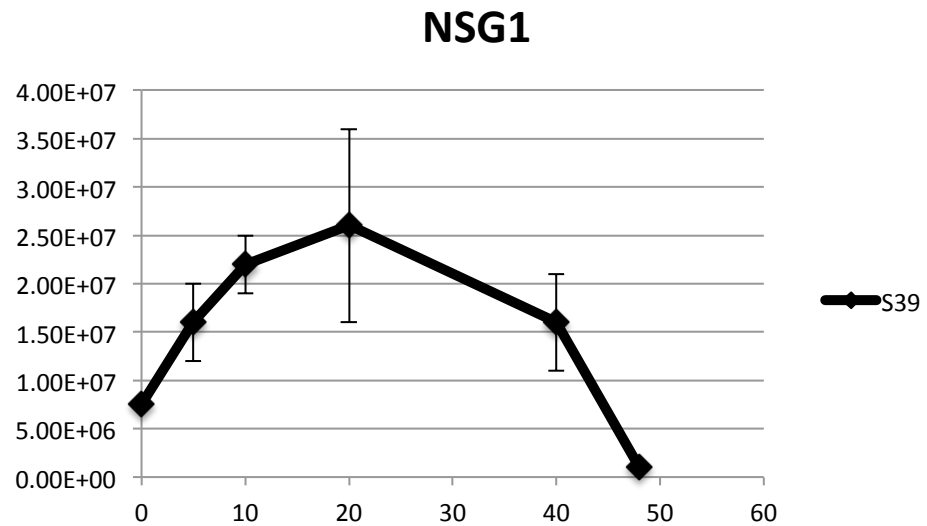
2 NM-PP1 (μM)



Min of derepression (48 = + 2 NM-PP1 (10 μM))

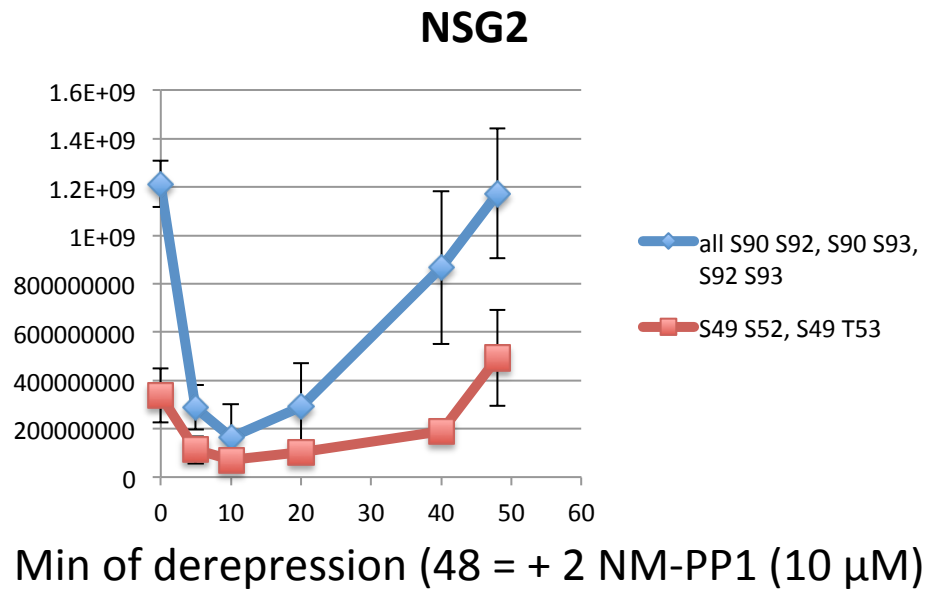
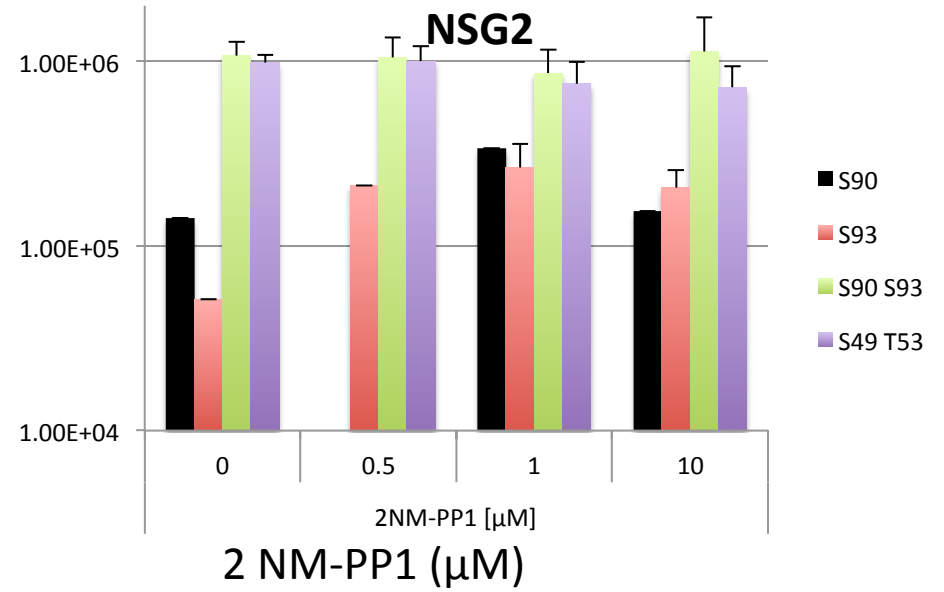
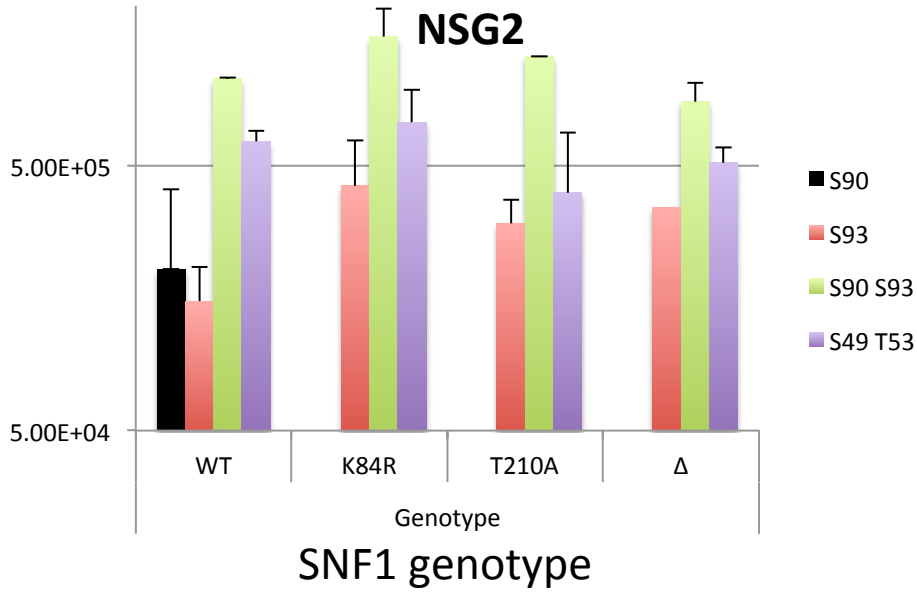
NSG1

No data from experiments 1 and 2



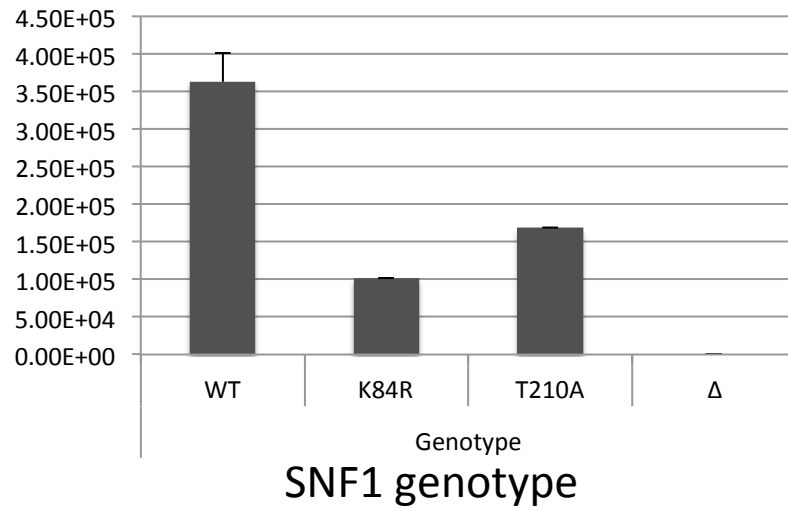
Min of derepression (48 = + 2 NM-PP1 (10 μ M))

NSG2

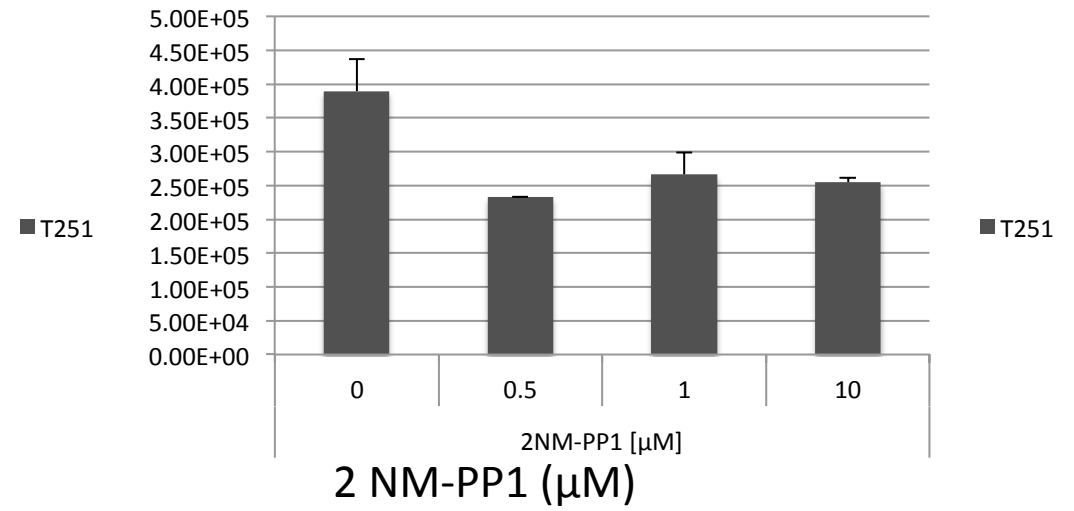


NVJ1

NVJ1



NVJ1

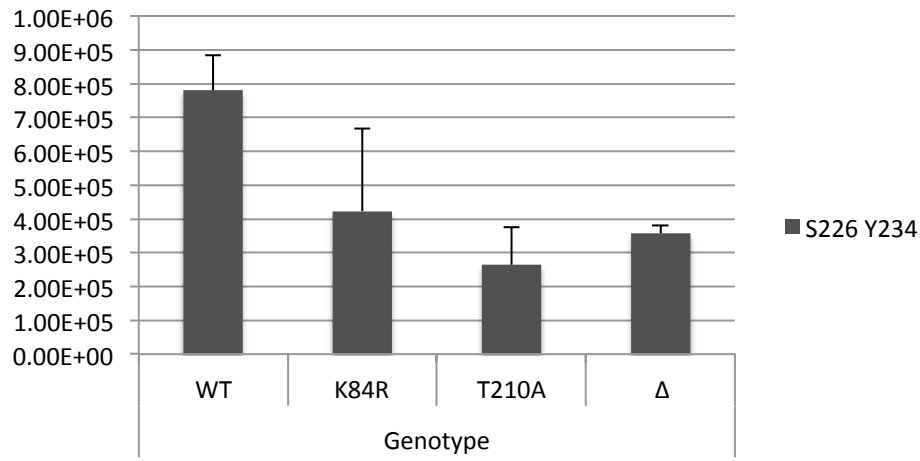


No data

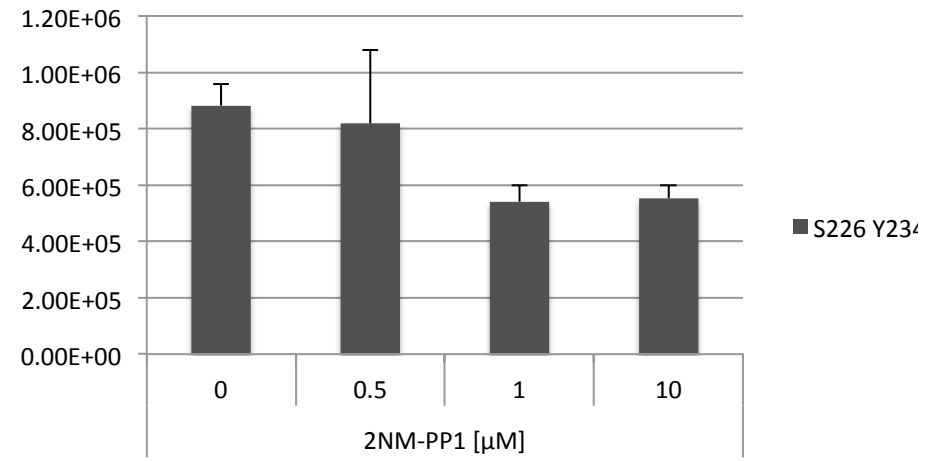
Min of derepression (48 = + 2 NM-PP1 (10 μM))

OAF3

OAF3



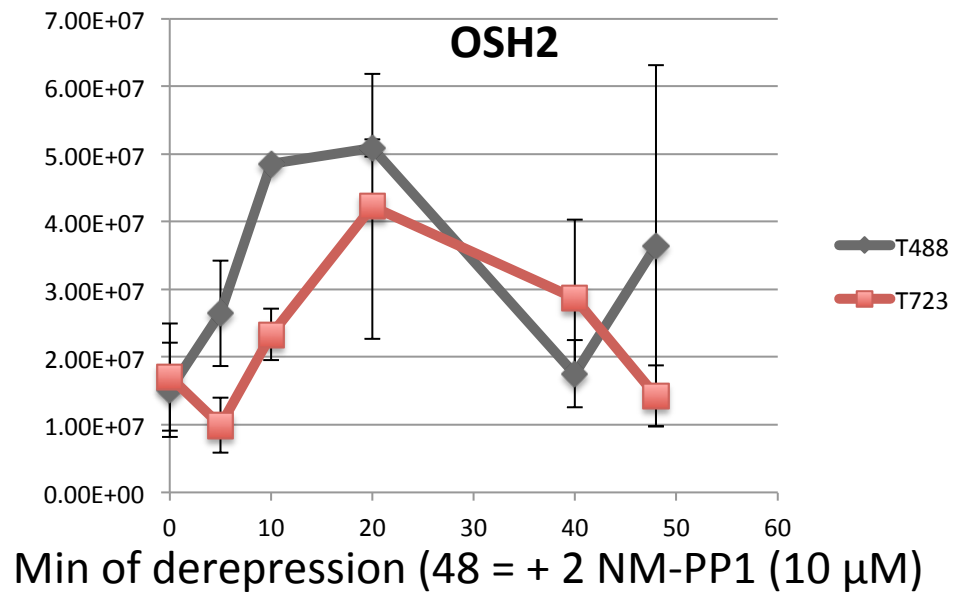
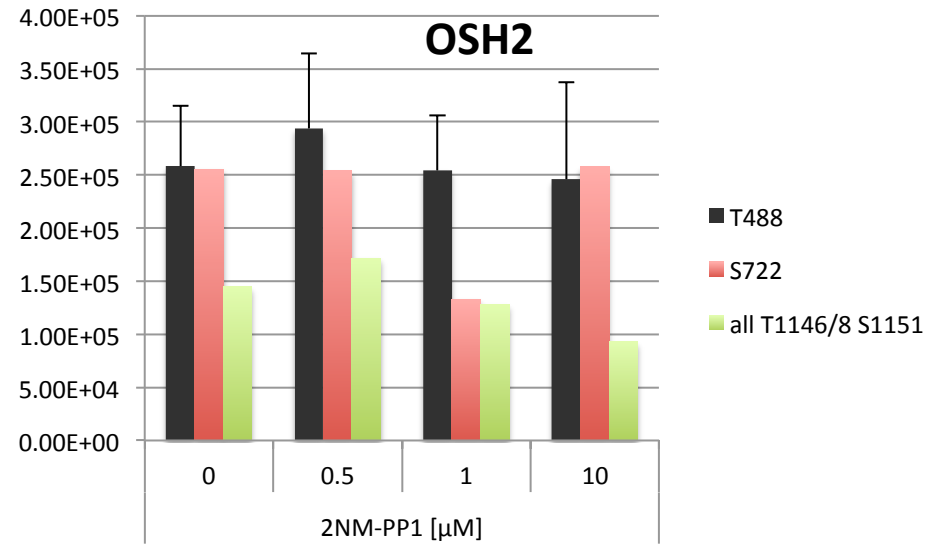
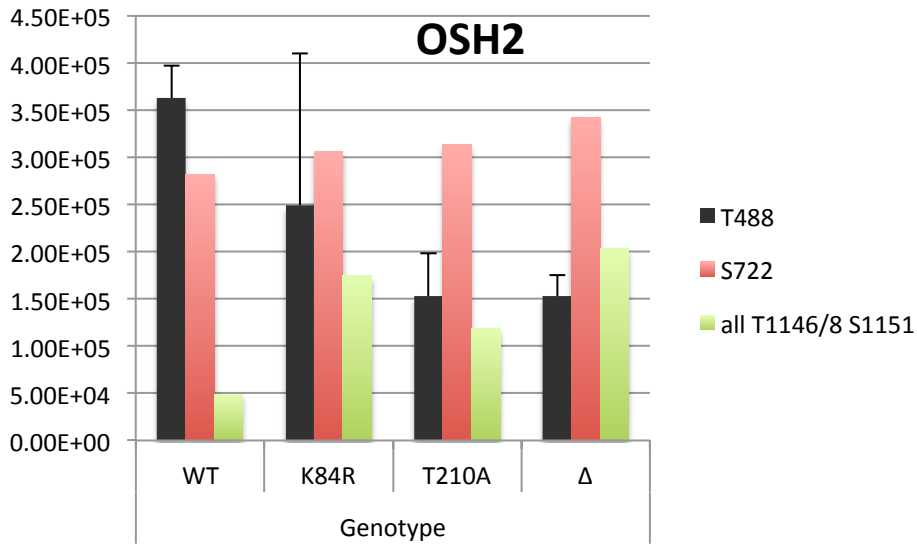
OAF3



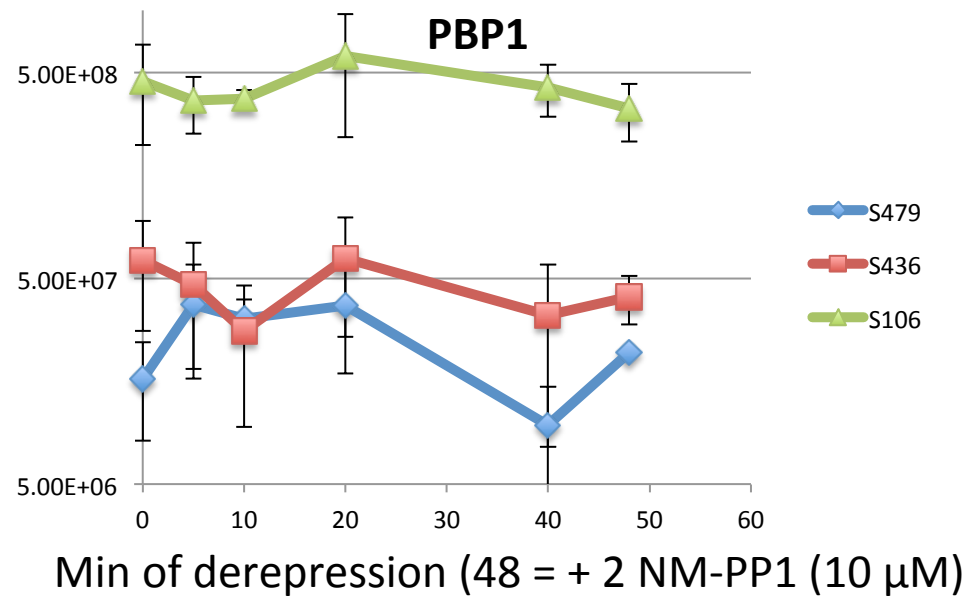
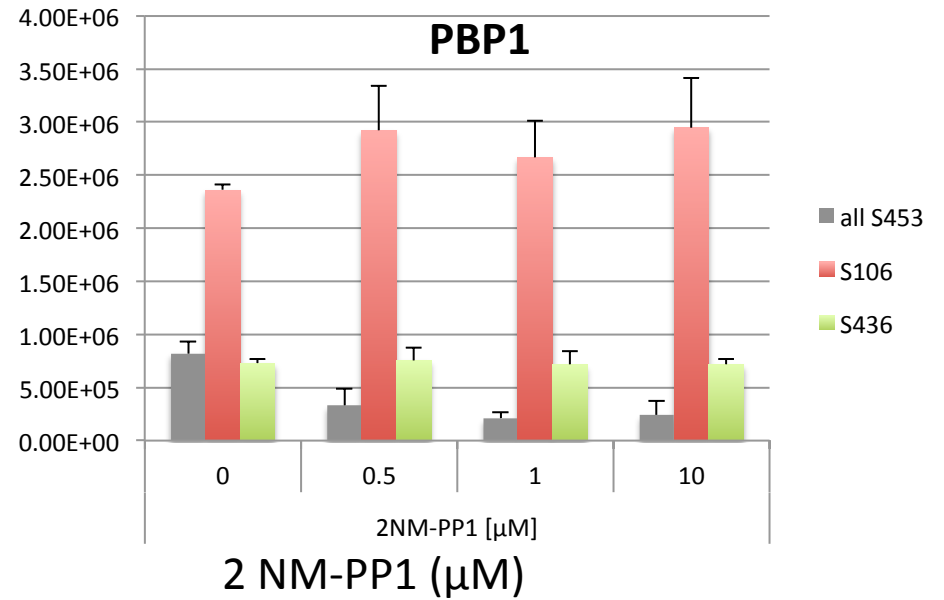
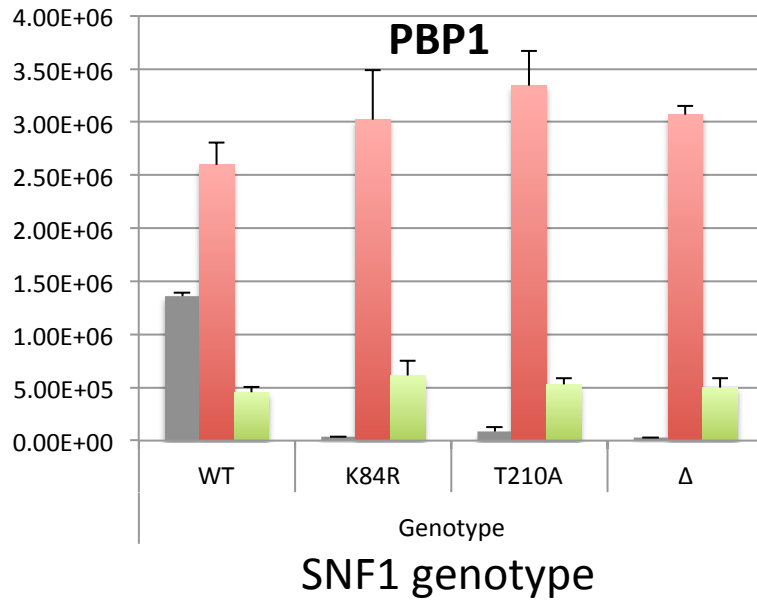
No data

Min of derepression (48 = + 2 NM-PP1 (10 μM))

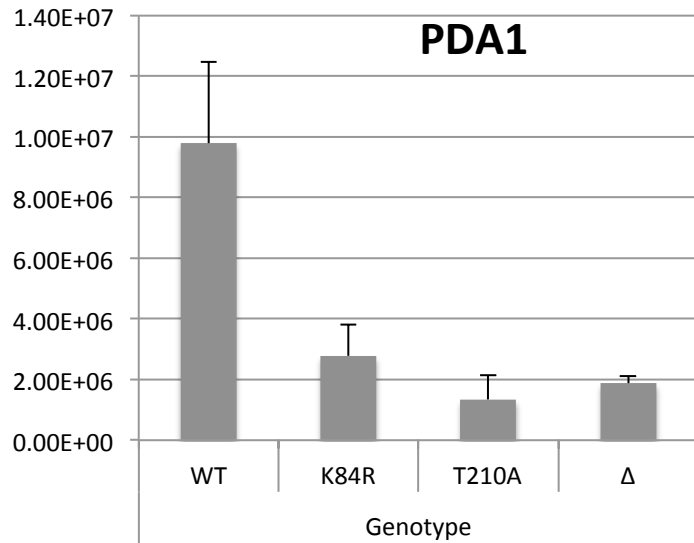
OSH2



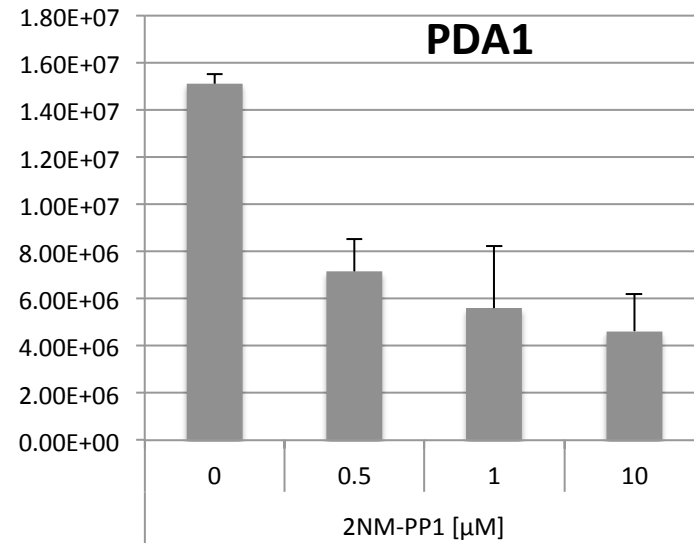
PBP1



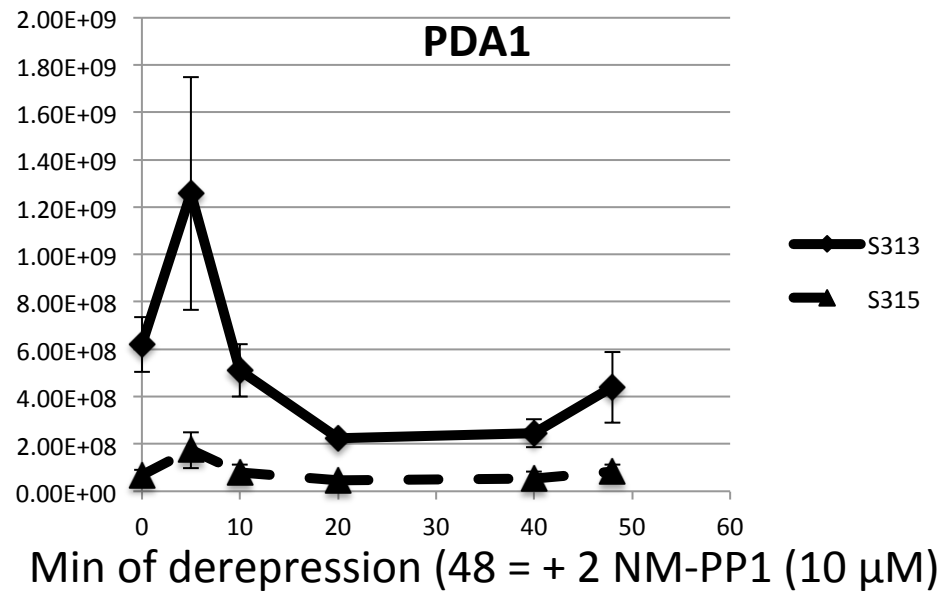
PDA1



SNF1 genotype

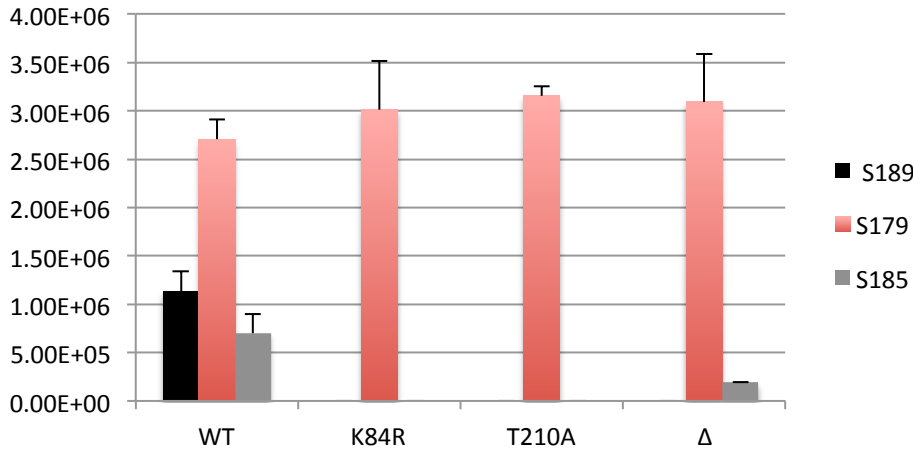


2 NM-PP1 (μM)



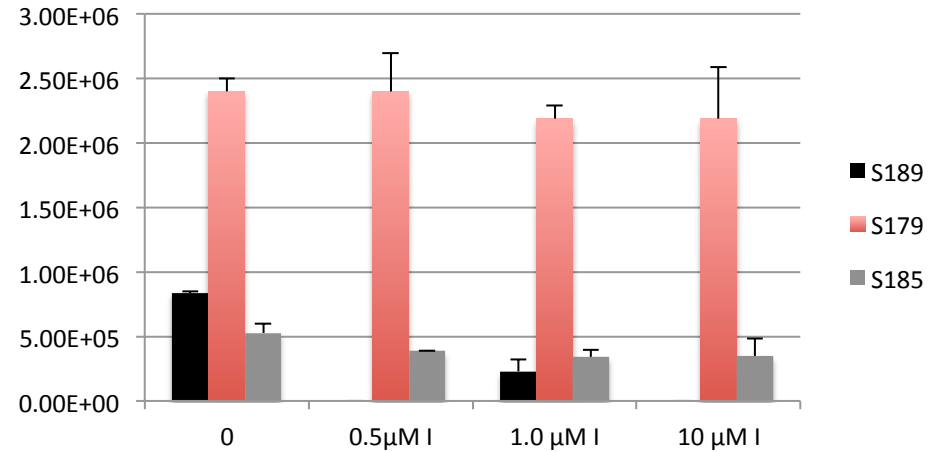
PFK1

PFK1

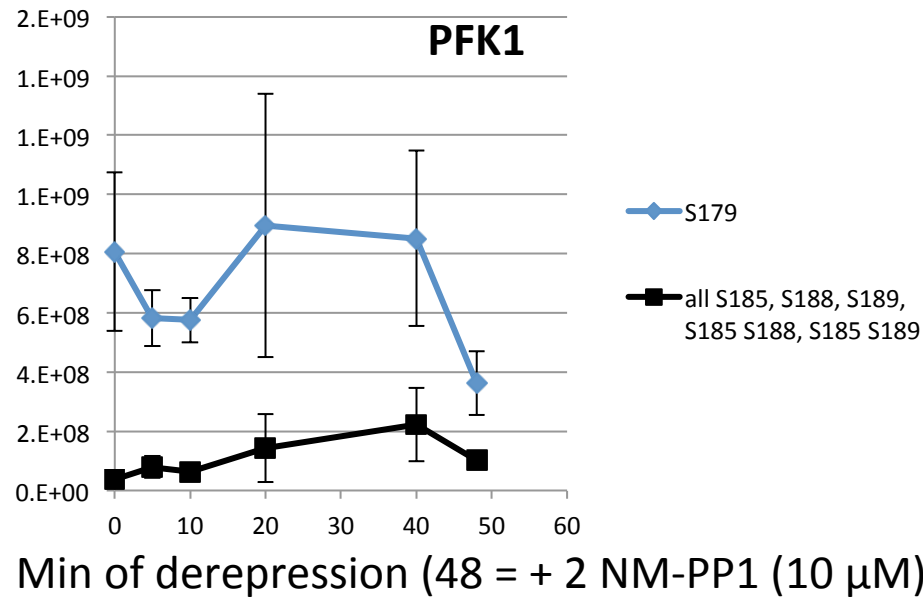


SNF1 genotype

PFK1

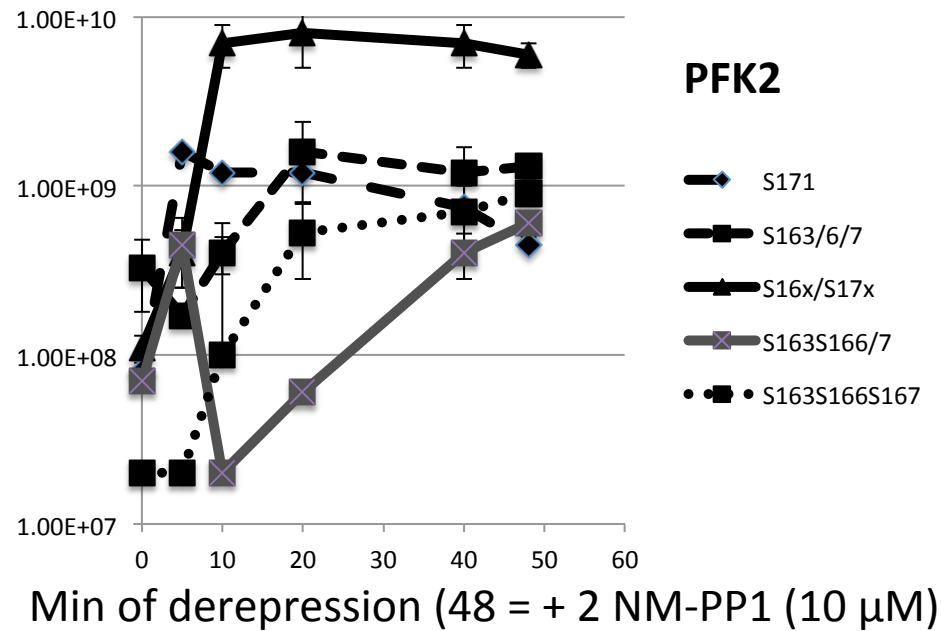
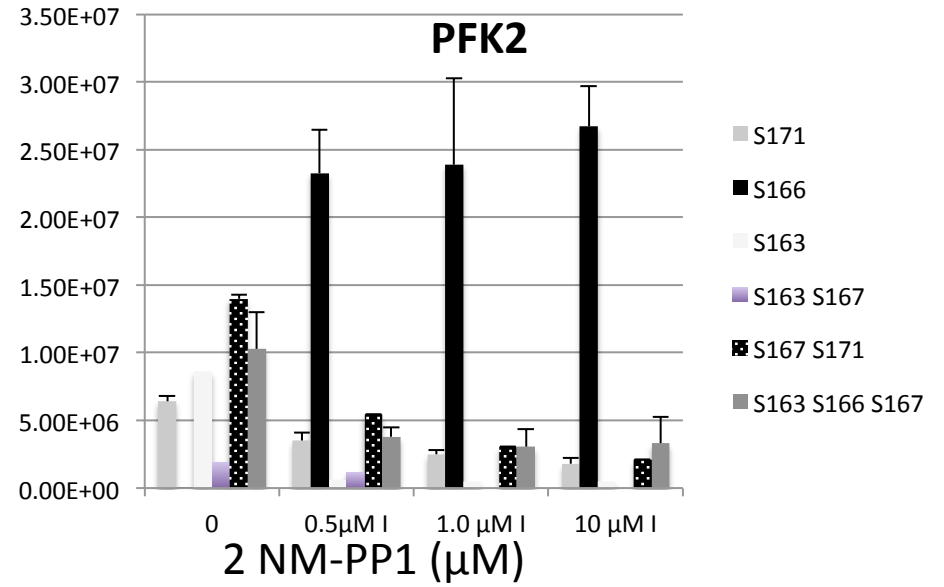
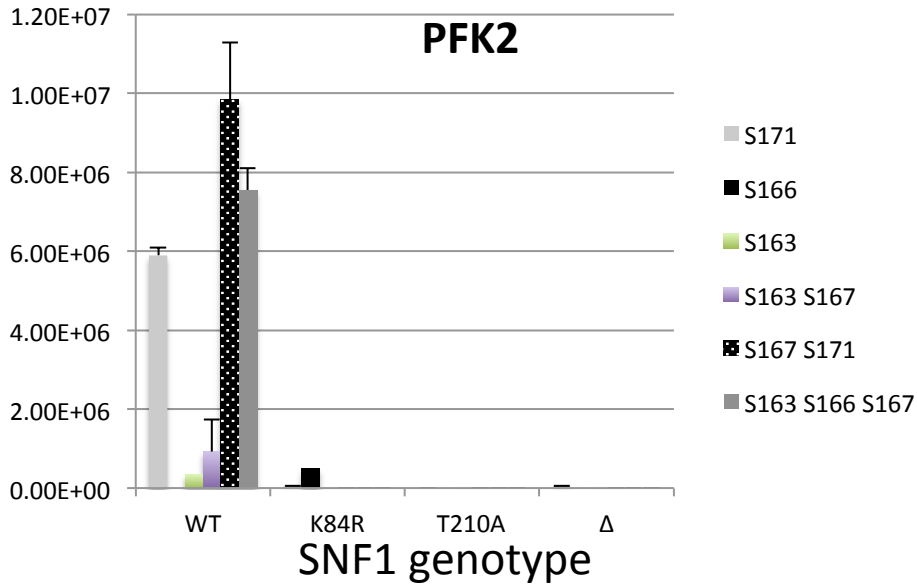


2 NM-PP1 (μM)

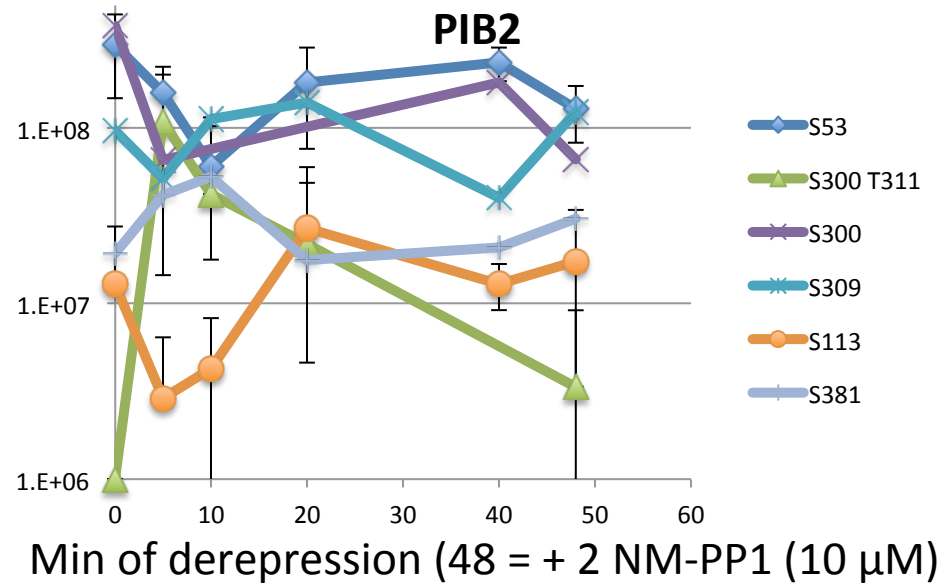
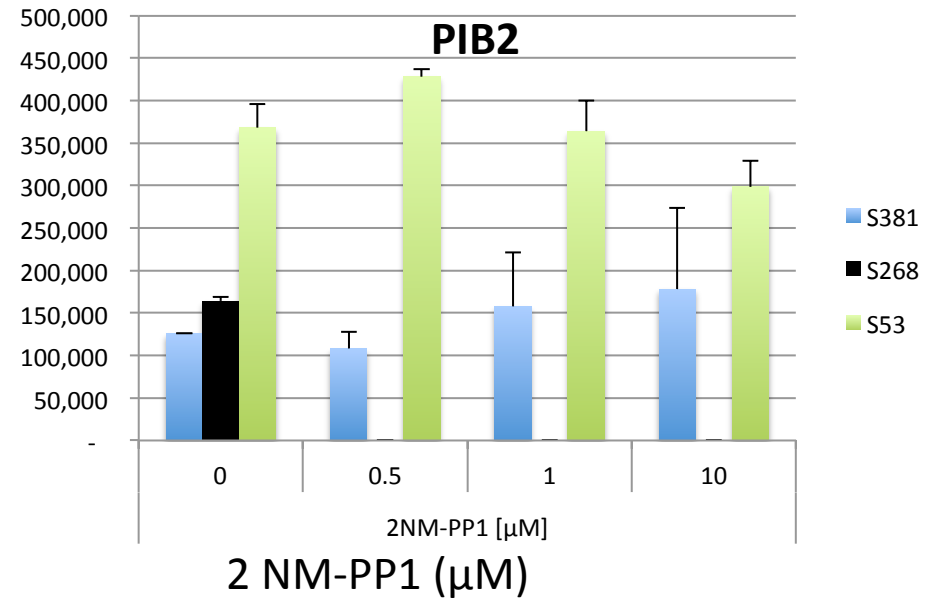
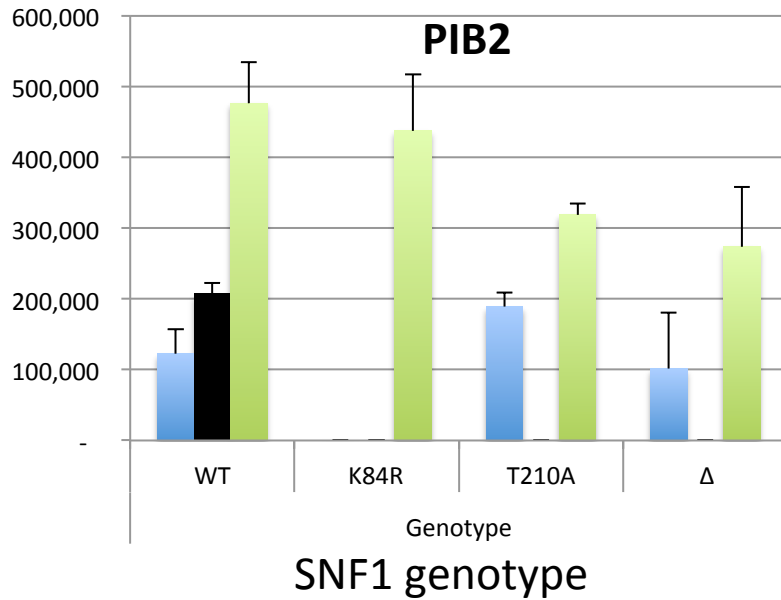


Min of derepression (48 = + 2 NM-PP1 (10 μM))

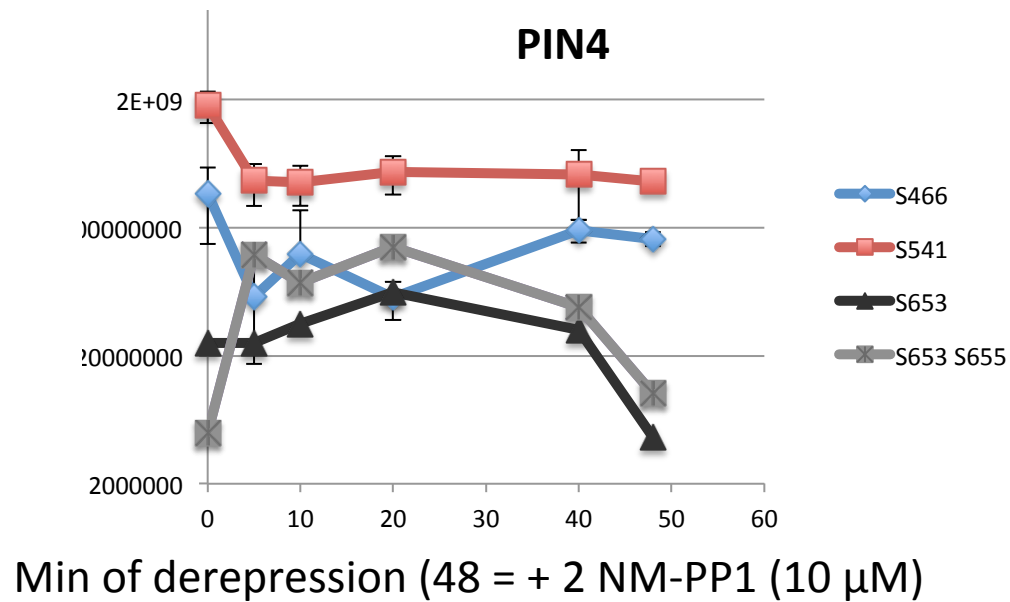
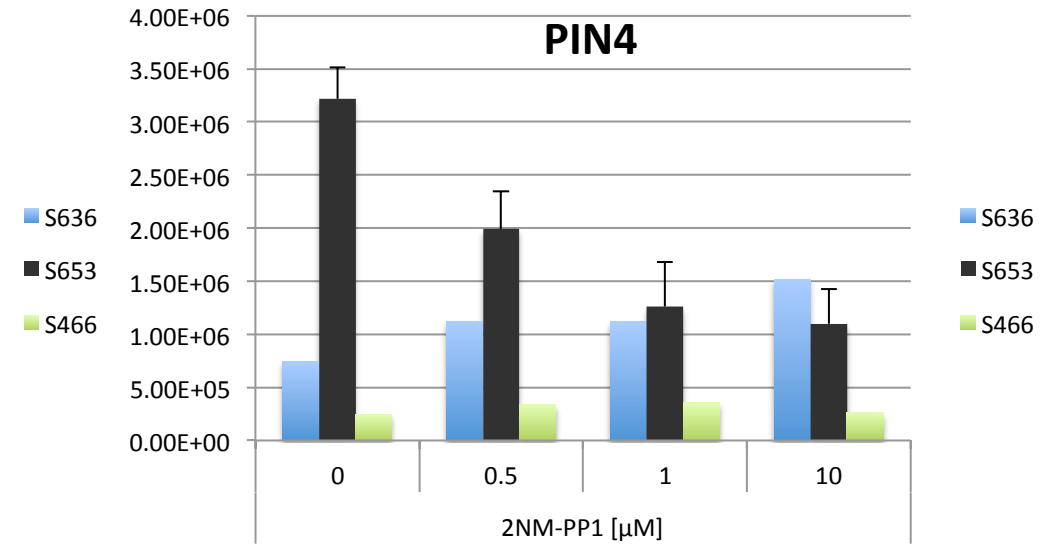
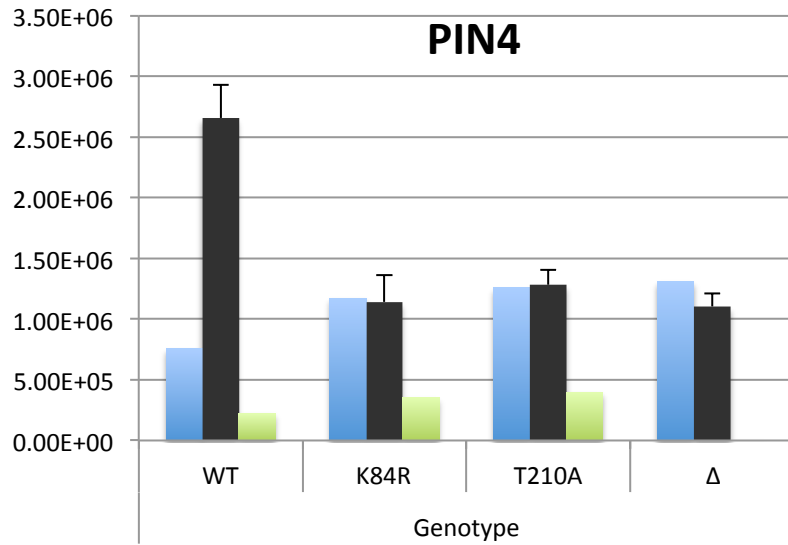
PFK2



PIB2

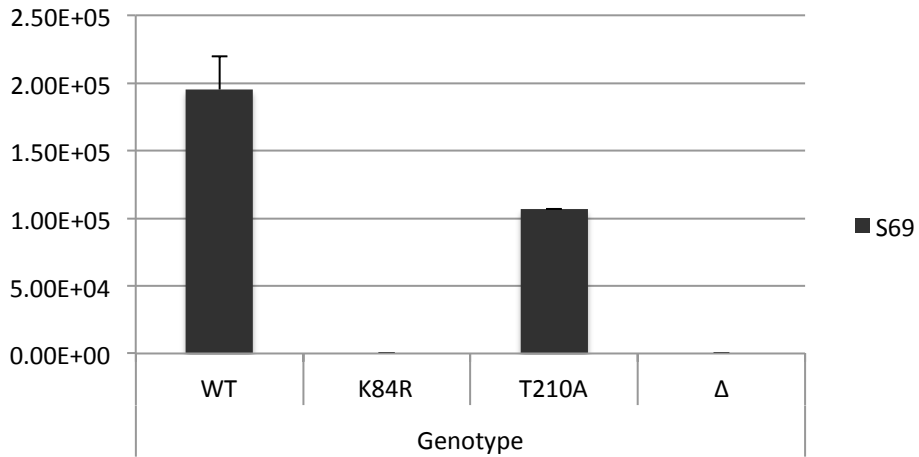


PIN4



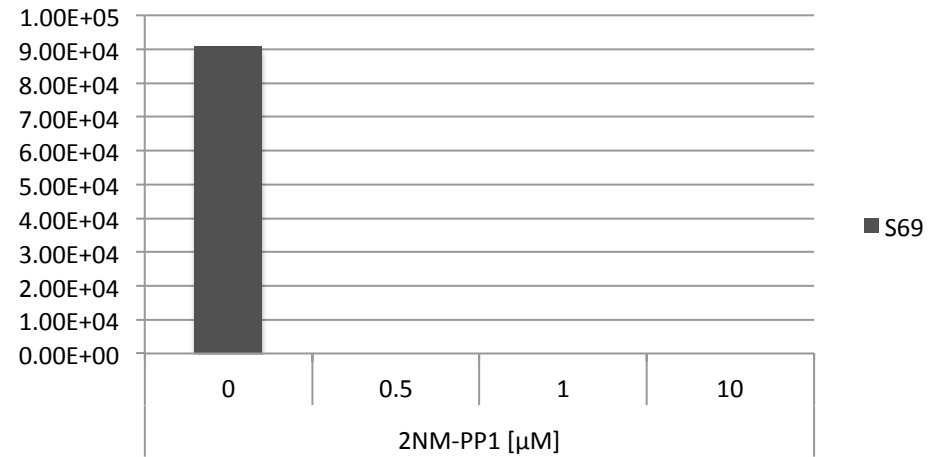
PRP28

PRP28



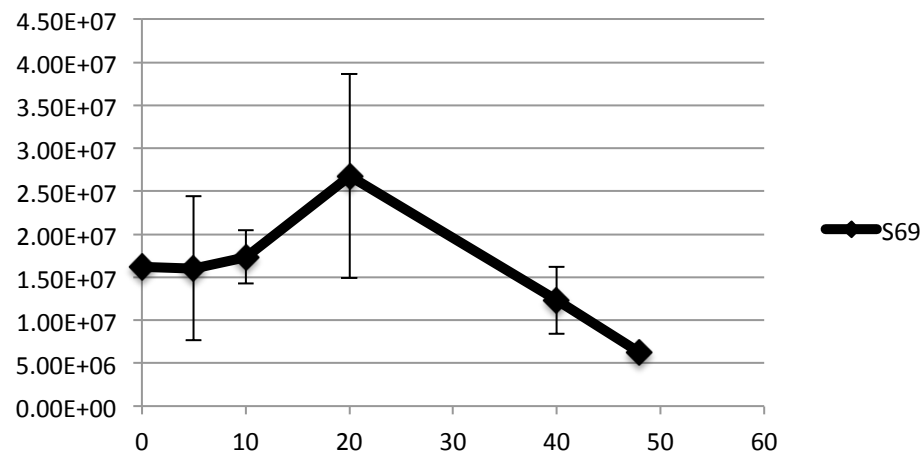
SNF1 genotype

PRP28



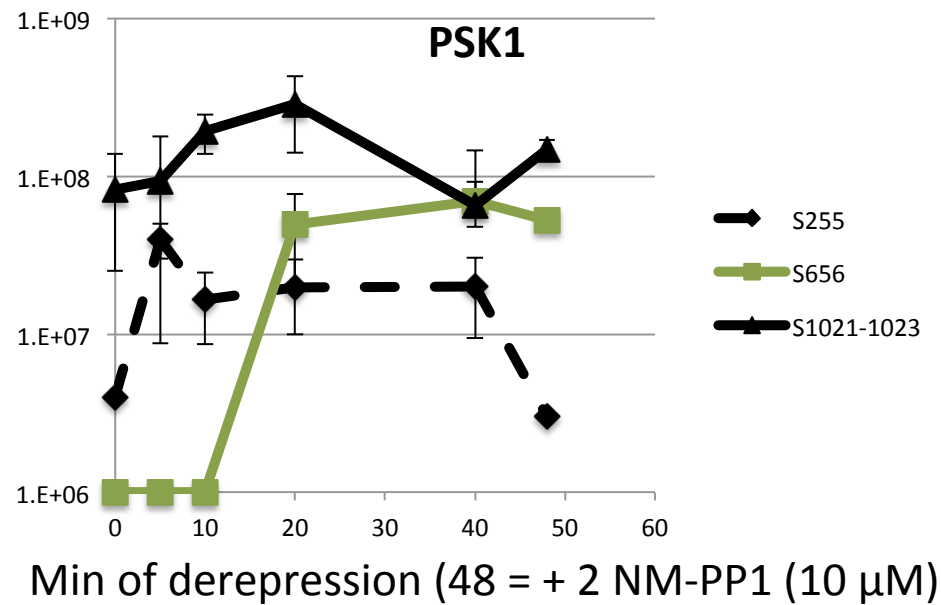
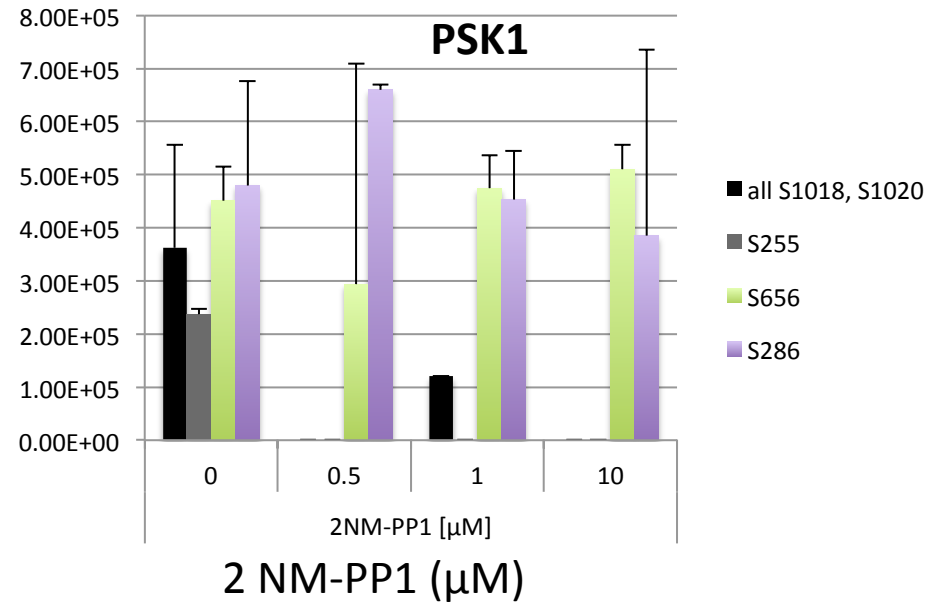
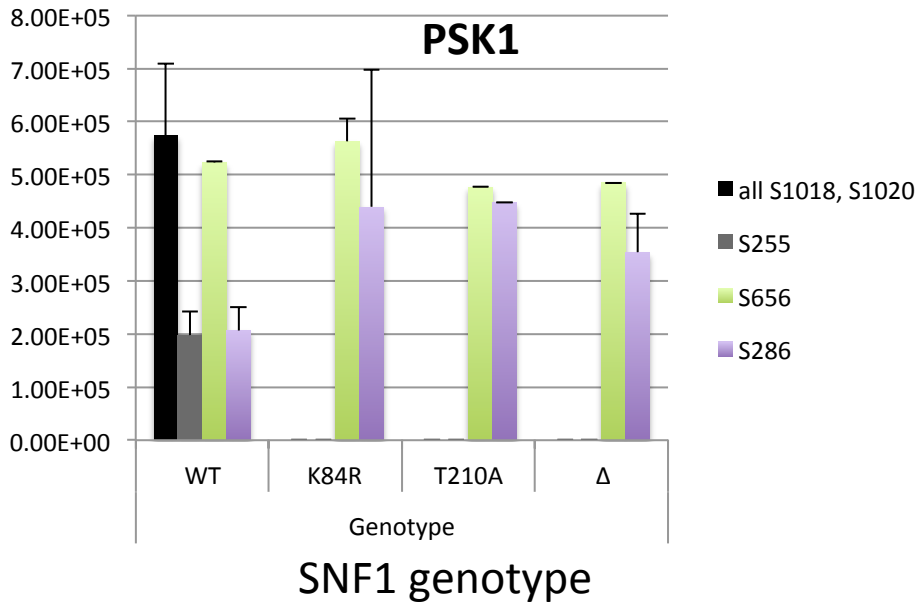
2NM-PP1 (μM)

PRP28

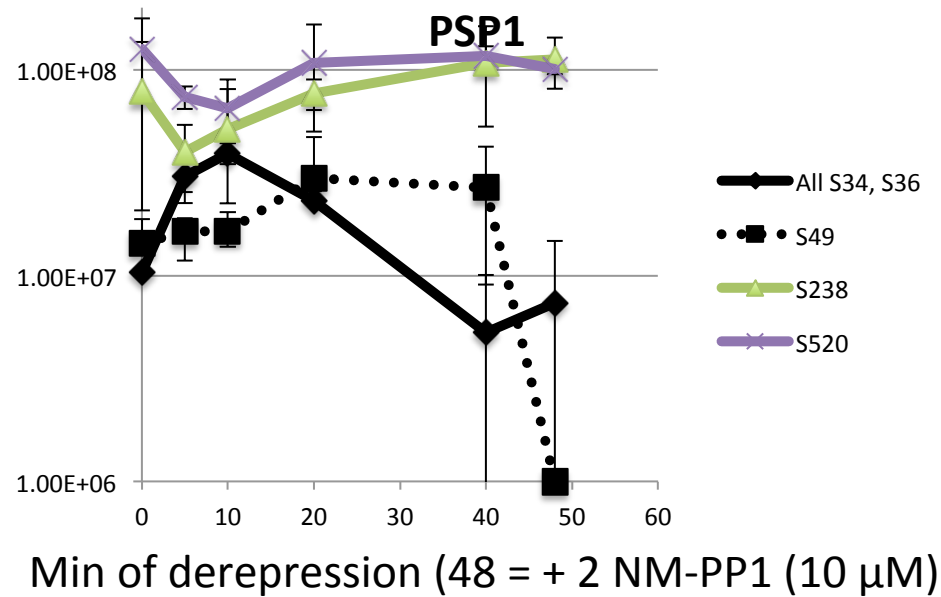
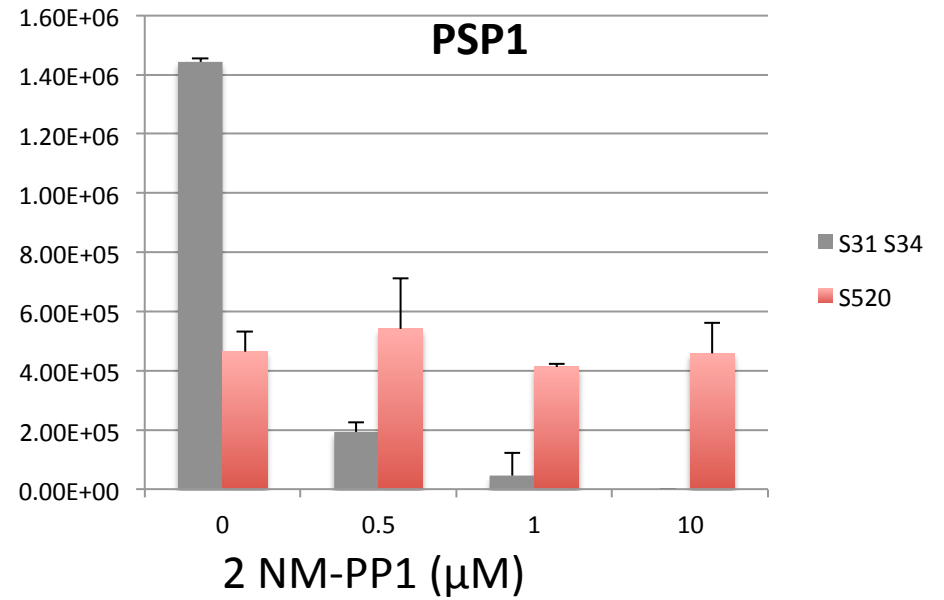
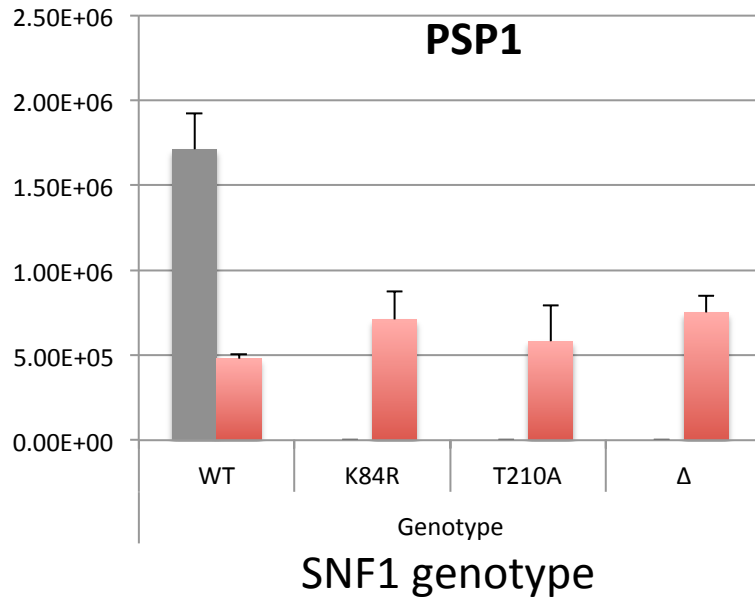


Min of derepression (48 = + 2 NM-PP1 (10 μM))

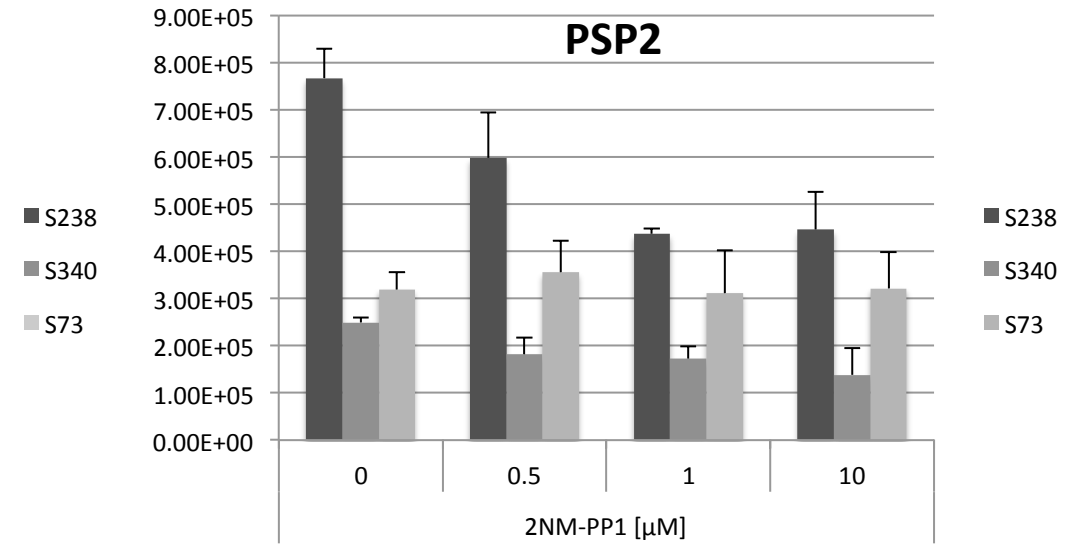
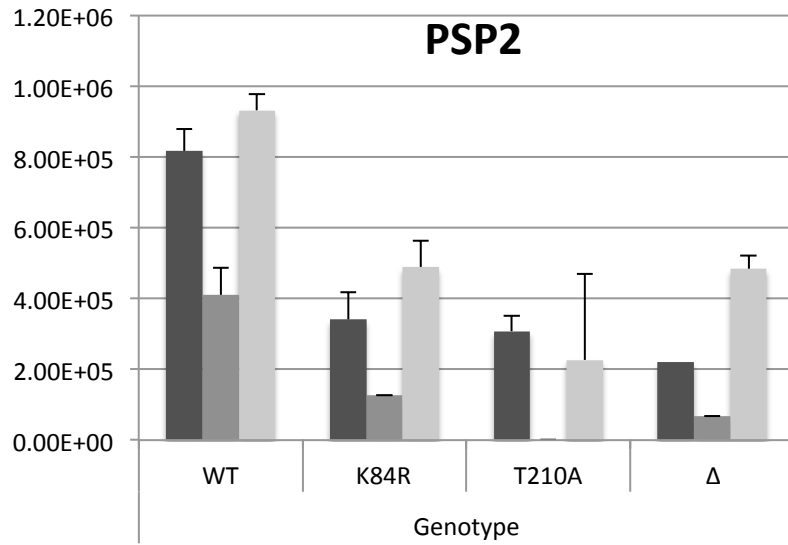
PSK1



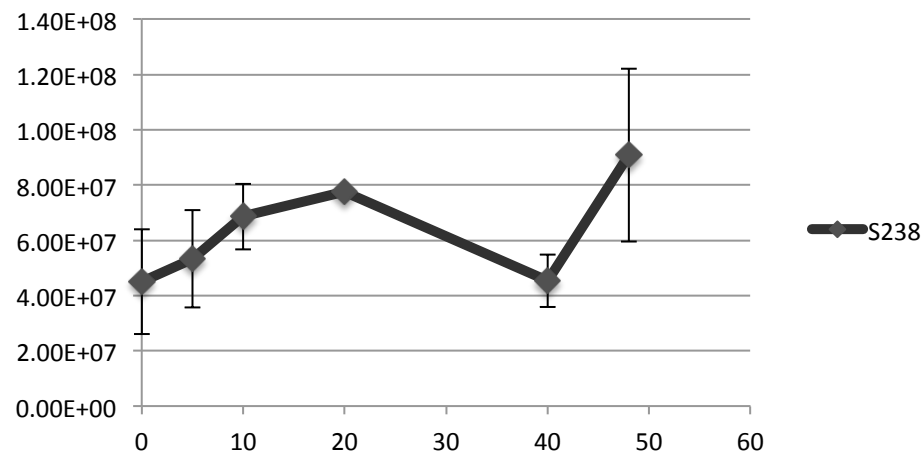
PSP1



PSP2

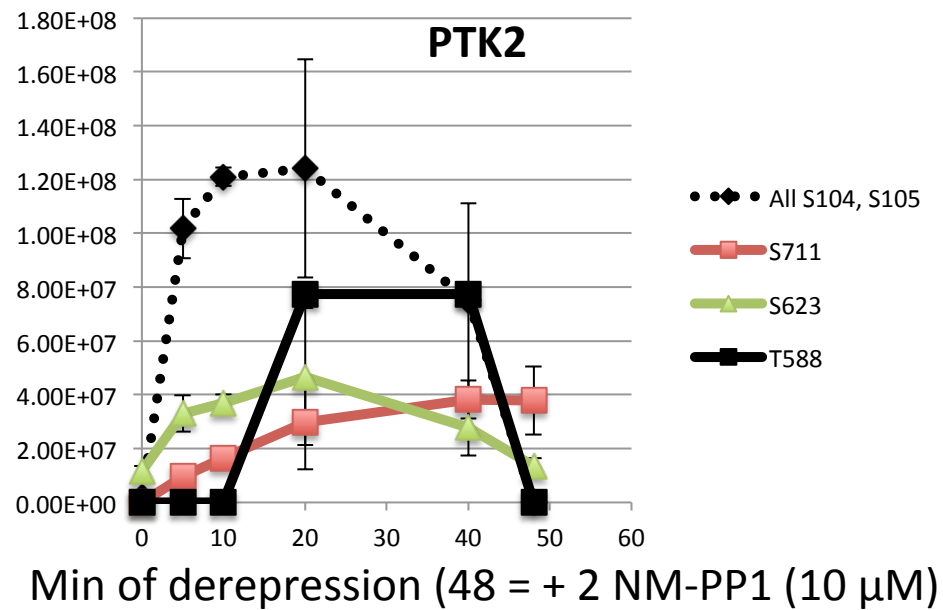
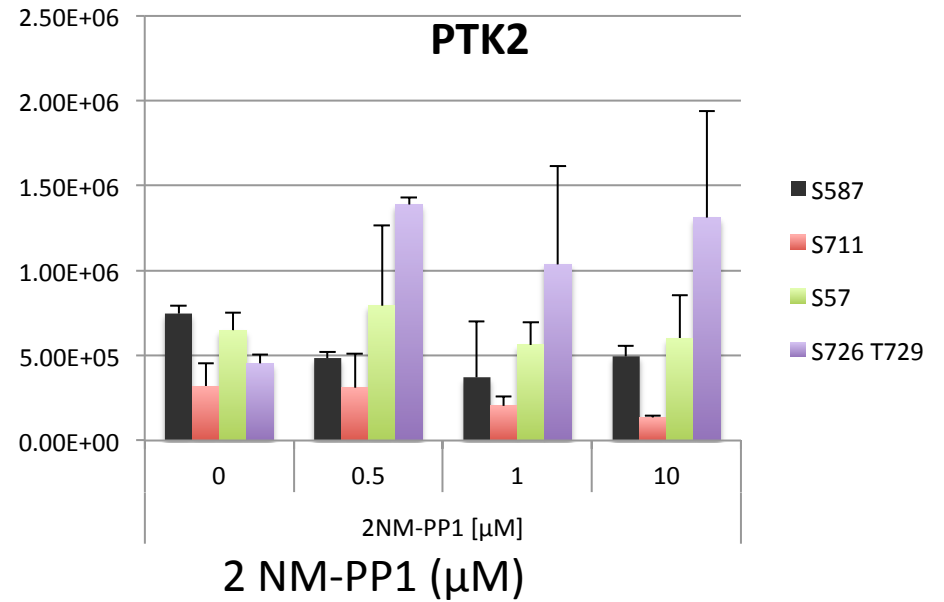
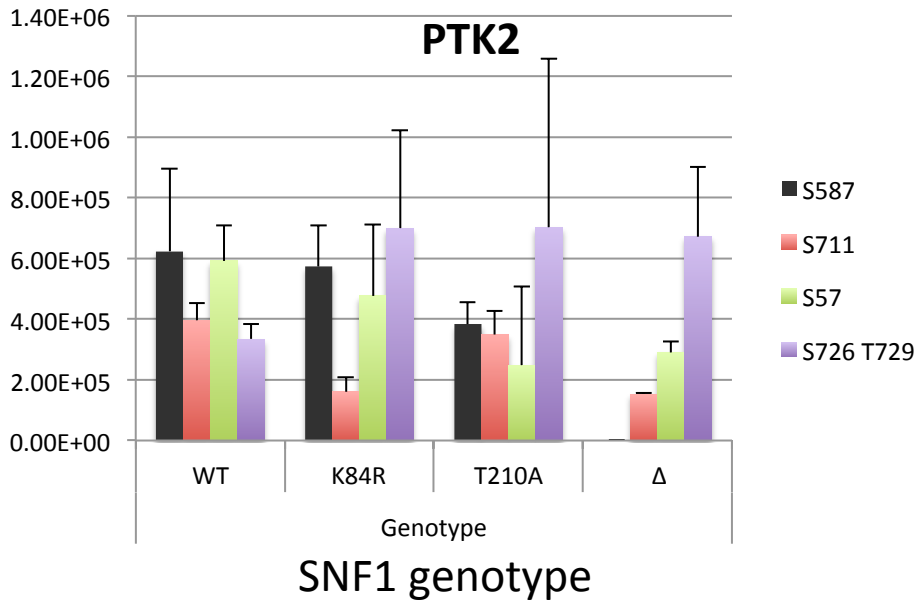


PSP2

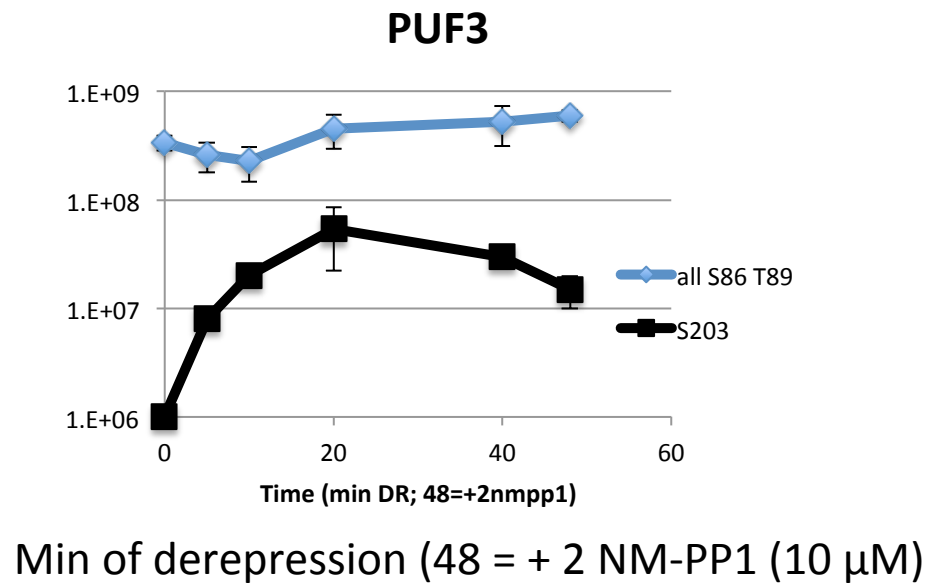
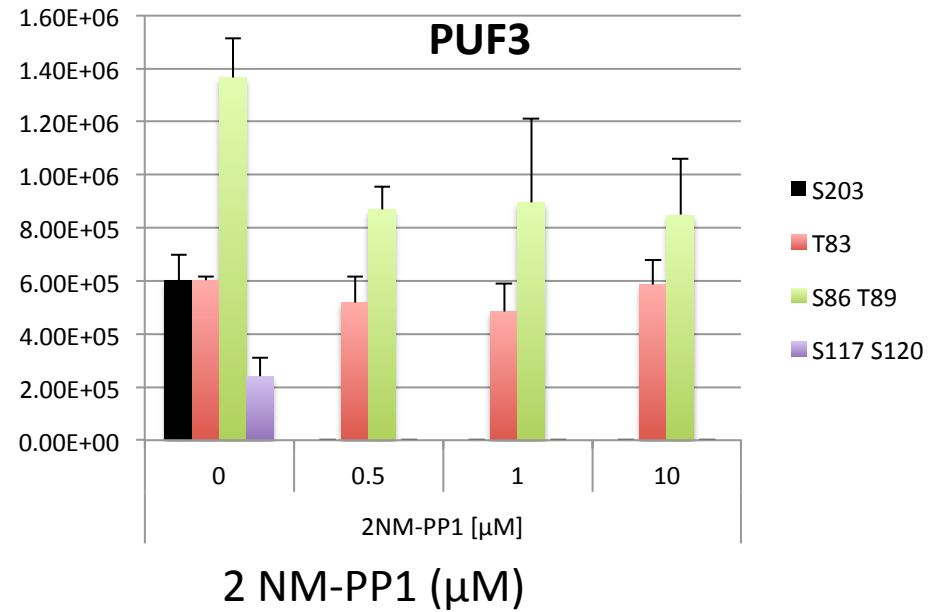
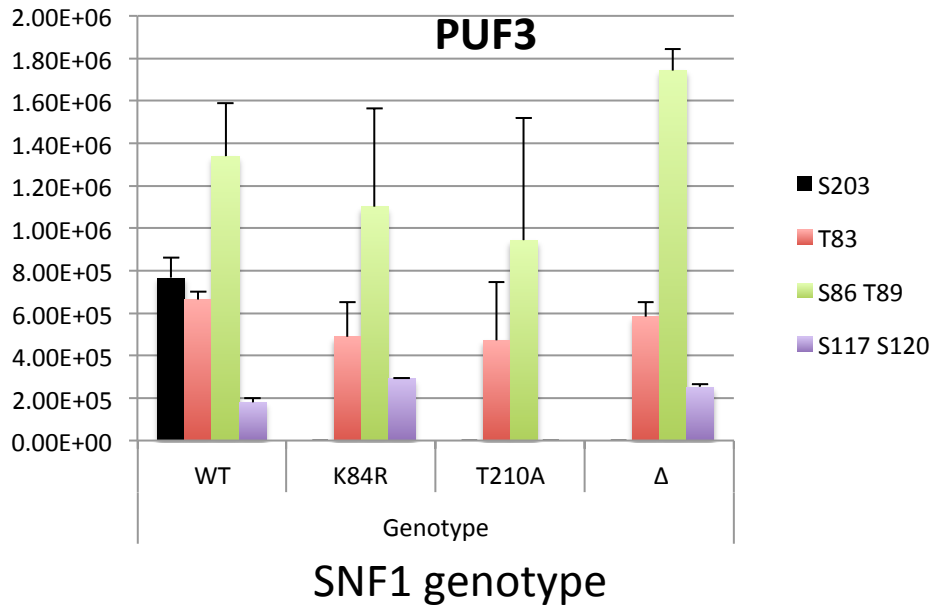


Min of derepression (48 = + 2 NM-PP1 (10 μM))

PTK2

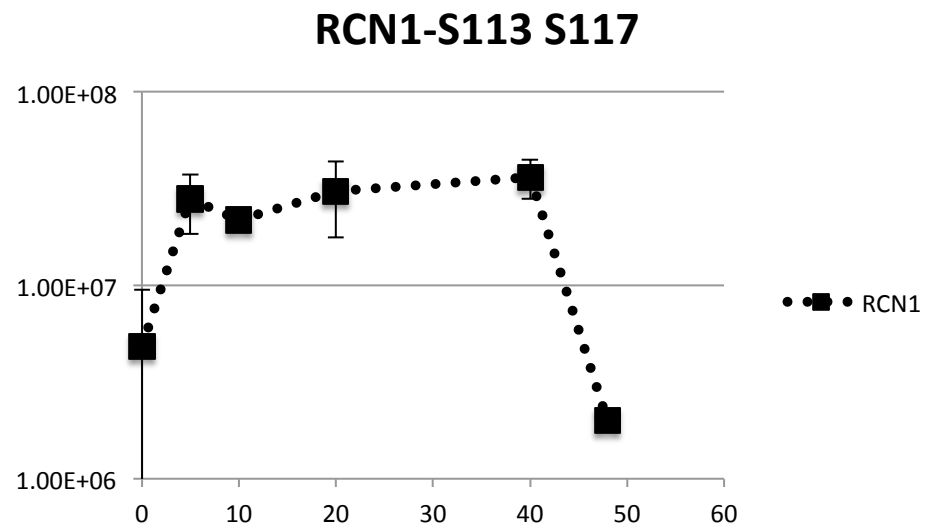


PUF3



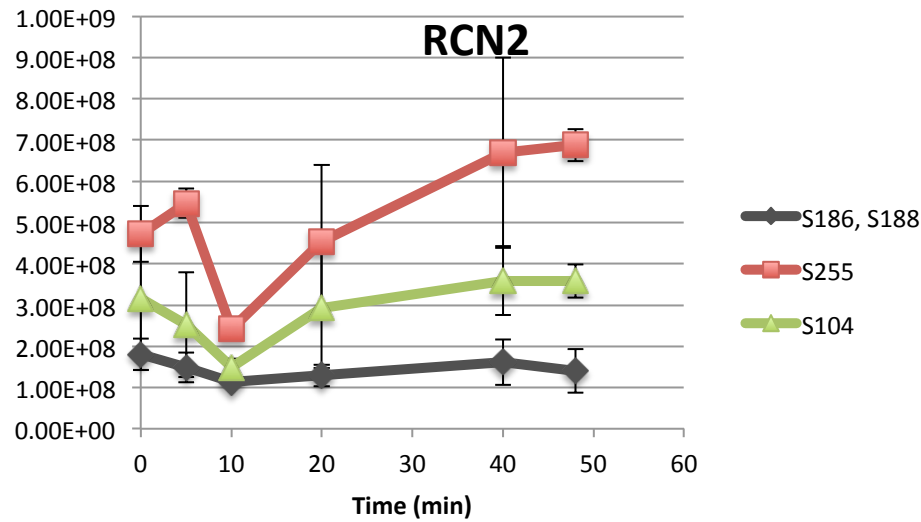
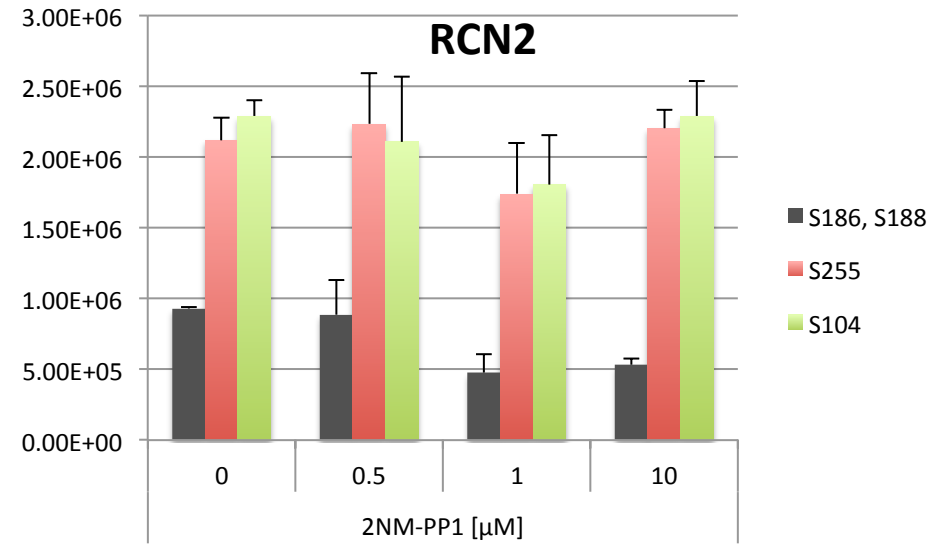
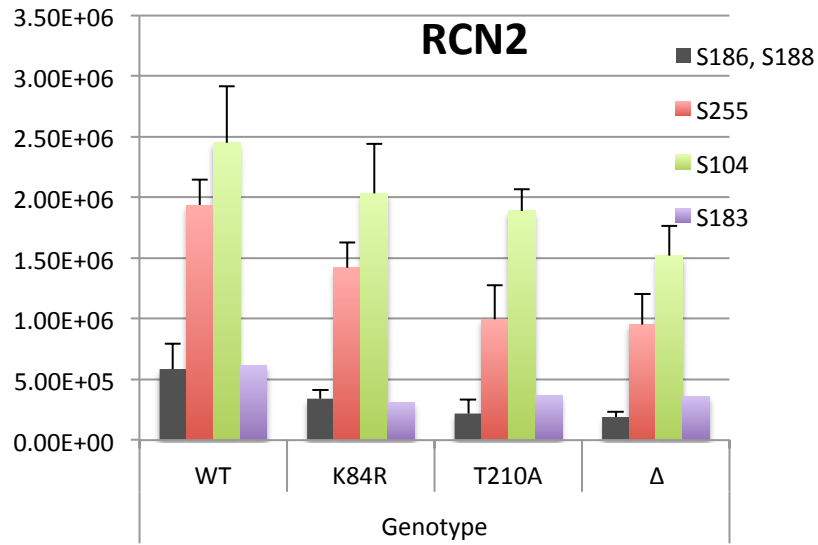
RCN1

No data in experiments 1 and 2



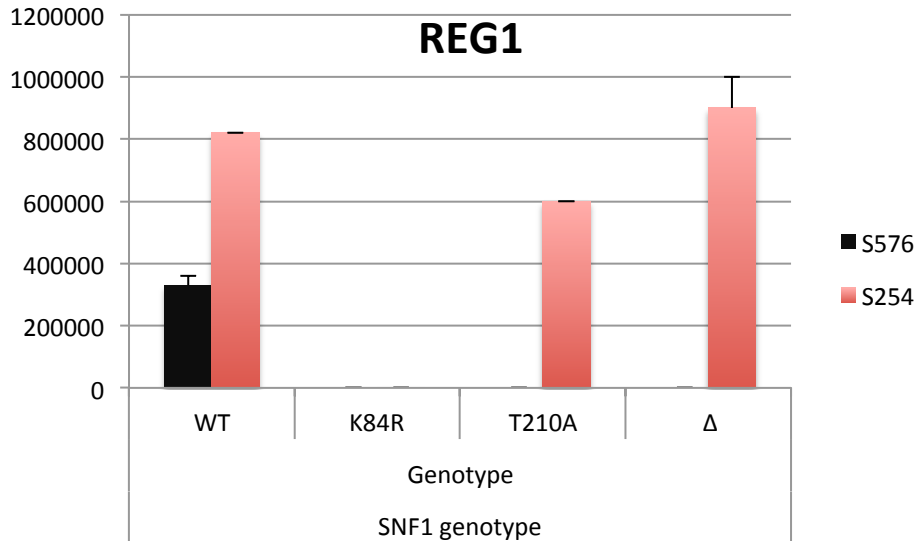
Min of derepression (48 = + 2 NM-PP1 (10 μ M))

RCN2



Min of derepression (48 = + 2 NM-PP1 (10 μ M))

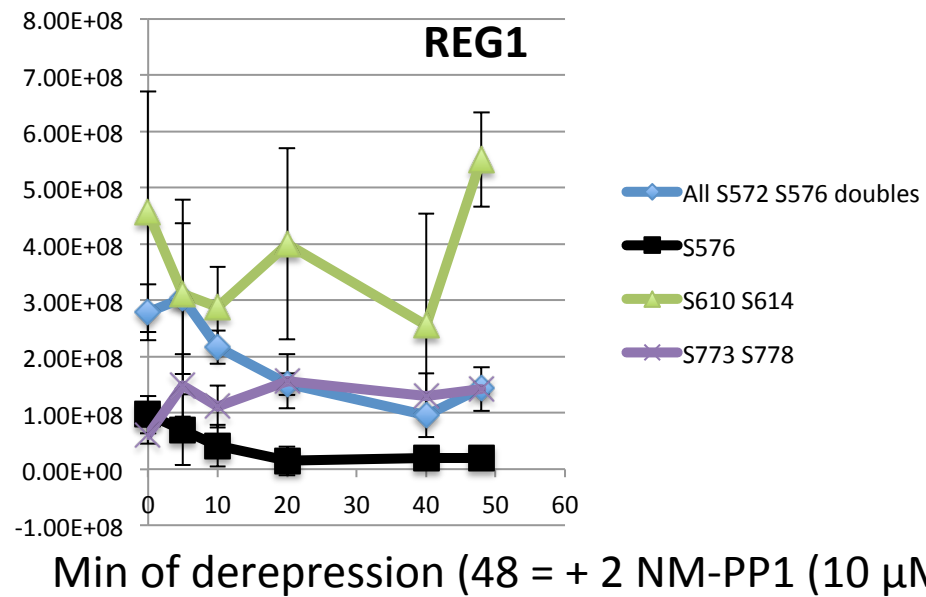
REG1



No data

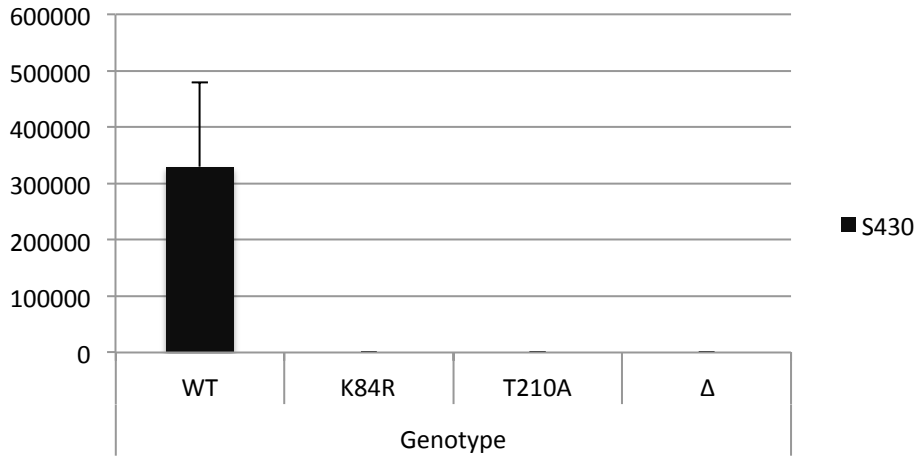
SNF1 genotype

2 NM-PP1 (μM)



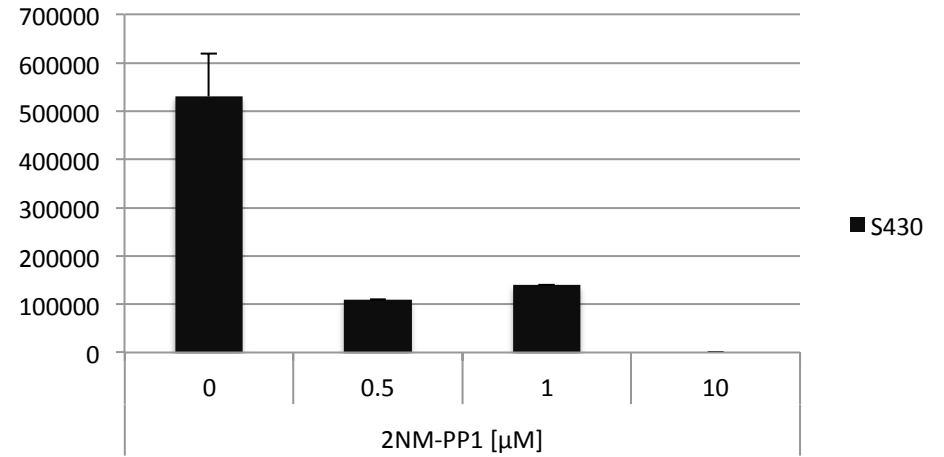
RIA1

RIA1



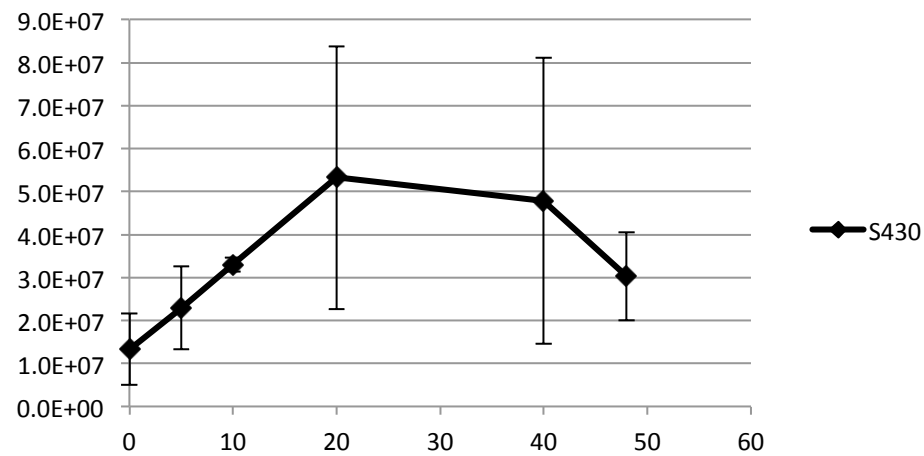
SNF1 genotype

RIA1



2 NM-PP1 (μM)

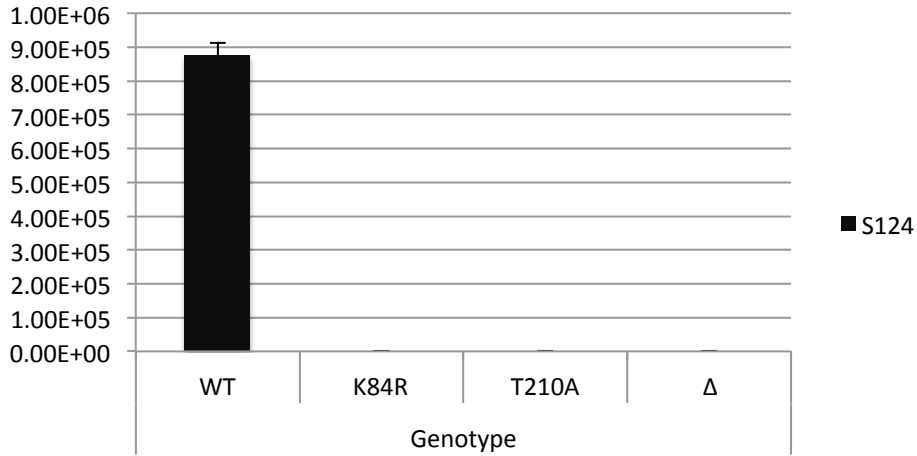
RIA1



Min of derepression (48 = + 2 NM-PP1 (10 μM))

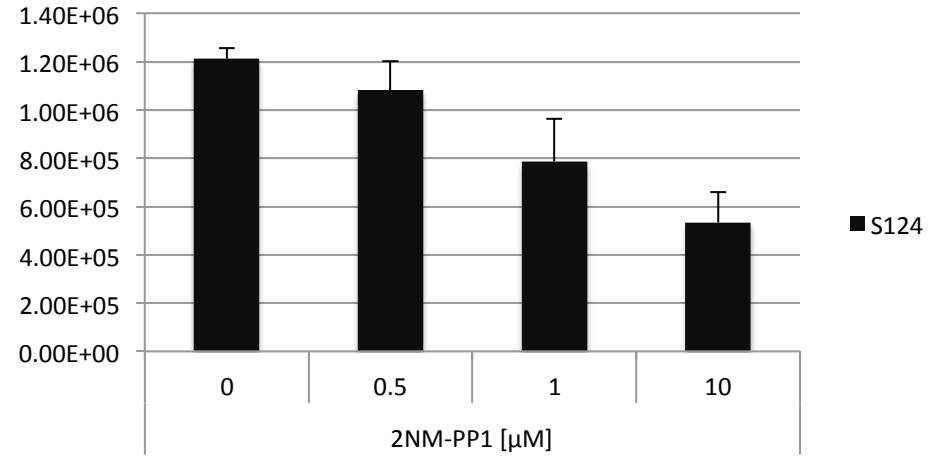
ROD1

ROD1

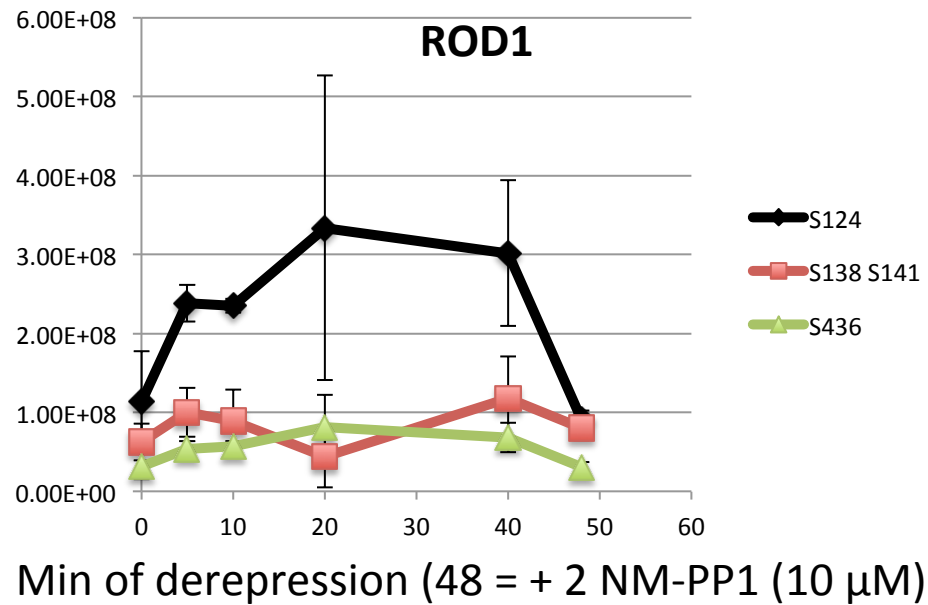


SNF1 genotype

ROD1



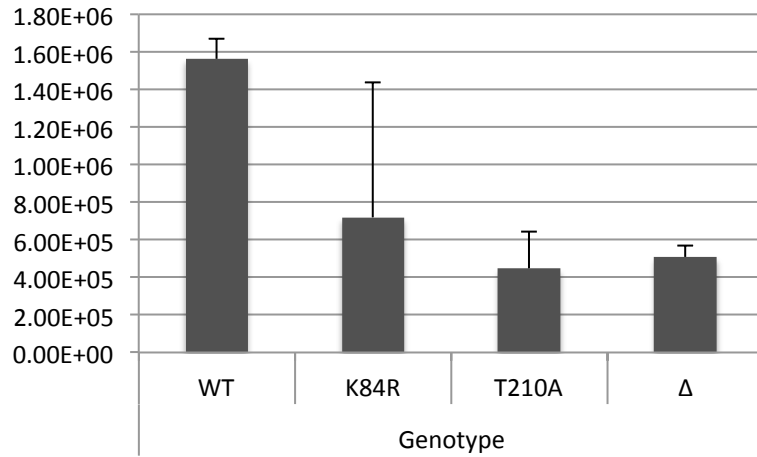
2 NM-PP1 (μM)



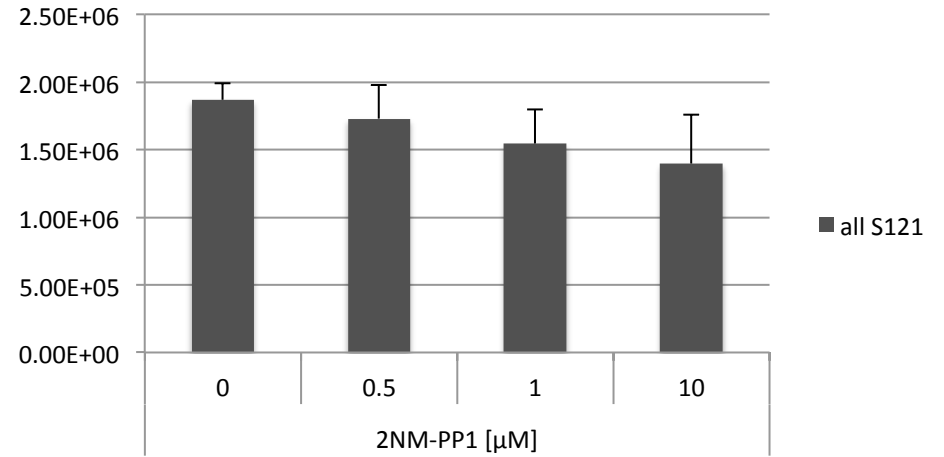
Min of derepression (48 = + 2 NM-PP1 (10 μM))

RPA14

RPA14



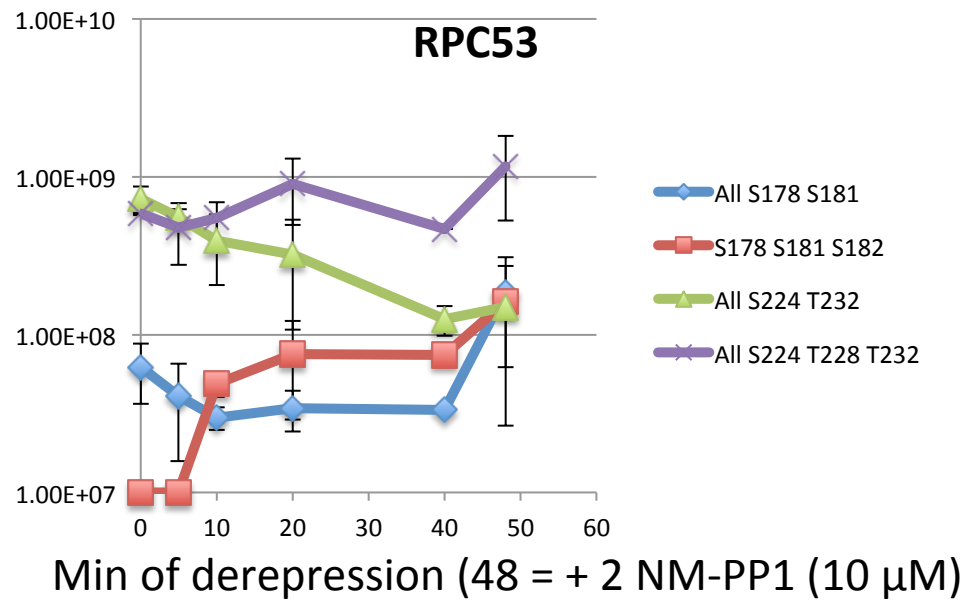
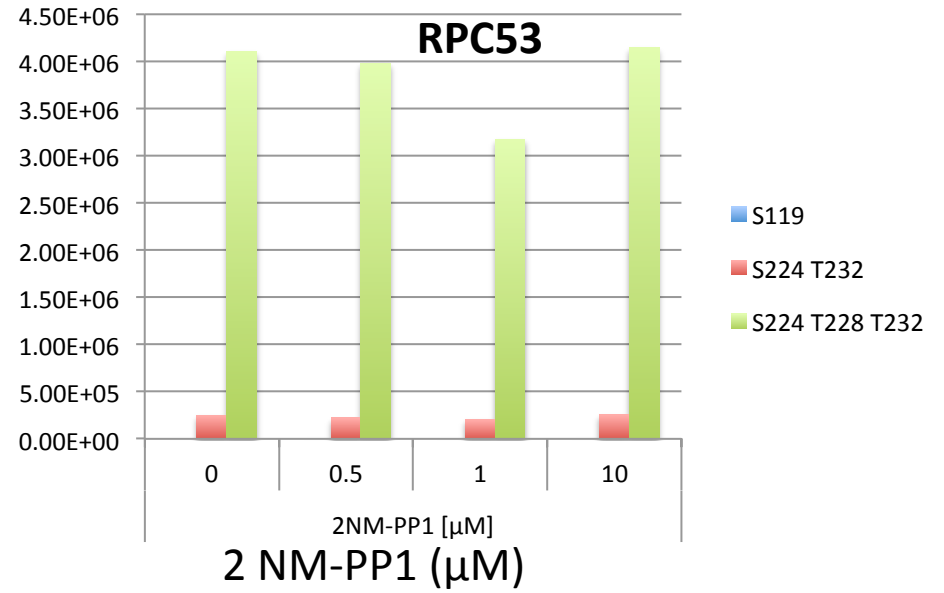
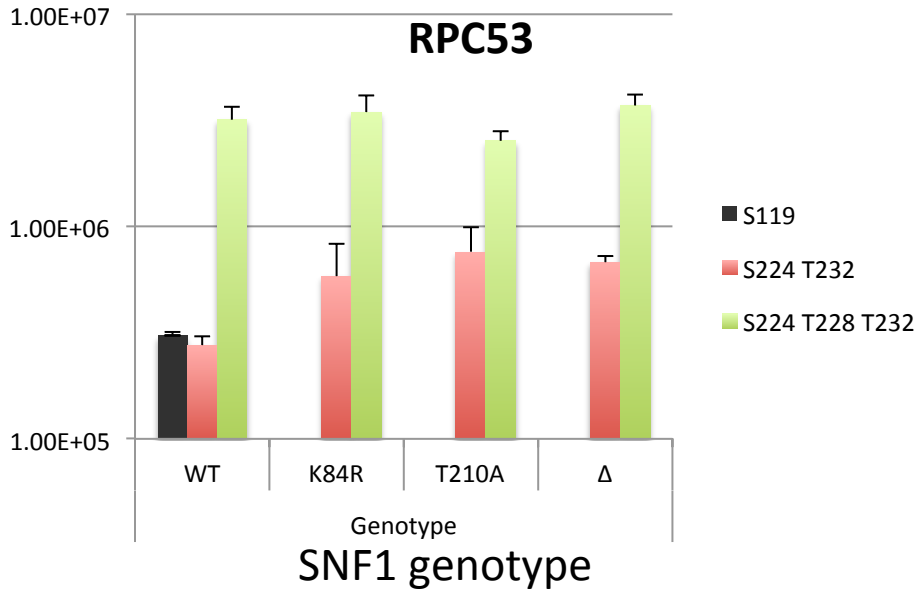
RPA14



No data

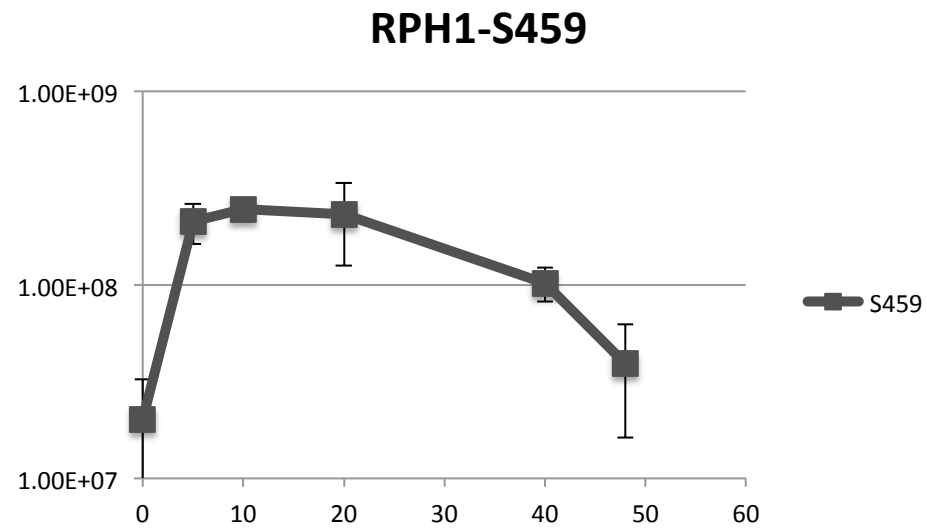
Min of derepression (48 = + 2 NM-PP1 (10 μM))

RPC53



RPH1

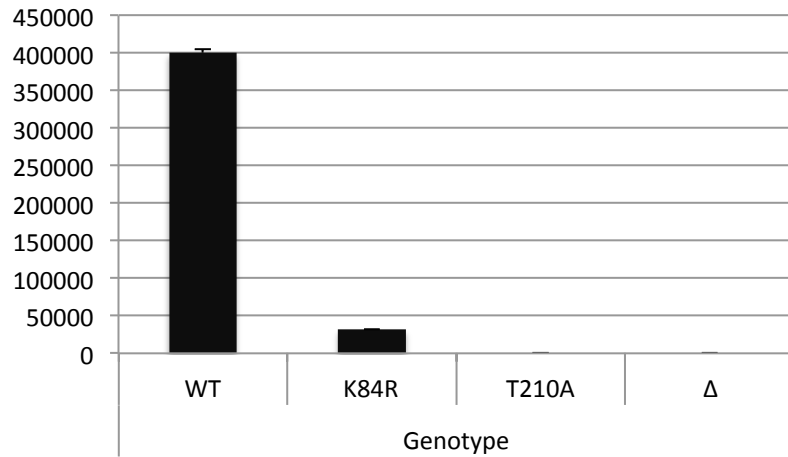
No data in experiments 1 and 2



Min of derepression (48 = + 2 NM-PP1 (10 μ M))

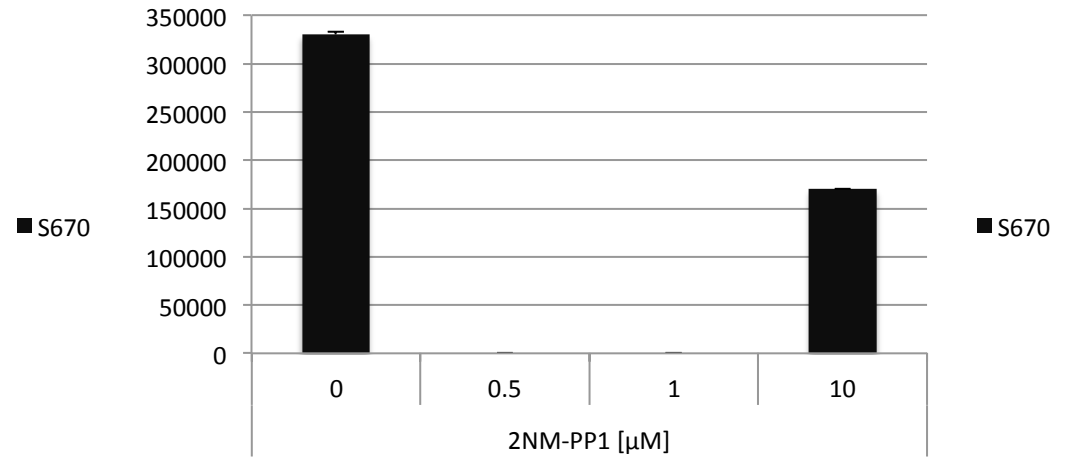
RSC1

RSC1



SNF1 genotype

RSC1



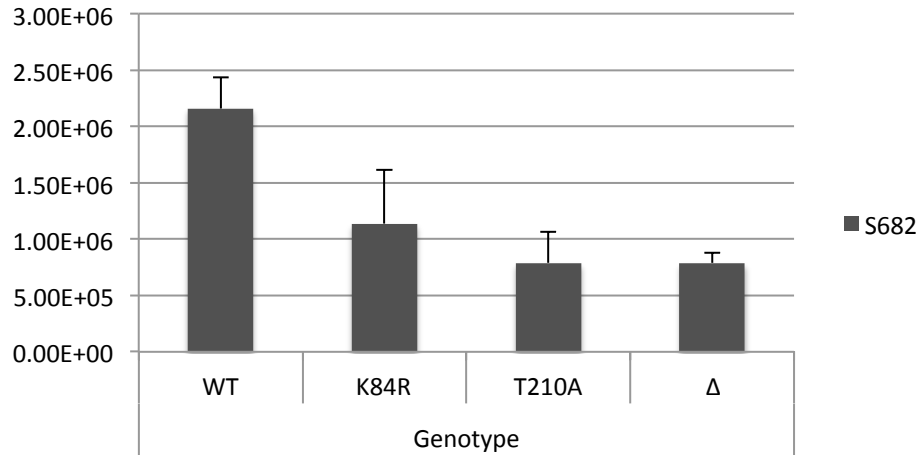
2 NM-PP1 (μM)

No data

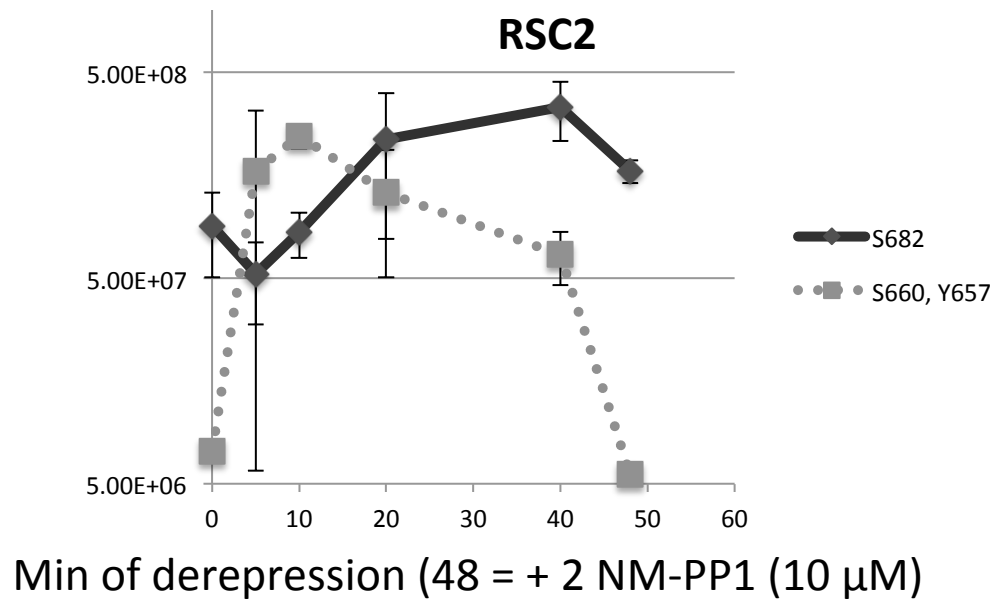
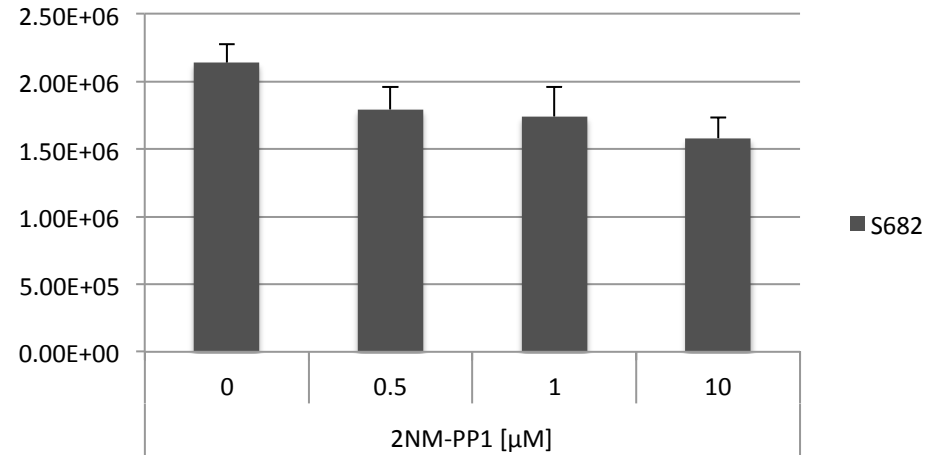
Min of derepression (48 = + 2 NM-PP1 (10 μM))

RSC2

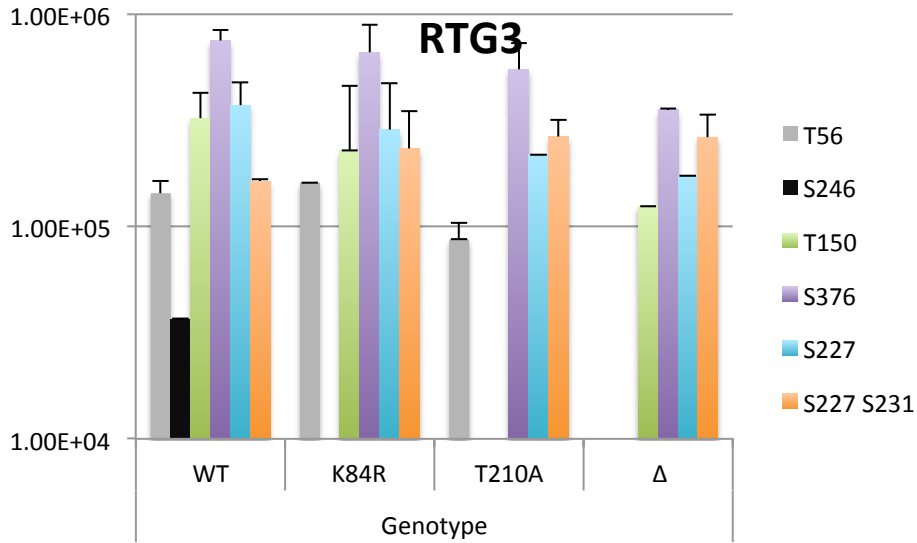
RSC2



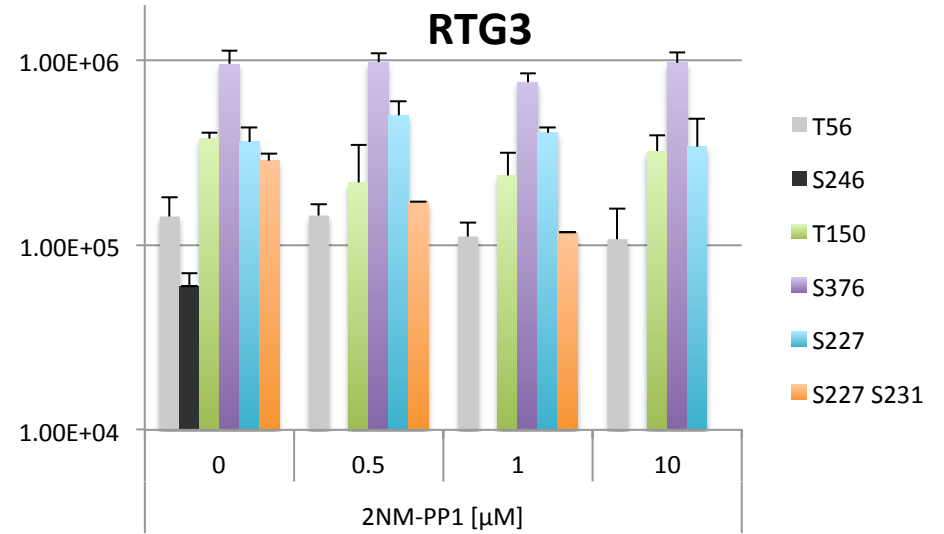
RSC2



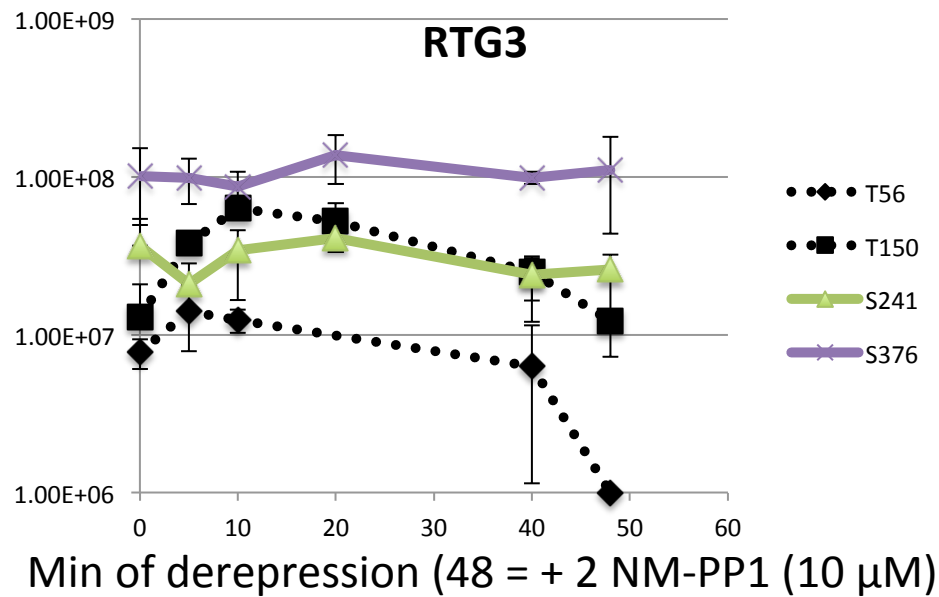
RTG3



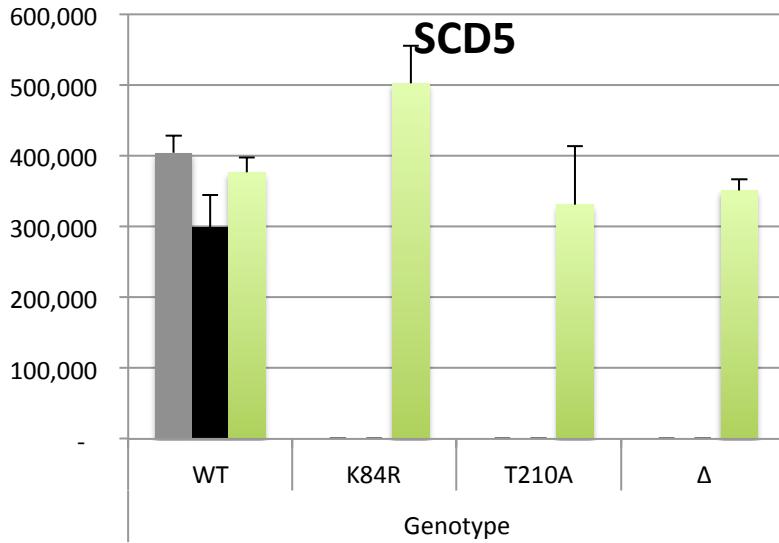
SNF1 genotype



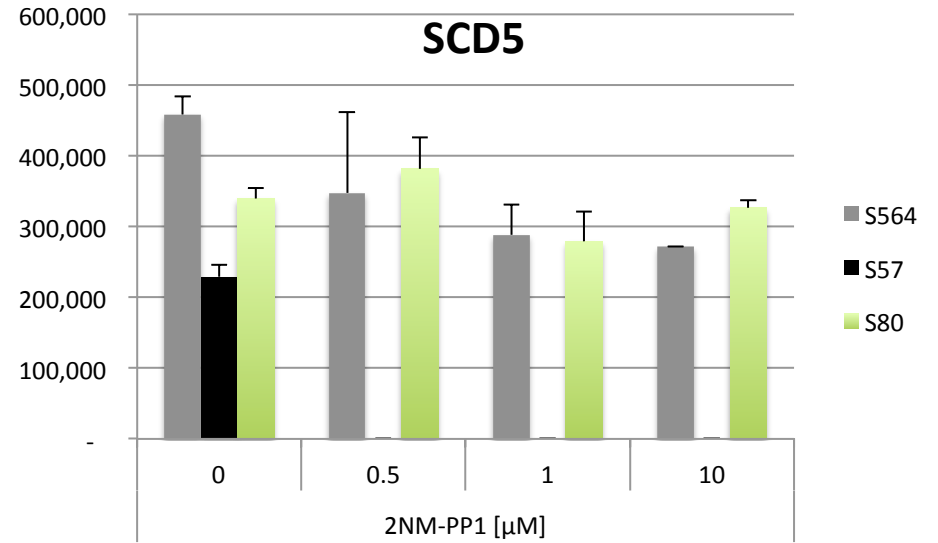
2 NM-PP1 (μ M)



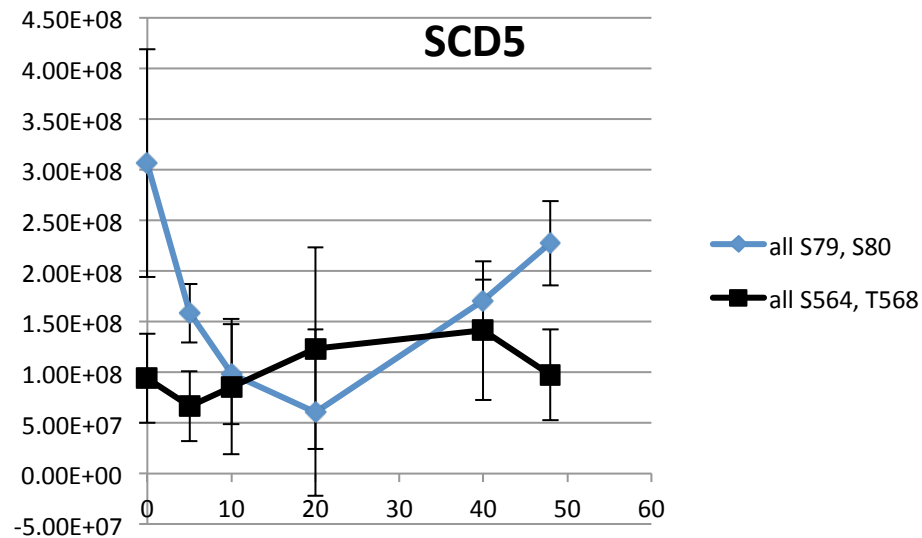
SCD5



SNF1 genotype



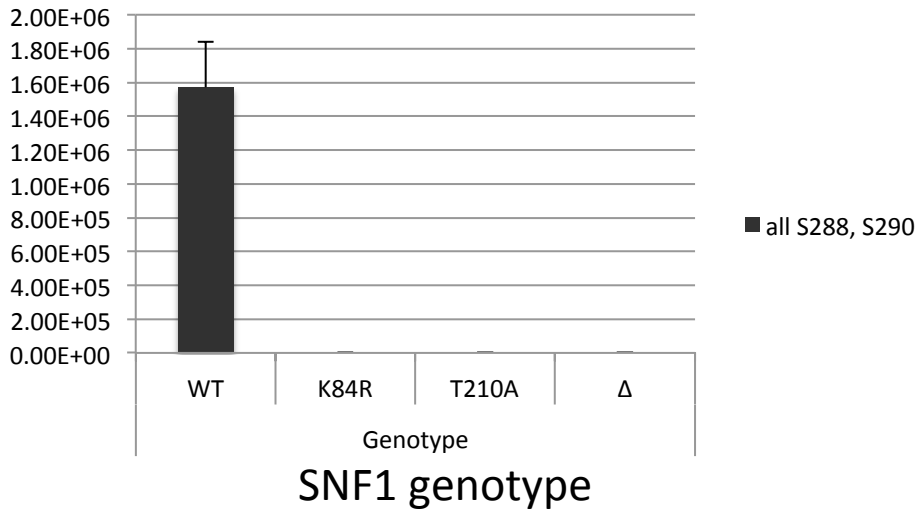
2 NM-PP1 (μM)



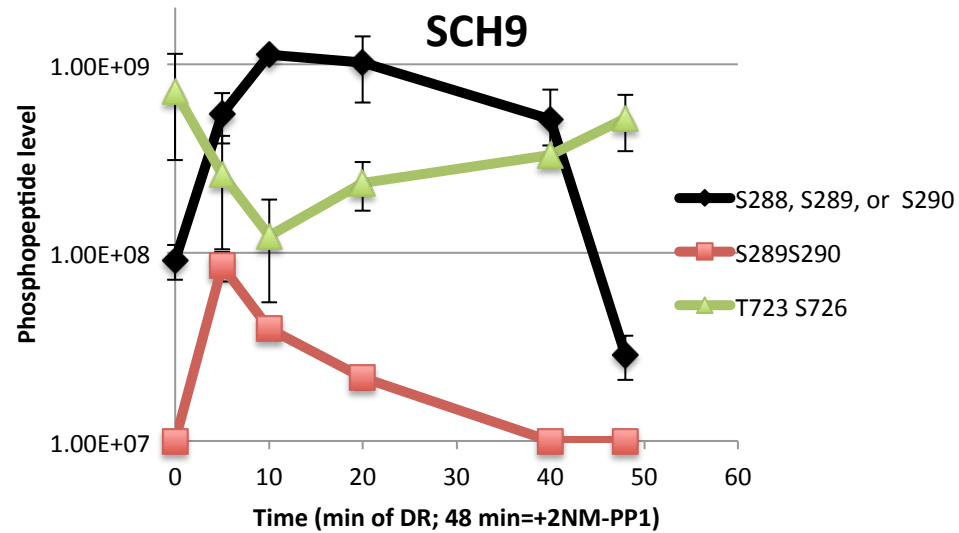
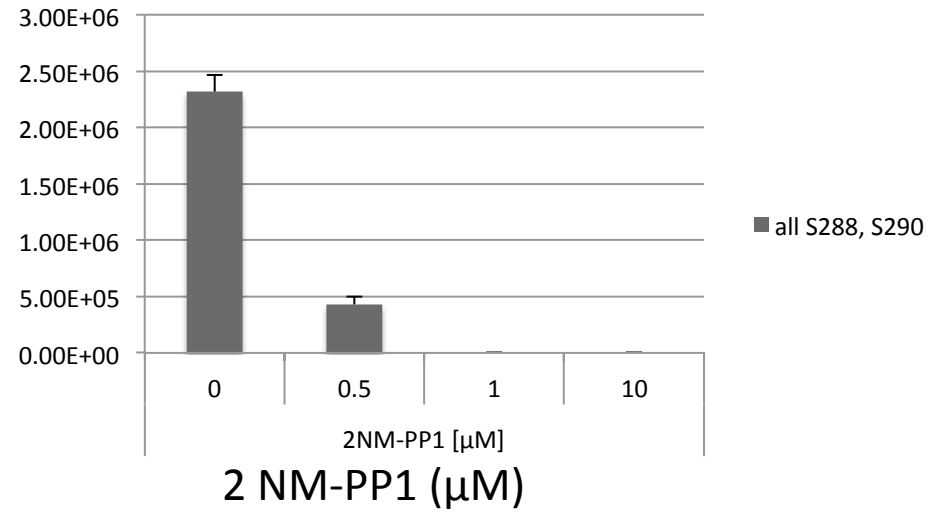
Min of derepression (48 = + 2 NM-PP1 (10 μM))

SCH9

SCH9

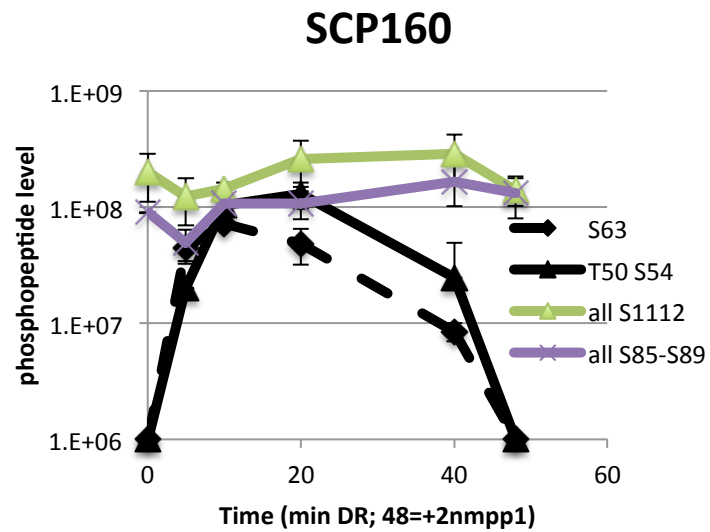
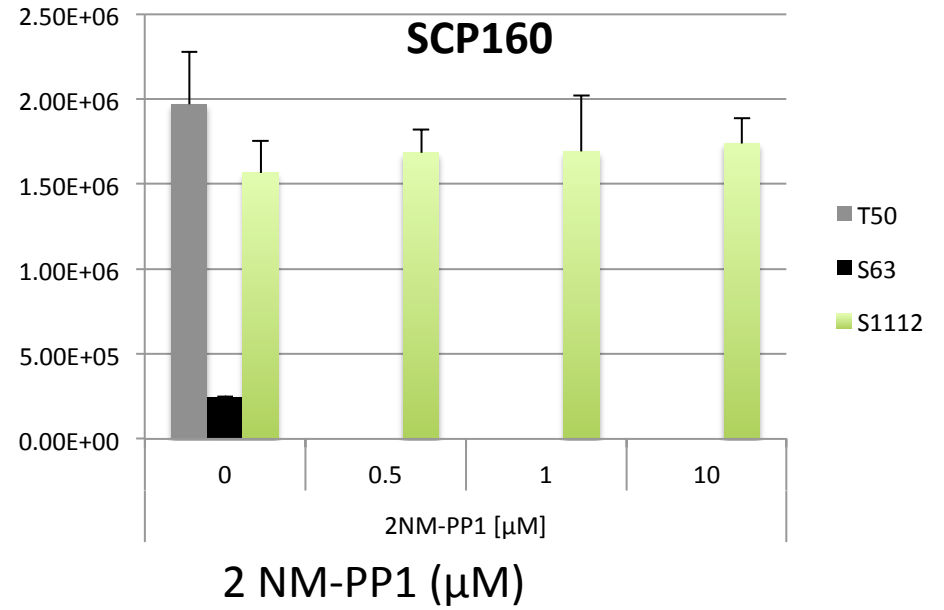
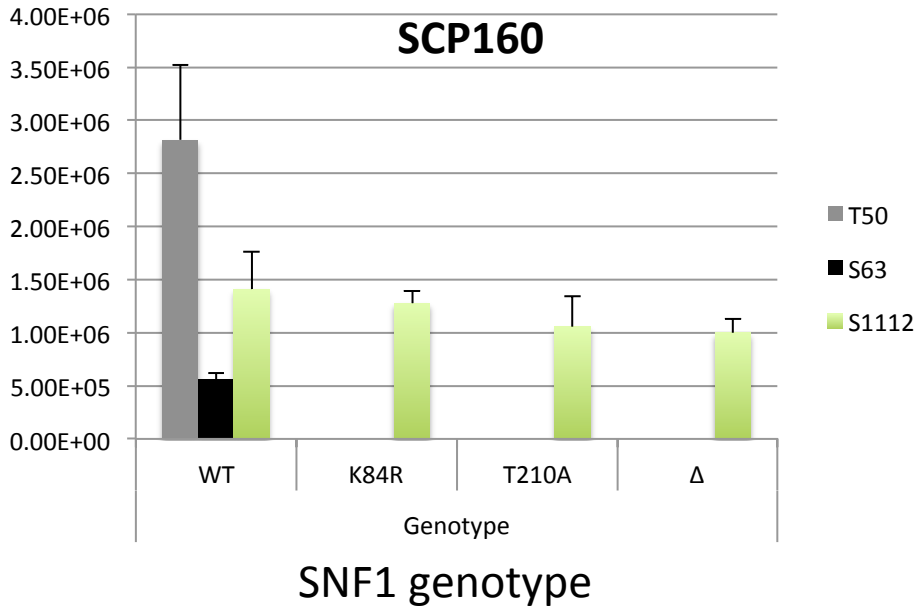


SCH9



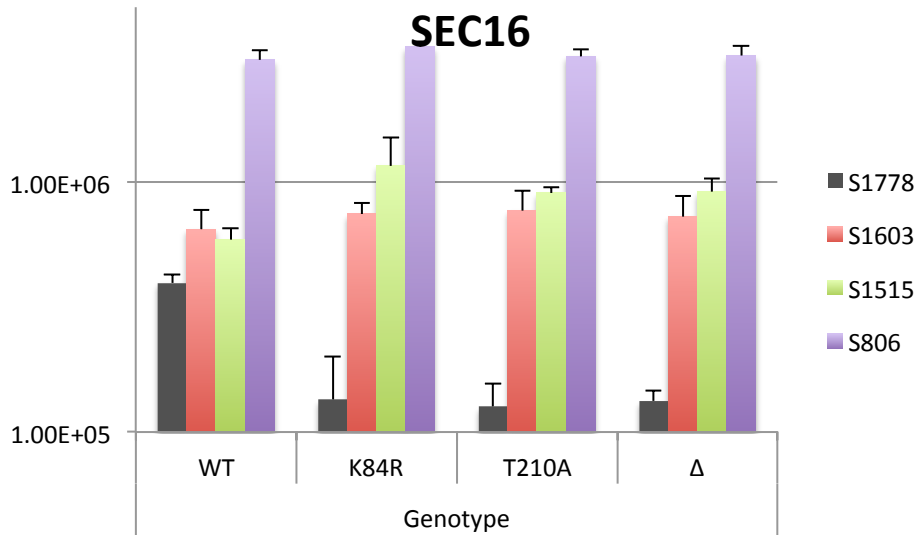
Min of derepression (48 = + 2 NM-PP1 (10 μM))

SCP160

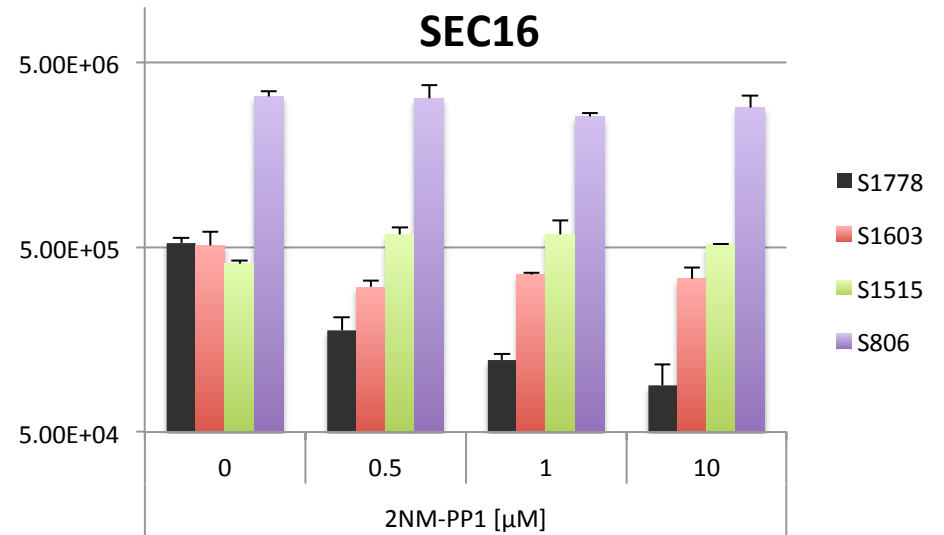


Min of derepression (48 = + 2 NM-PP1 (10 μM))

SEC16



SNF1 genotype

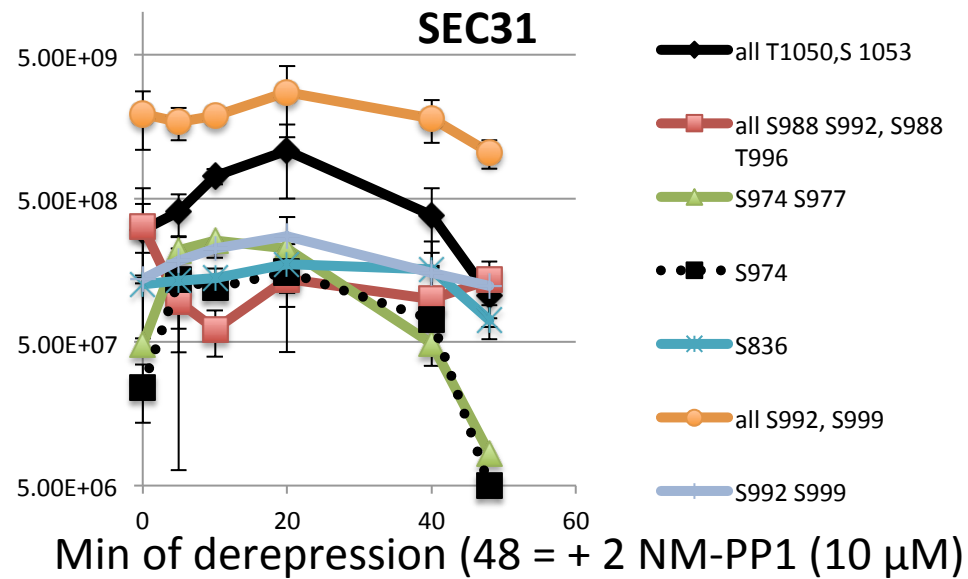
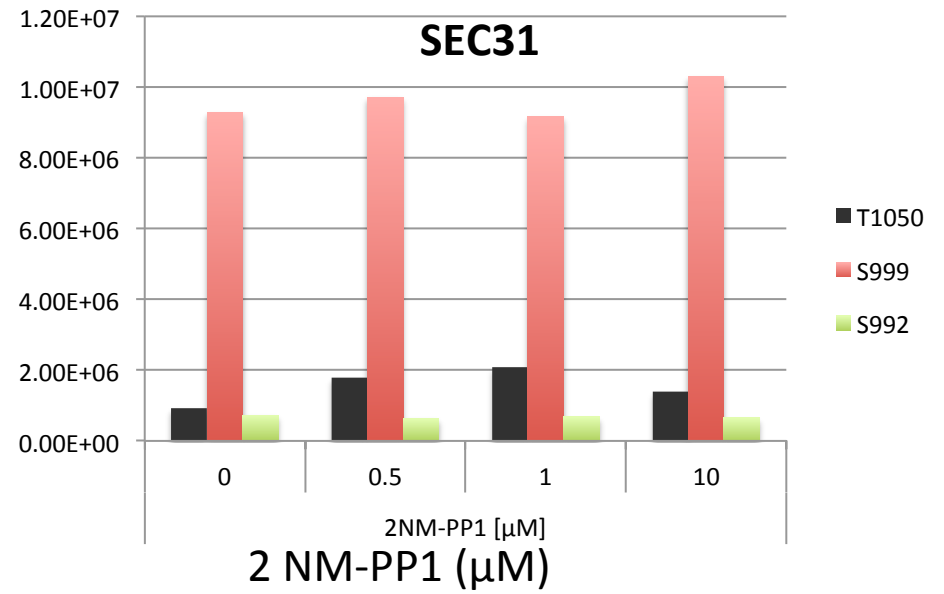
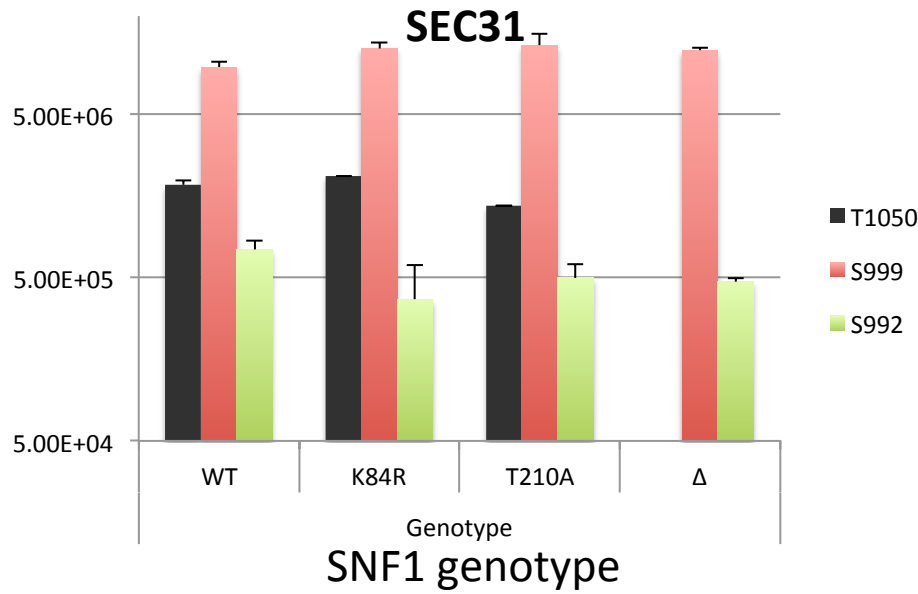


2 NM-PP1 (μ M)

No data for S1778

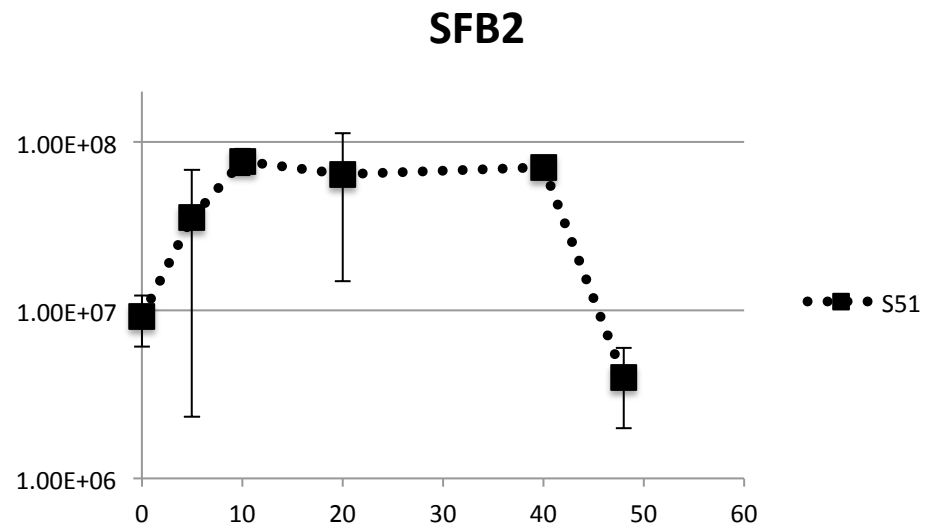
Min of derepression (48 = + 2 NM-PP1 (10 μ M))

SEC31



SFB2

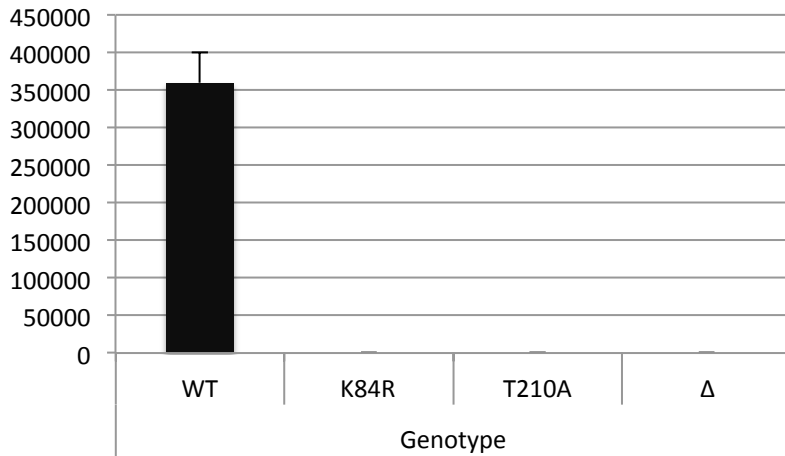
No data in experiments 1 and 2



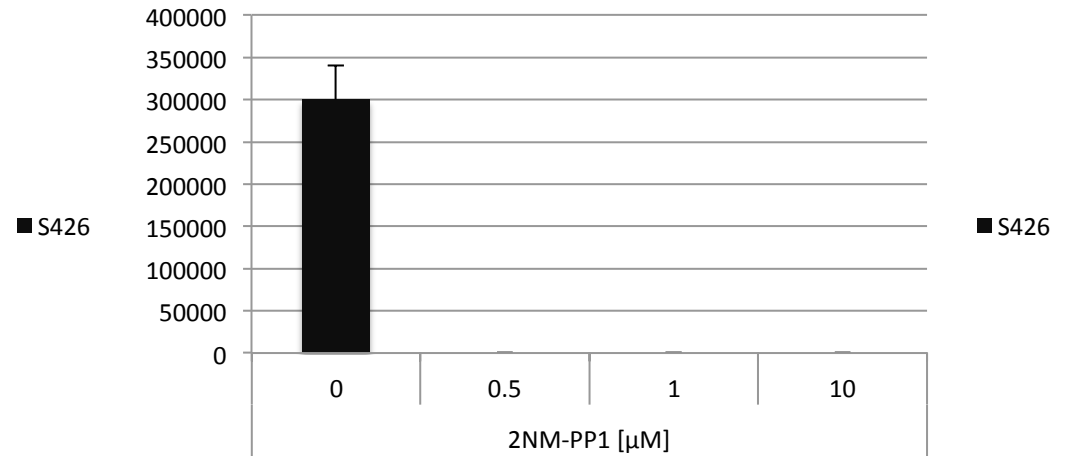
Min of derepression (48 = + 2 NM-PP1 (10 μ M))

SFL1

SFL1



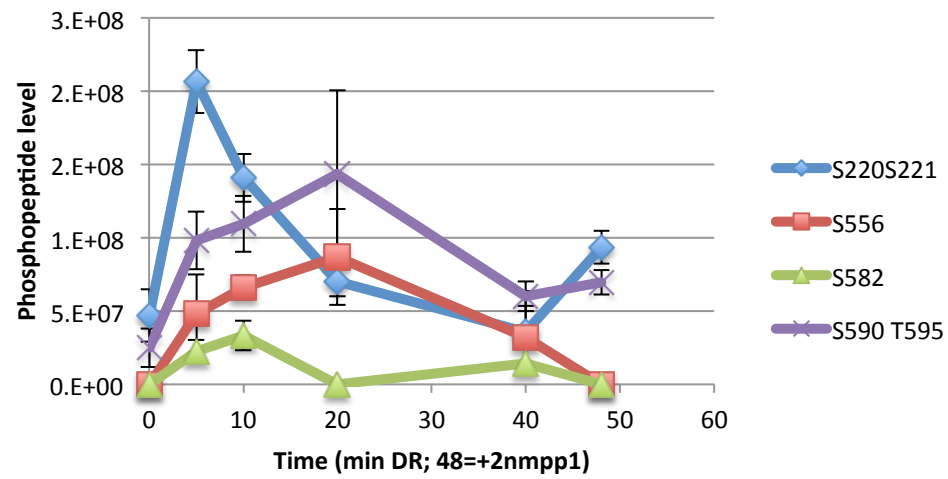
SFL1



SNF1 genotype

SFL1

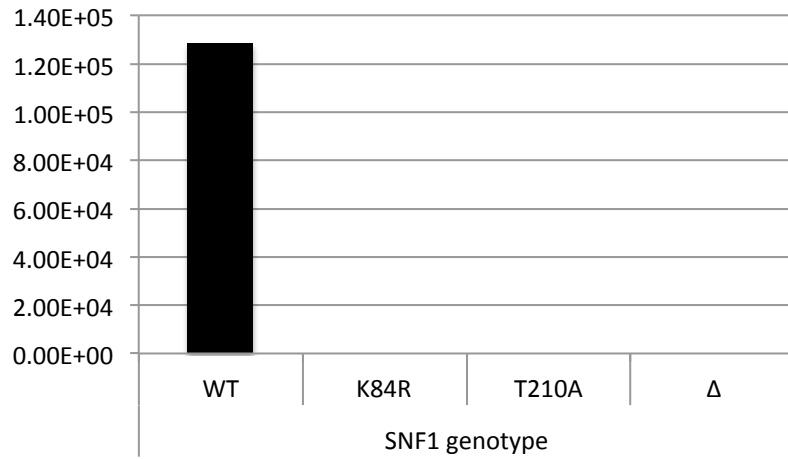
2 NM-PP1 (μM)



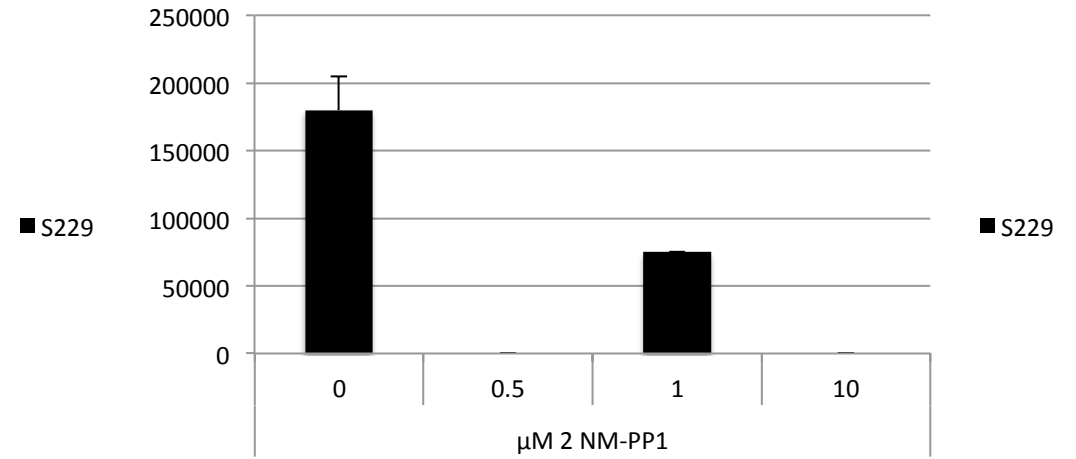
Min of derepression (48 = + 2 NM-PP1 (10 μM))

SGN1

SGN1

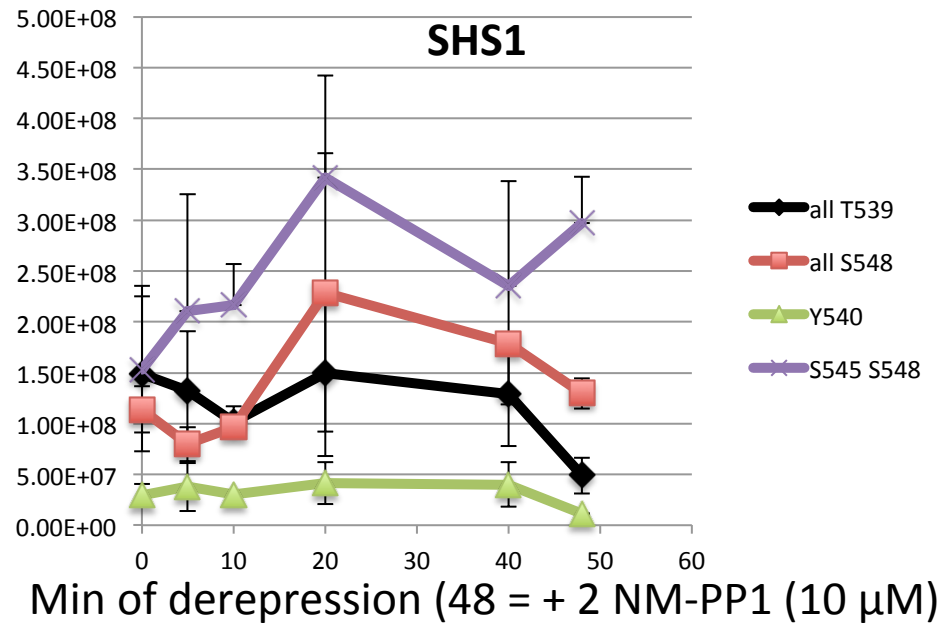
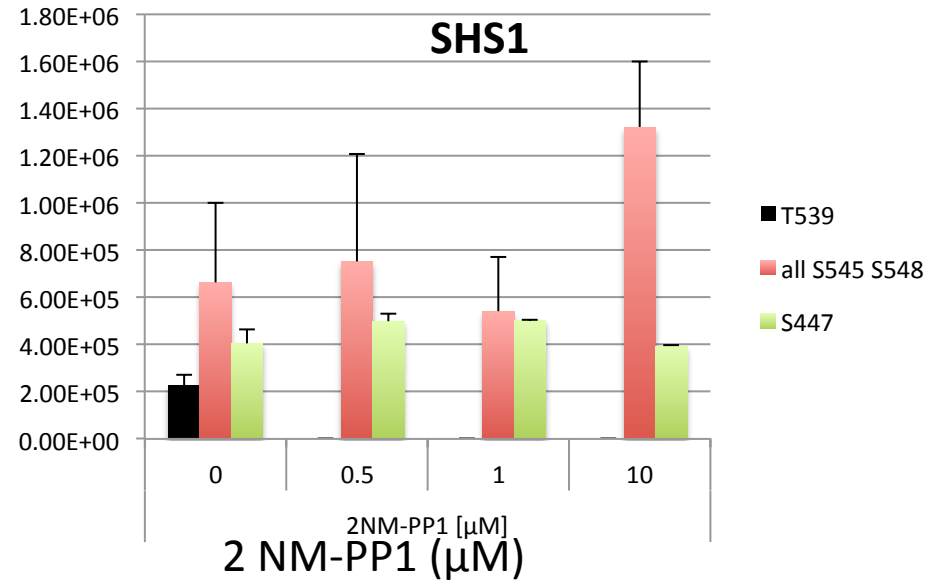
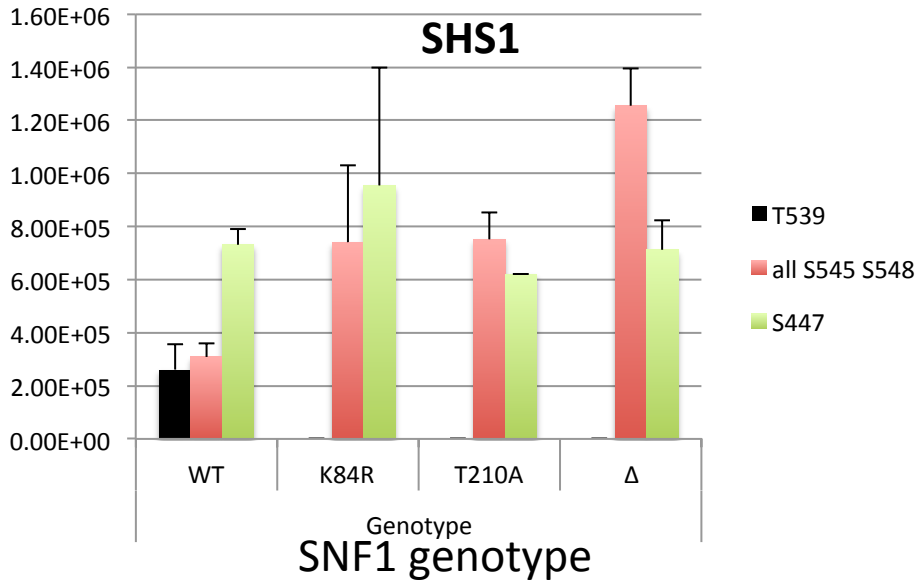


SGN1



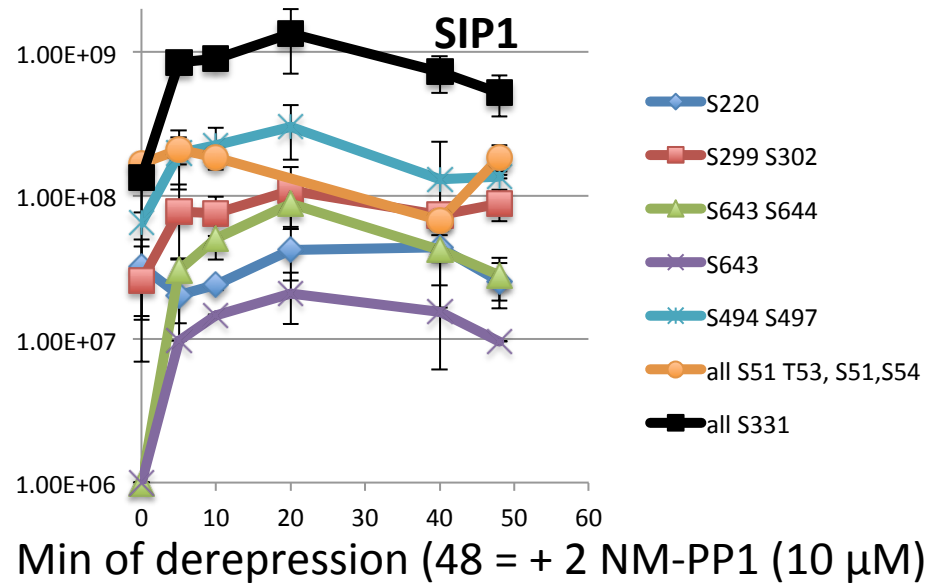
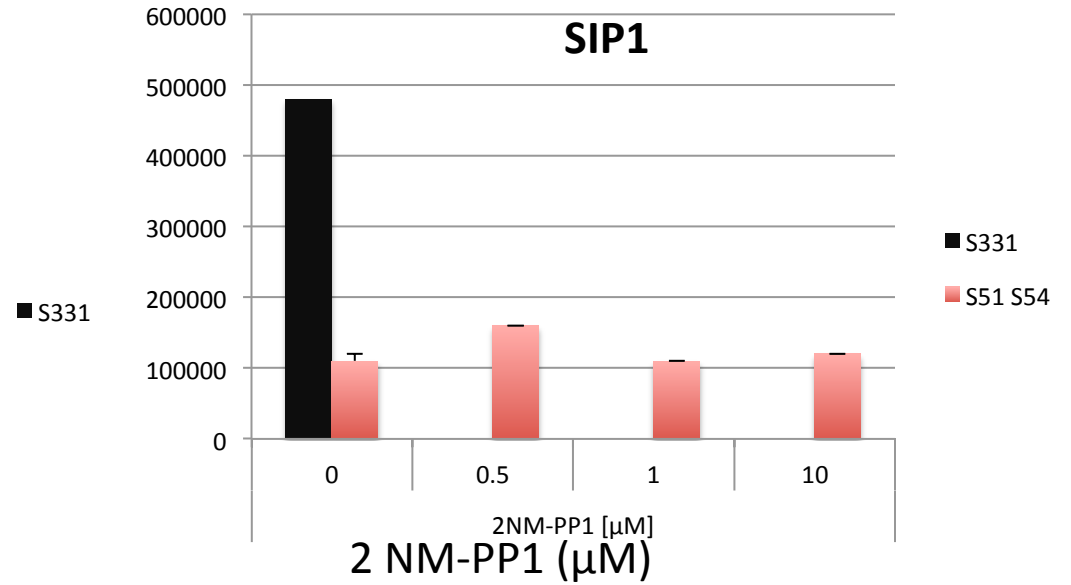
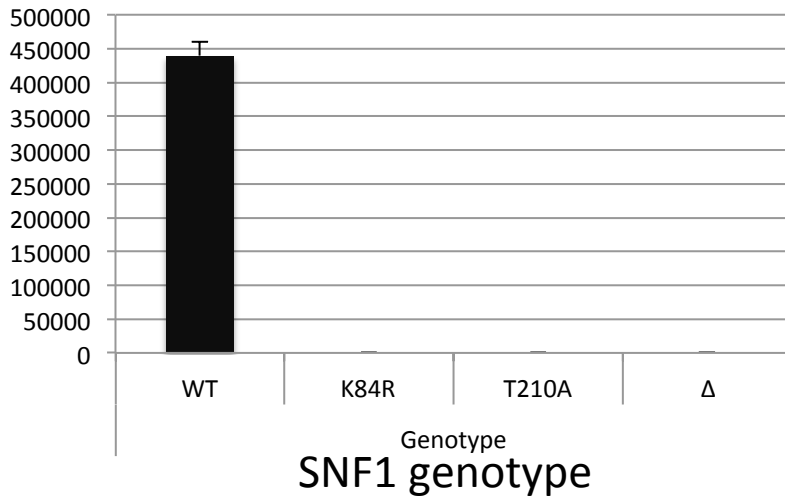
No data in time course experiment

SHS1



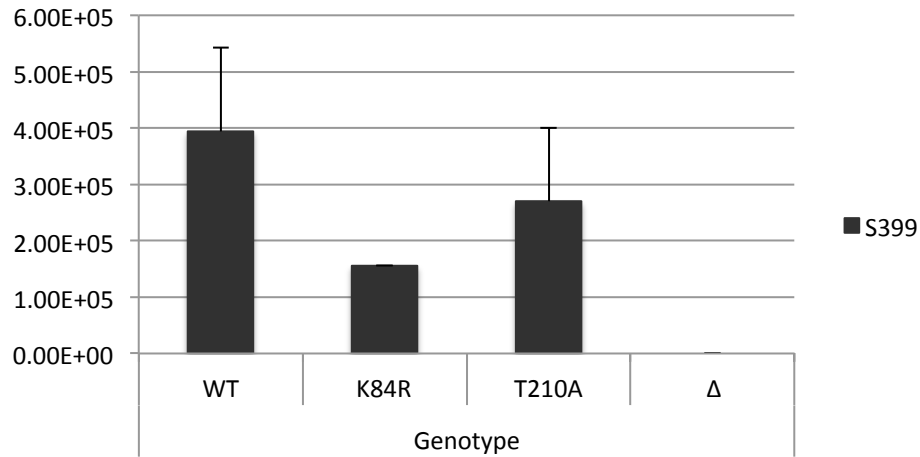
SIP1

SIP1

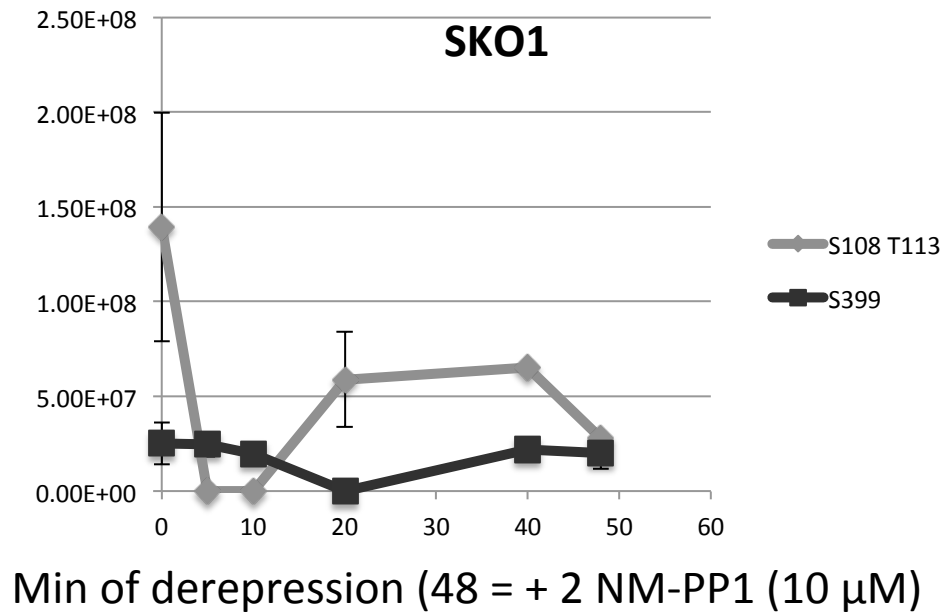
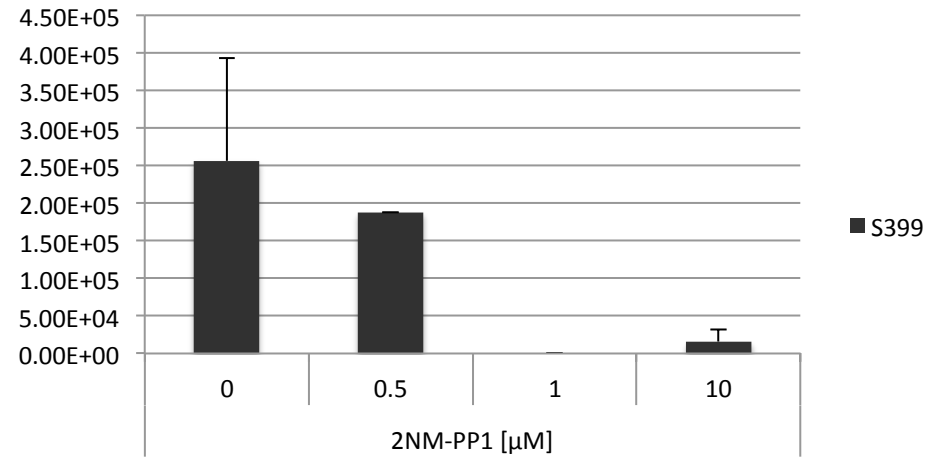


SKO1

SKO1

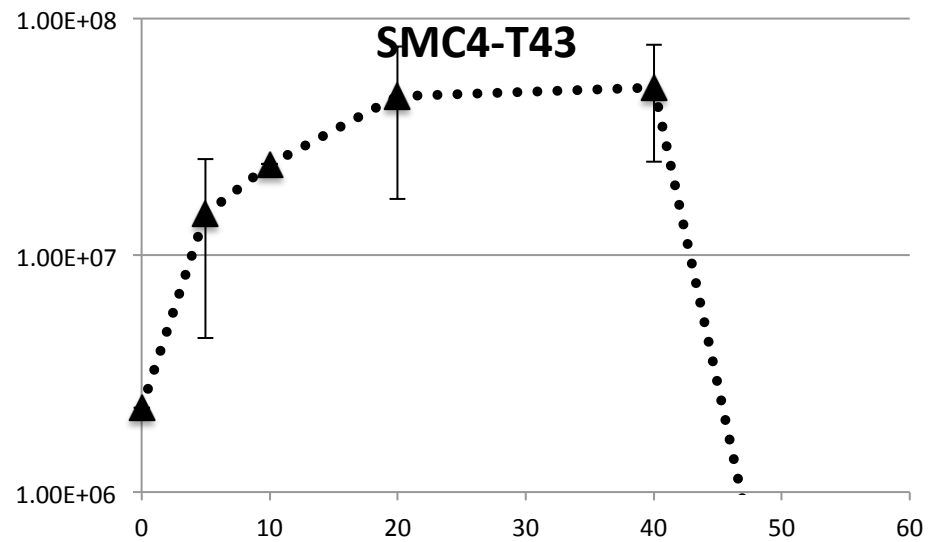


SKO1



SMC4

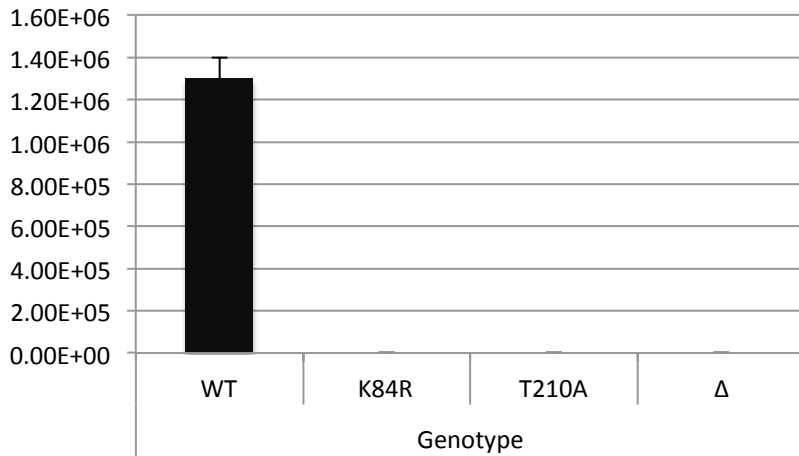
No data in experiments 1 and 2



Min of derepression (48 = + 2 NM-PP1 (10 μ M))

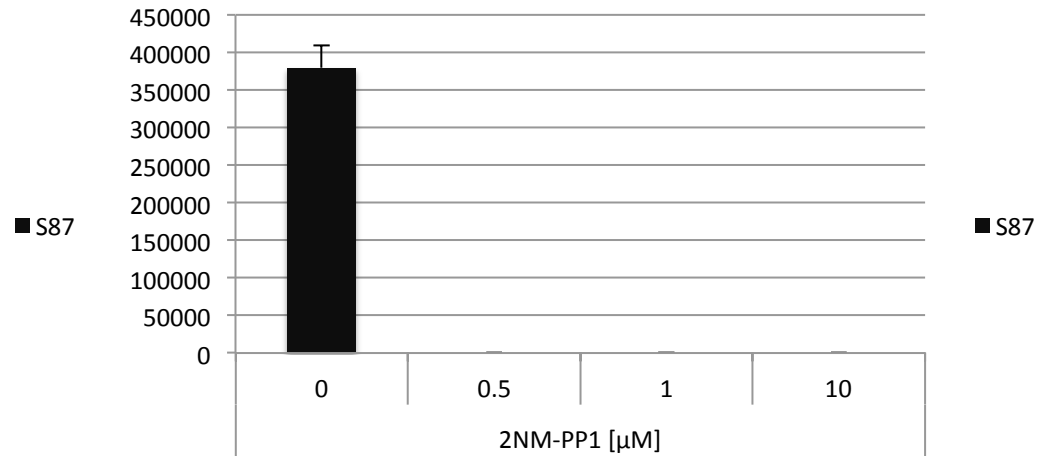
SML1

SML1



SNF1 genotype

SML1

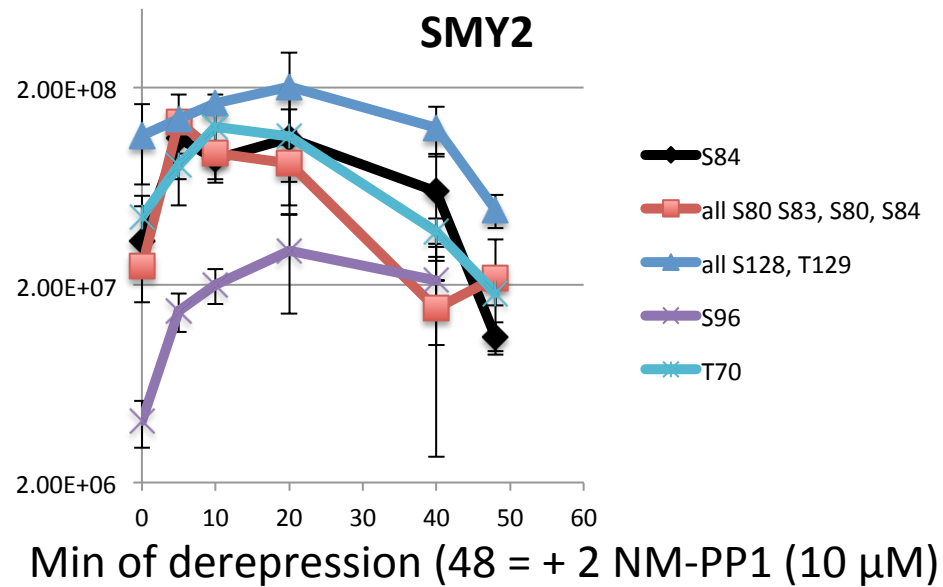
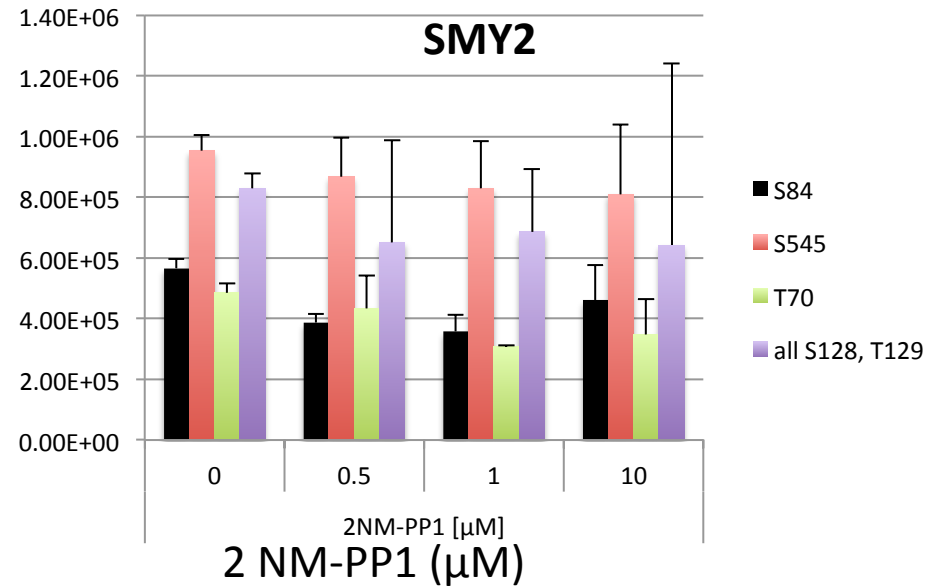
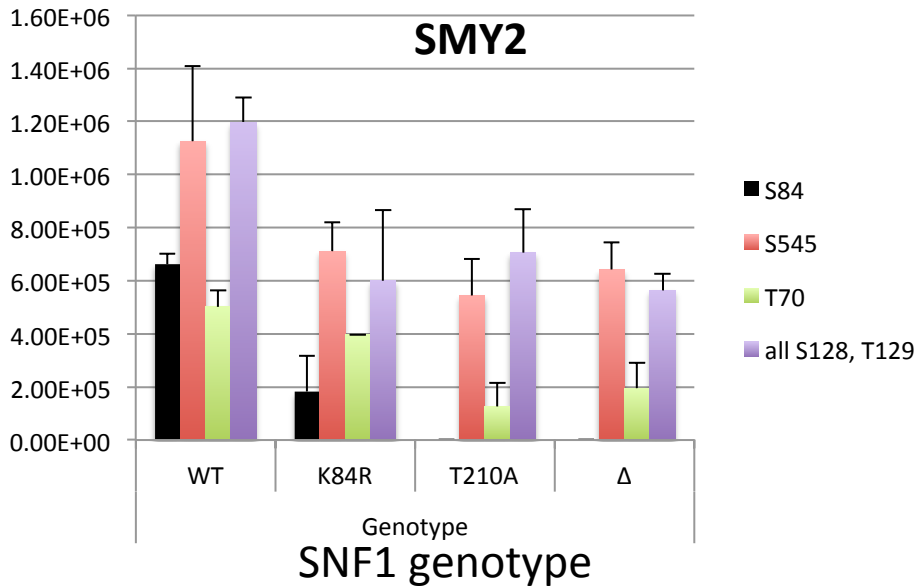


2 NM-PP1 (μM)

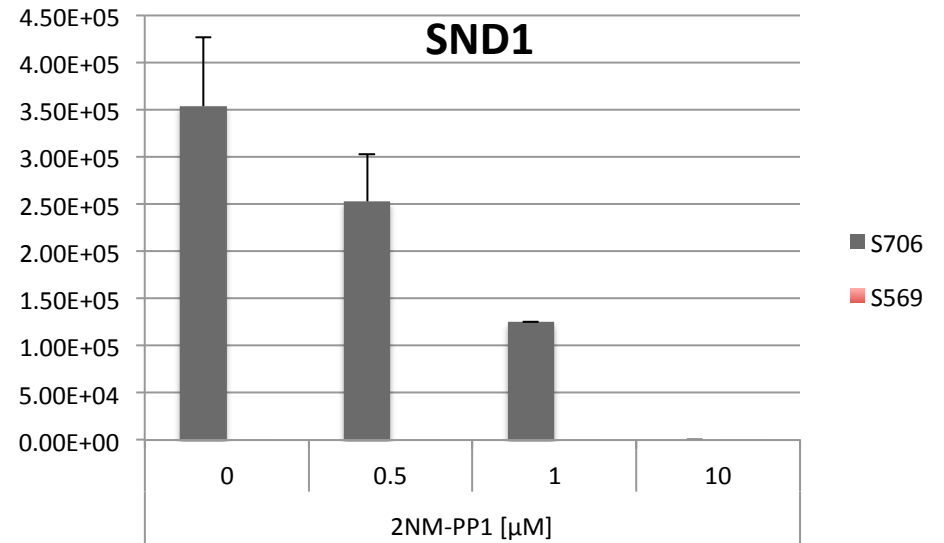
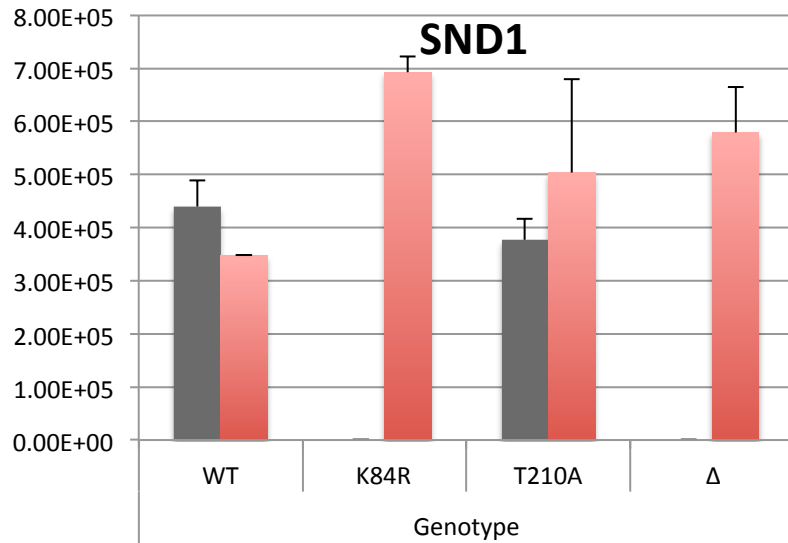
No data

Min of derepression (48 = + 2 NM-PP1 (10 μM))

SMY2

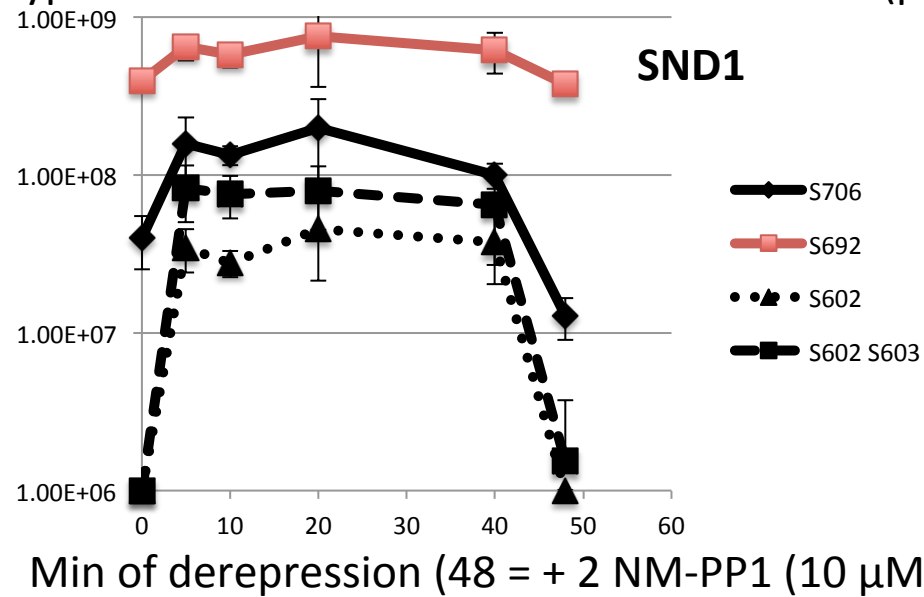


SND1

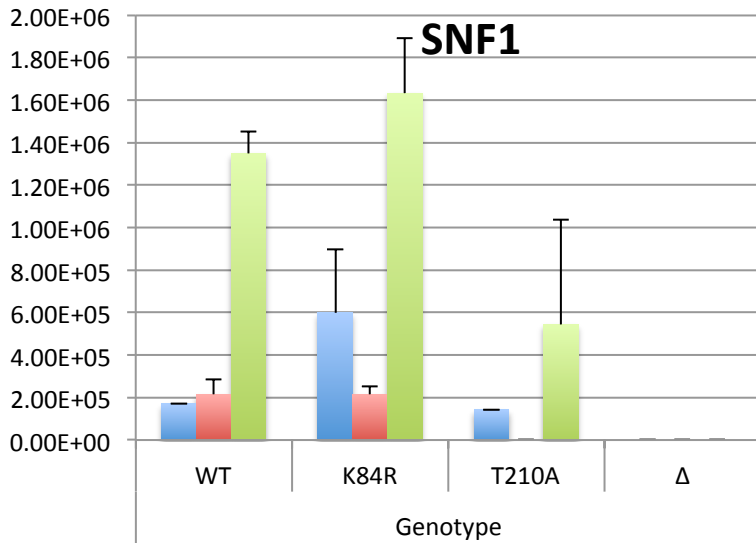


SNF1 genotype

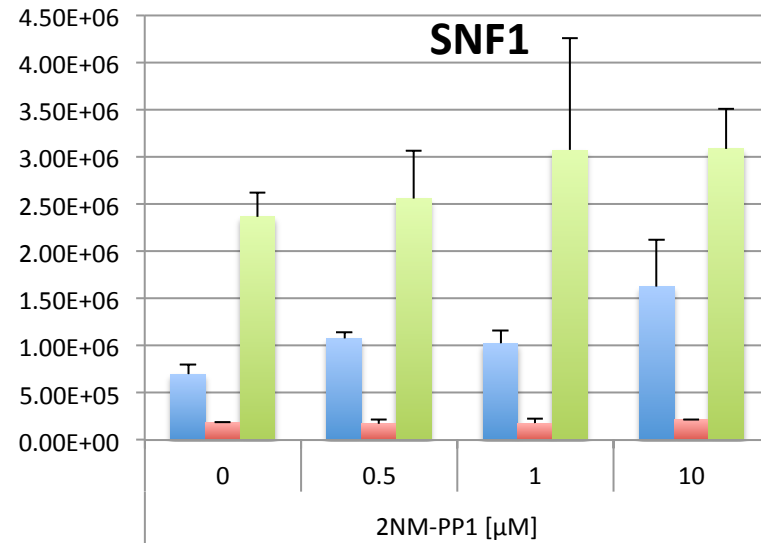
2 NM-PP1 (μM)



SNF1

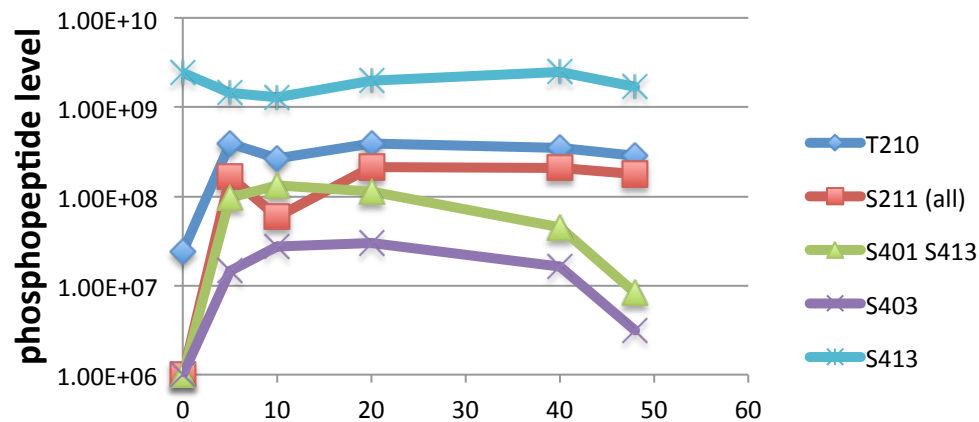


SNF1 genotype



2 NM-PP1 (μM)

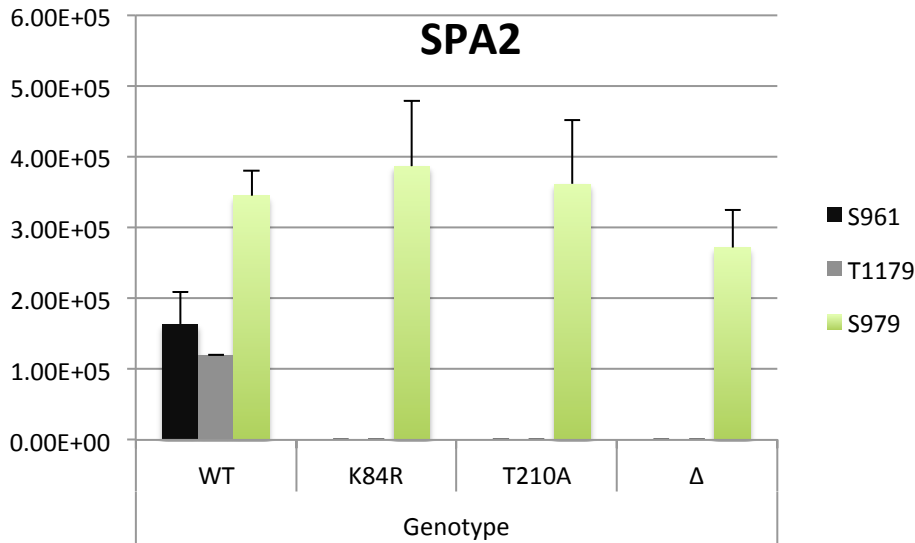
SNF1



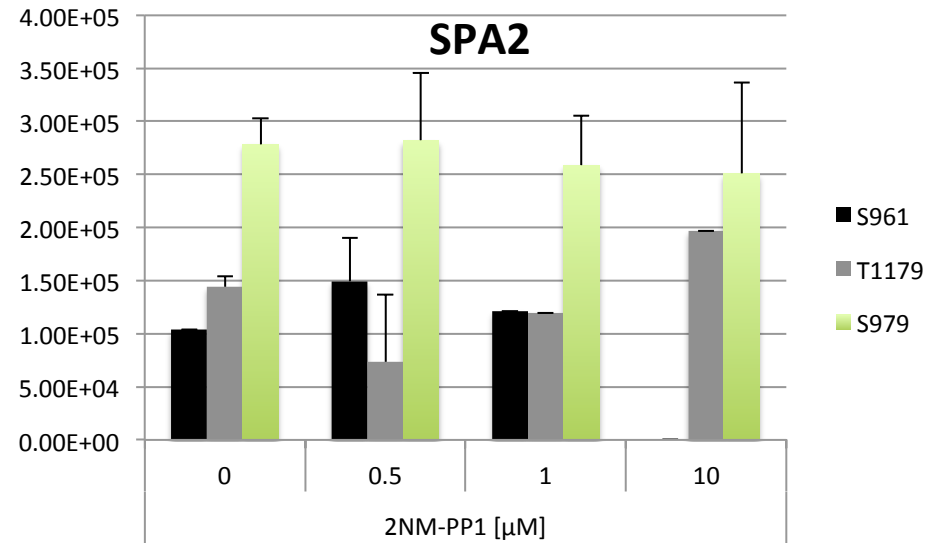
Min of DR (48 min=+2NM-PP1)

Min of derepression (48 = + 2 NM-PP1 (10 μM))

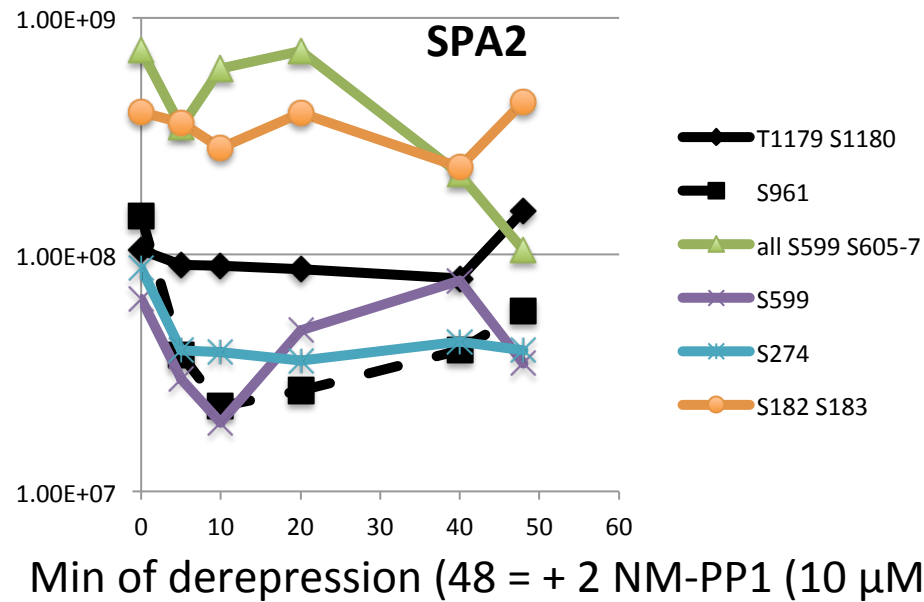
SPA2



SNF1 genotype



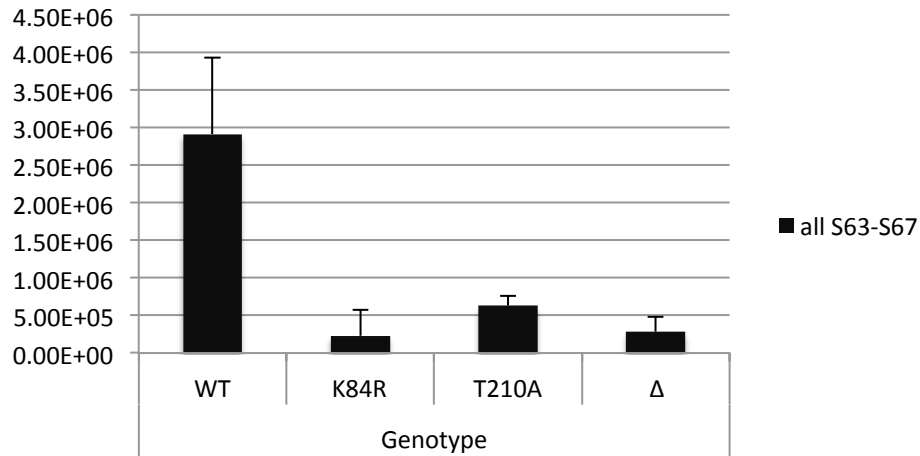
2 NM-PP1 (μM)



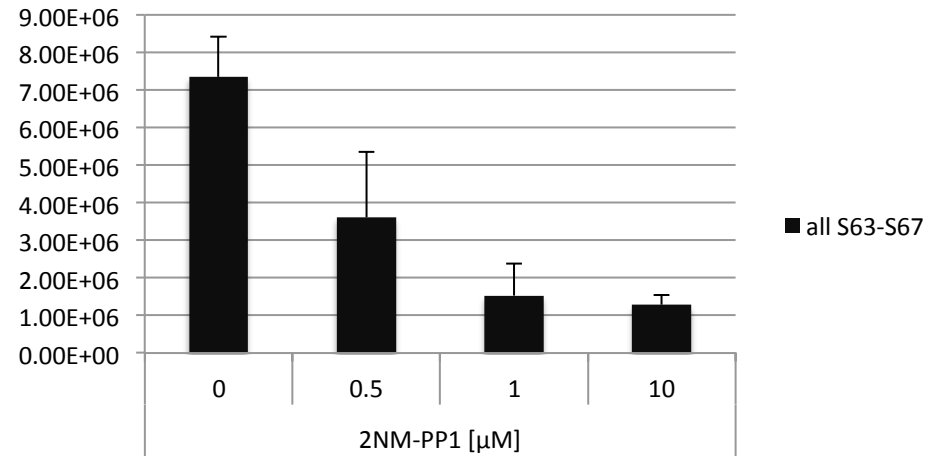
Min of derepression (48 = + 2 NM-PP1 (10 μM))

SPG4

SPG4

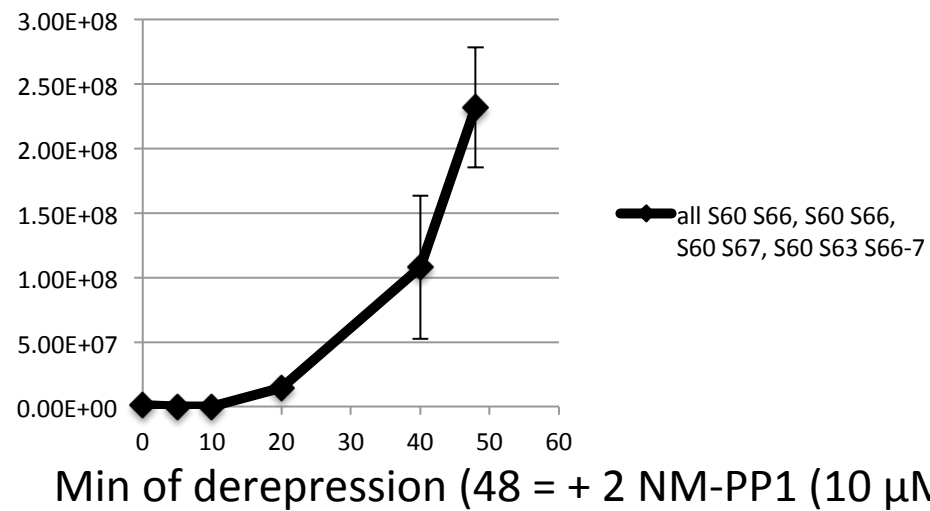


SPG4



SNF1 genotype

SPG4

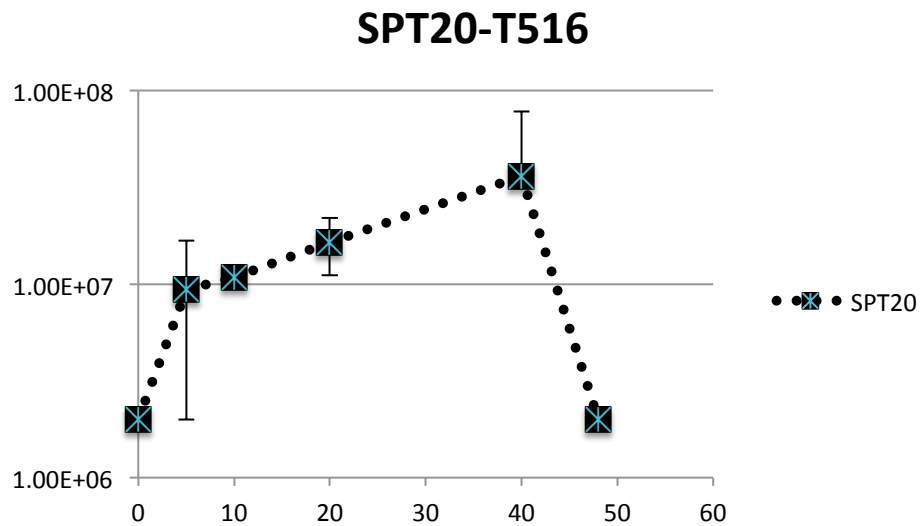


2 NM-PP1 (μM)

Min of derepression (48 = + 2 NM-PP1 (10 μM))

SPT20

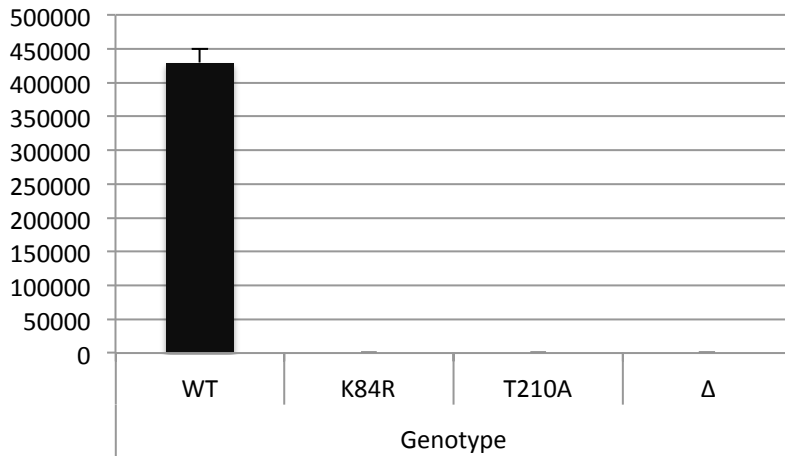
No data in experiments 1 and 2



Min of derepression (48 = + 2 NM-PP1 (10 μ M))

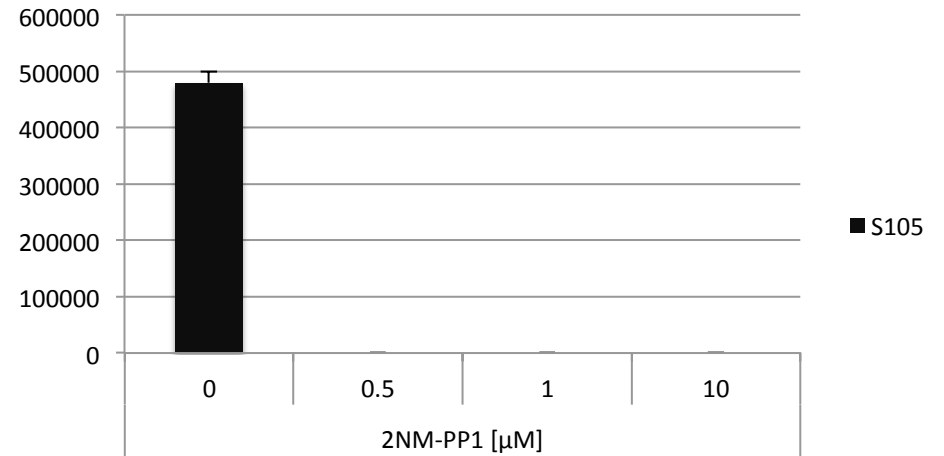
SQS1

SQS1

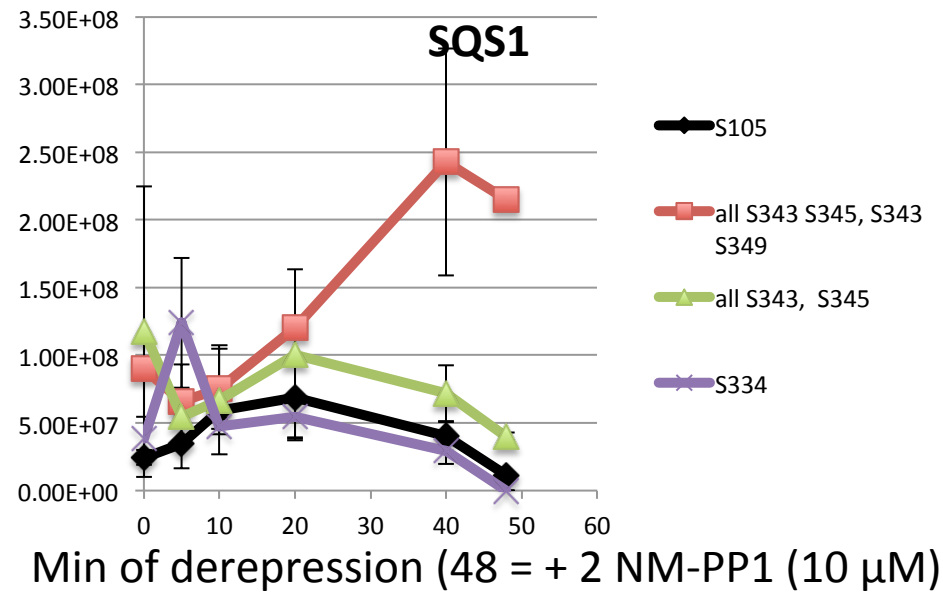


SNF1 genotype

SQS1

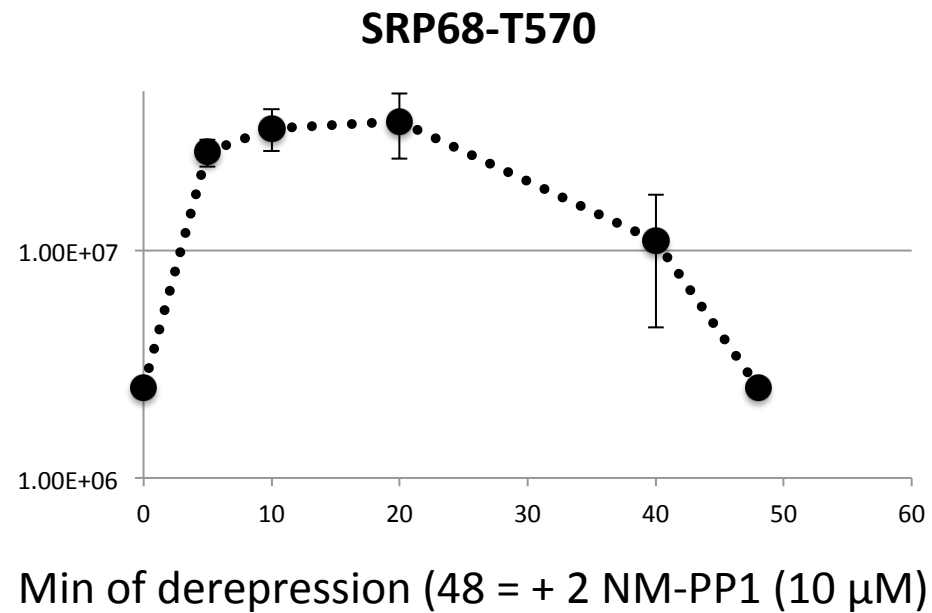


2 NM-PP1 (μM)

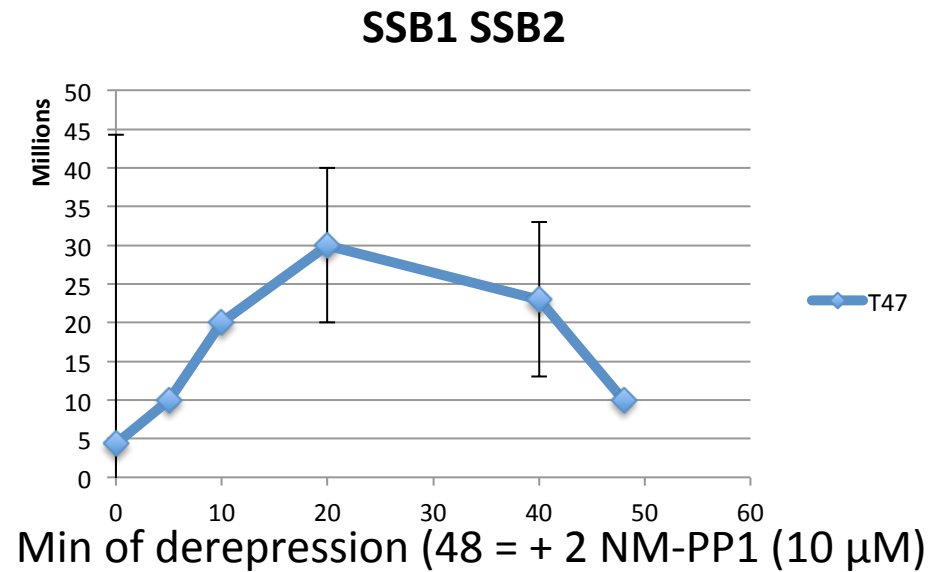
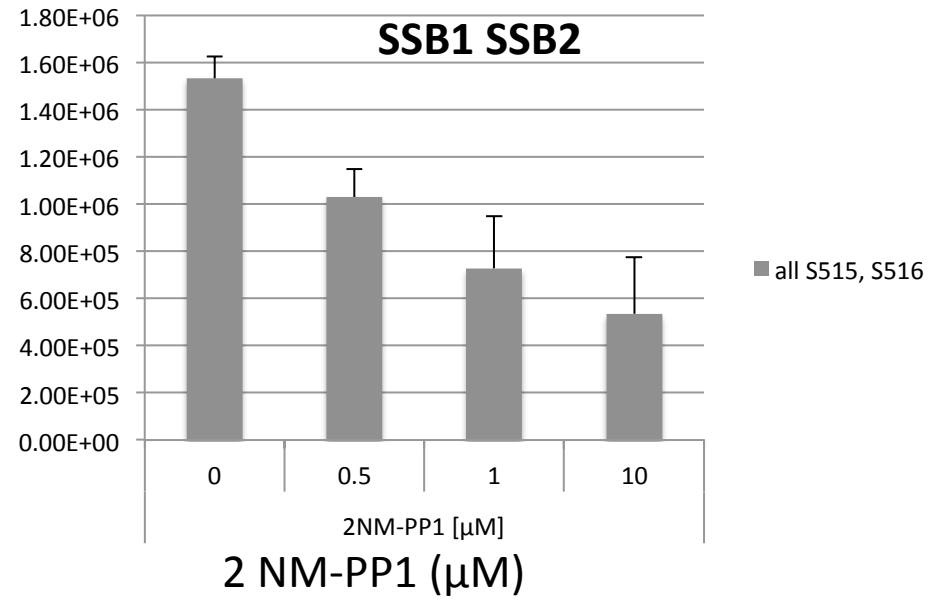
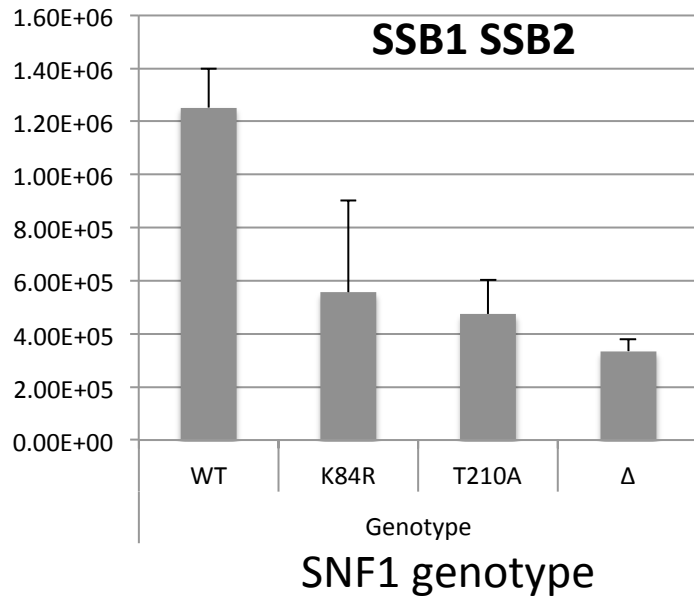


SRP68

No data in experiments 1 and 2

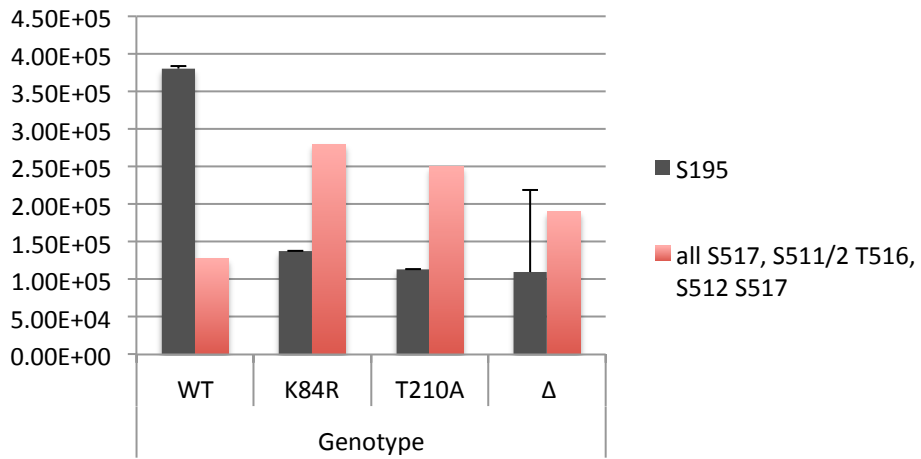


SSB1 SSB2

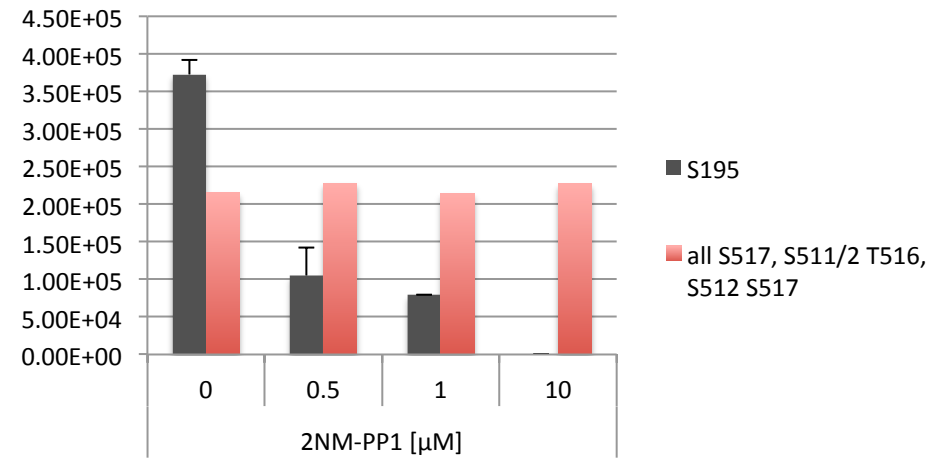


STE20

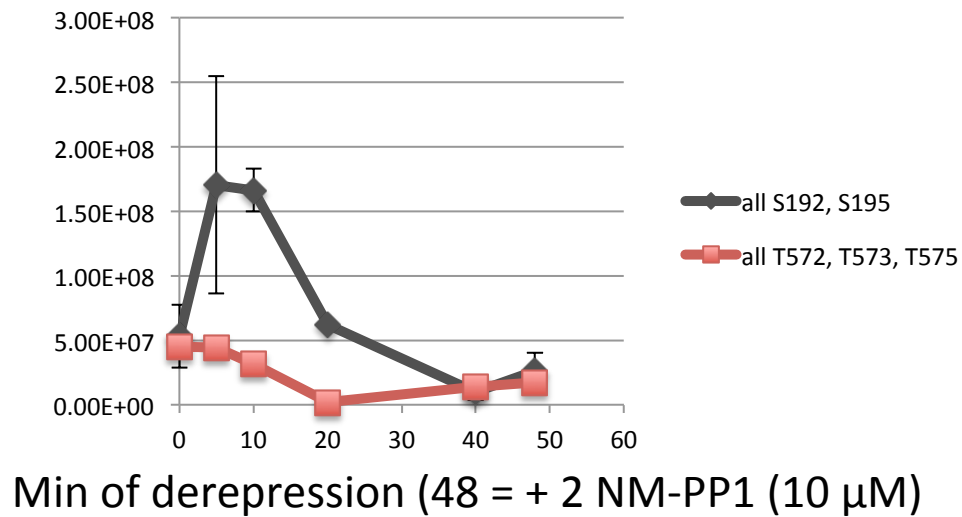
STE20



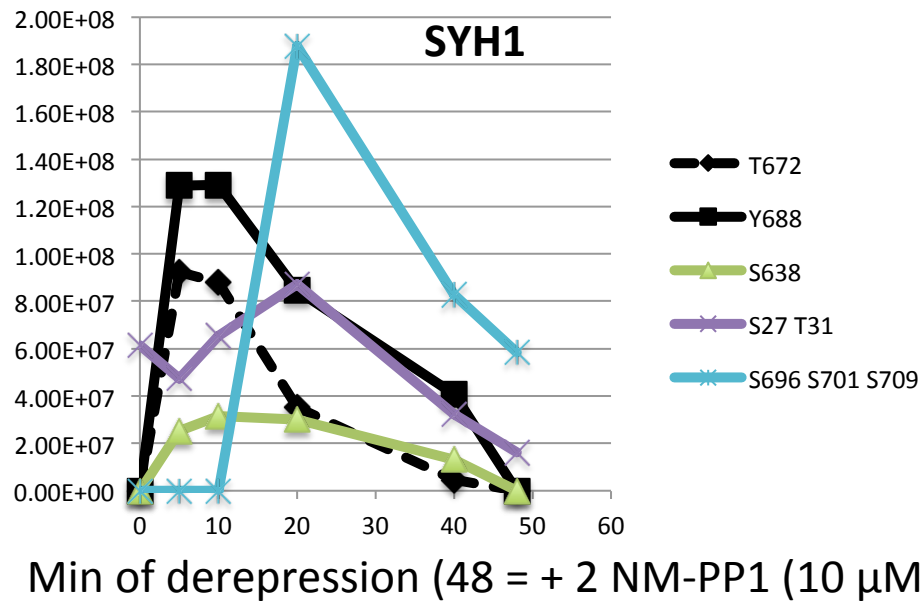
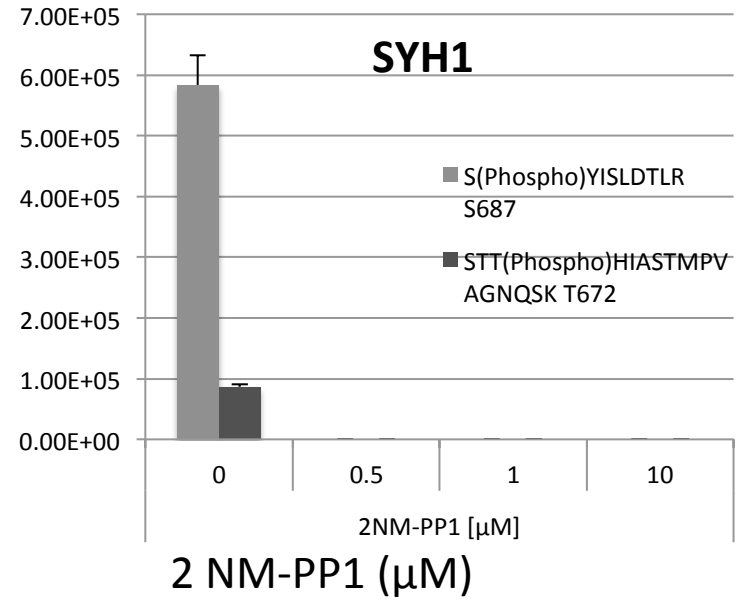
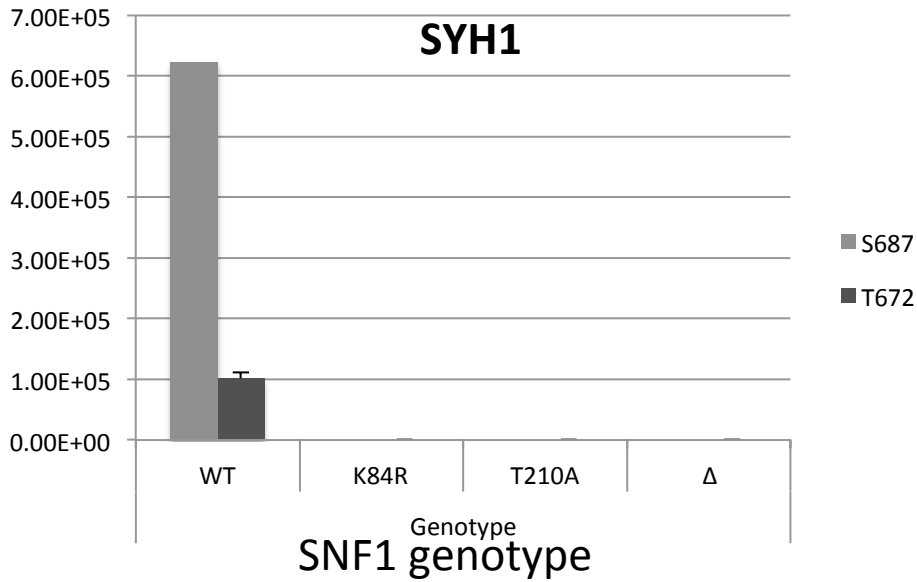
STE20



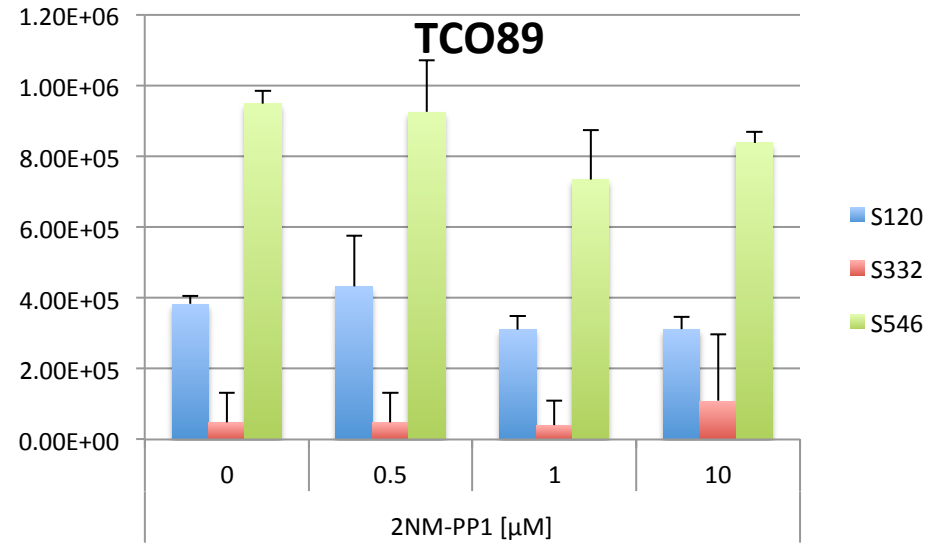
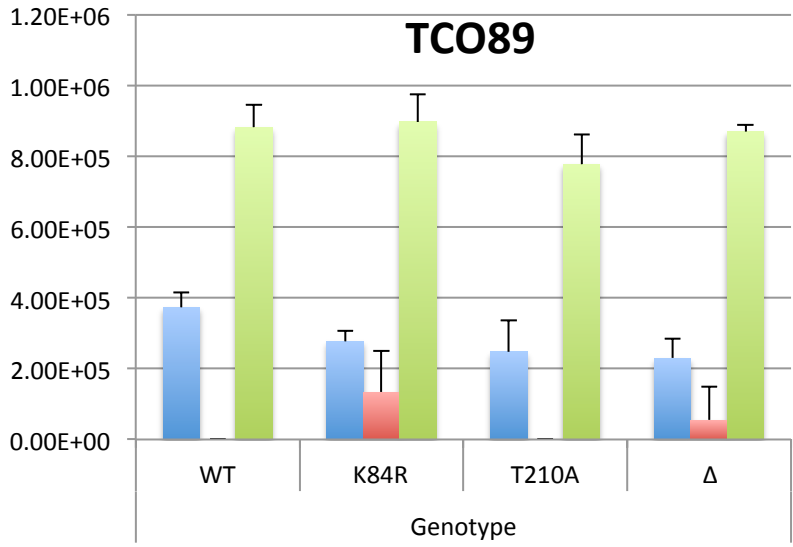
STE20



SYH1

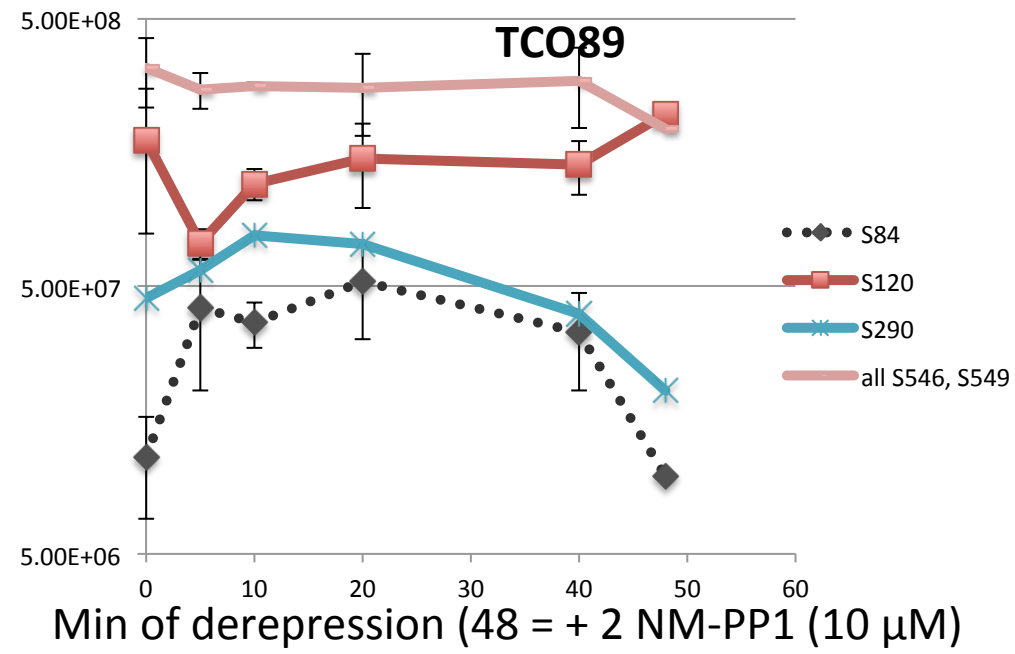


TCO89

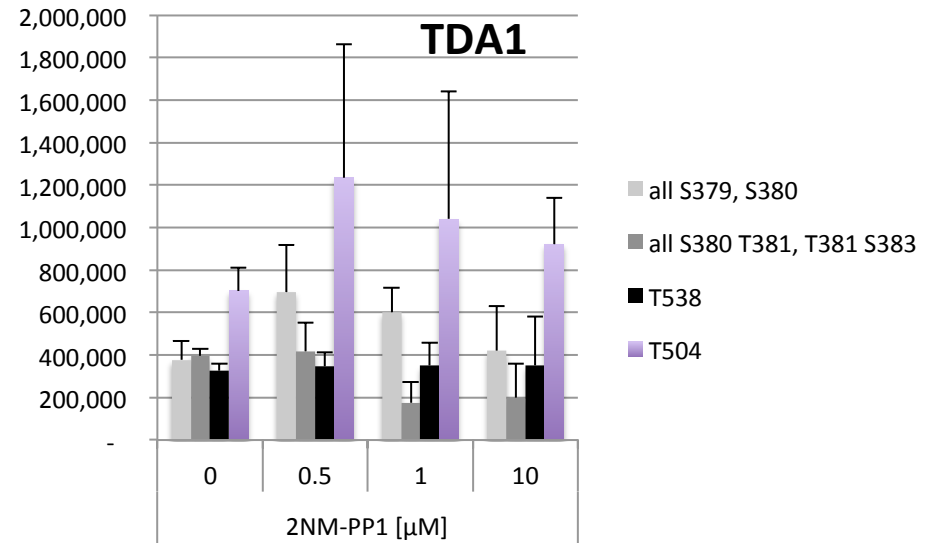
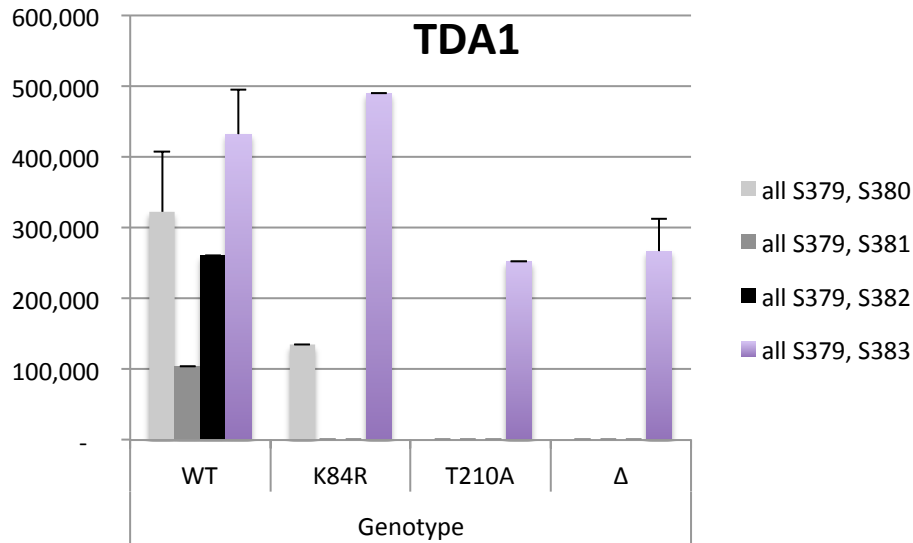


SNF1 genotype

2 NM-PP1 (μM)

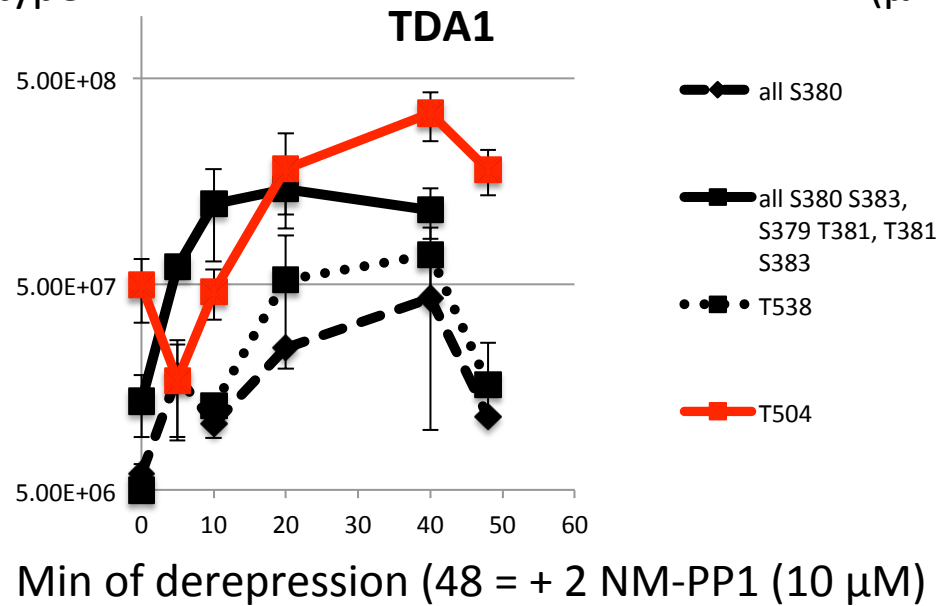


TDA1

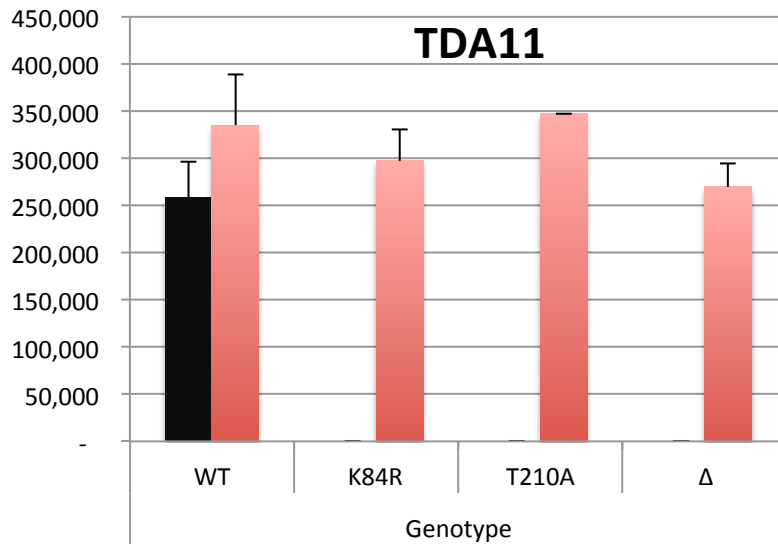


SNF1 genotype

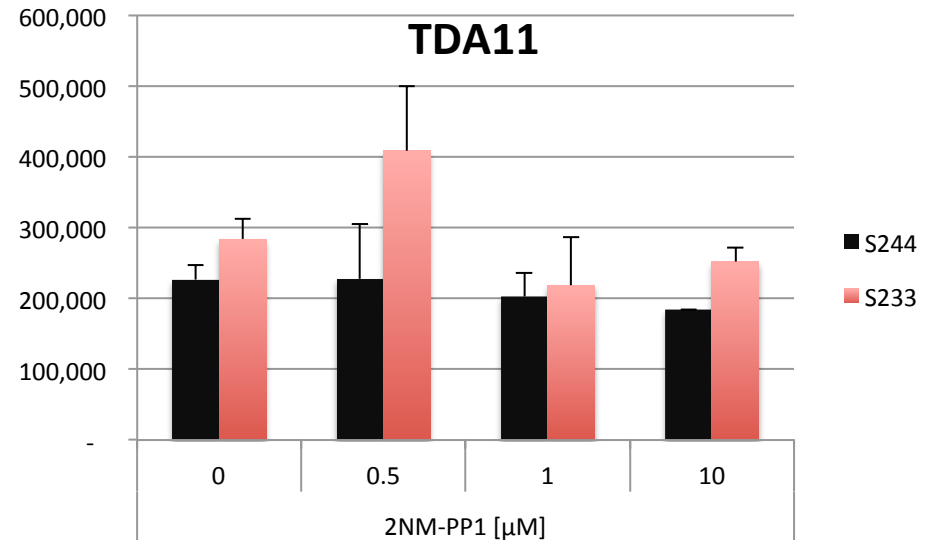
2 NM-PP1 (μM)



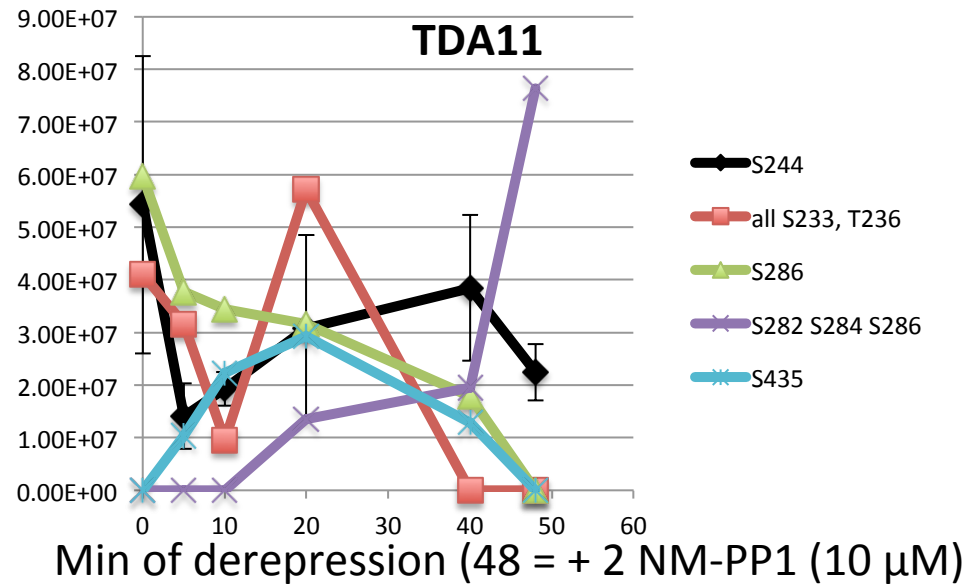
TDA1



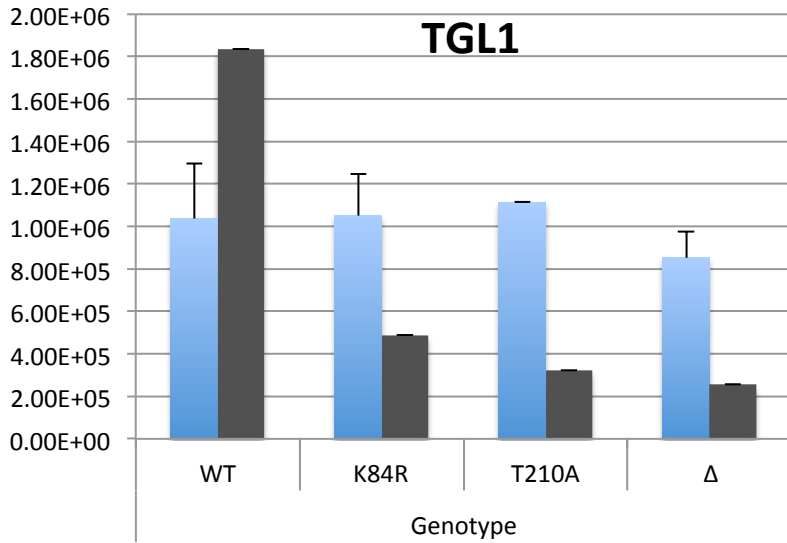
SNF1 genotype



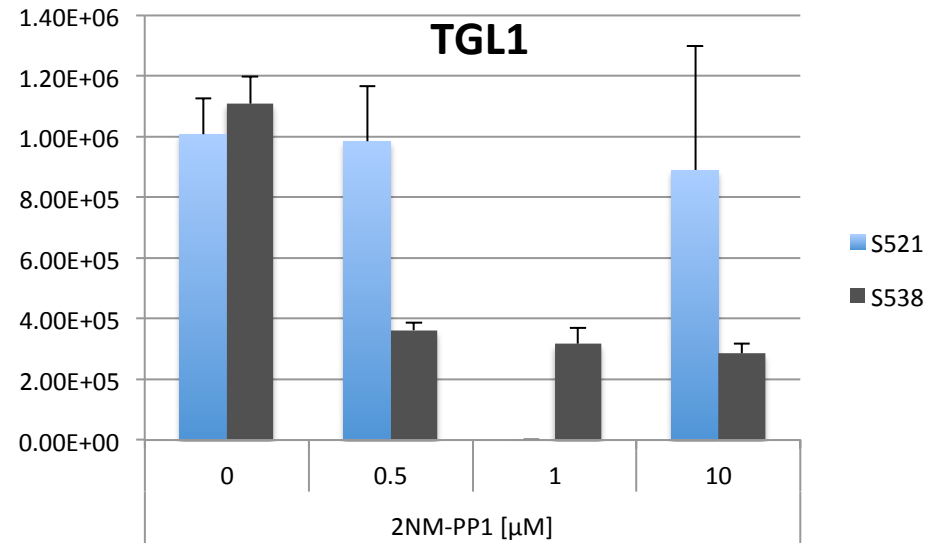
2 NM-PP1 (μM)



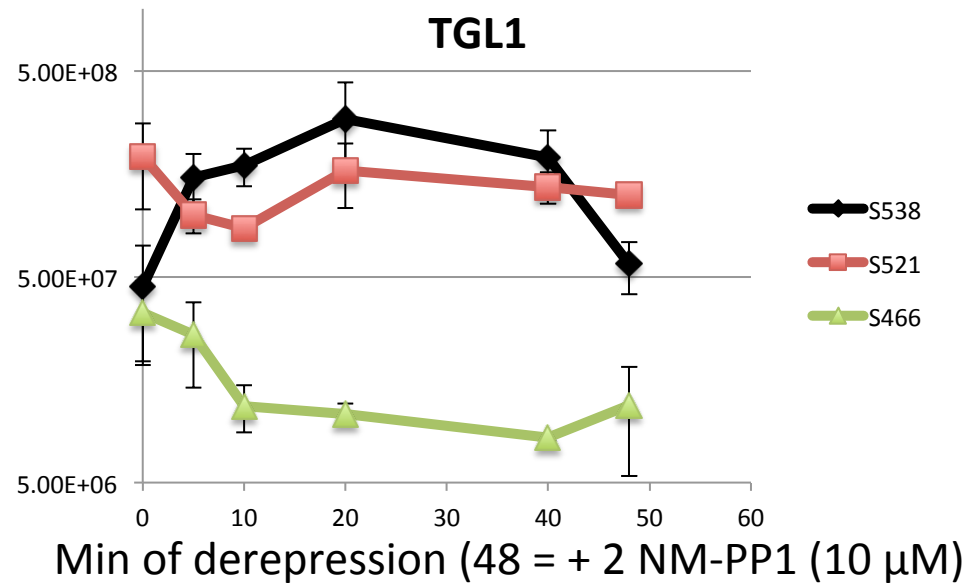
TGL1



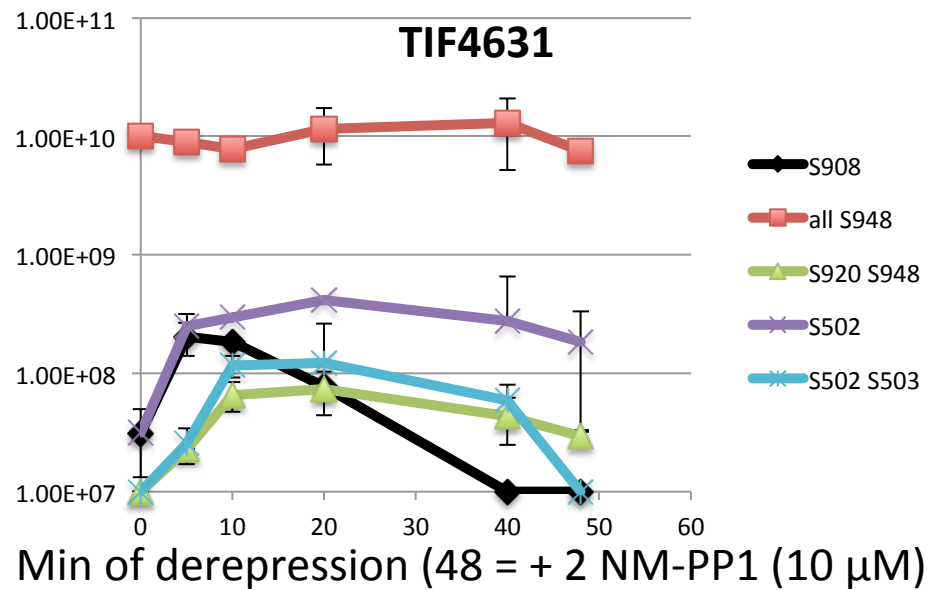
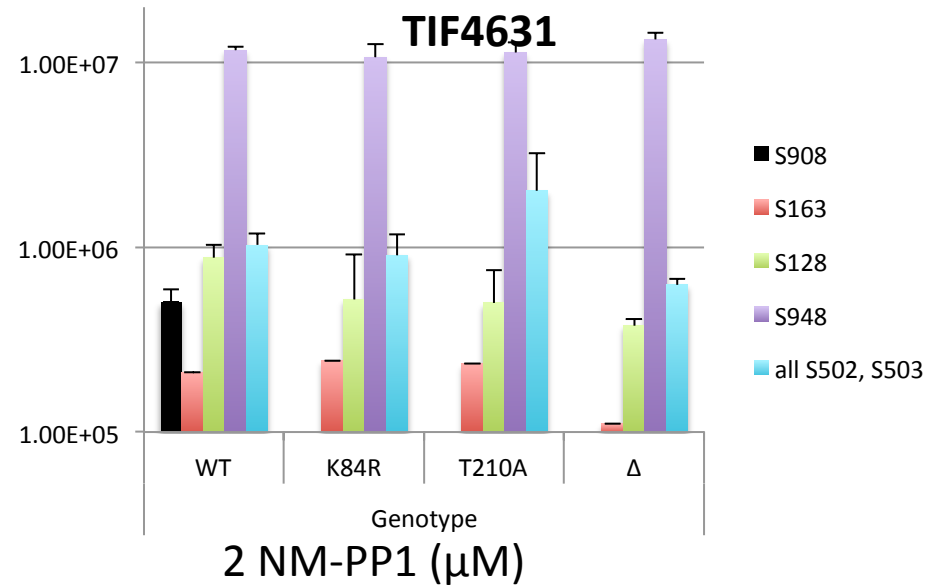
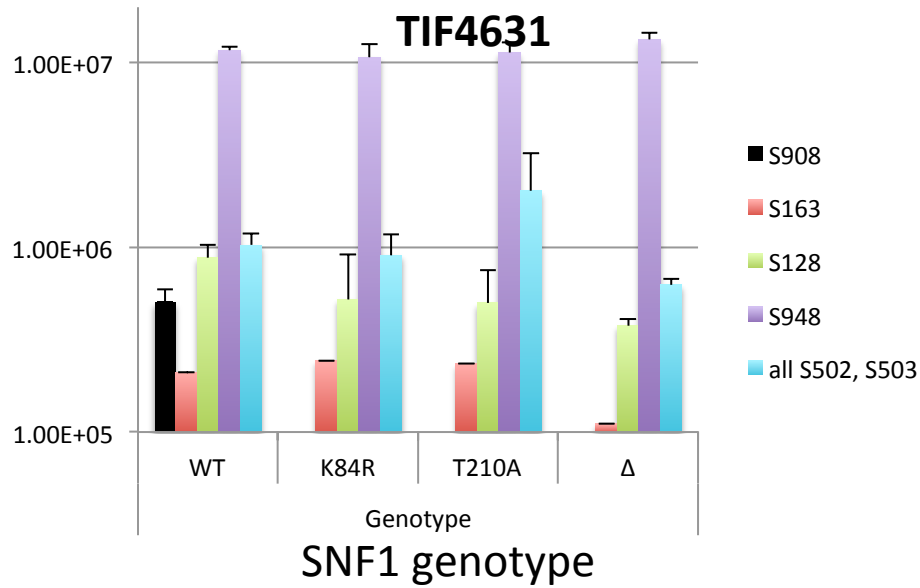
SNF1 genotype



2 NM-PP1 (μ M)

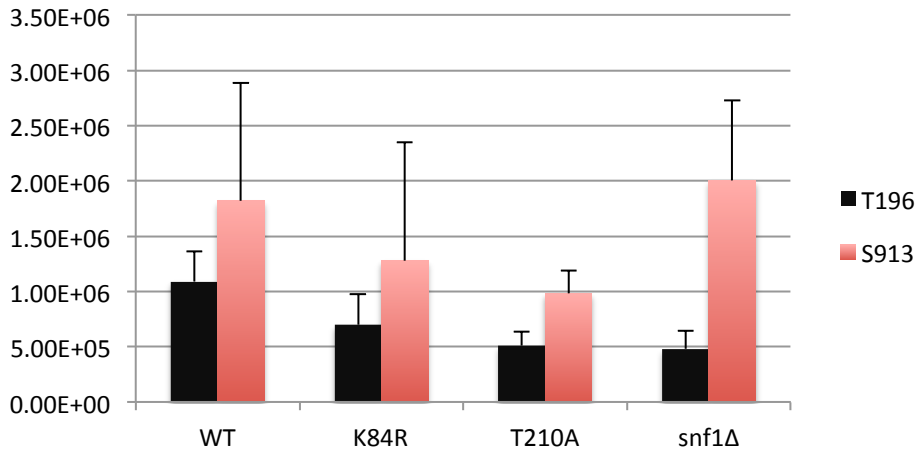


TIF4631



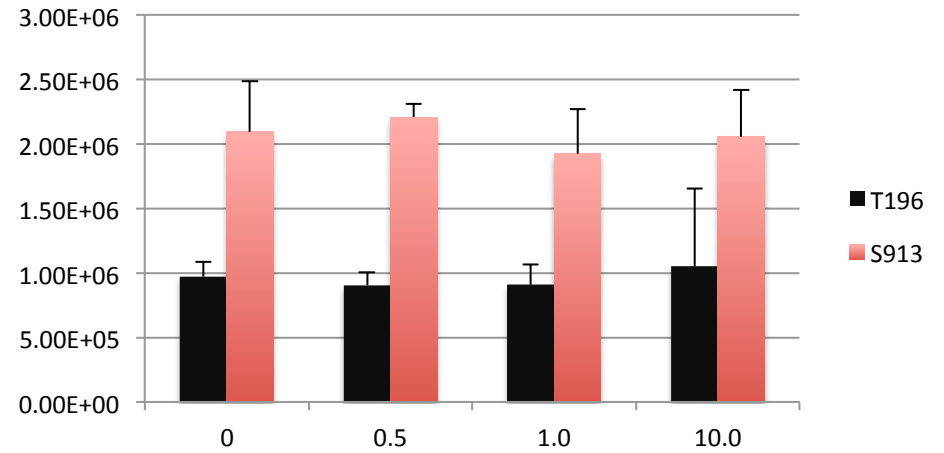
TIF4632

TIF4632



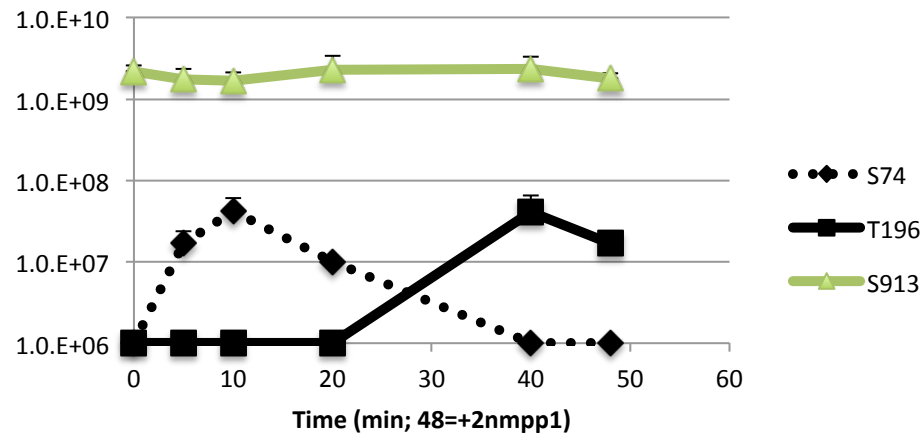
SNF1 genotype

TIF4632



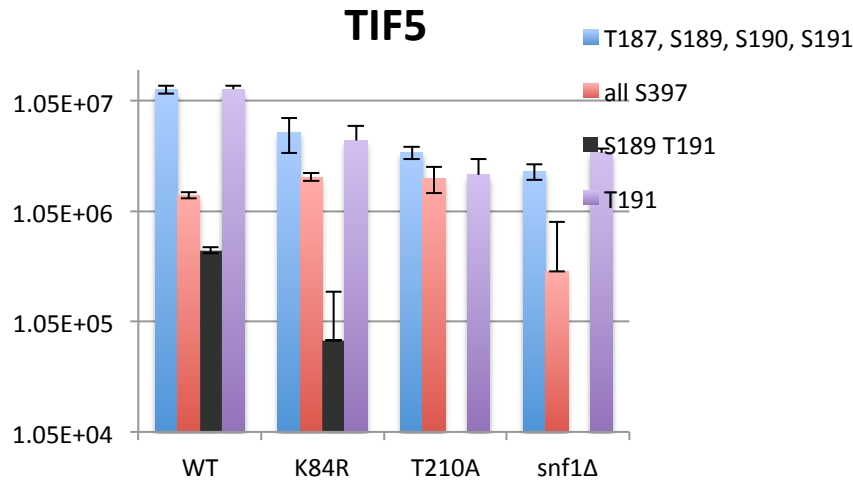
2 NM-PP1 (μM)

TIF4632

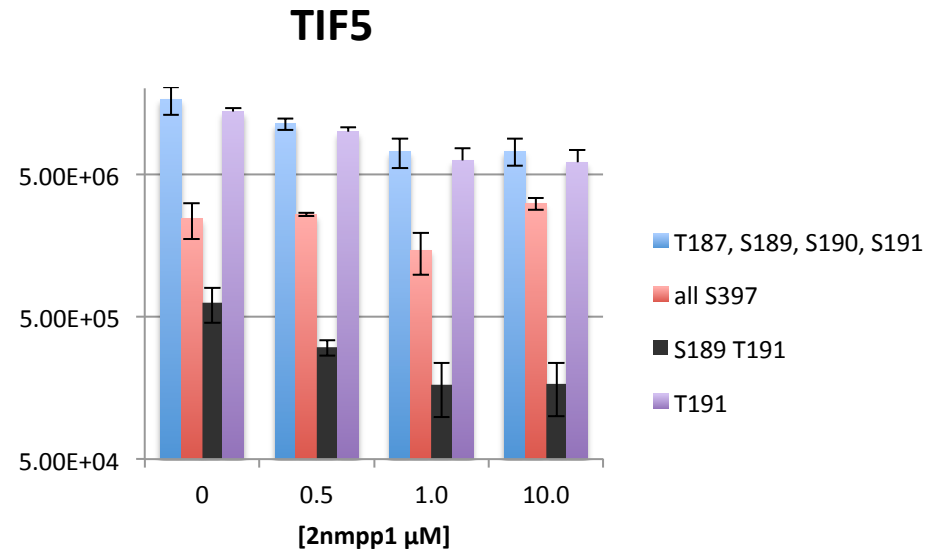


Min of derepression (48 = + 2 NM-PP1 (10 μM))

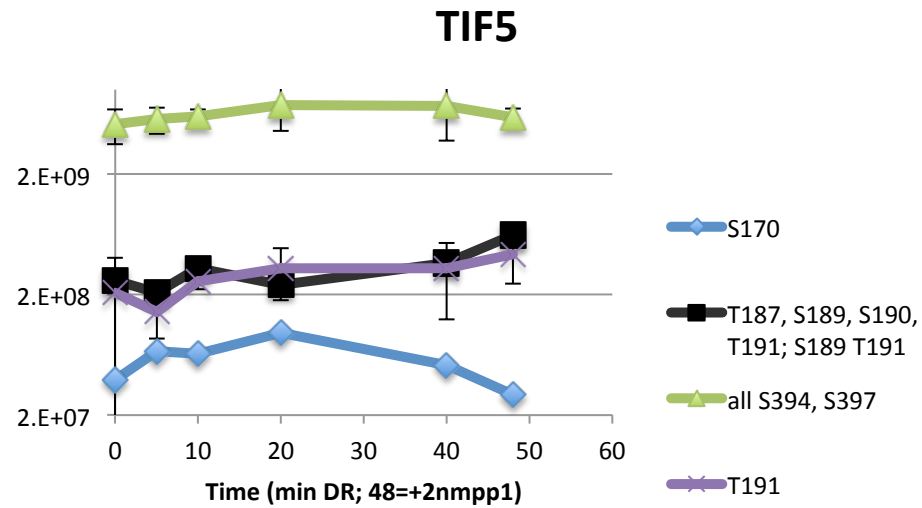
TIF5



SNF1 genotype



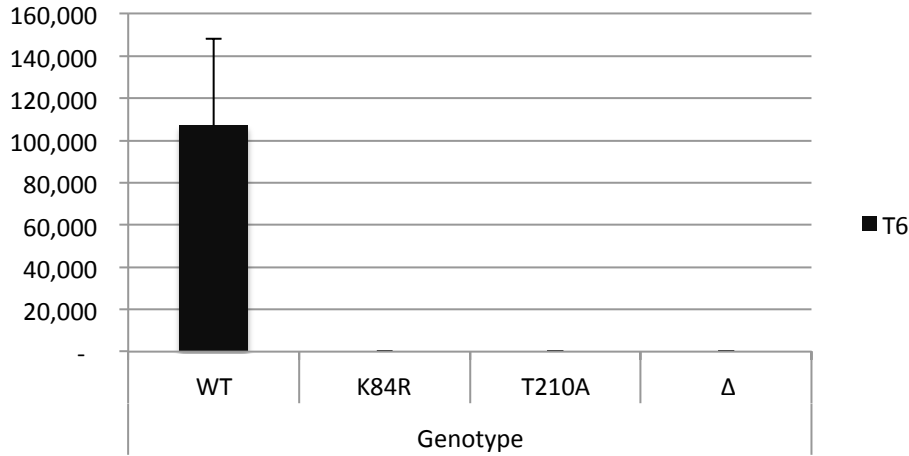
2NM-PP1 (μM)



Min of derepression (48 = + 2 NM-PP1 (10 μM))

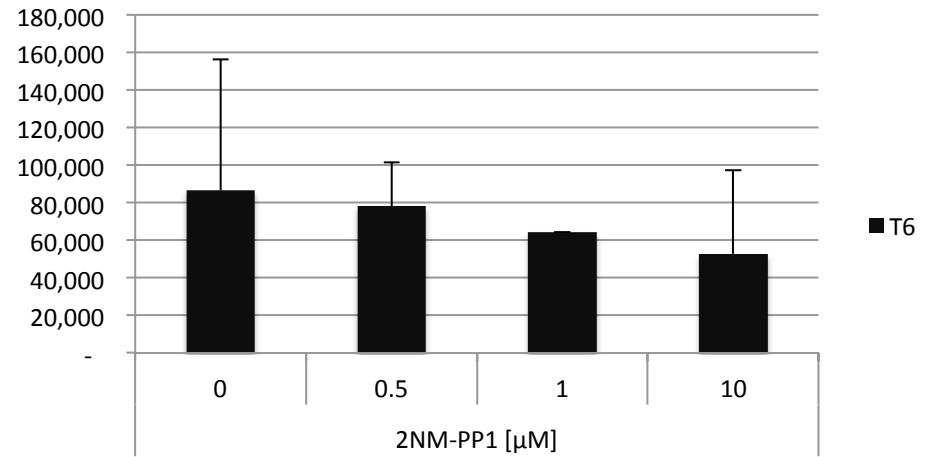
TIP41

TIP41



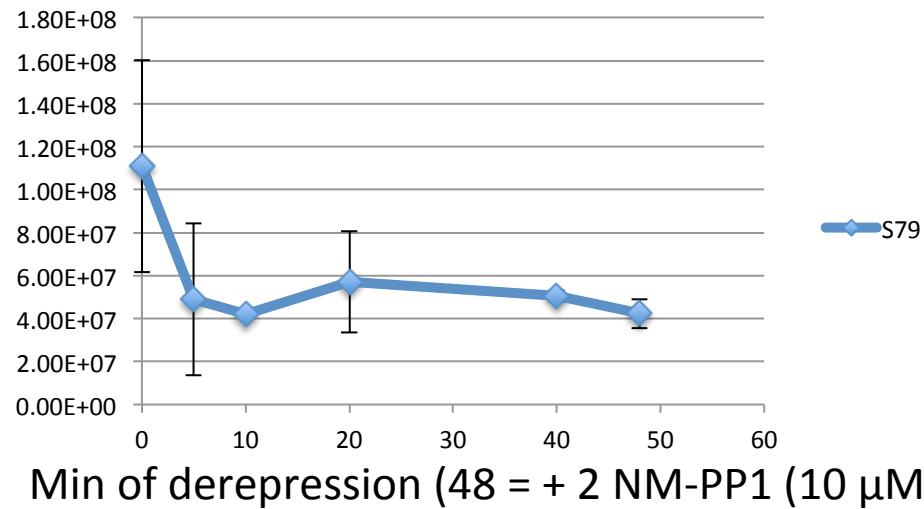
SNF1 genotype

TIP41



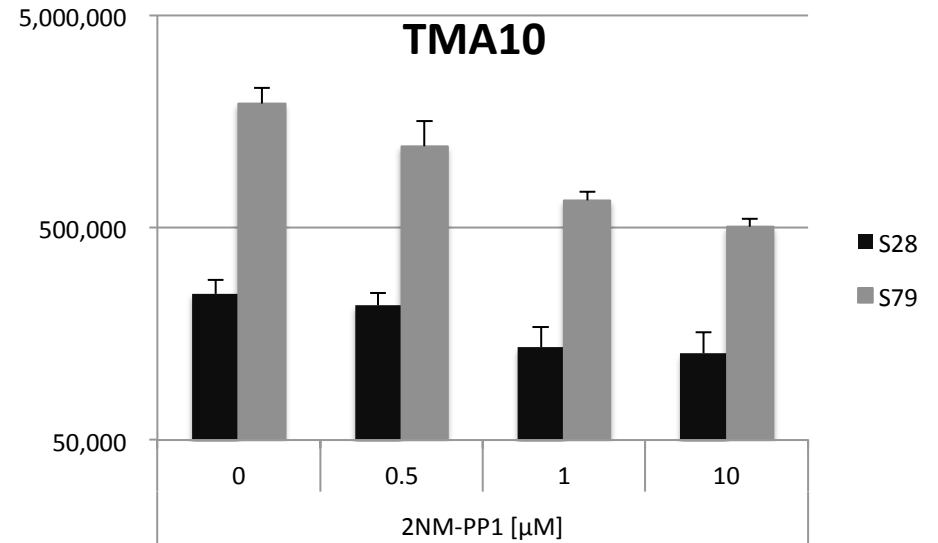
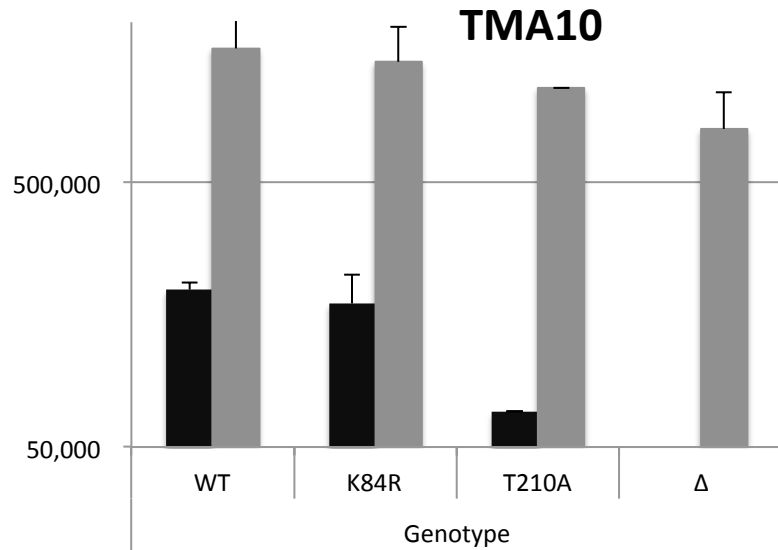
2 NM-PP1 (μM)

TIP41



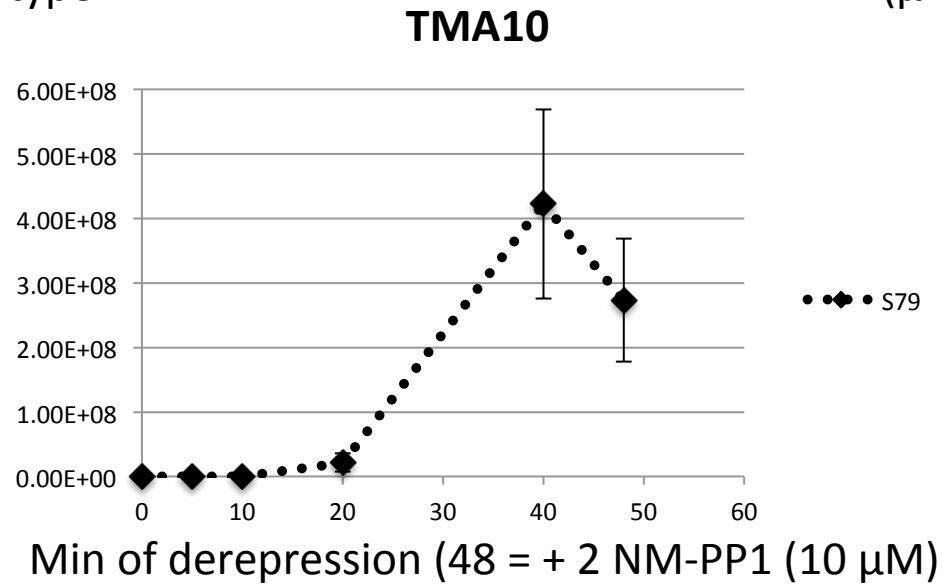
Min of derepression (48 = + 2 NM-PP1 (10 μM))

TMA10

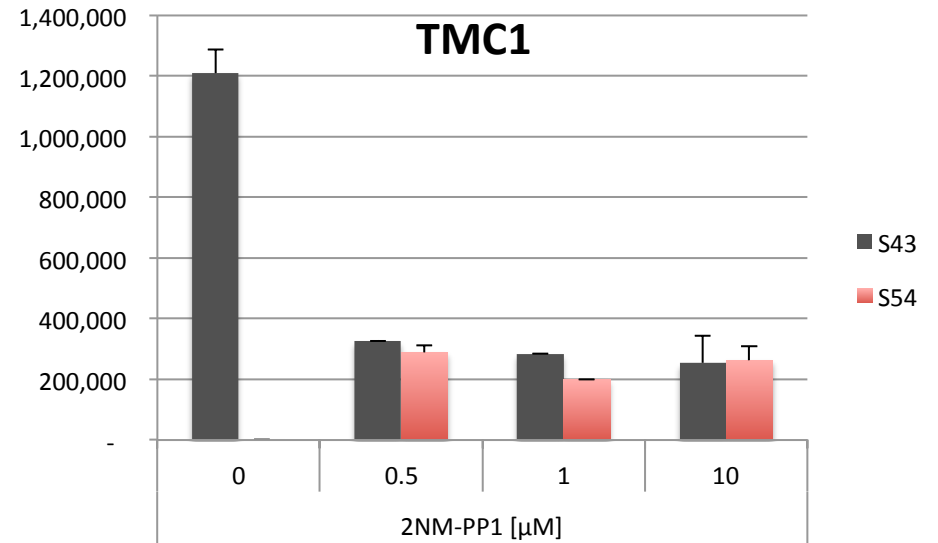
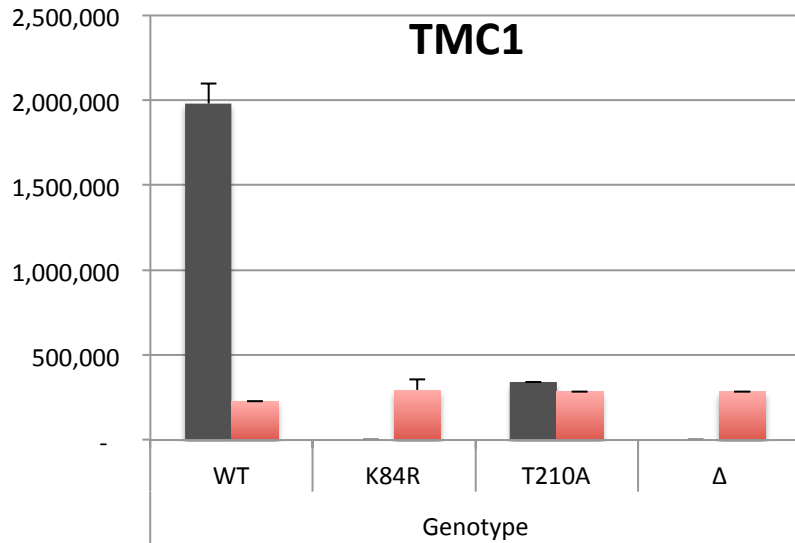


SNF1 genotype

2 NM-PP1 (μM)

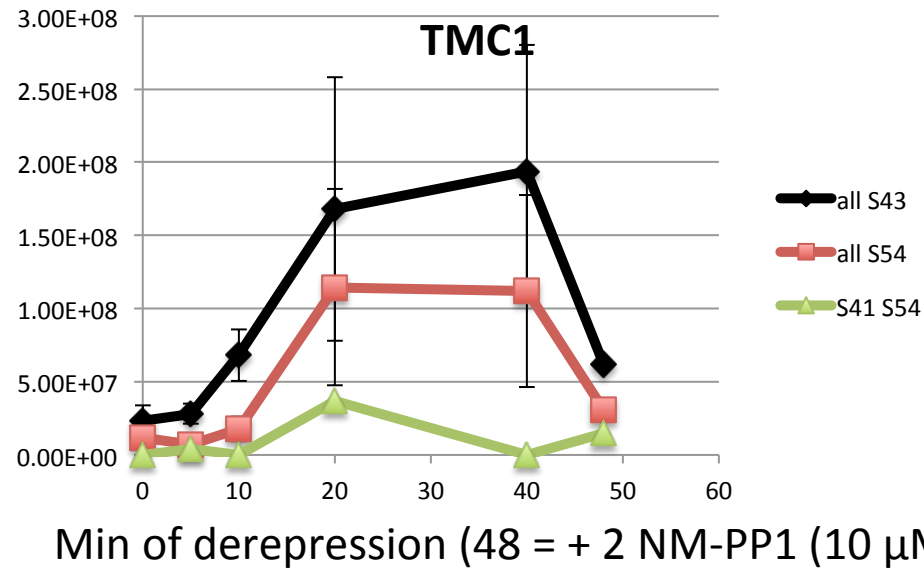


TMC1



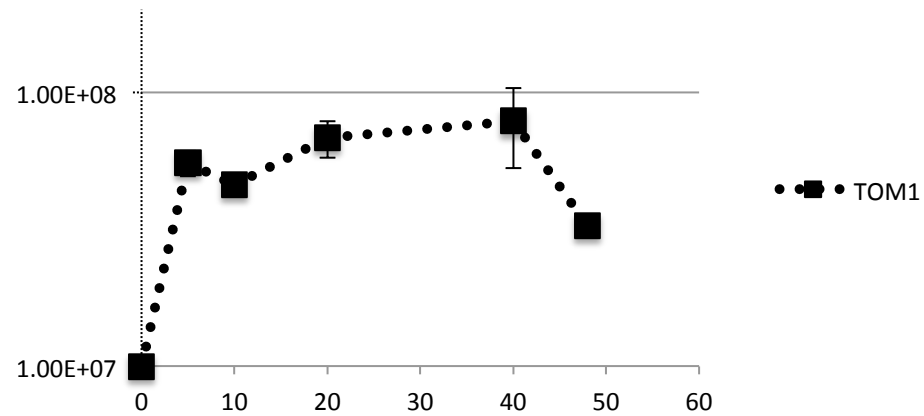
SNF1 genotype

2 NM-PP1 (μM)



TOM10

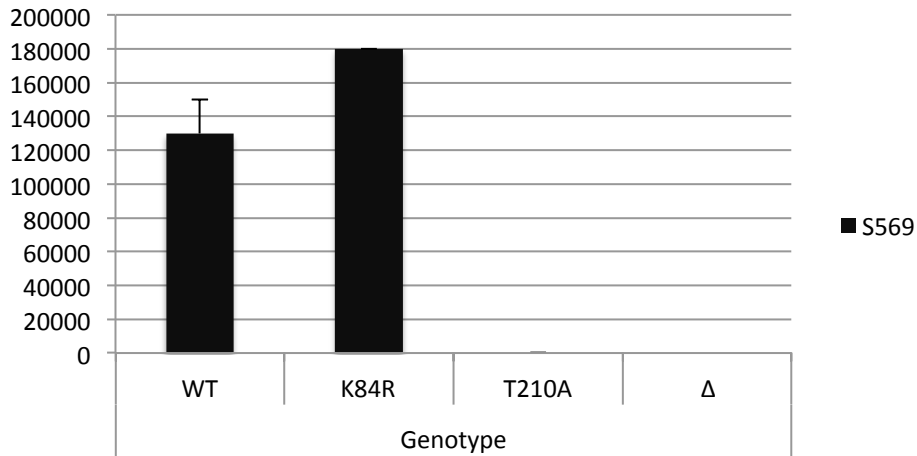
TOM1-S1890



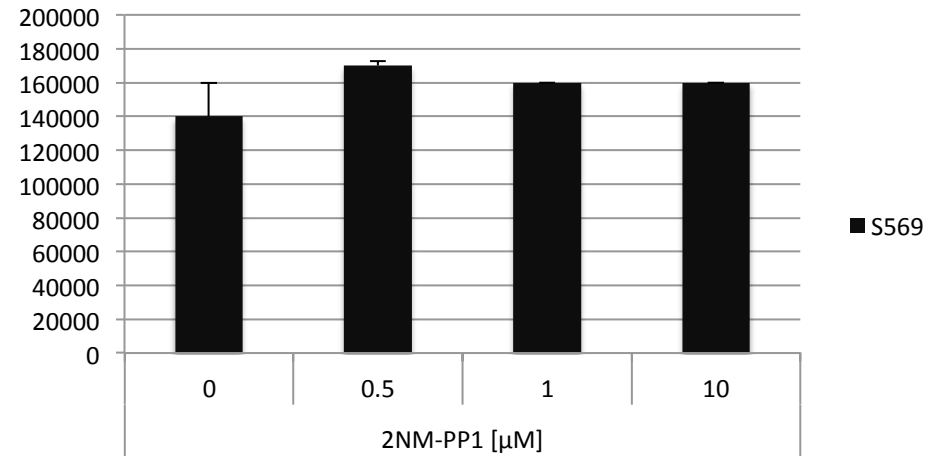
Min of derepression (48 = + 2 NM-PP1 (10 μ M))

TPO5

TPO5



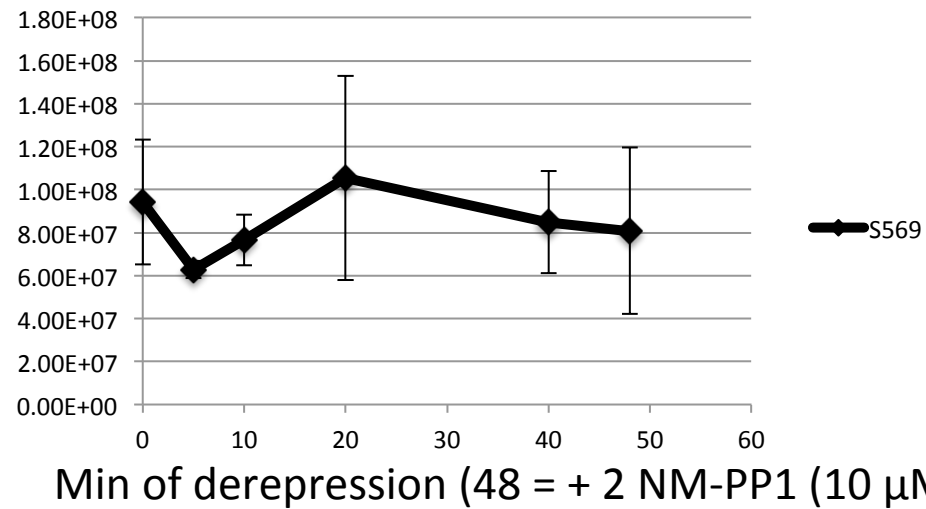
TPO5



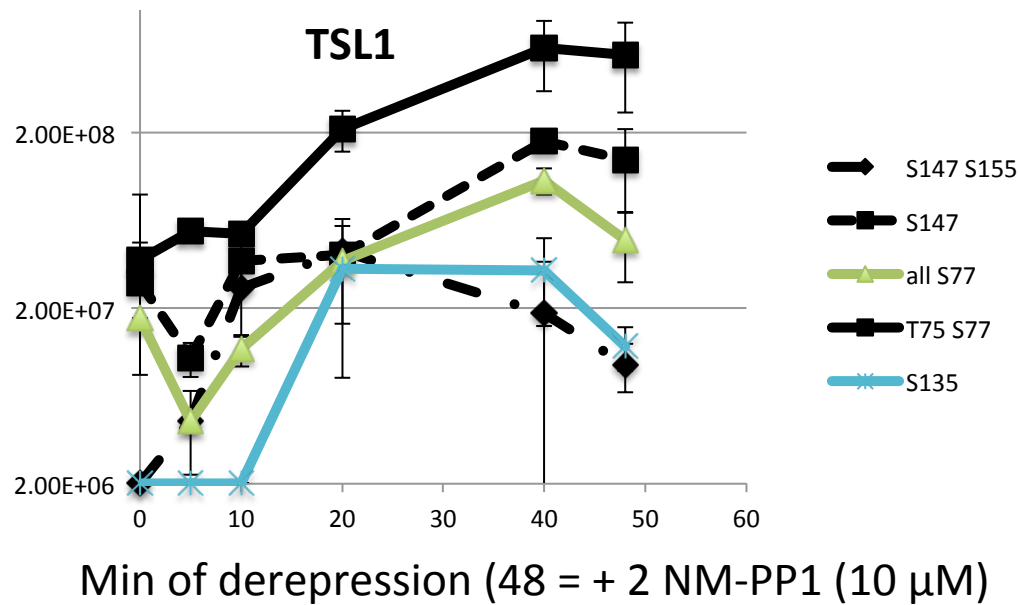
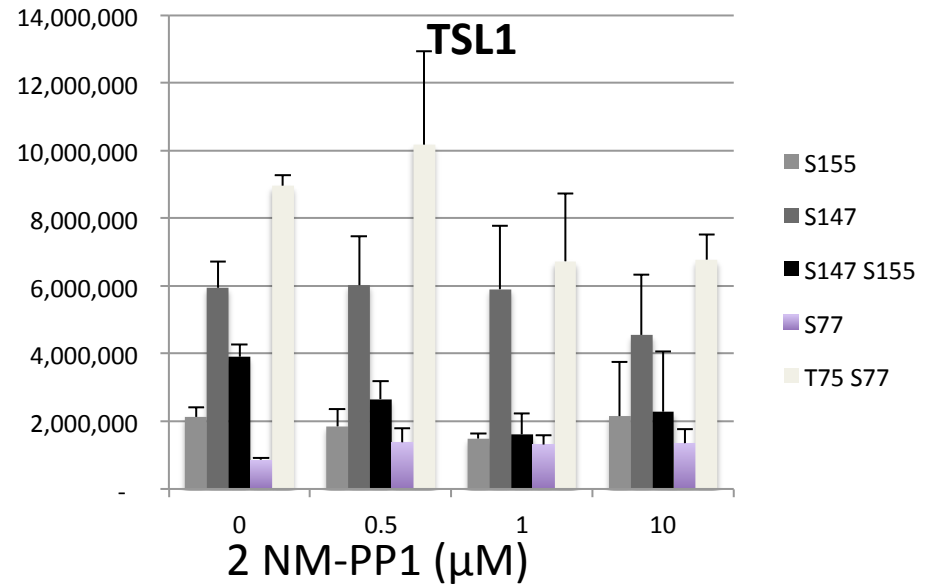
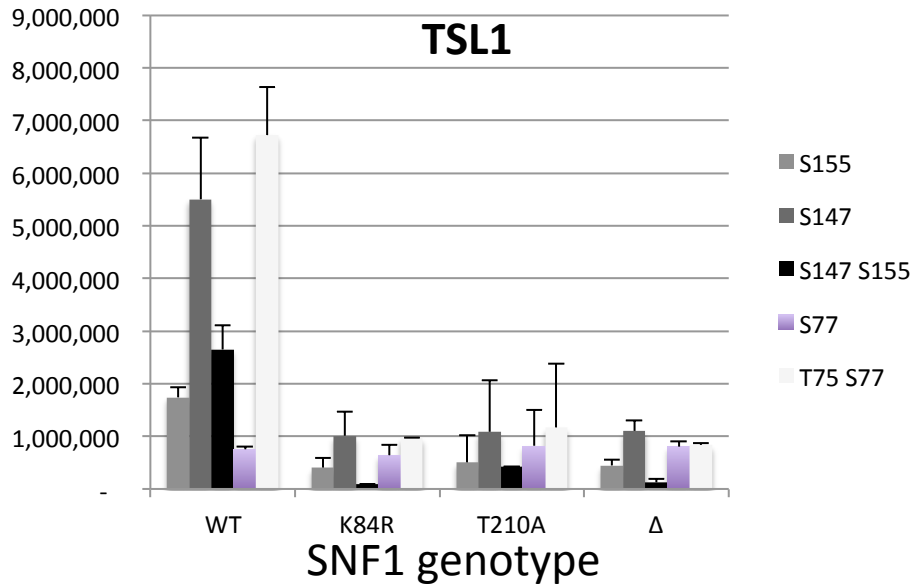
SNF1 genotype

2 NM-PP1 (μM)

TPO5

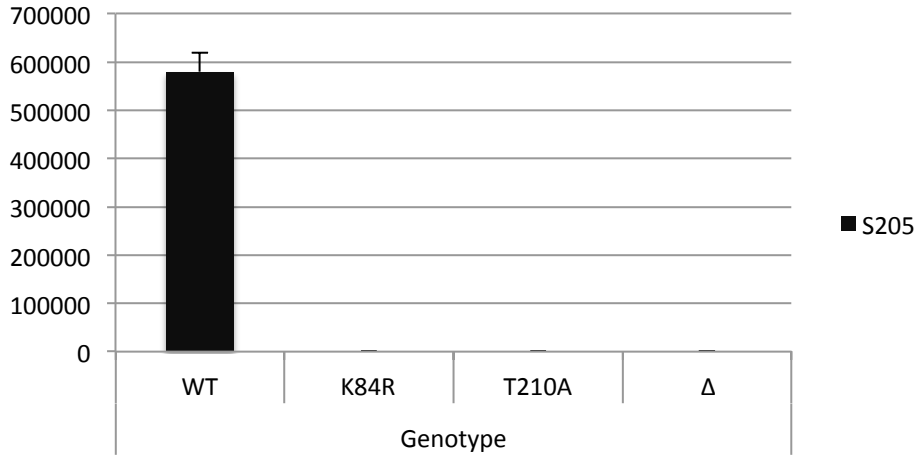


TSL1



UIP4

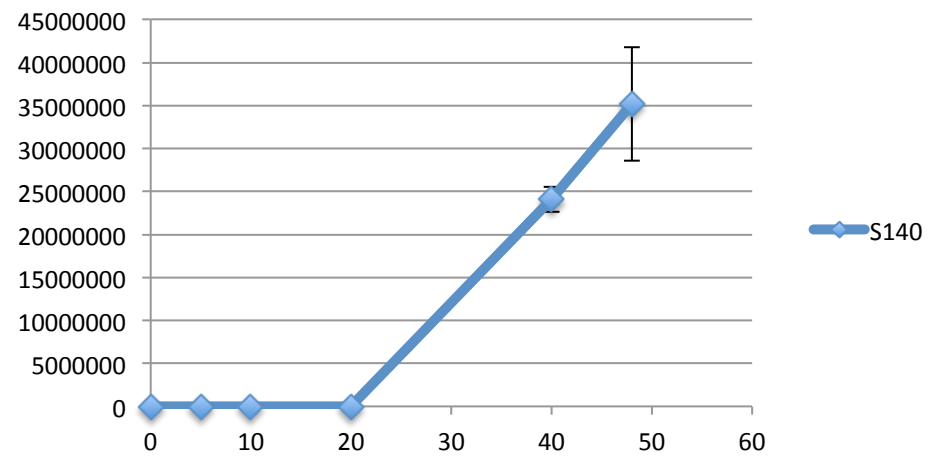
UIP4



No data

SNF1 genotype

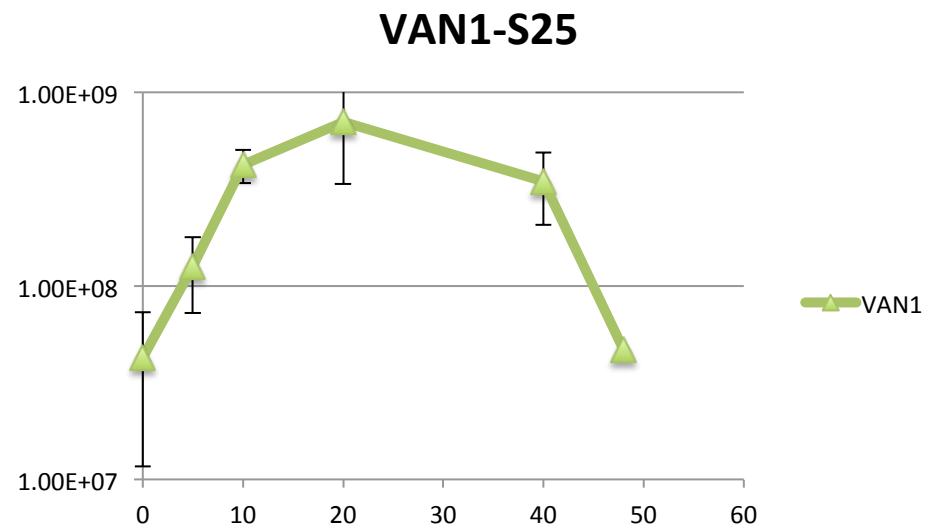
UIP4



Min of derepression (48 = + 2 NM-PP1 (10 μM))

2 NM-PP1 (μM)

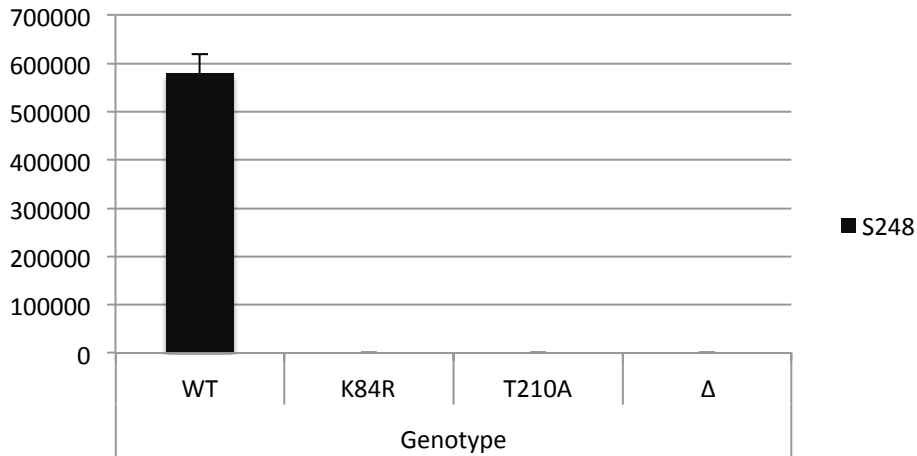
VAN1



Min of derepression (48 = + 2 NM-PP1 (10 μ M))

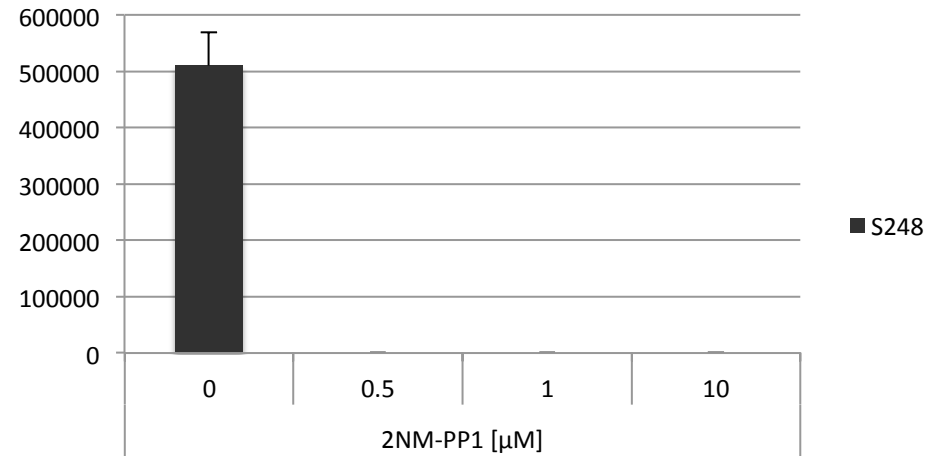
VID30

VID30



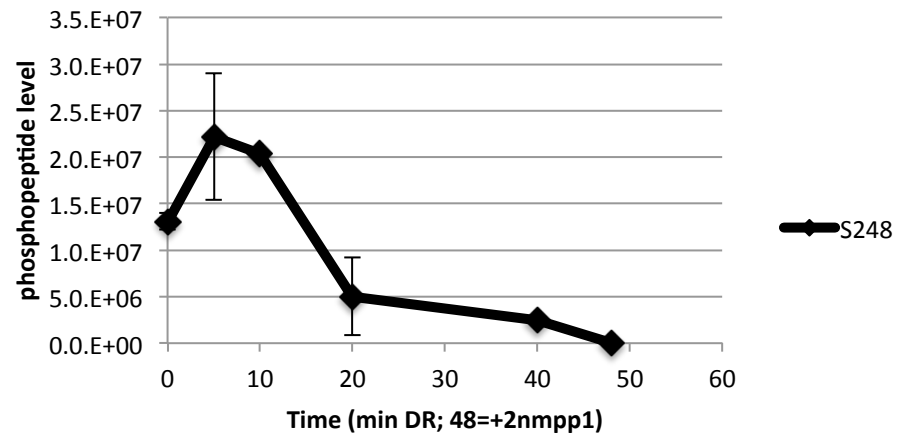
SNF1 genotype

VID30



2 NM-PP1 (μM)

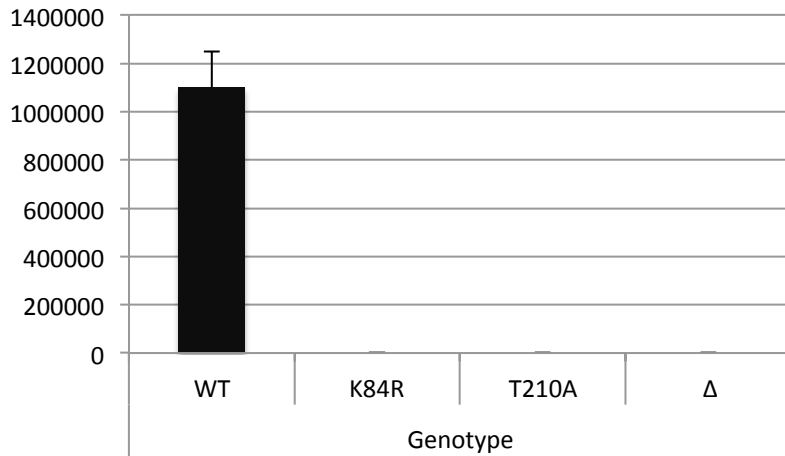
VID30



Min of derepression (48 = + 2 NM-PP1 (10 μM))

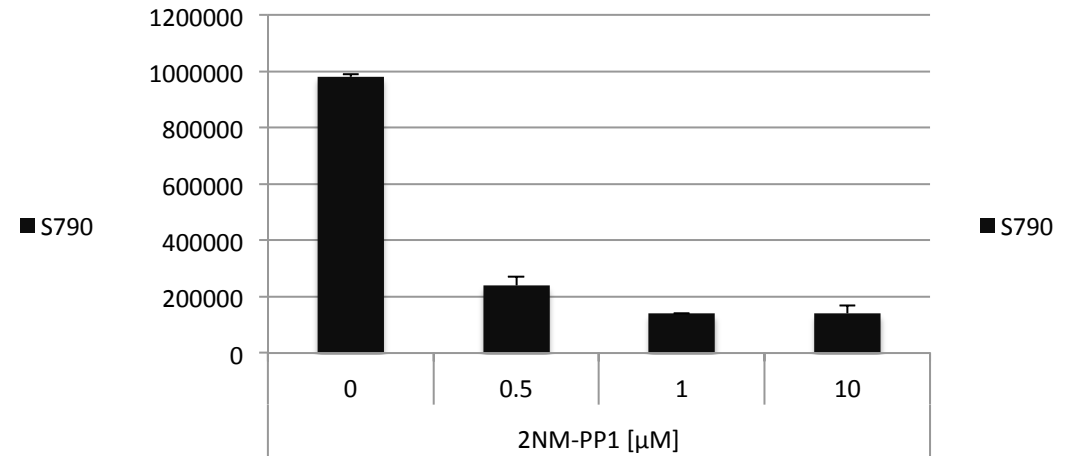
VPS53

VPS53



SNF1 genotype

VPS53



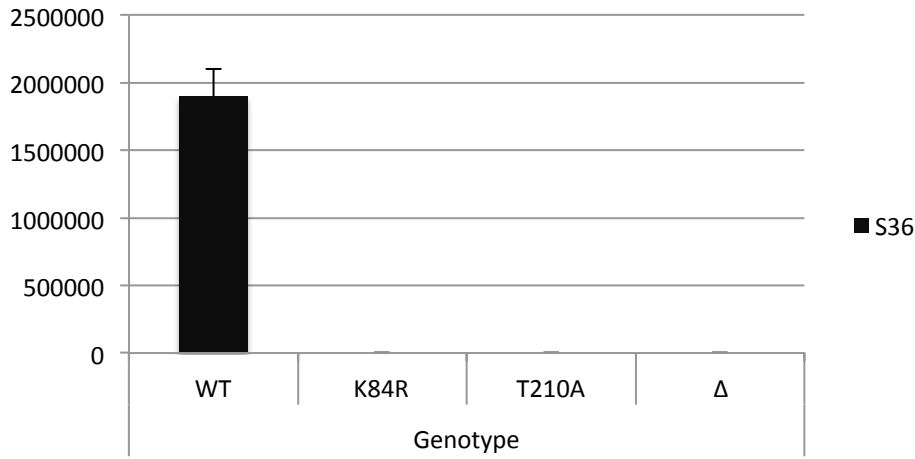
2 NM-PP1 (μM)

No data

Min of derepression (48 = + 2 NM-PP1 (10 μM))

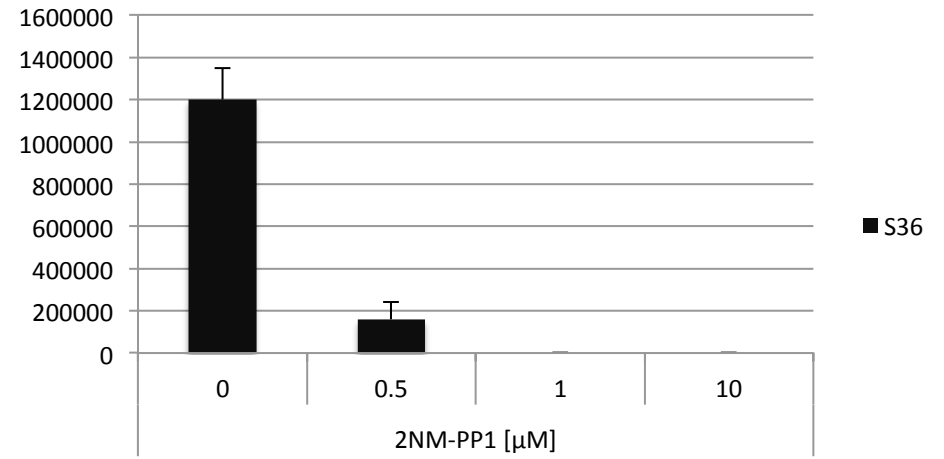
WHI3

WHI3



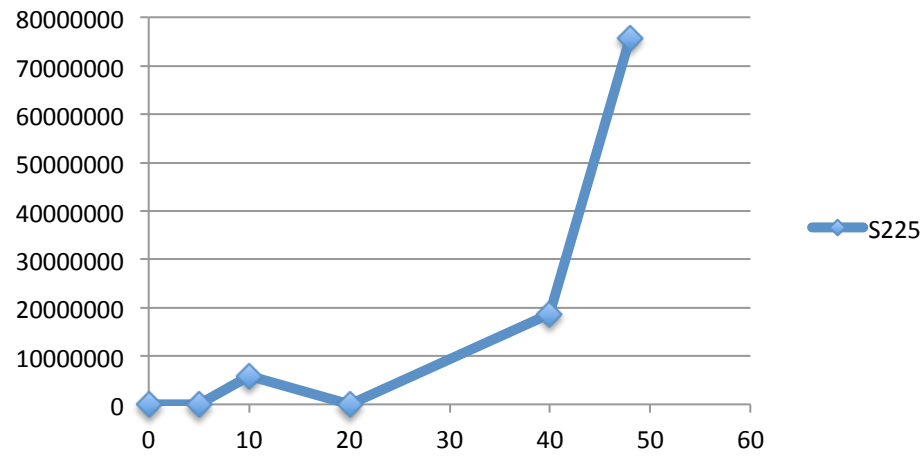
SNF1 genotype

WHI3



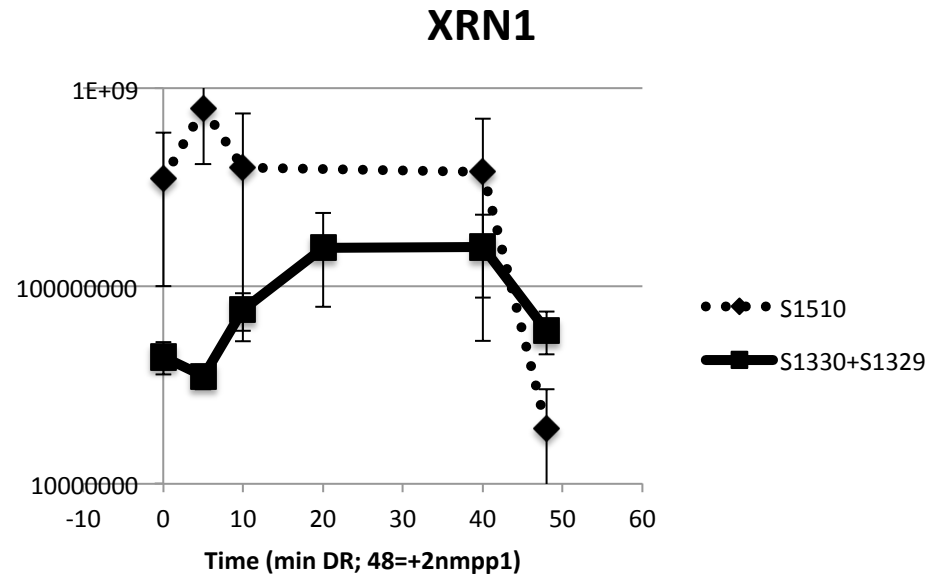
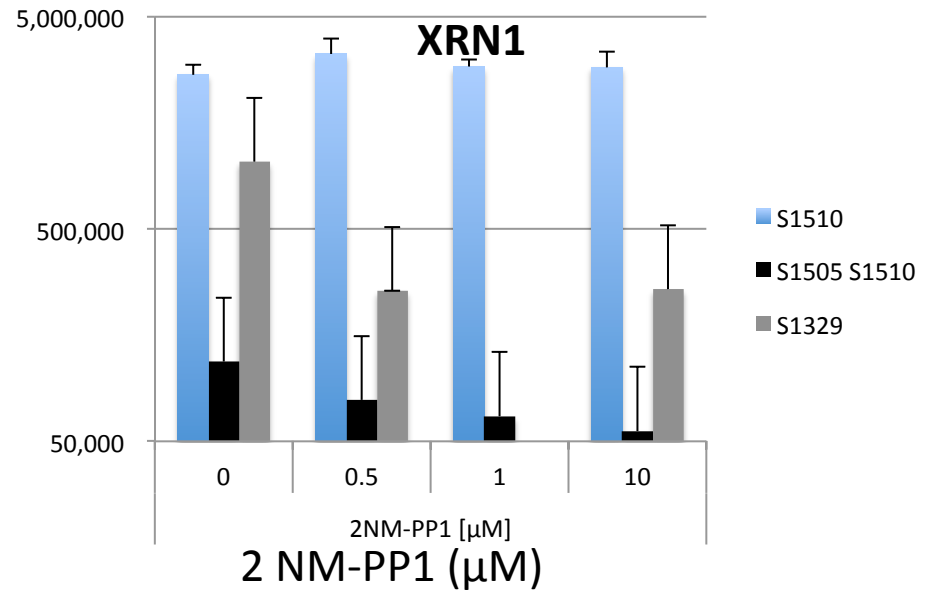
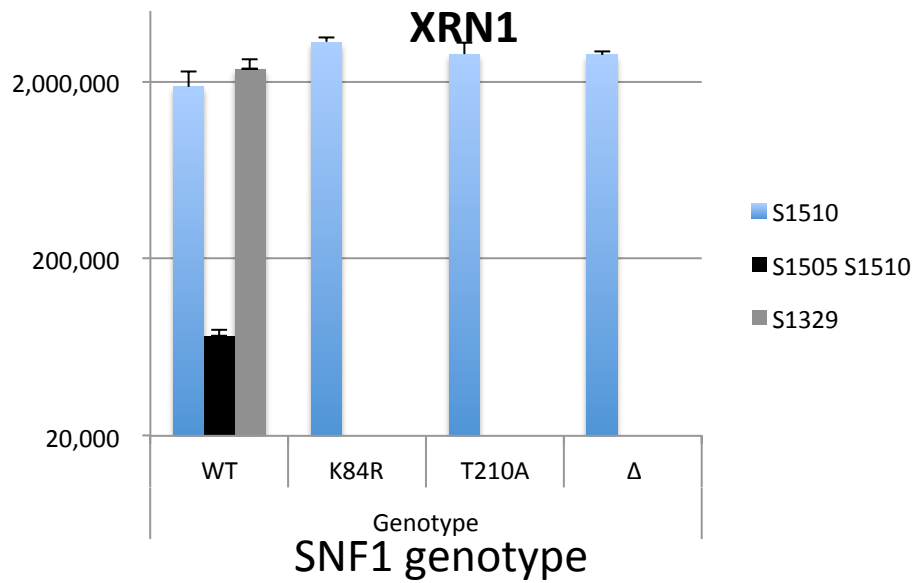
2 NM-PP1 (μM)

WHI3



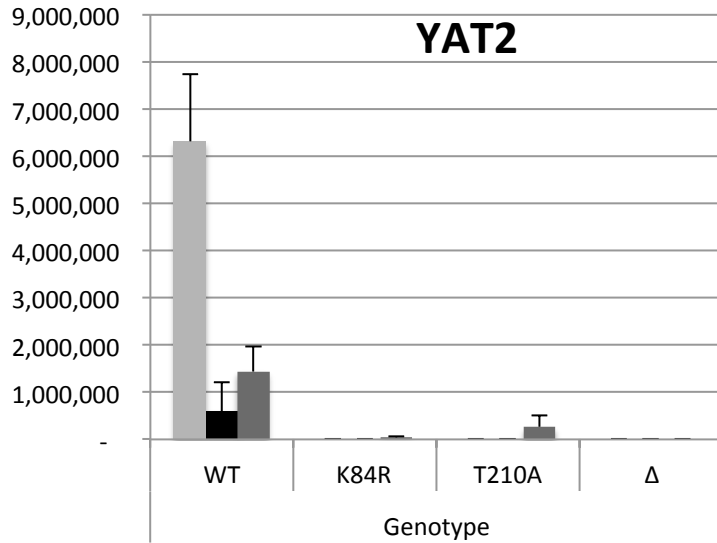
Min of derepression (48 = + 2 NM-PP1 (10 μM))

XRN1

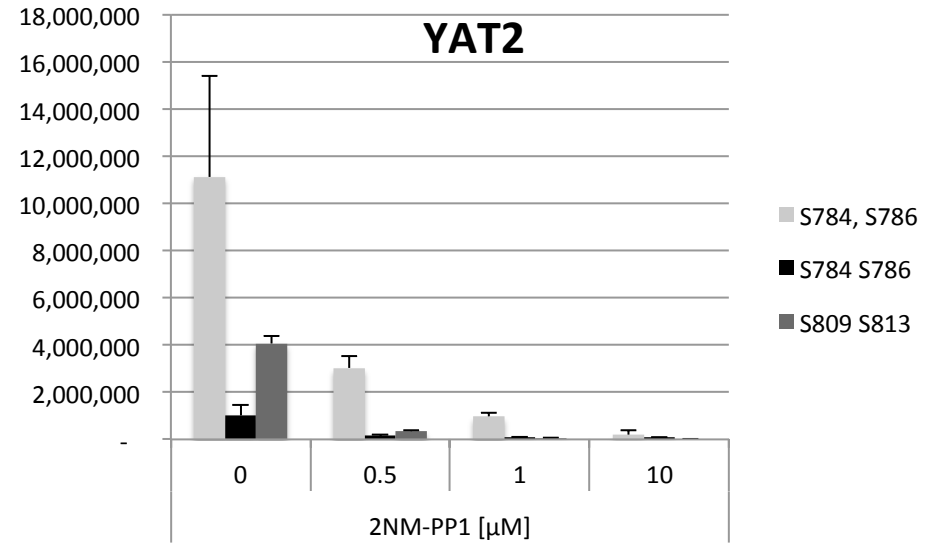


Min of derepression (48 = + 2 NM-PP1 (10 μM))

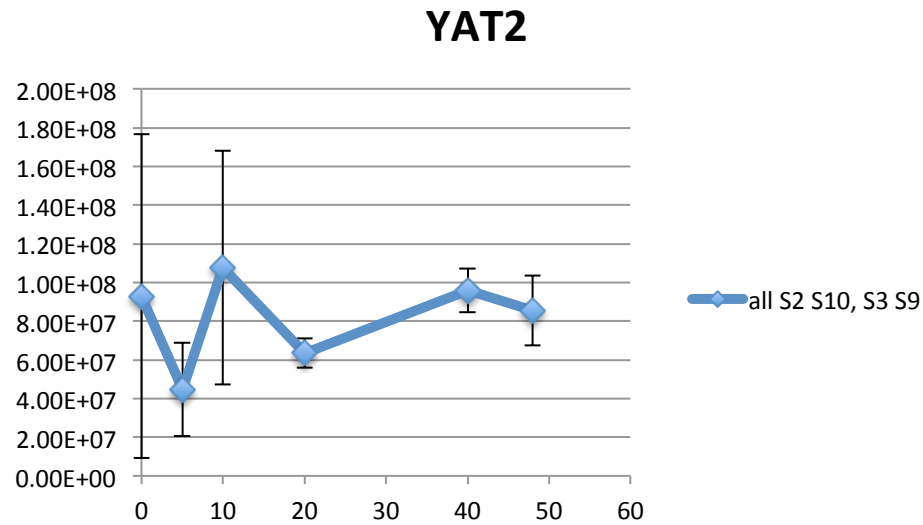
YAT2



SNF1 genotype

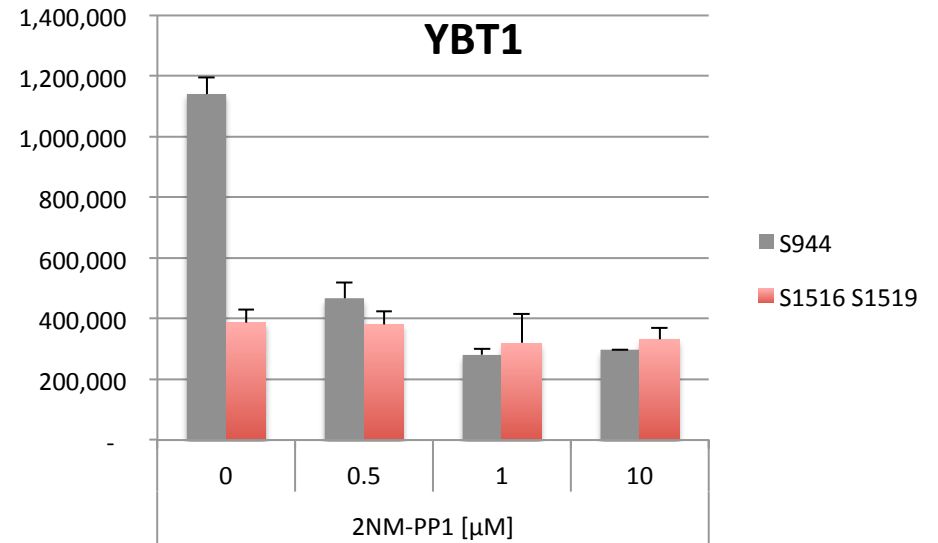
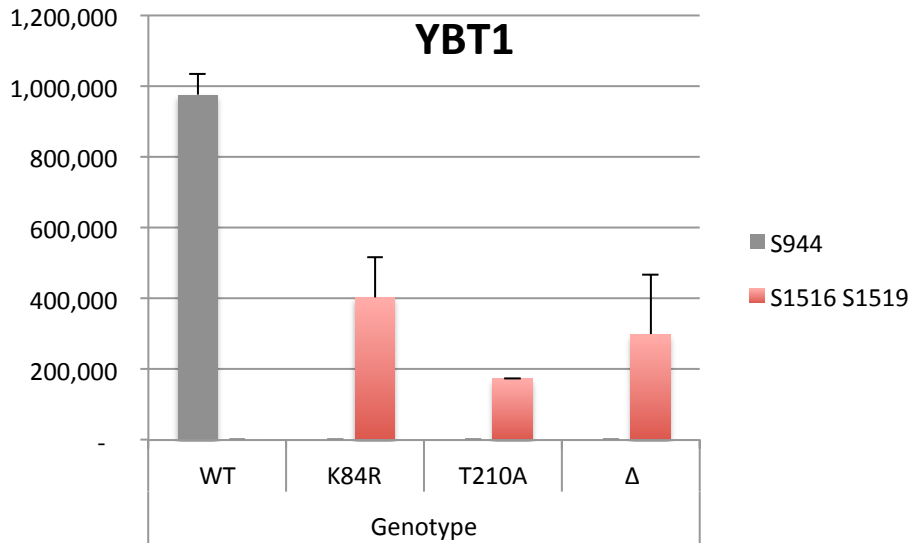


2 NM-PP1 (μM)



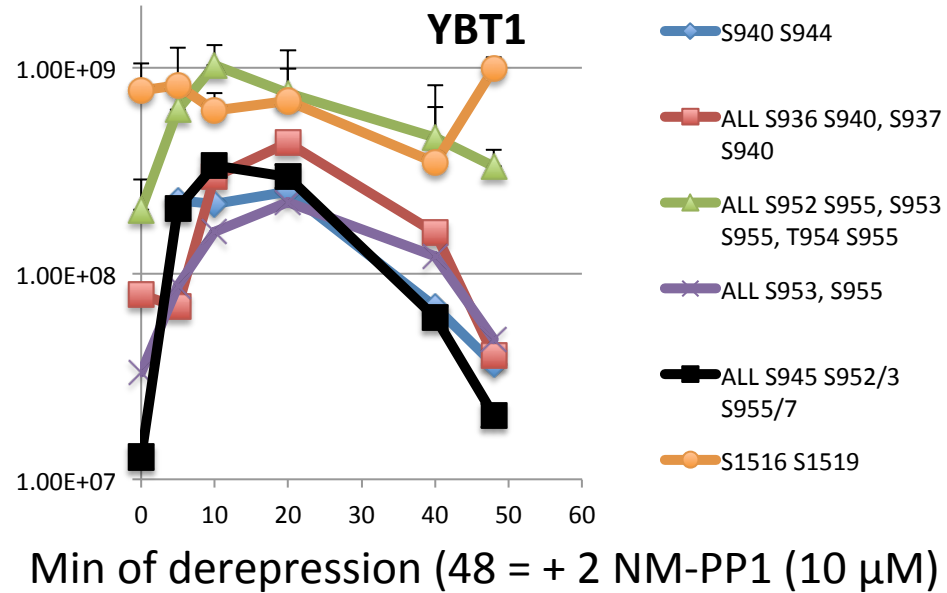
Min of derepression (48 = + 2 NM-PP1 (10 μM))

YBT1

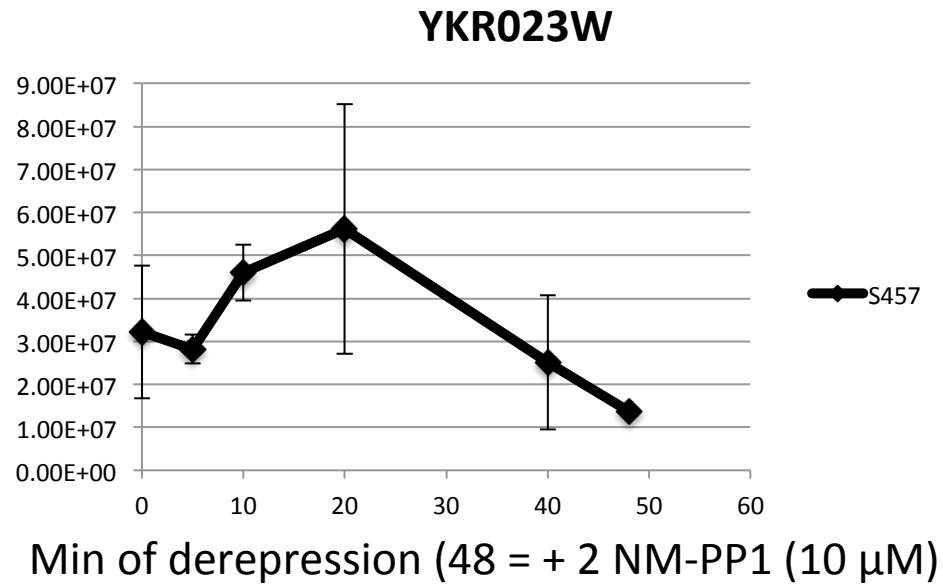
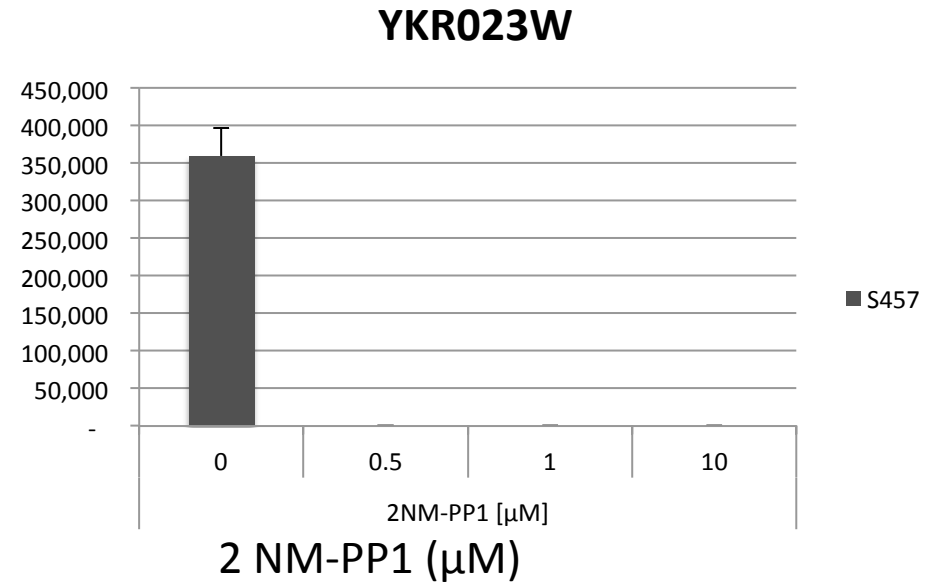
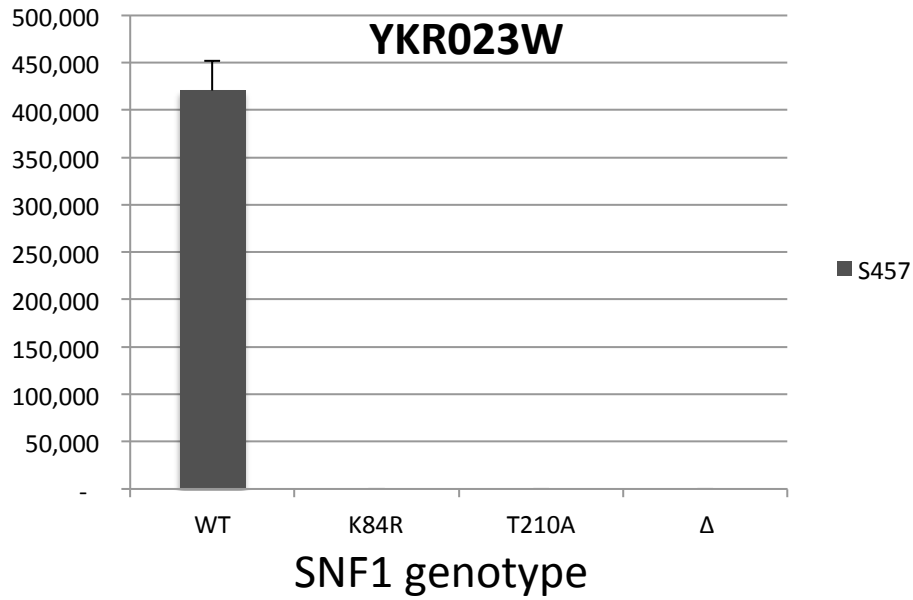


SNF1 genotype

2 NM-PP1 (μM)

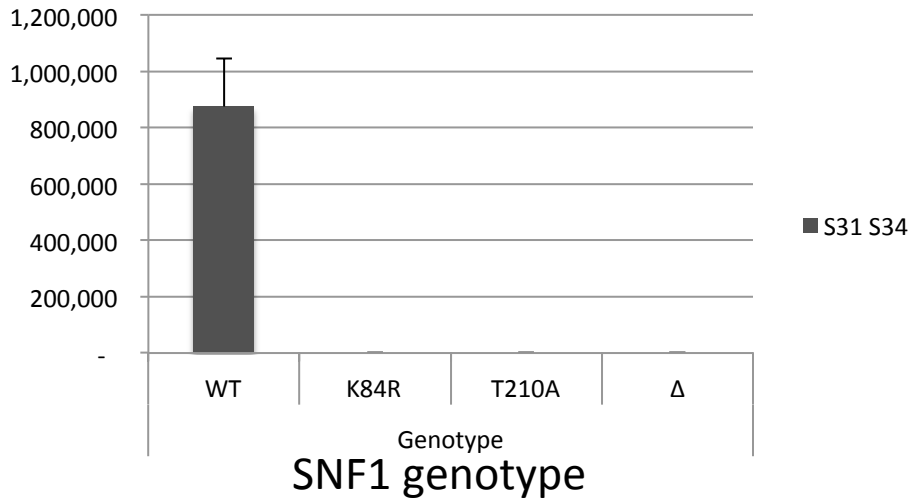


YKR023W

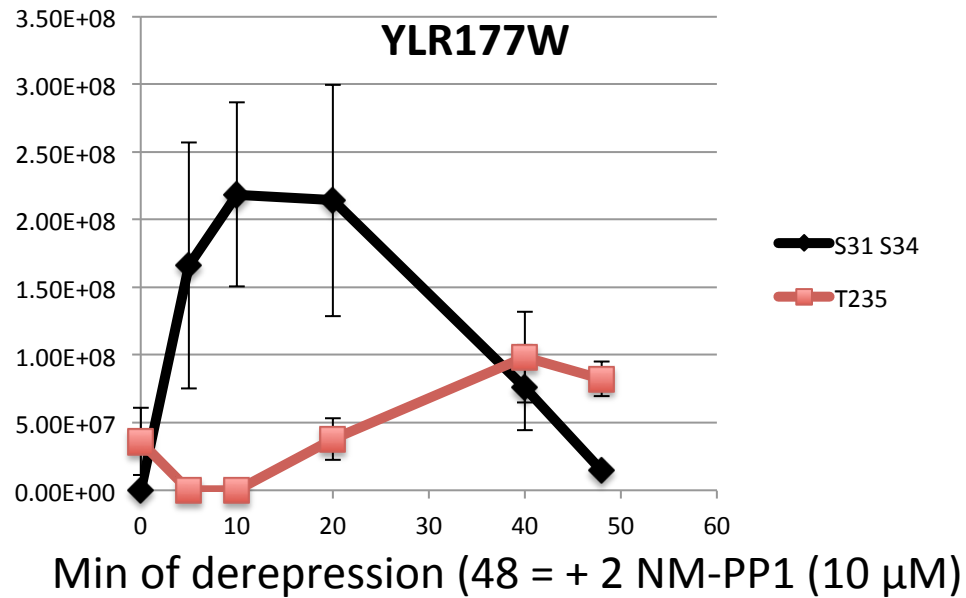
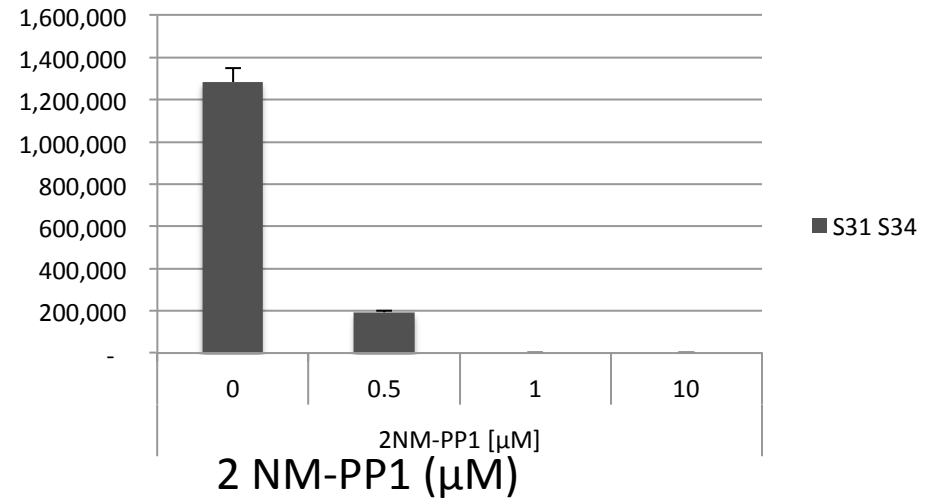


YLR177W

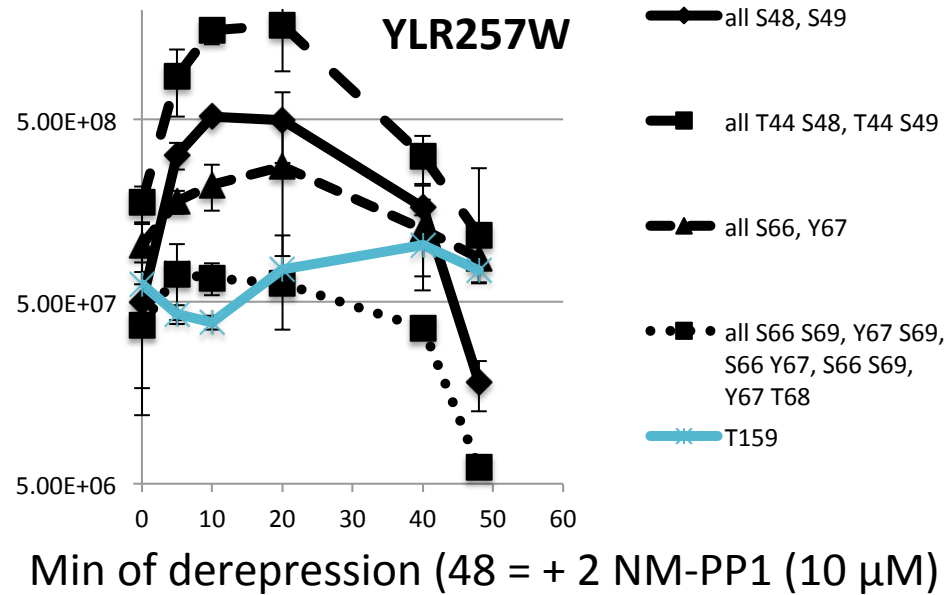
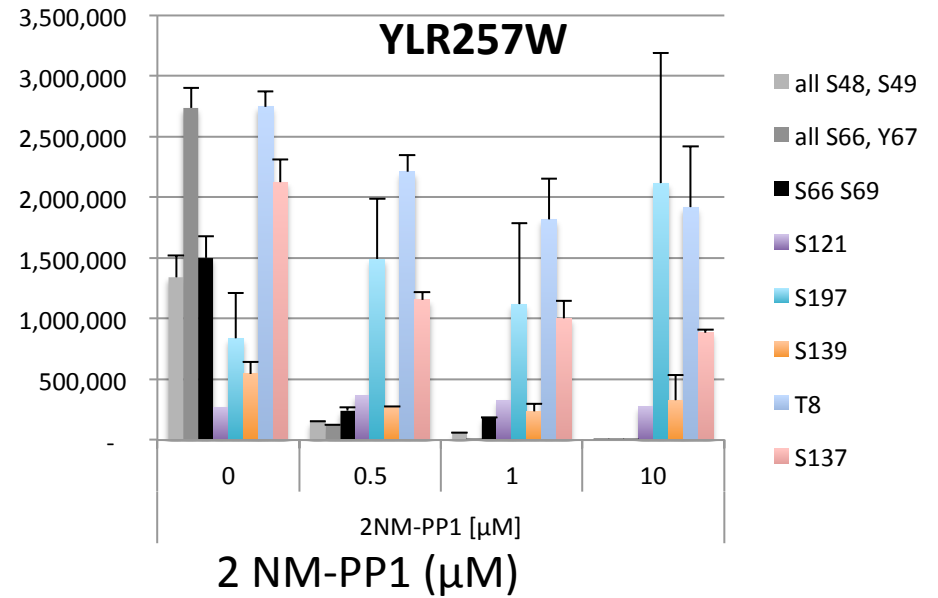
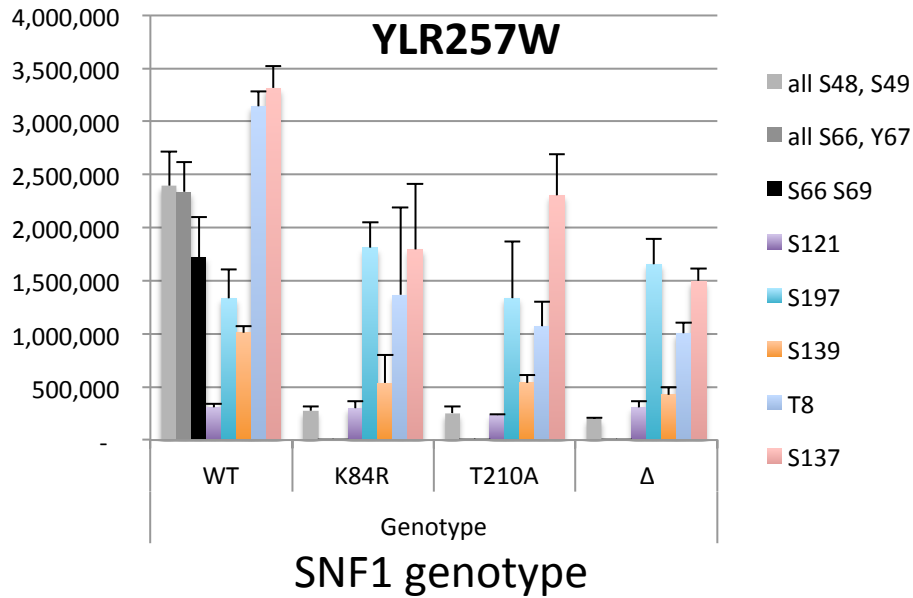
YLR177W



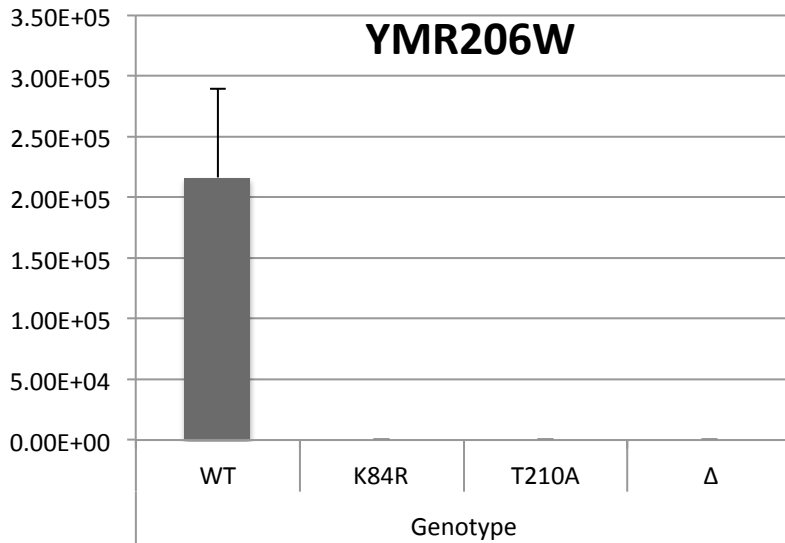
YLR177W



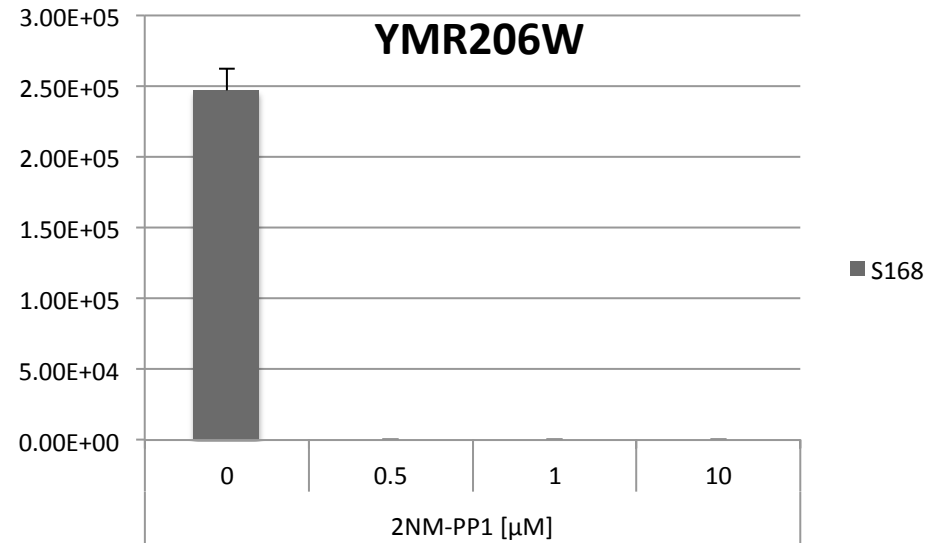
YLR257W



YMR206W



SNF1 genotype

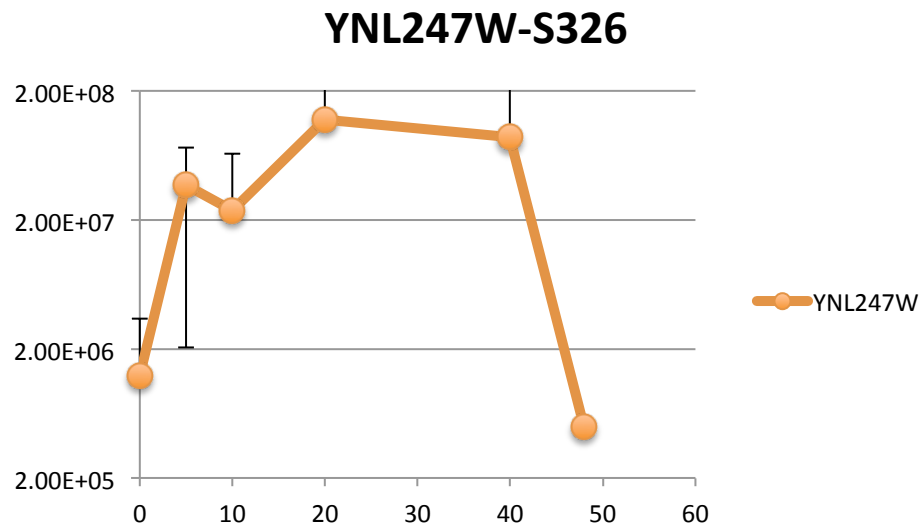


2 NM-PP1 (μM)

No data

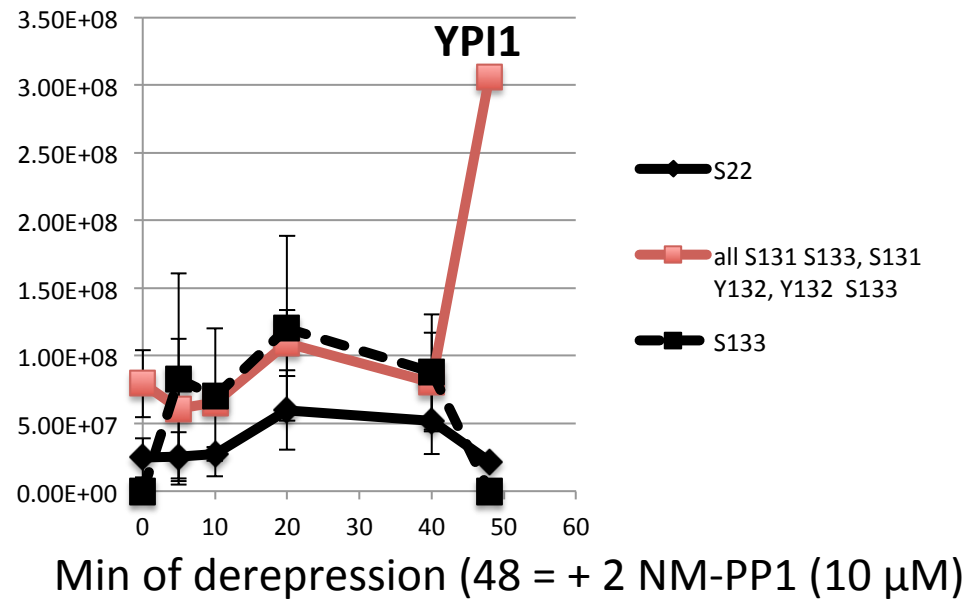
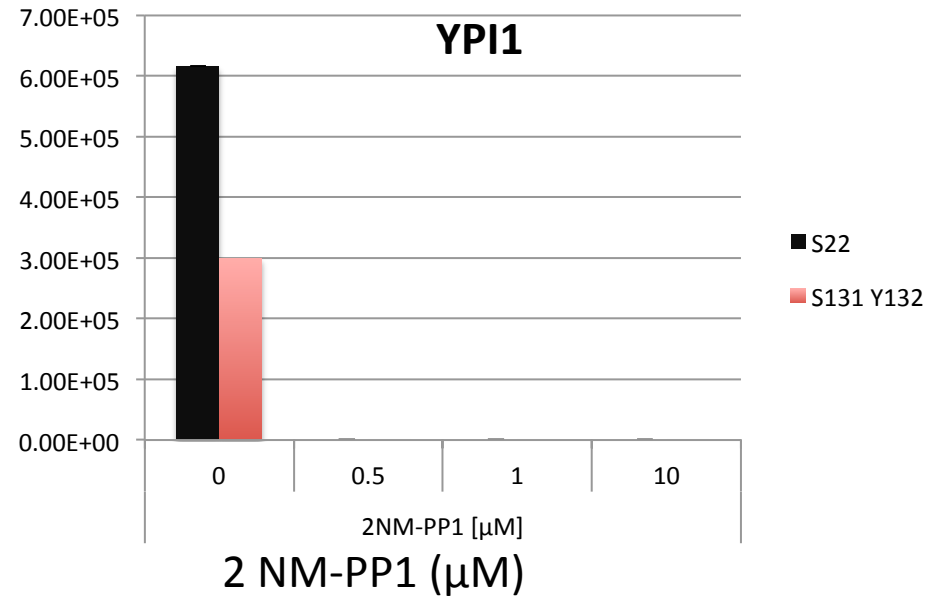
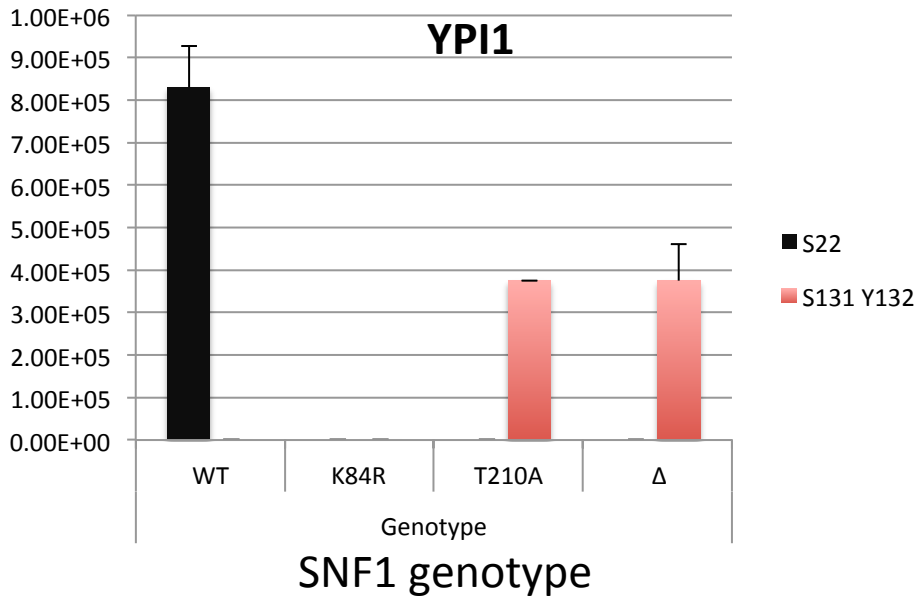
Min of derepression (48 = + 2 NM-PP1 (10 μM))

YNL247W

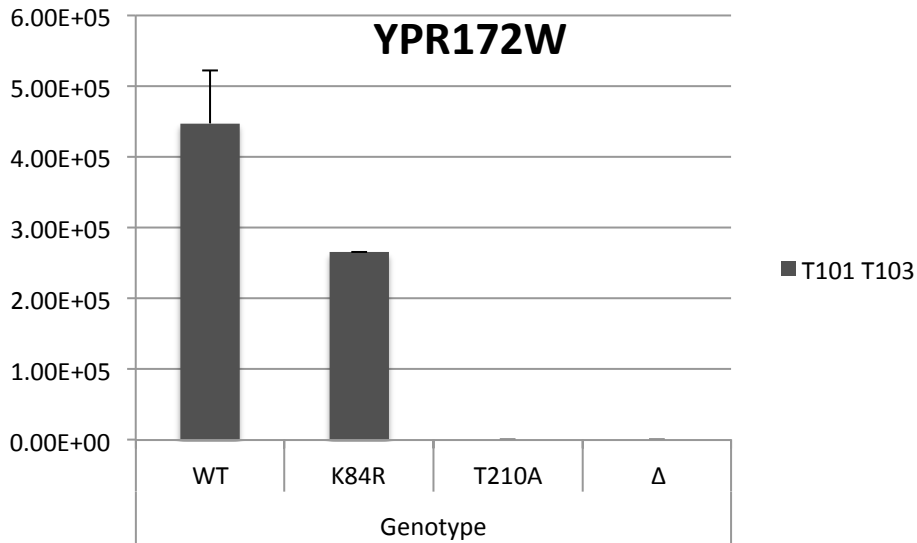


Min of derepression (48 = + 2 NM-PP1 (10 μ M))

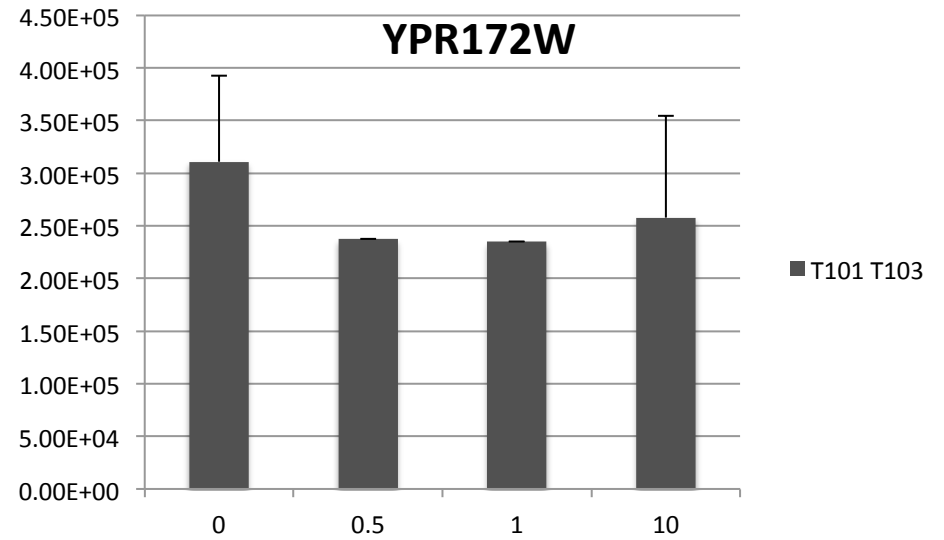
YPI1



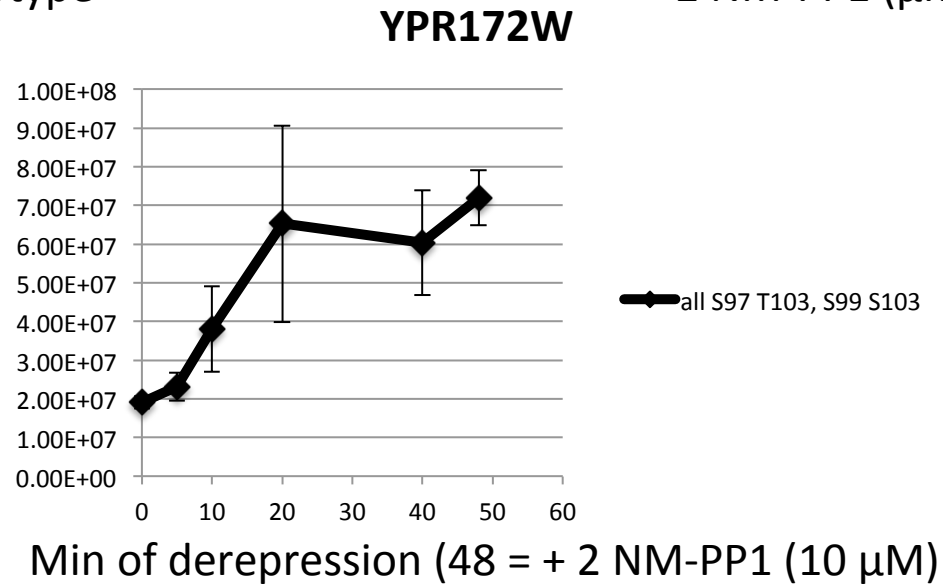
YPR172W



SNF1 genotype



2 NM-PP1 (μM)



Min of derepression (48 = + 2 NM-PP1 (10 μM))