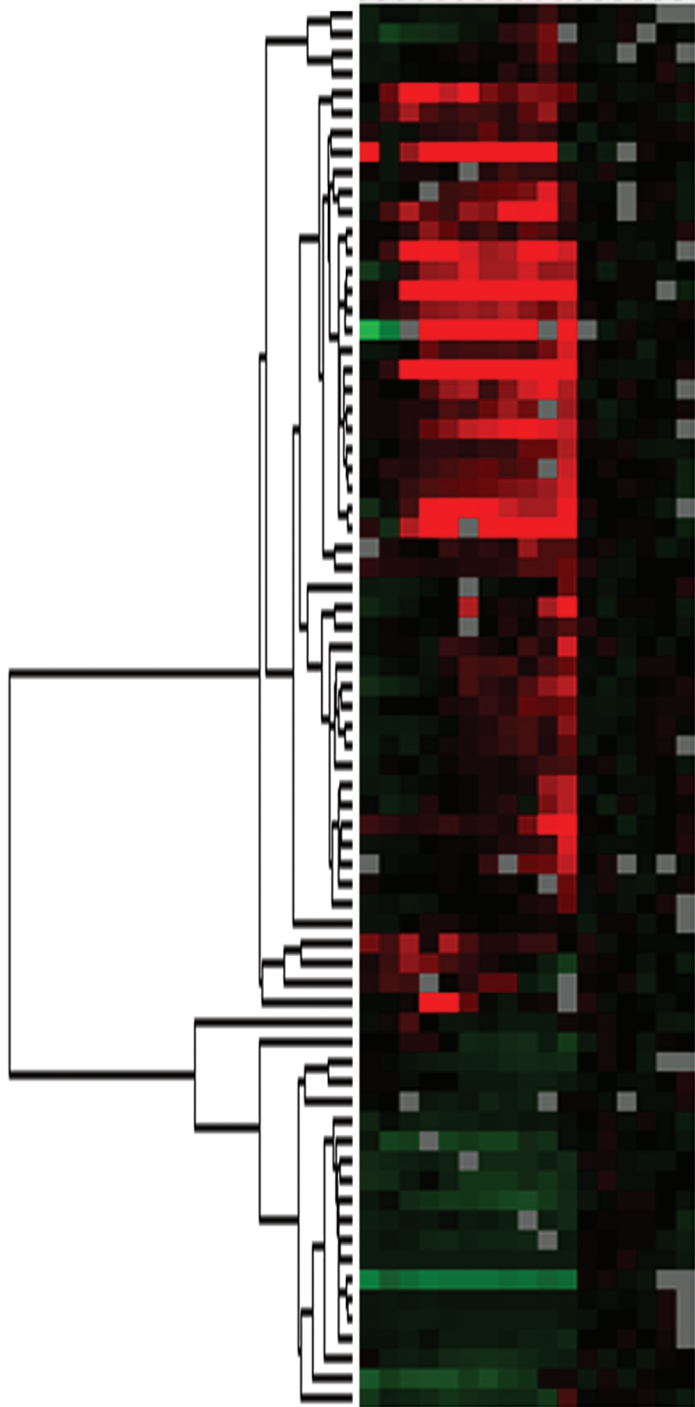


MEJA 0
MEJA 1
MEJA 2
MEJA 4
MEJA 6
MEJA 8
MEJA 10
MEJA 12
MEJA 16
MEJA 24
MEJA 48
DMSO 0
DMSO 4
DMSO 8
DMSO 12
DMSO 24
DMSO 48



T211-BT2M3-3-No hit
C112-BC1M4-2=putative protein At3g11810 [Arabidopsis thaliana]
T131-BT12M3-6-No hit
C408-BC4M2-1=beta-glucan binding protein
T434-BT42M4-5=putative SGPI monomeric G-protein
C209-BC24M2-3=N.tabacum DNA (recombination breakpoint between T-DNA and plant DNA)
T314-BT32M1-1=clonazepam factor-1 alpha
C168-BC14M1-4-No hit
C22-BC21M4-3=ribonucleotide reductase
C419-BC41M4-2=putative potassium transporter
C137-BC13M1-3=retrotransposon copia-like
T315-BT32M1-2=probable glutathione S-transferase PARA
T164-BT14M1-3=probable glutathione S-transferase PARA
T402-BT41M2-2=phenylalanine ammonia lyase
C449-BC43M1-1=vacuolar H(+)-ATPase
C228-BC24M4-3=arginine decarboxylase
T32-BC22M2-1-No hit
C359-BC33M4-2=putative protein kinase
T357-BT34M2-3=endo-1,3-1,4-beta-D-glucanase
T114-BT12M1-8-No hit
T9-BT21M2-1=putative adenosine kinase
T345-BT33M4-6-No hit
T351-BT34M1-3-No hit
T362-BT34M3-2-No hit
T358-BT34M2-4-No hit
T305-BT31M2-1=arabinoxylan-protein
C18-BC21M4-1=RNA-binding-like protein
C364-BC34M1-1=heme oxygenase
C146-BC13M4-2-No hit
C207-BC24M2-1=N.tabacum DNA (recombination breakpoint between T-DNA and plant DNA)
MT112-M3BT1-3-No hit
C304-BC31M1-4-F1-ATPase alpha subunit
C333-BC34M4-3=dihydrofolinamide dehydrogenase
C358-BC33M4-1=putative potassium transporter
C430-BC42M2-1=vacuolar H(+)-ATPase subunit-like protein
C425-BC42M1-2=vacuolar H(+)-ATPase subunit-like protein
T421-BT43M3-3=high-affinity nitrate transporter
C445-BC42M4-5=putative inorganic pyrophosphatase
C410-BC41M2-3=40S ribosomal protein S3a
C331-BC34M4-1=putative protein At3g62270 [Arabidopsis thaliana]
C338-BC33M3-1=putative protein P0401G10.6 [Orzya sativa]
C328-BC34M3-1=putative protein P0401G10.6 [Orzya sativa]
C134-BC12M4-5-No hit
C485-BC44M4-4=putative vacuolar ATP synthase proteolipid subunit
T327-BT33M2-1=expansin
C443-BC42M4-3=potassium transporter
T113-BT12M1-7=probable glutathione S-transferase PARA
MT407-M4BT4-1=probable glutathione S-transferase PARA
C206-BC24M1-2=N.tabacum DNA (recombination breakpoint between T-DNA and plant DNA)
C102-BC11M1-2=mitochondrial ribosomal protein
MT401-M1BT4-1=probable glutathione S-transferase PARA
T28-BT22M1-4-No hit
C350-BC32M3-1=putative protein At3g52110 [Arabidopsis thaliana]
T226-BT24M4-1-No hit
T18-BT21M4-1=pathogenesis-related protein
C201-BC23M2-1=putative protein At5g44670 [Arabidopsis thaliana]
C178-BC14M3-5=auxin-induced protein IAA
MT207-M2BT2-2=ooly(A)-binding protein
T337-BT33M3-3=putative protein At3g52110 [Arabidopsis thaliana]
MC402-M1BC4-2=putative beta-1,3-glucanase
C157-BC12M4-6=zinc-binding protein
C7-BC21M1-7-No hit
T331-BT33M2-5=RNA polymerase beta''
C216-BC24M3-3=N.tabacum DNA (recombination breakpoint between T-DNA and plant DNA)
T204-BT23M2-2-No hit
T153-BT13M3-3=hyoscyamine 6 beta-hydroxylase like protein
T109-BT12M1-3=small subunit ribosomal RNA
C323-BC34M2-2=amino acid transport protein AAT1
C340-BC33M3-3-No hit
C150-BC12M1-4=putative aminotransferase-like protein
T26-BT22M1-2-No hit