Figure S1

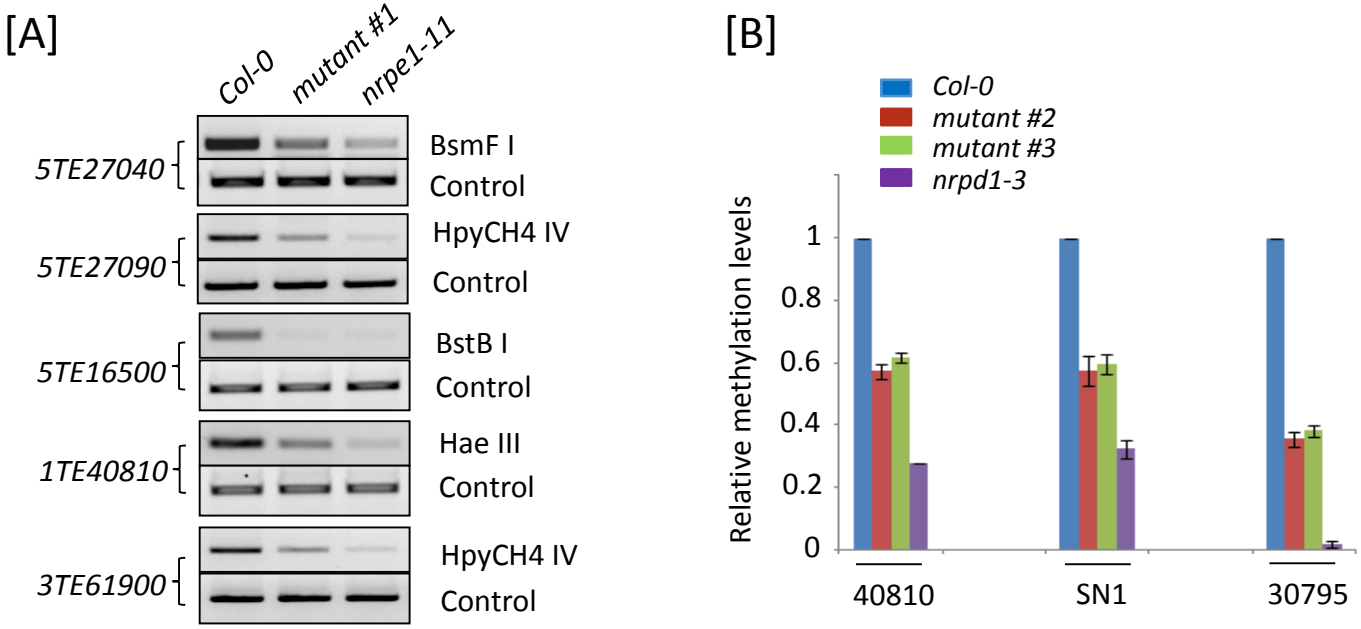
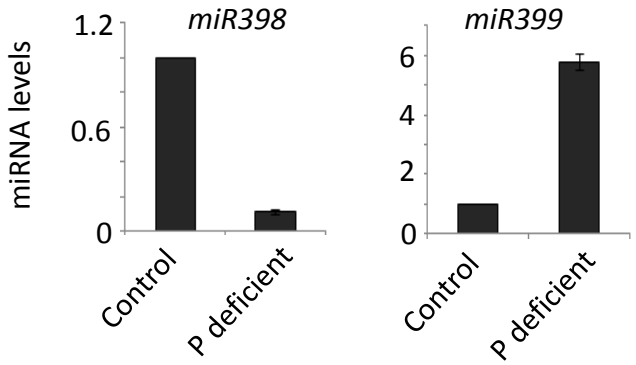

[A]
Col-0
mutant #1
nrpe1-11
5TE27040
BsmF I
Control
5TE27090
HpyCH4 IV
Control
5TE16500
BstB I
Control
1TE40810
Hae III
Control
3TE61900
HpyCH4 IV
Control
[B]
Col-0
mutant #2
mutant #3
nrpd1-3
Relative methylation levels
1
0.8
0.6
0.4
0.2
0
40810
SN1
30795

Figure S2


miRNA levels
1.2
0.6
0
miR398
Control
P deficient
6
4
2
0
miR399
Control
P deficient

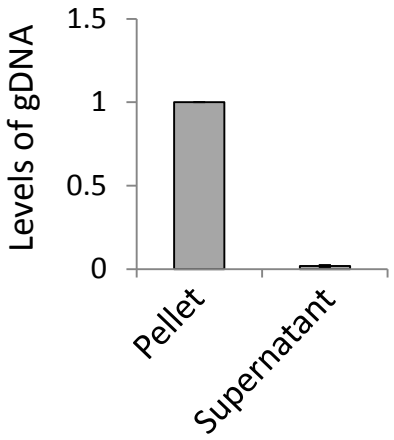



Figure S3