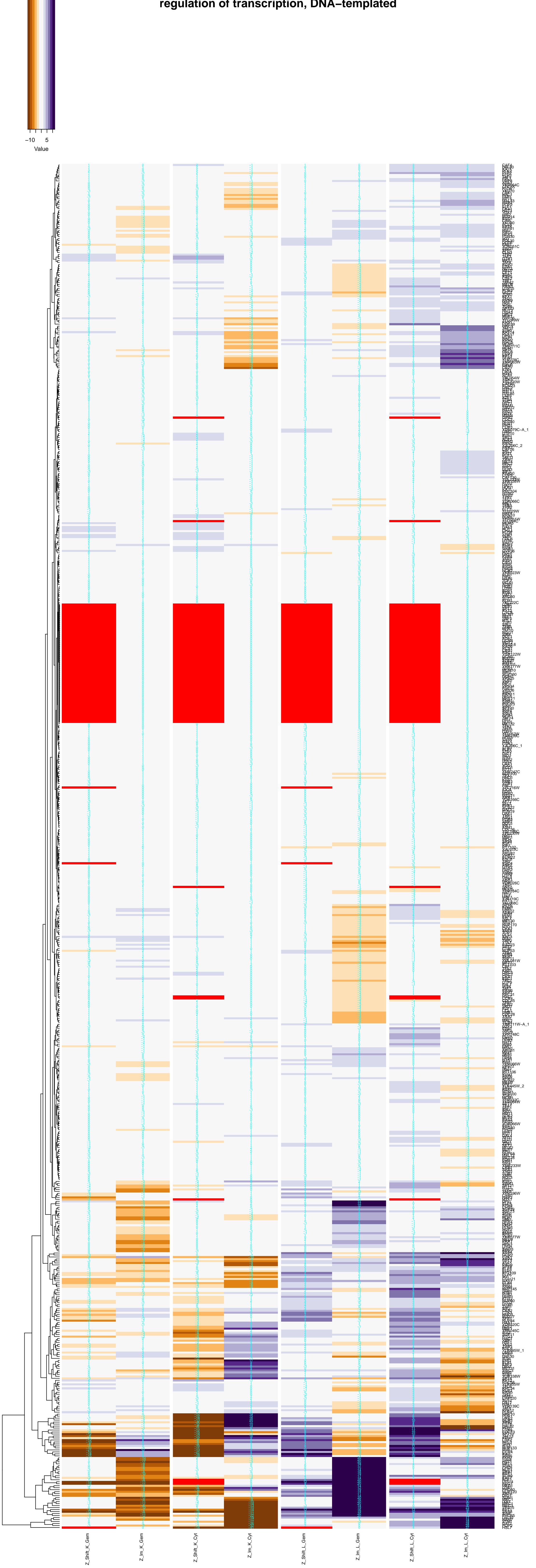
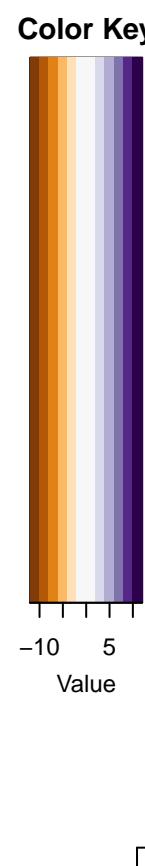


Color Key

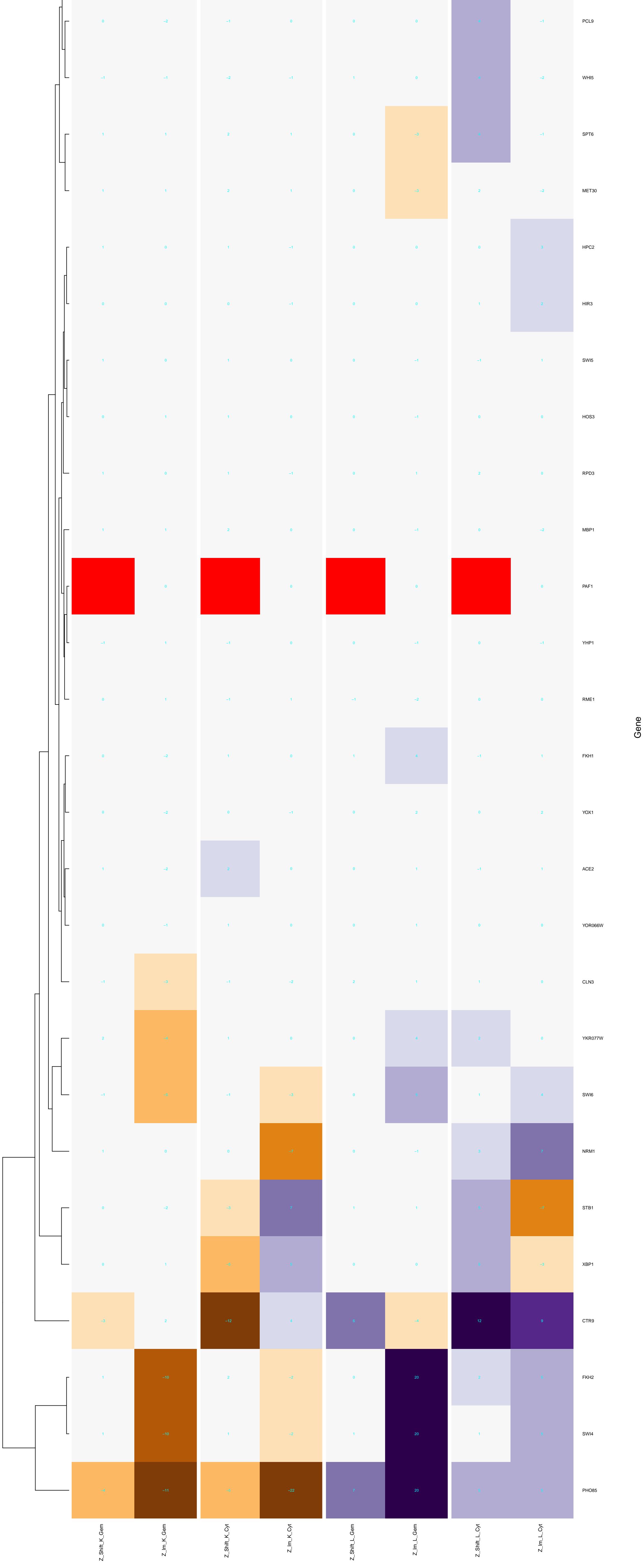


regulation of transcription, DNA-templated





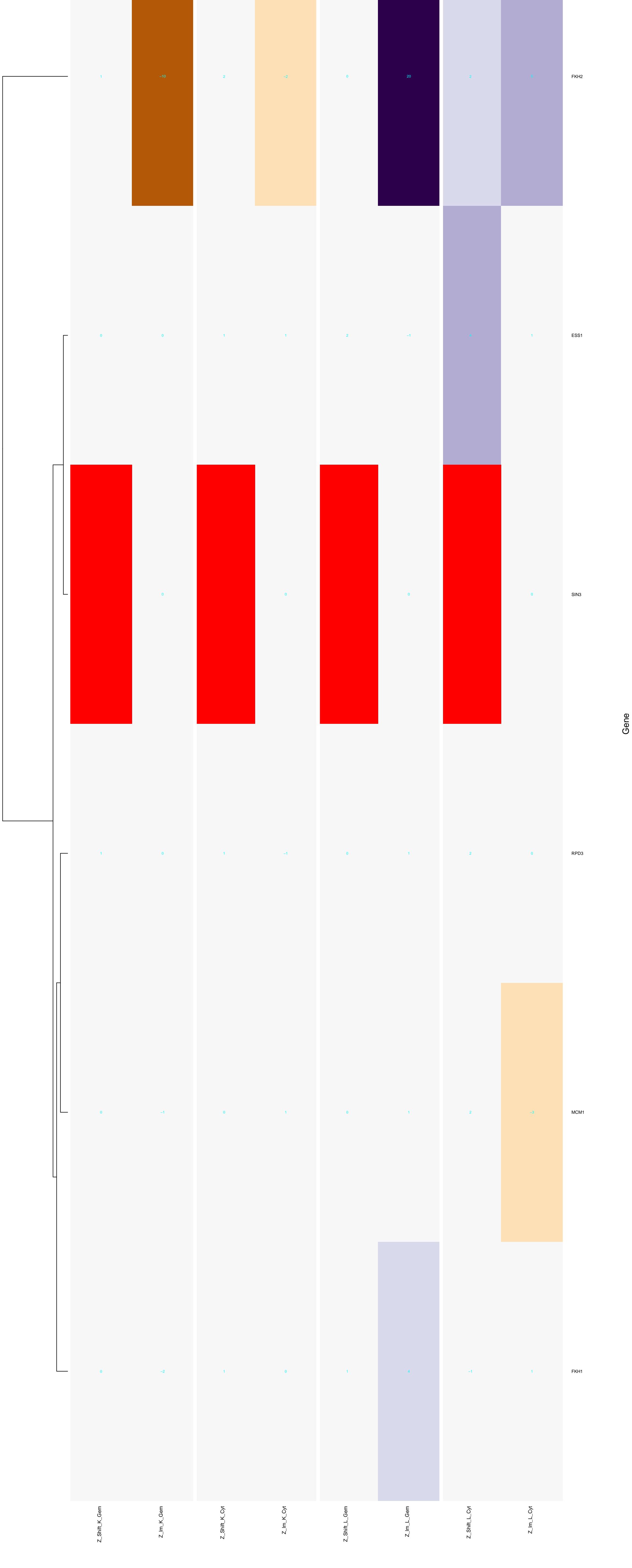
regulation of transcription involved in G1/S transition of mitotic cell cycle



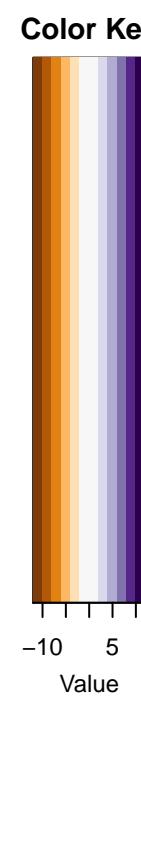
Color Key



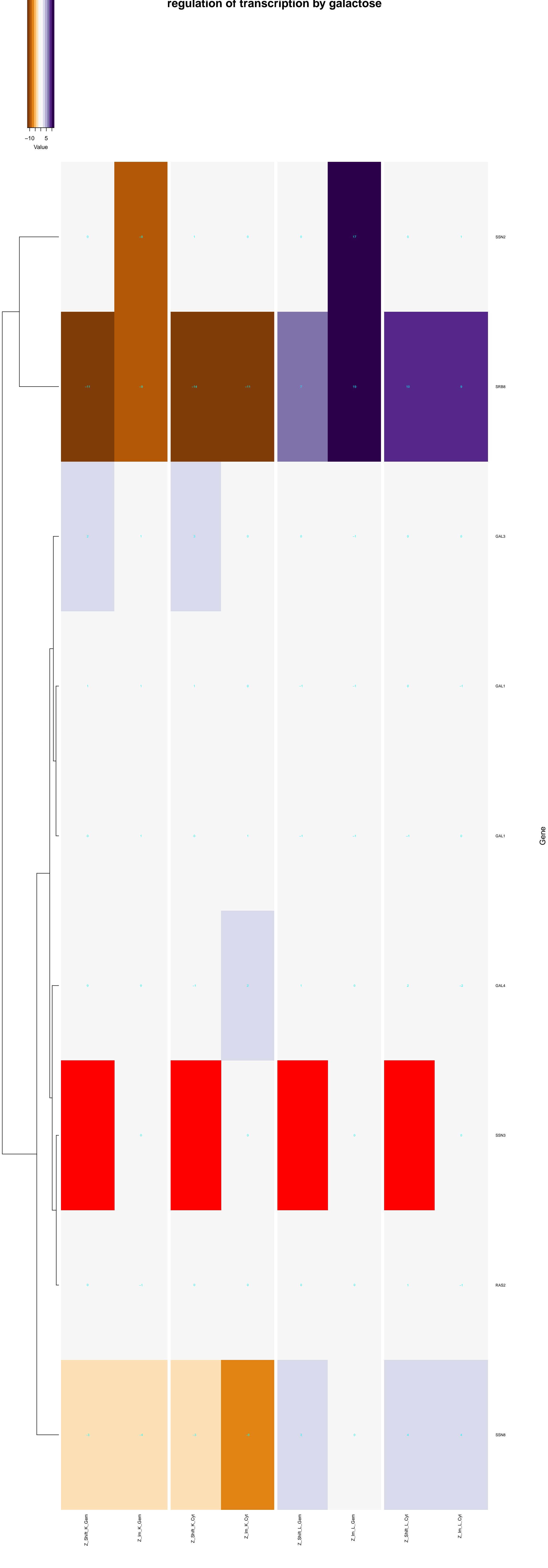
regulation of transcription involved in G2/M transition of mitotic cell cycle

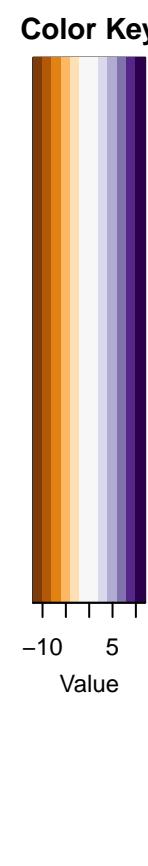


Color Key

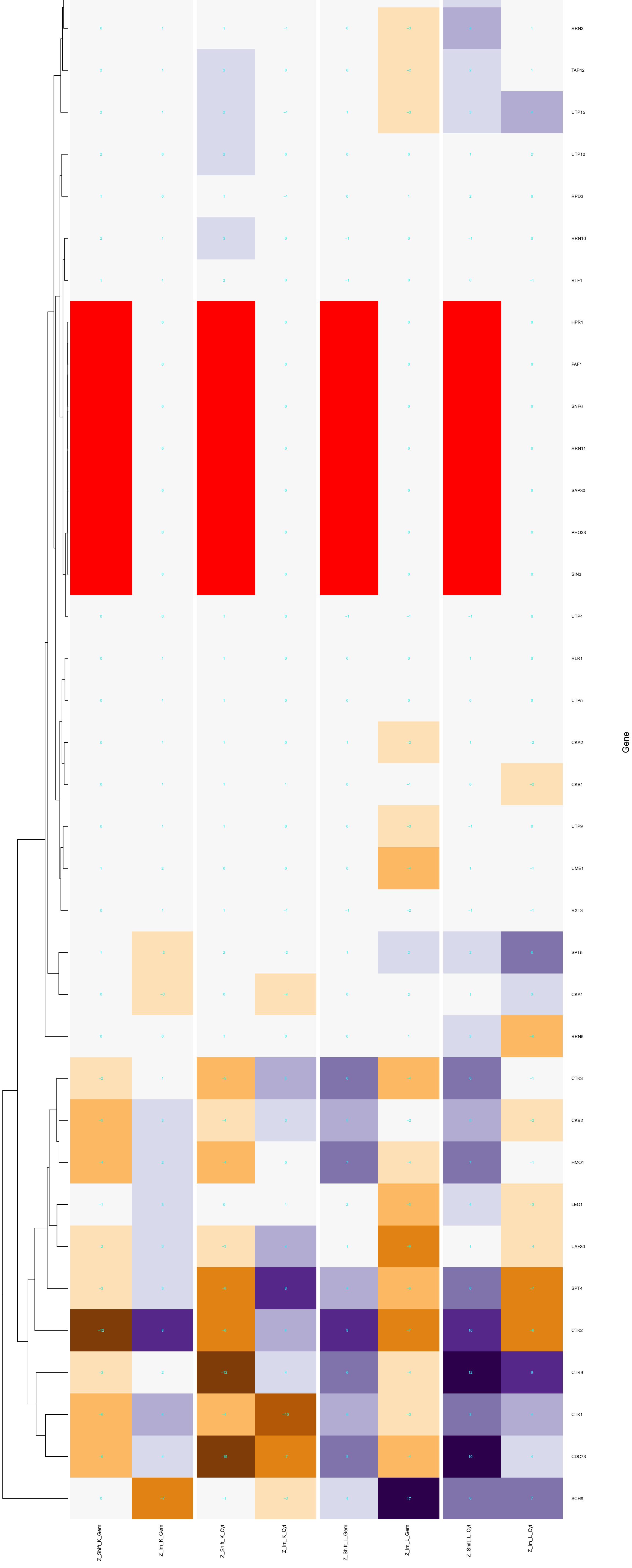


regulation of transcription by galactose





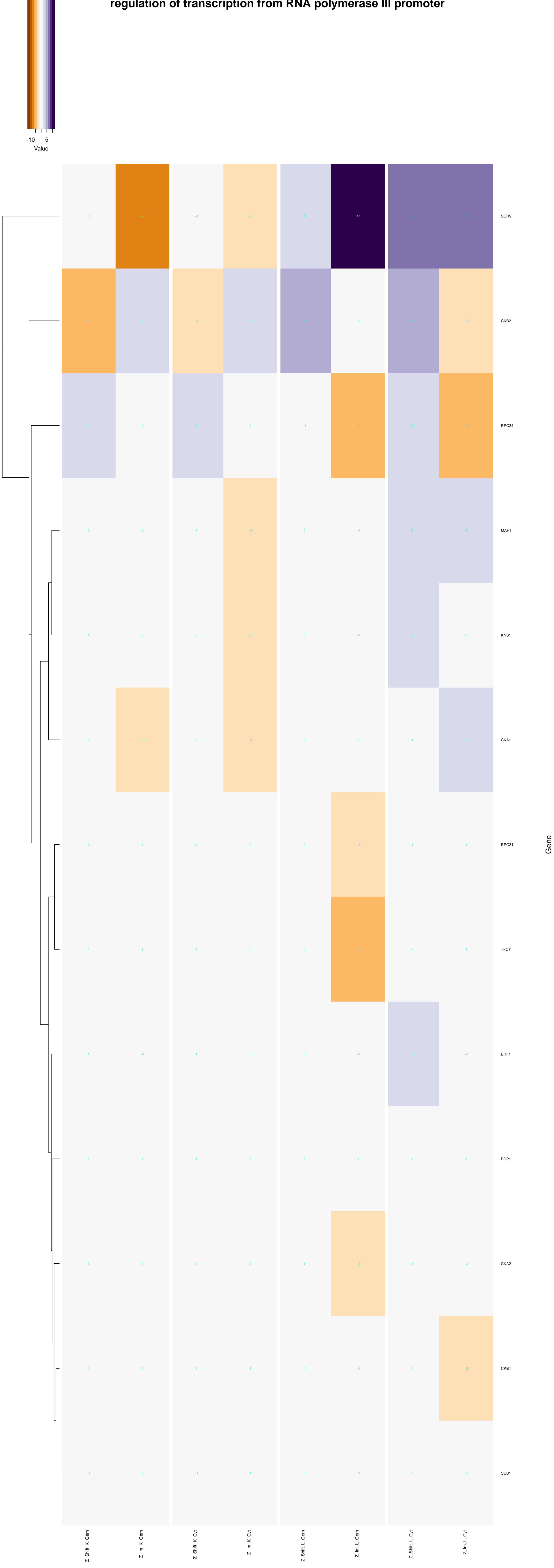
regulation of transcription from RNA polymerase I promoter



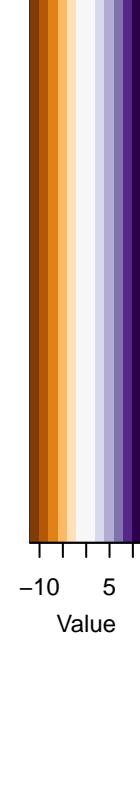
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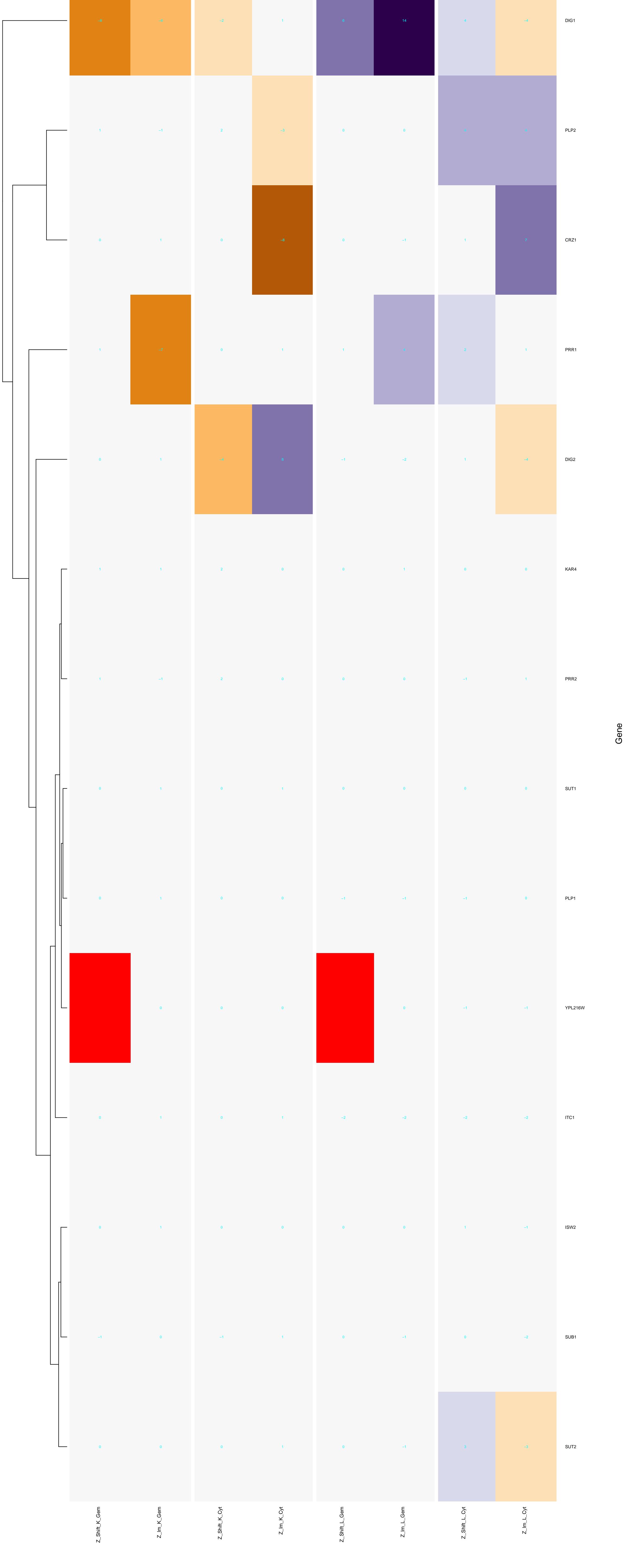
regulation of transcription from RNA polymerase III promoter



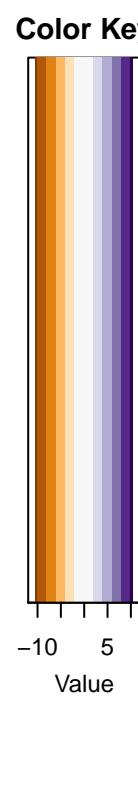
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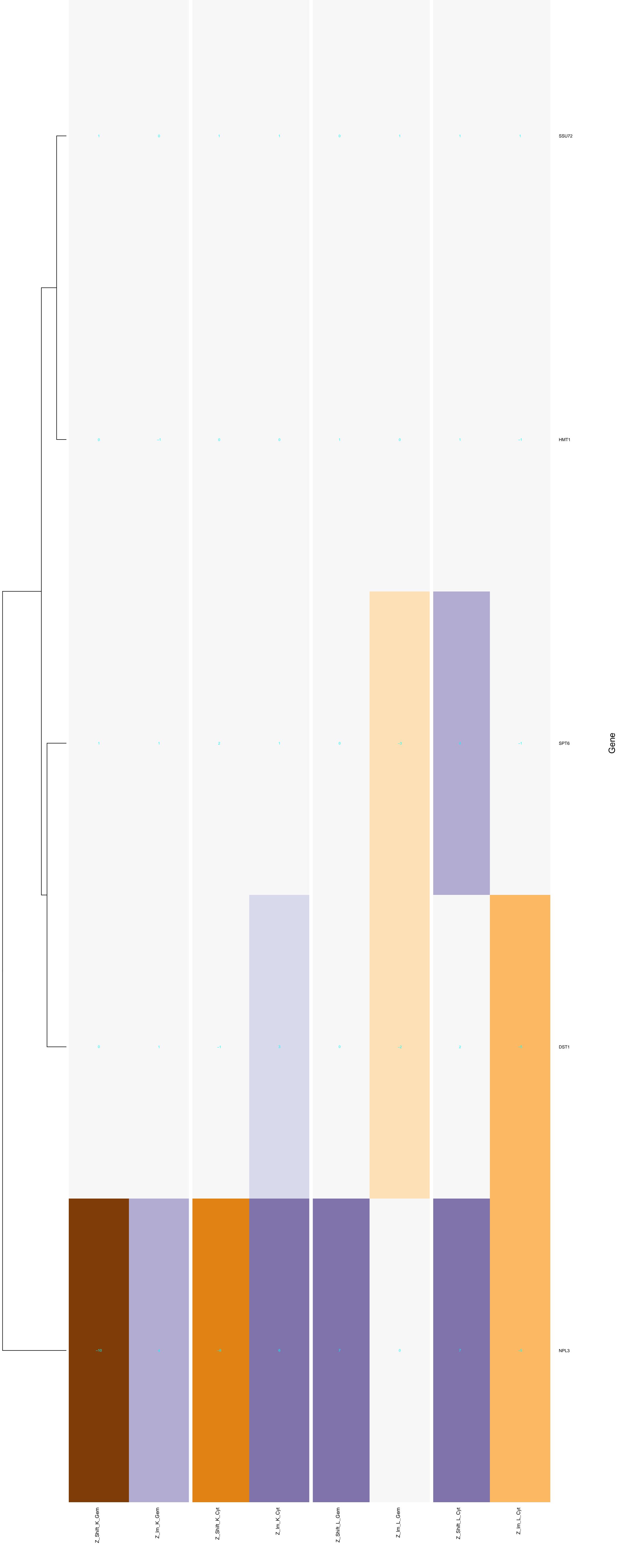
regulation of transcription by pheromones

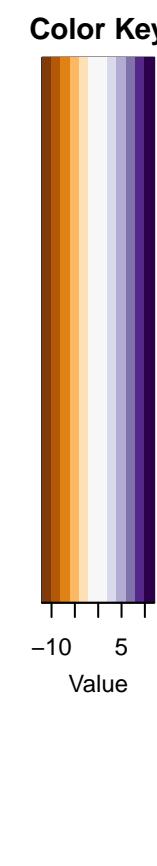


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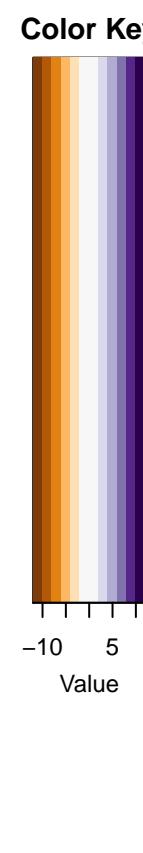
regulation of DNA-templated transcription, termination



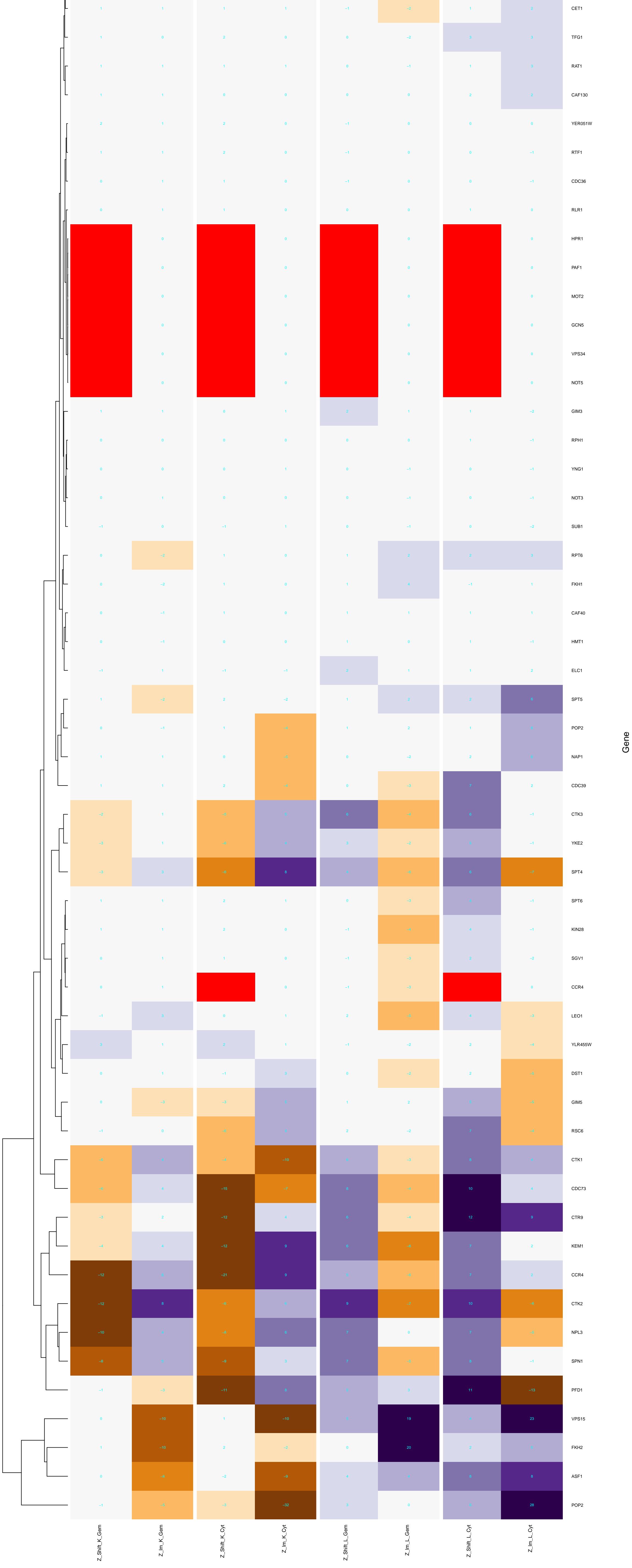


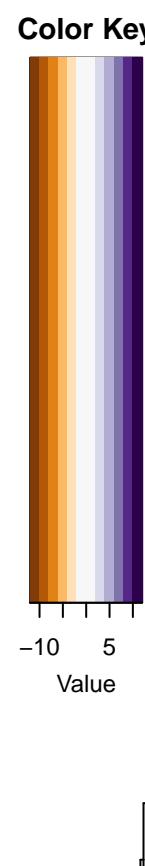
regulation of chromatin silencing



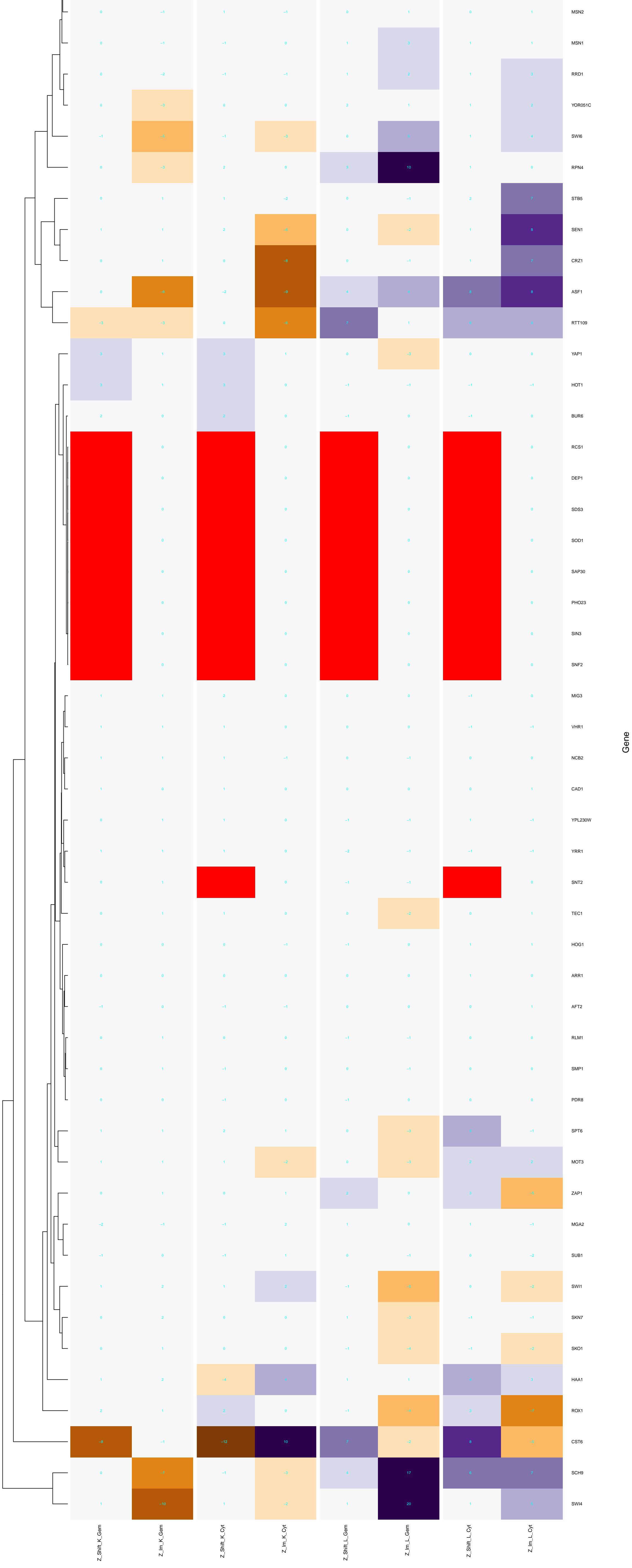


regulation of DNA-templated transcription, elongation



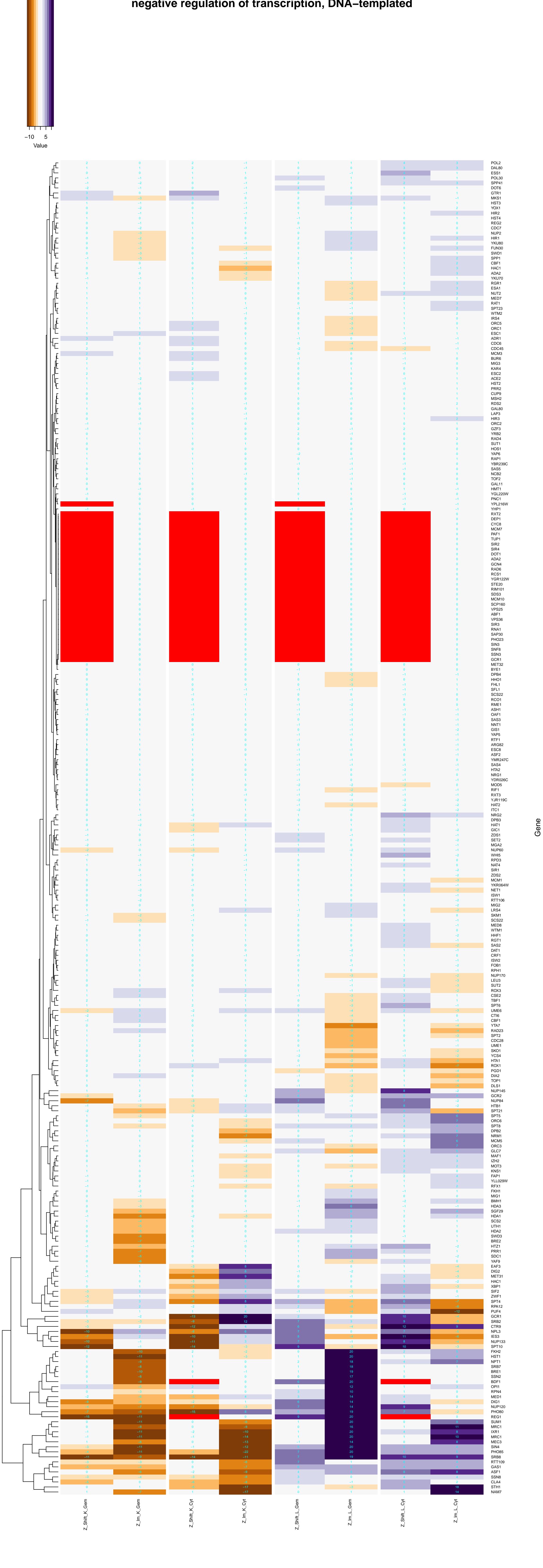


regulation of DNA-templated transcription in response to stress



Color Key

negative regulation of transcription, DNA-templated

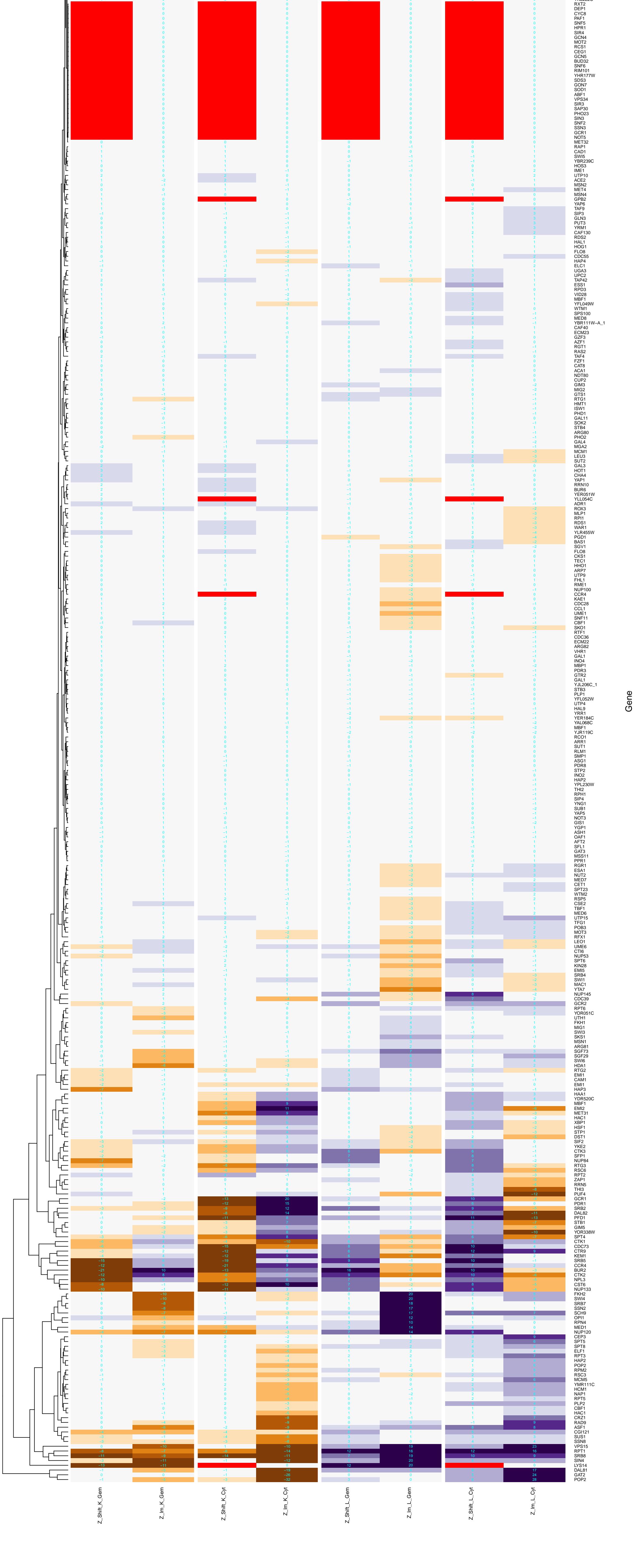


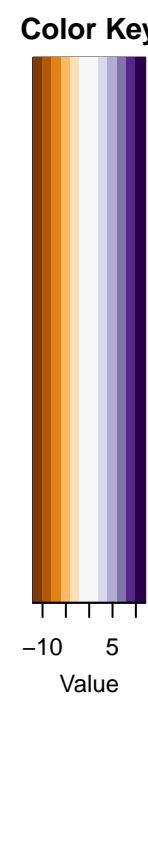
Color Key

positive regulation of transcription, DNA-templated

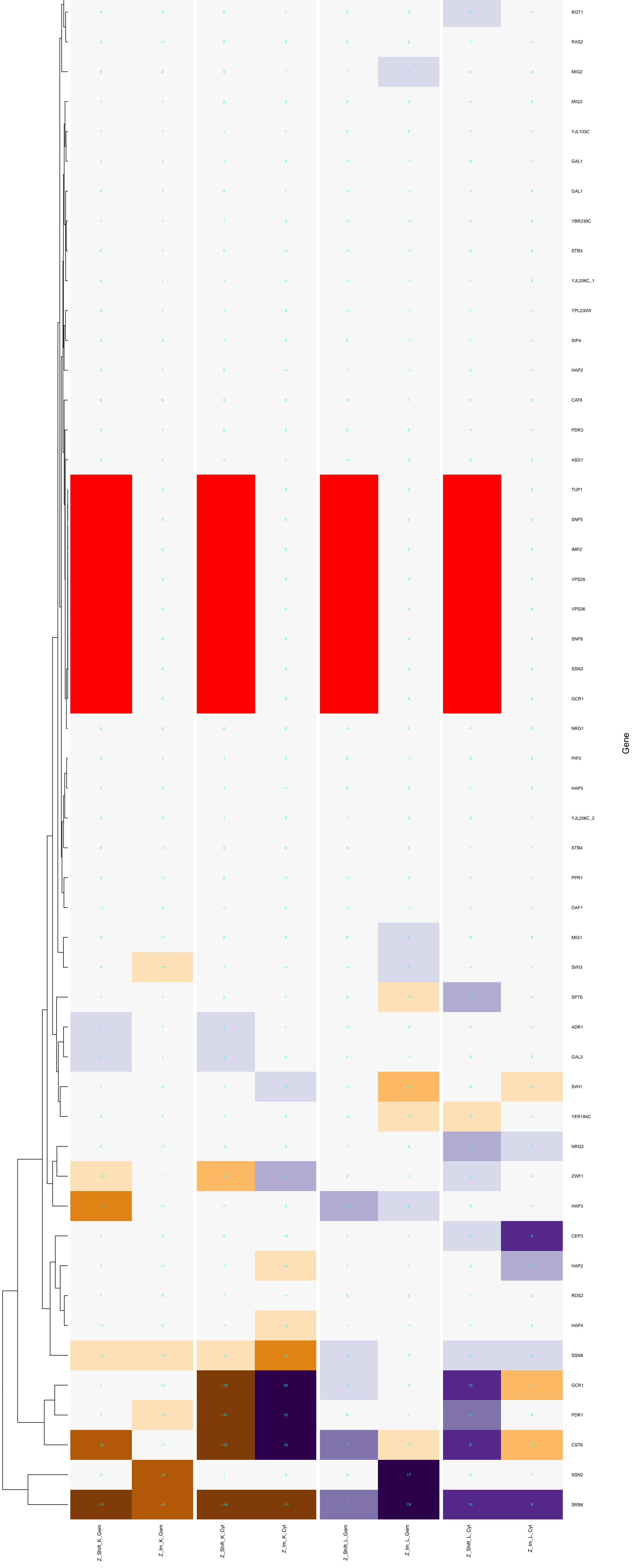


Value





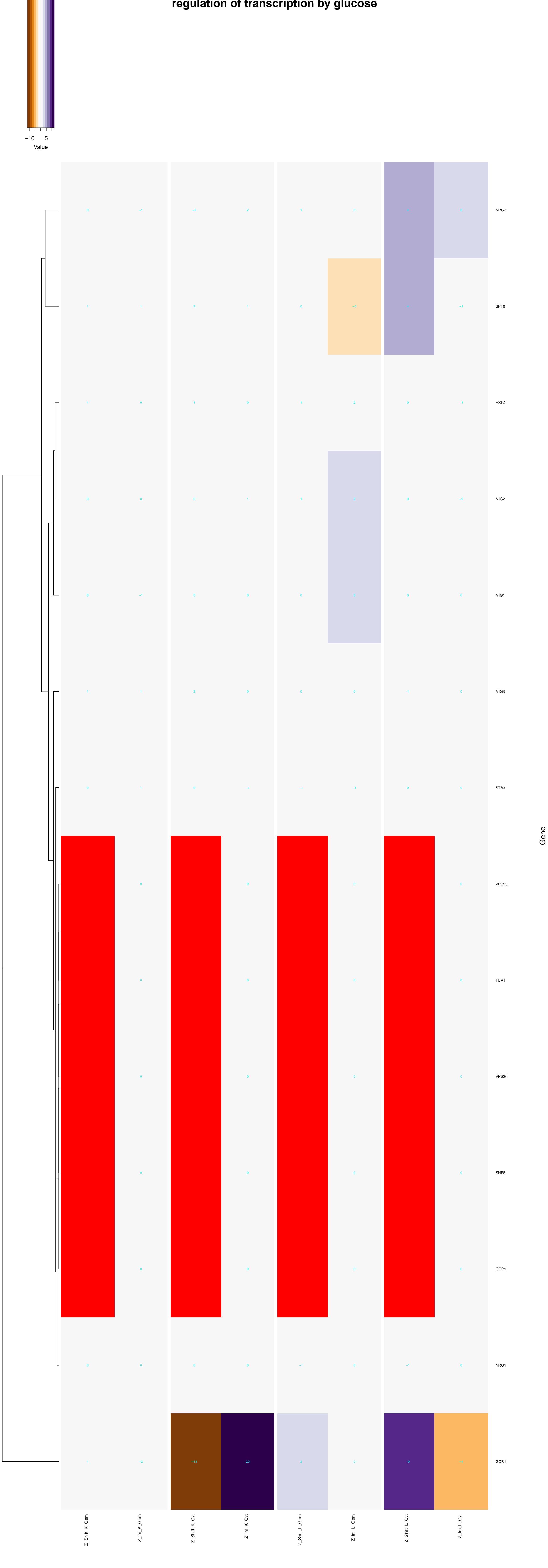
carbon catabolite regulation of transcription



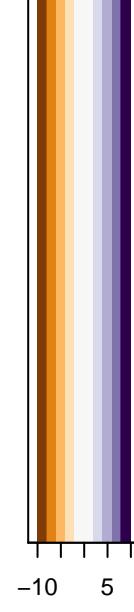
Color Key



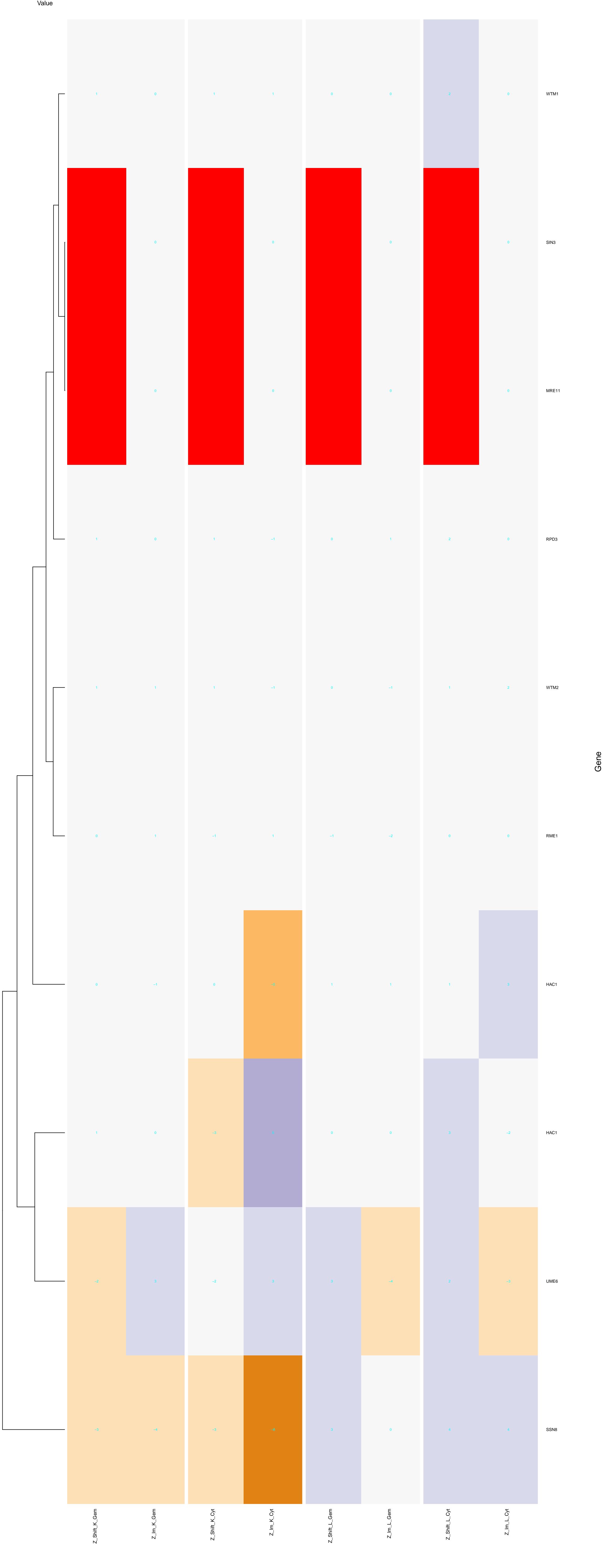
regulation of transcription by glucose



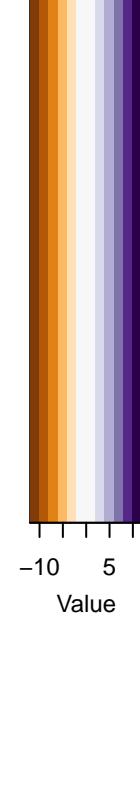
Color Key



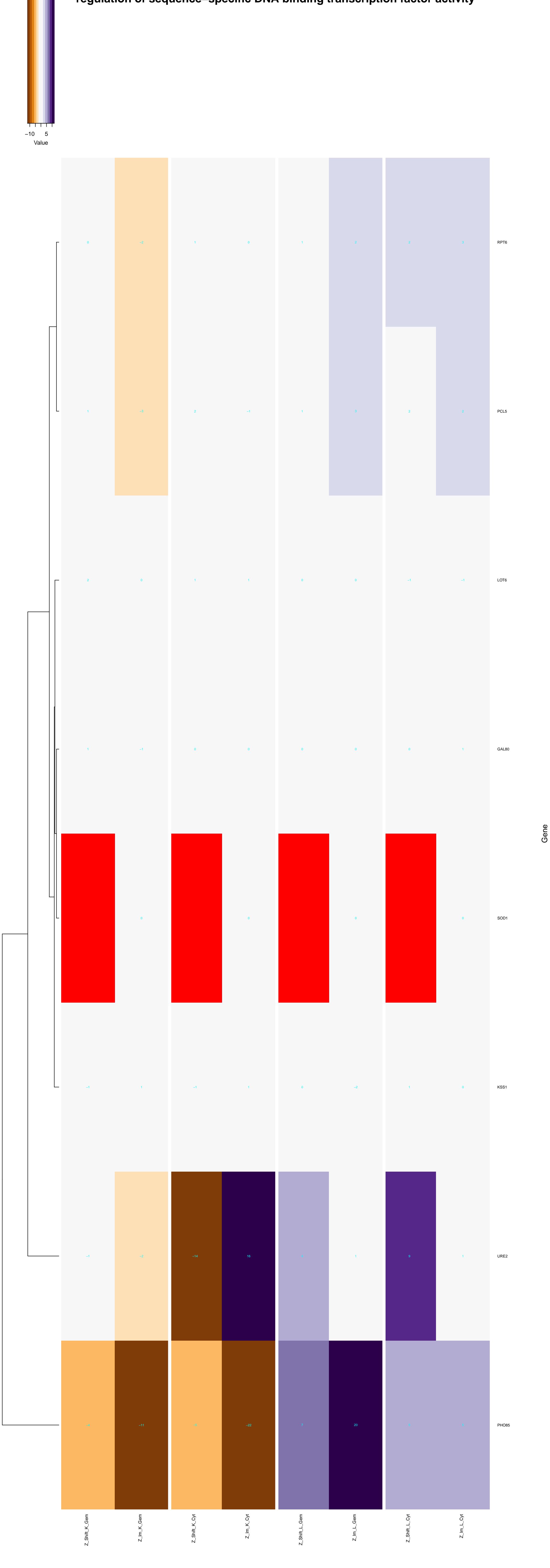
regulation of transcription involved in meiotic cell cycle



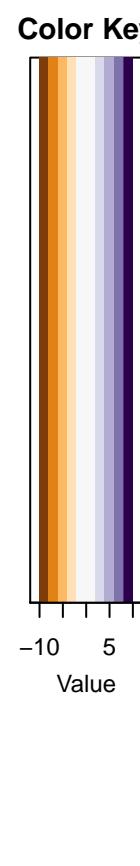
Color Key



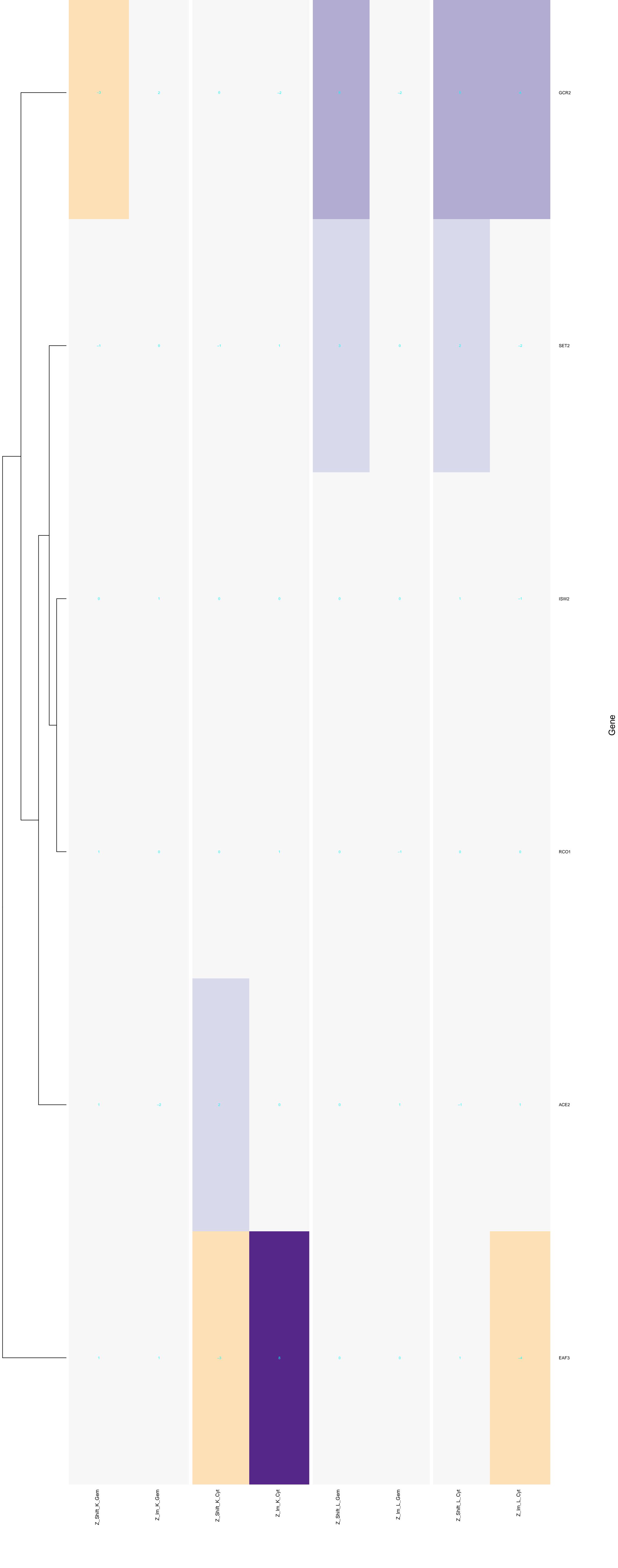
regulation of sequence-specific DNA binding transcription factor activity



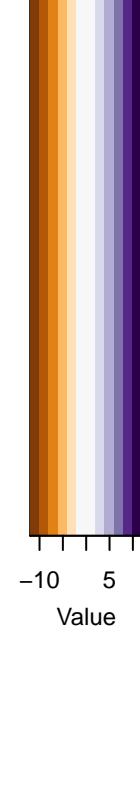
Color Key



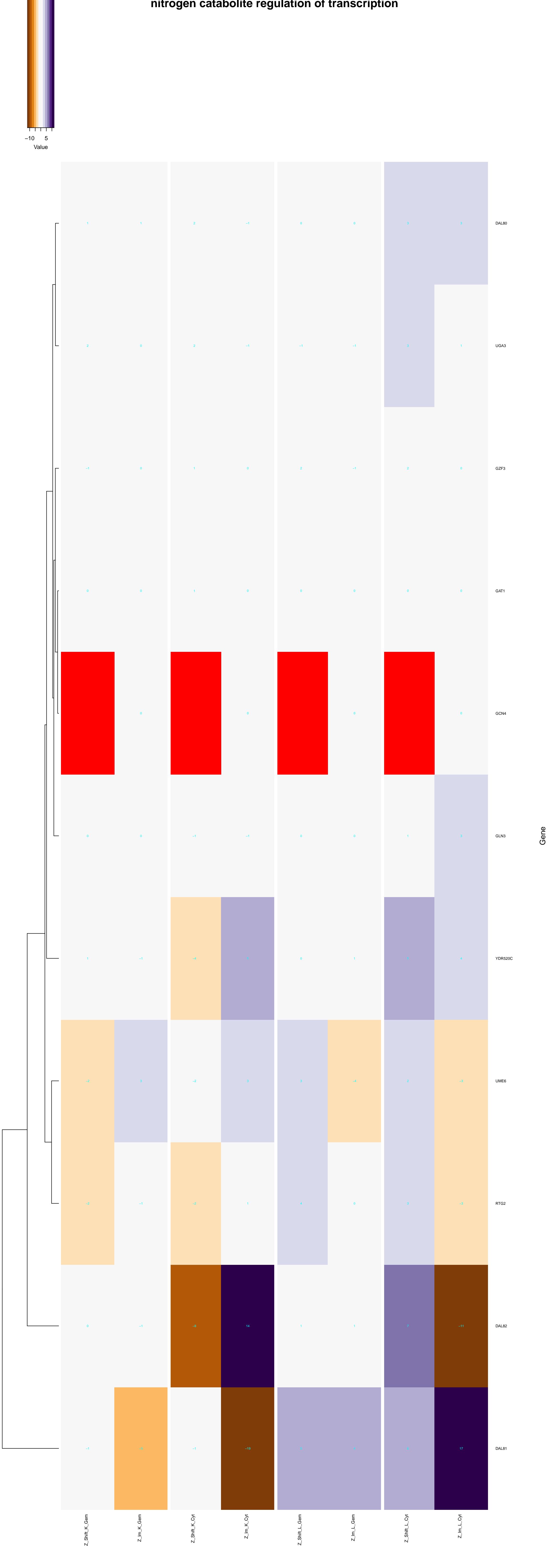
regulation of antisense RNA transcription



Color Key



nitrogen catabolite regulation of transcription



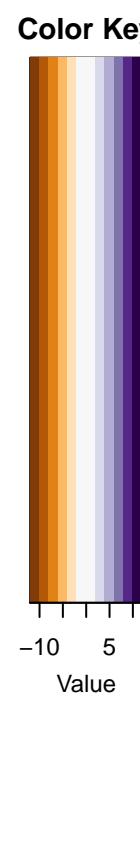
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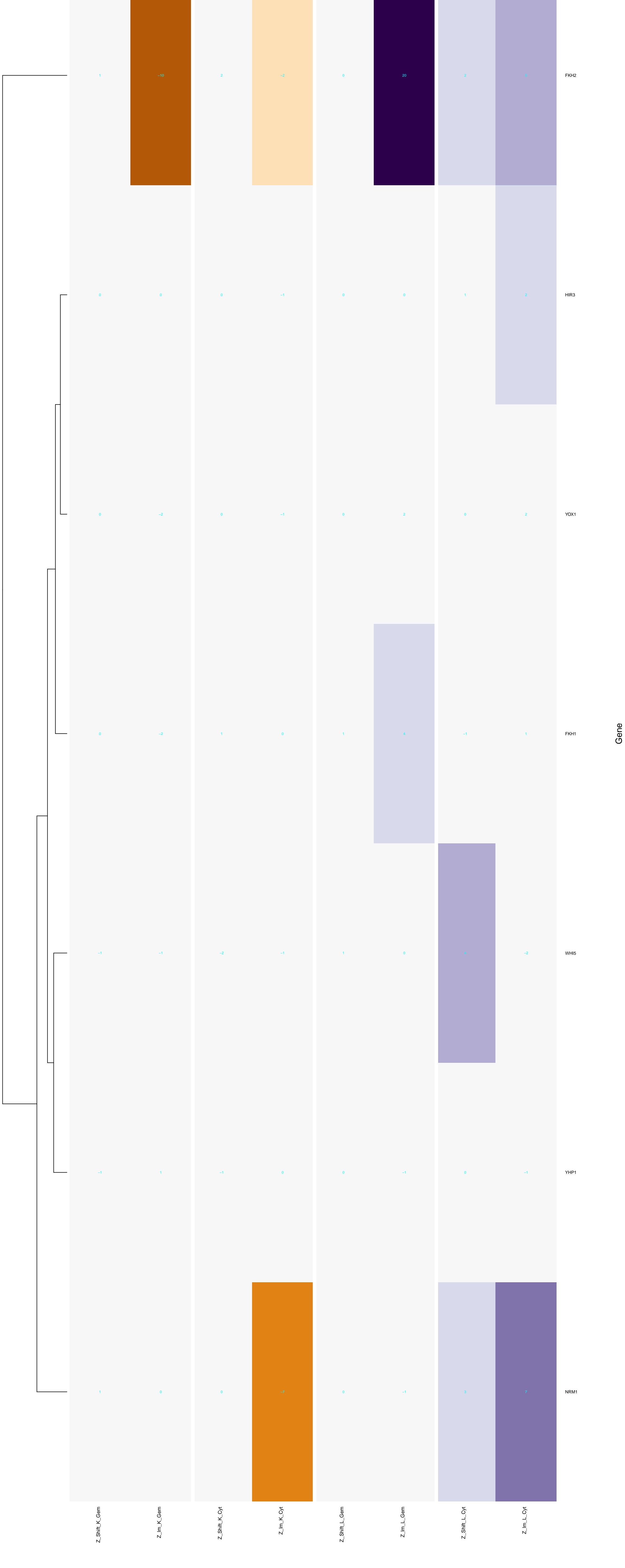
regulation of DNA-templated transcription, initiation



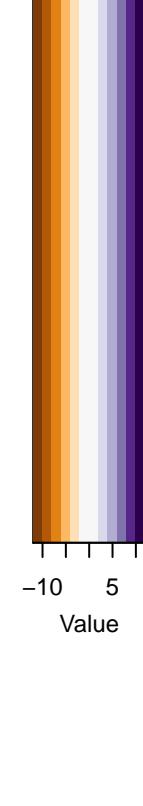
Color Key



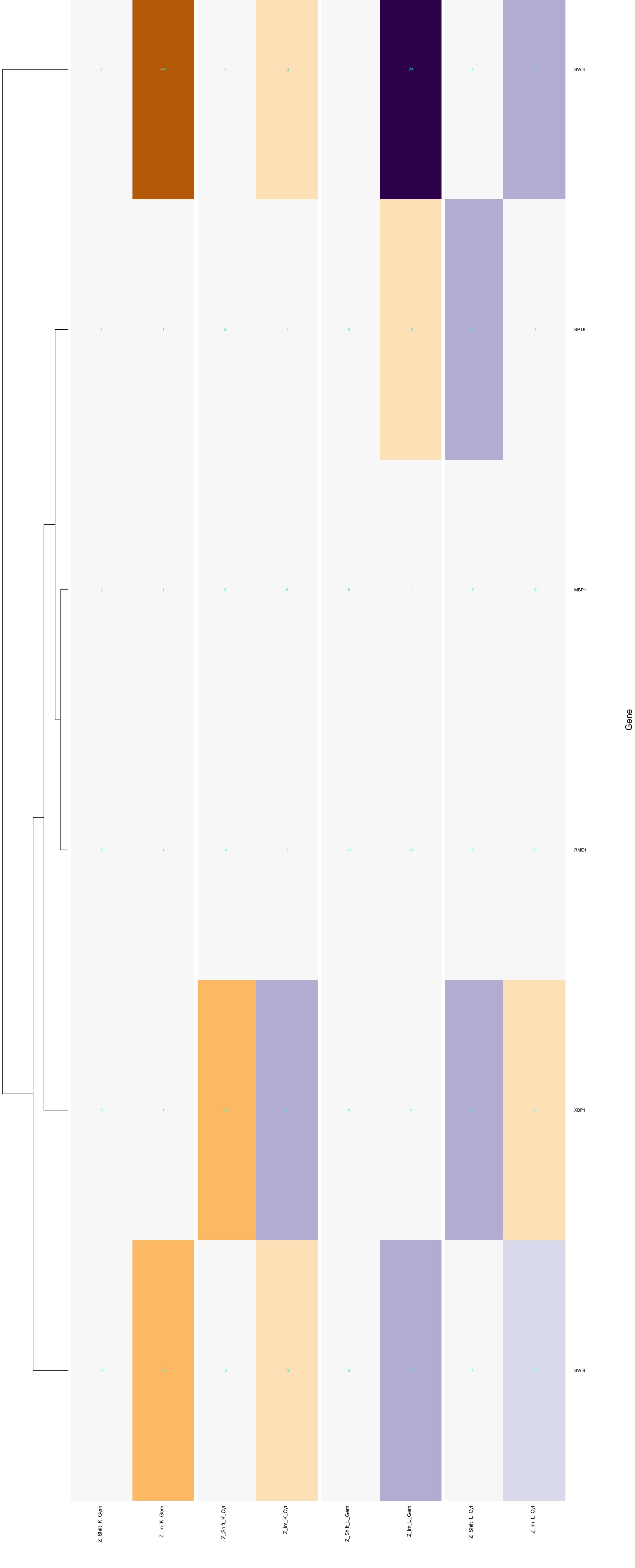
Negative regulation of transcription involved in G1/S transition of mitotic cell cycle



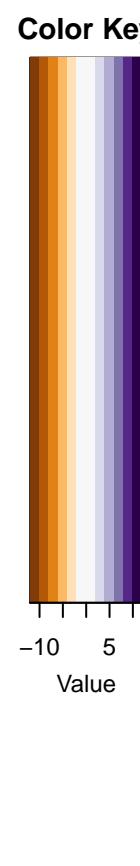
Color Key



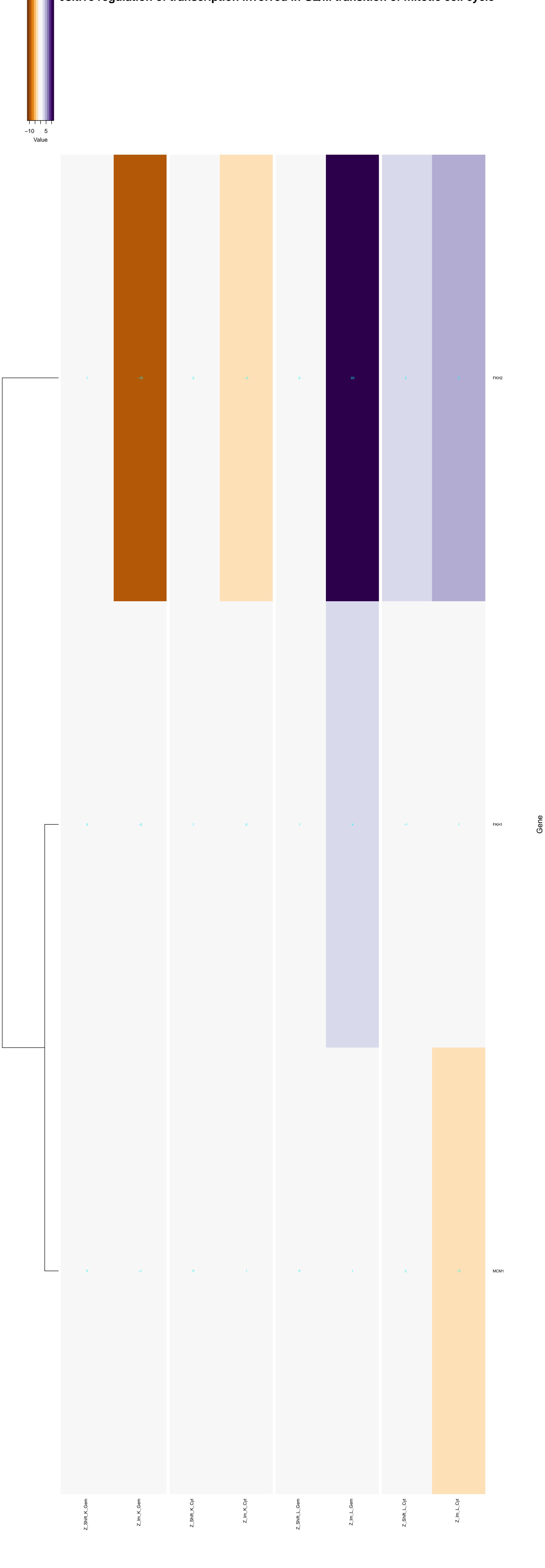
Positive regulation of transcription involved in G1/S transition of mitotic cell cycle



Color Key



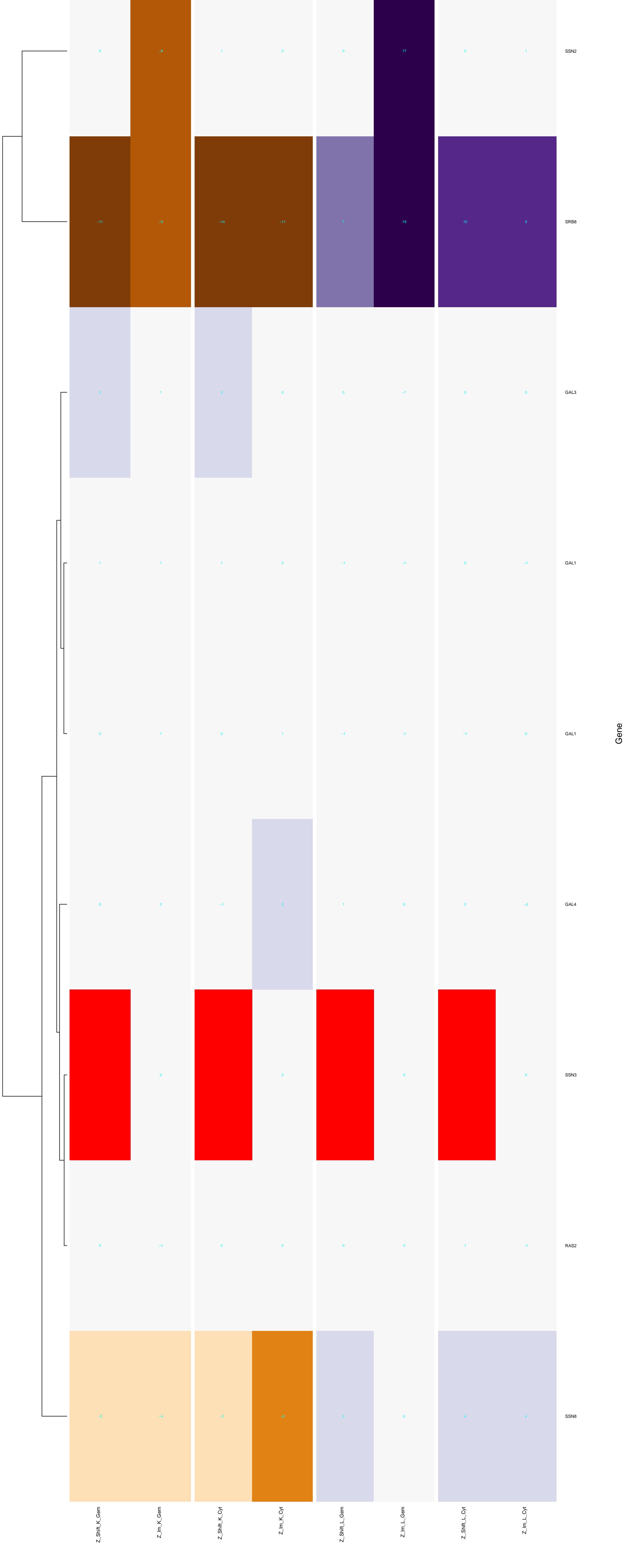
Positive regulation of transcription involved in G2/M transition of mitotic cell cycle



Color Key



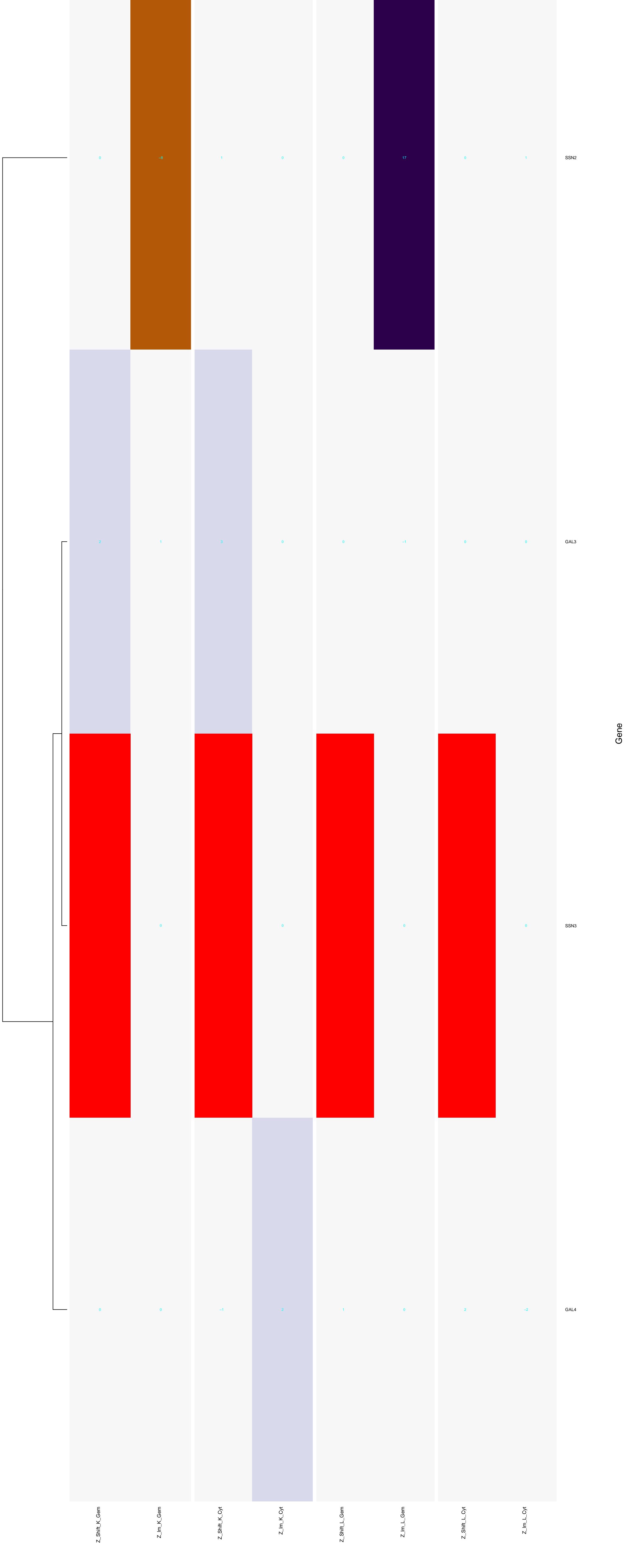
positive regulation of transcription by galactose



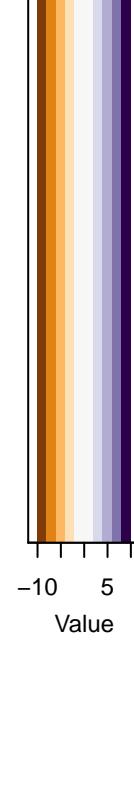
Color Key



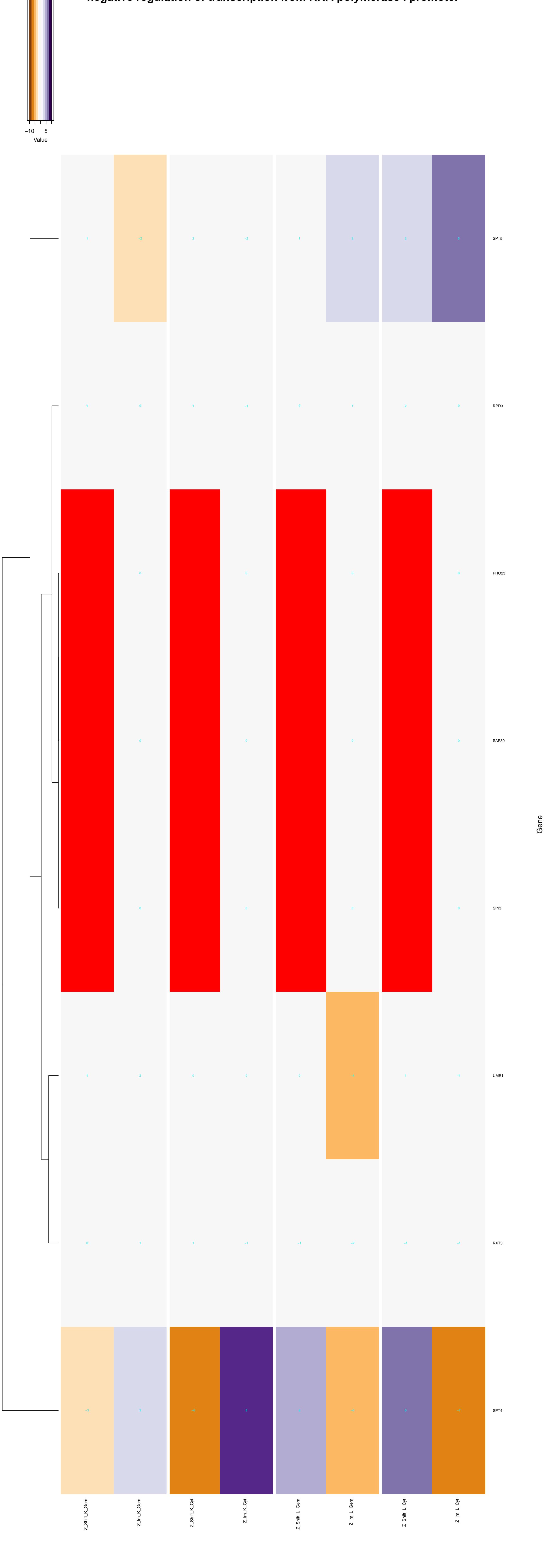
regulation of transcription from RNA polymerase II promoter by galactose



Color Key

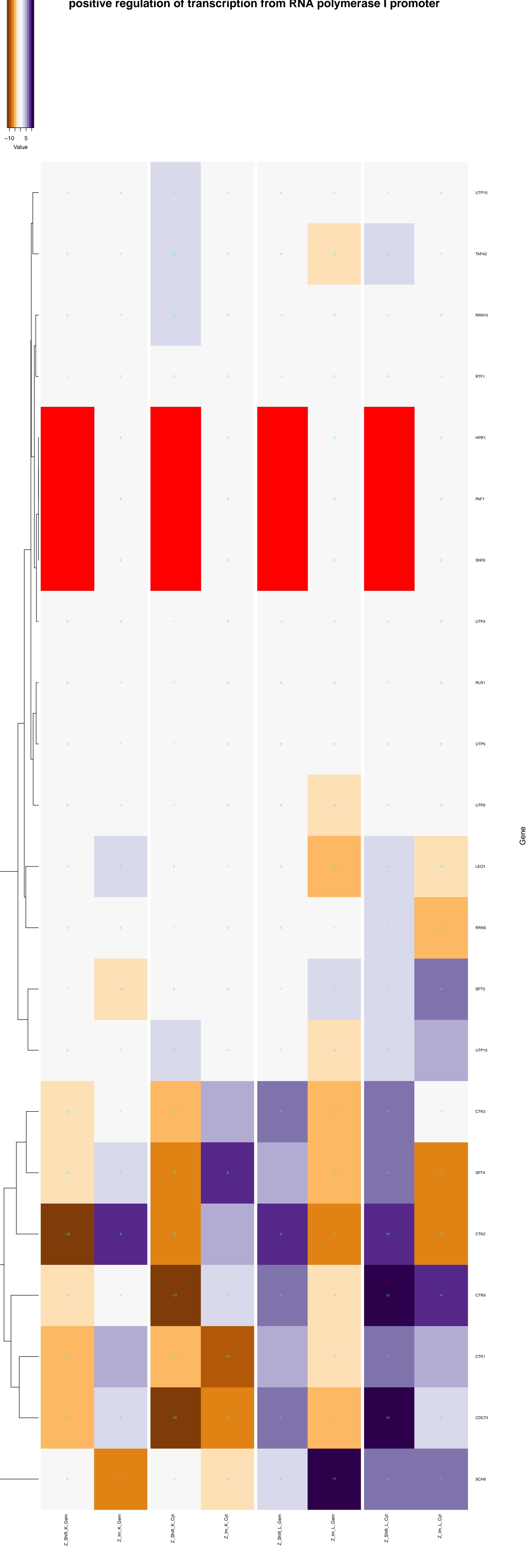


negative regulation of transcription from RNA polymerase I promoter

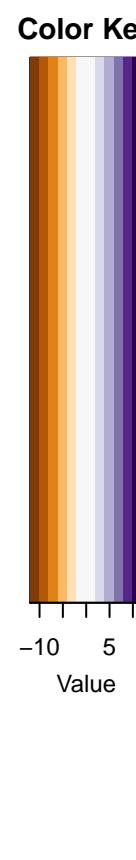


Color Key

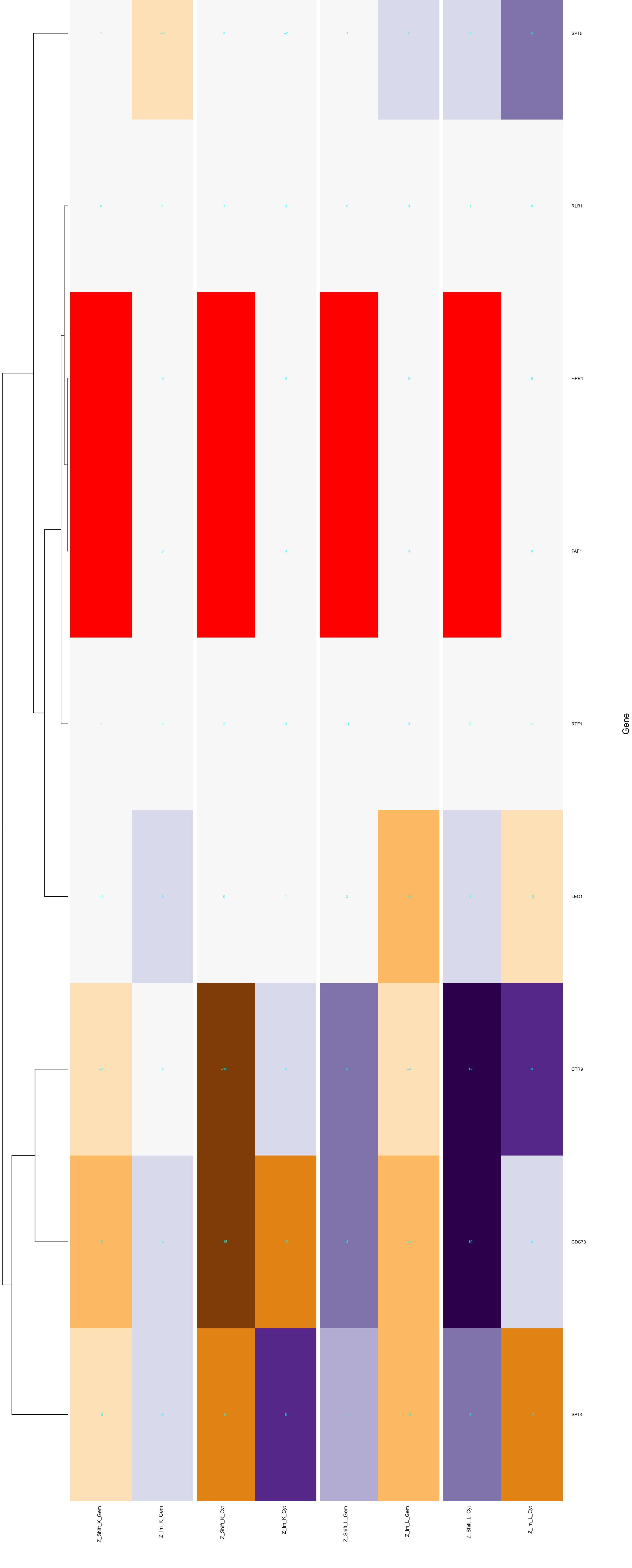
positive regulation of transcription from RNA polymerase I promoter



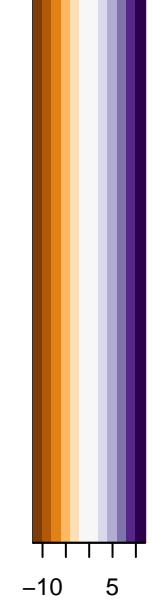
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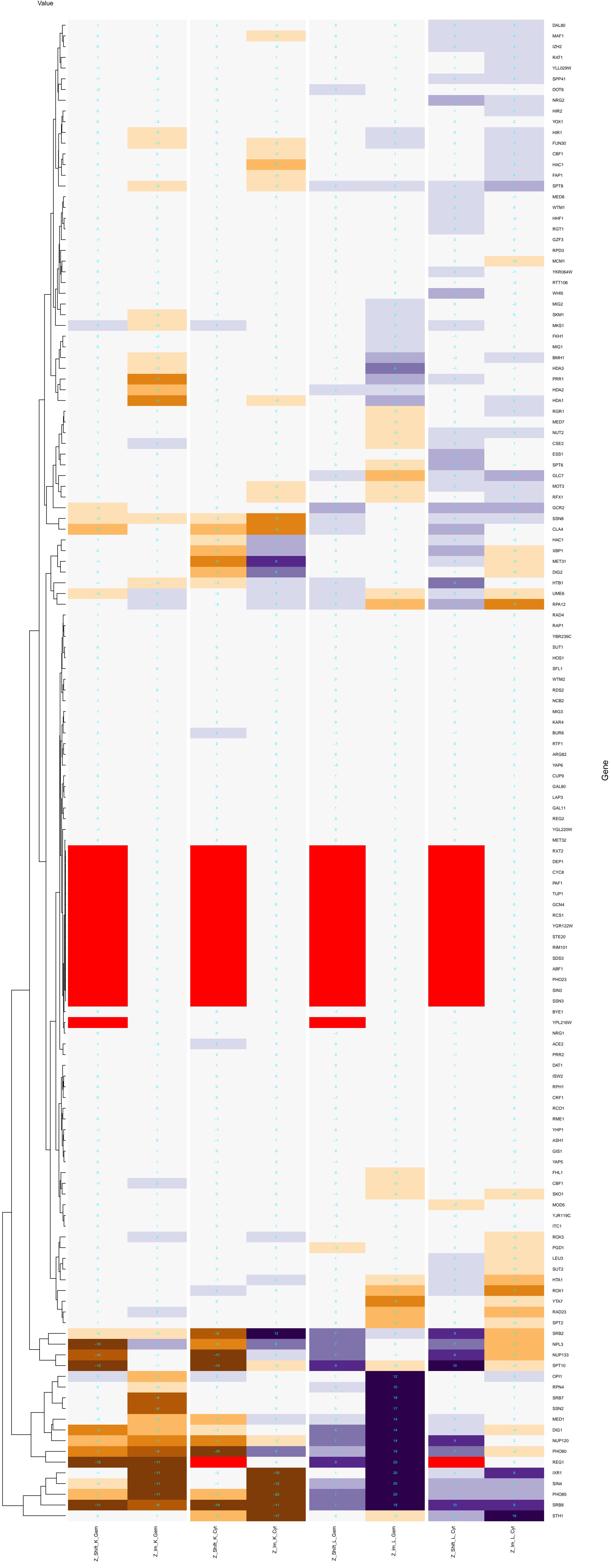
regulation of transcription elongation from RNA polymerase I promoter

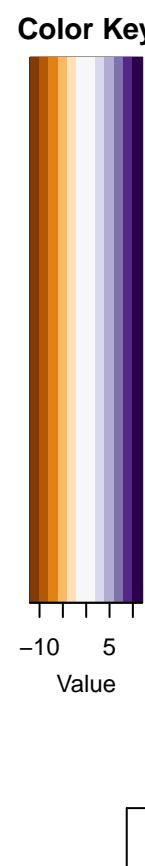


Color Key

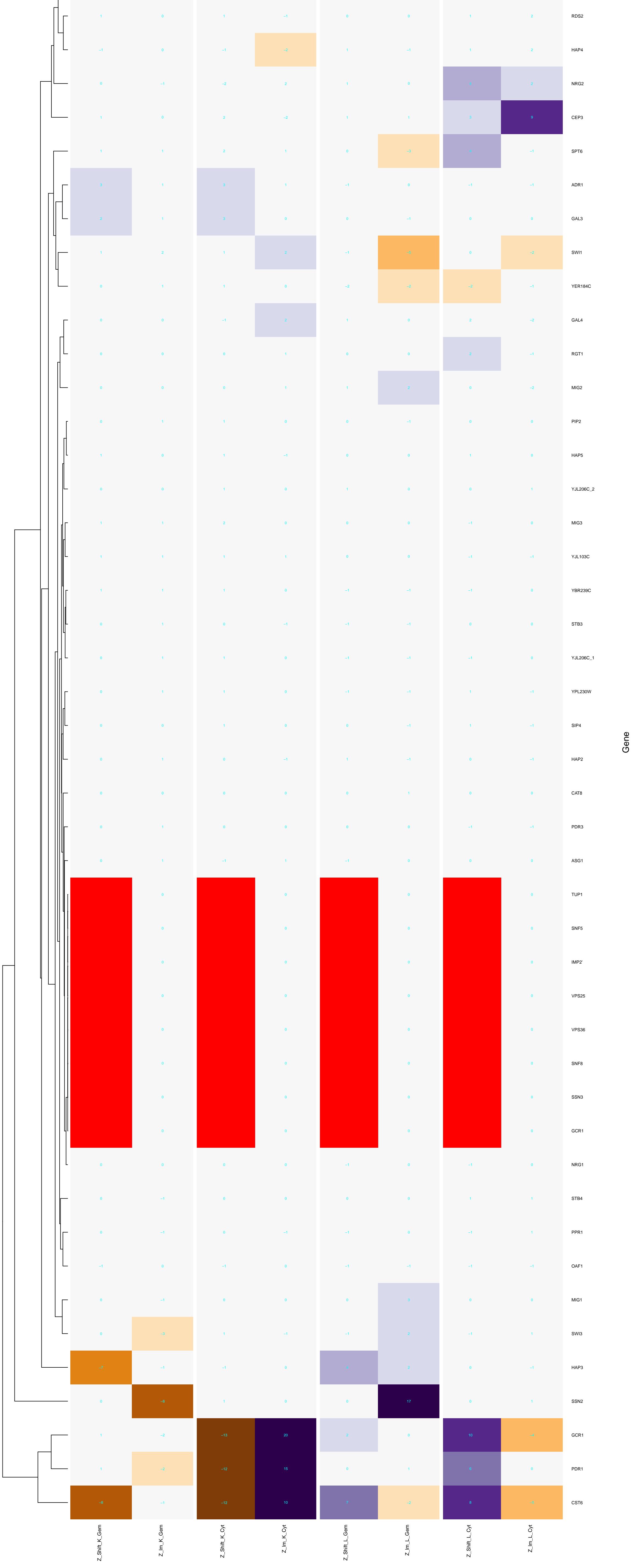


negative regulation of transcription from RNA polymerase II promoter





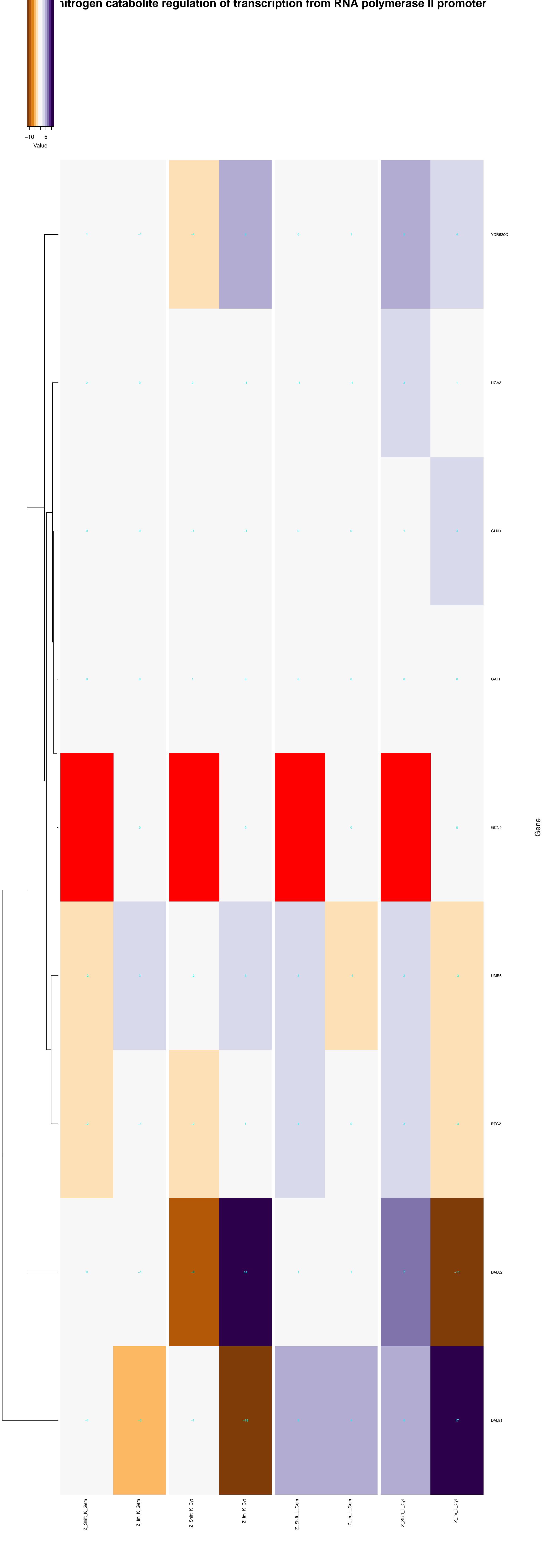
carbon catabolite regulation of transcription from RNA polymerase II promoter



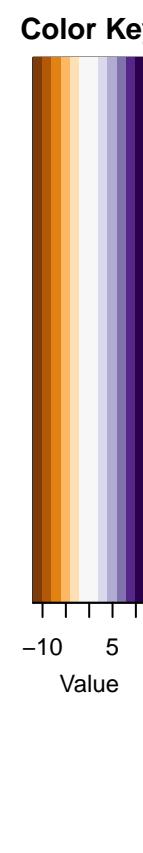
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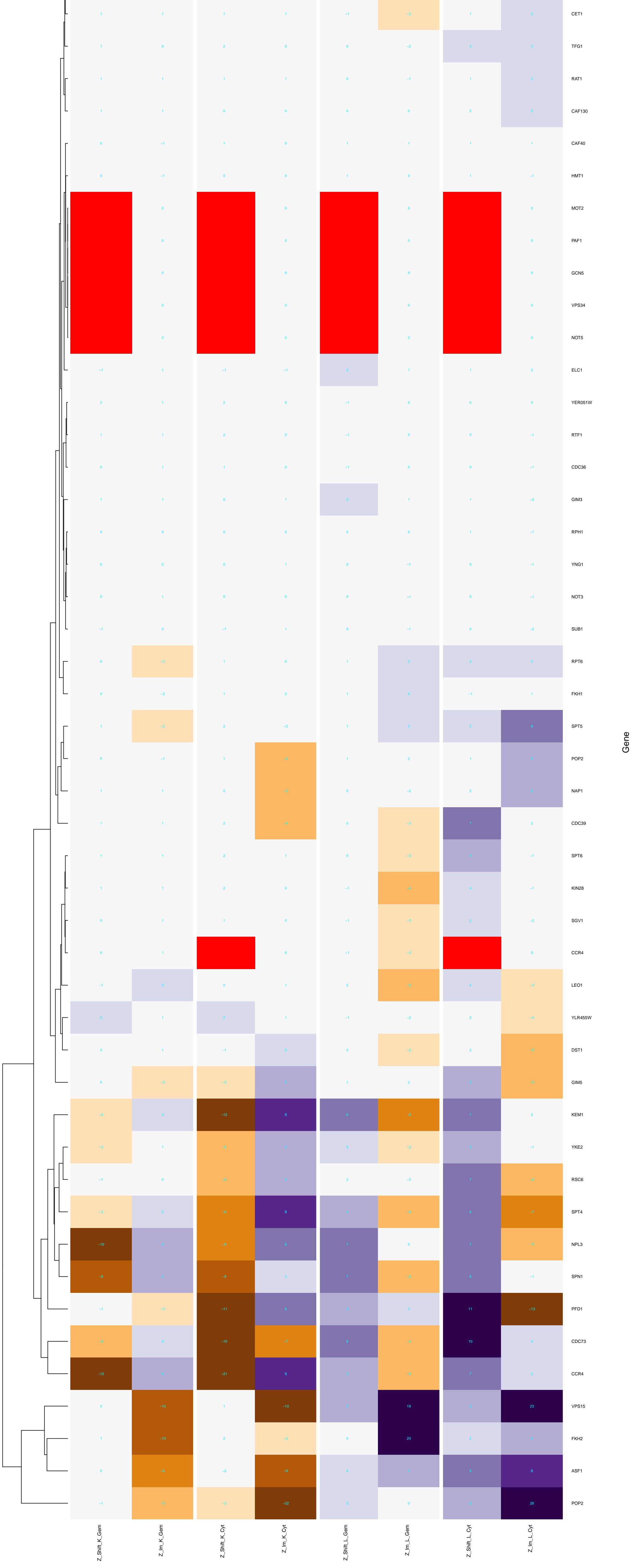
nitrogen catabolite regulation of transcription from RNA polymerase II promoter



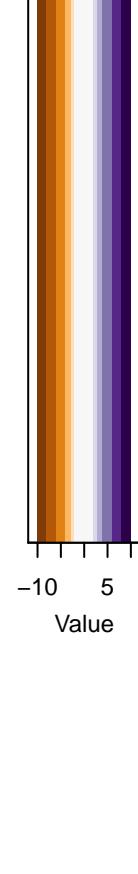




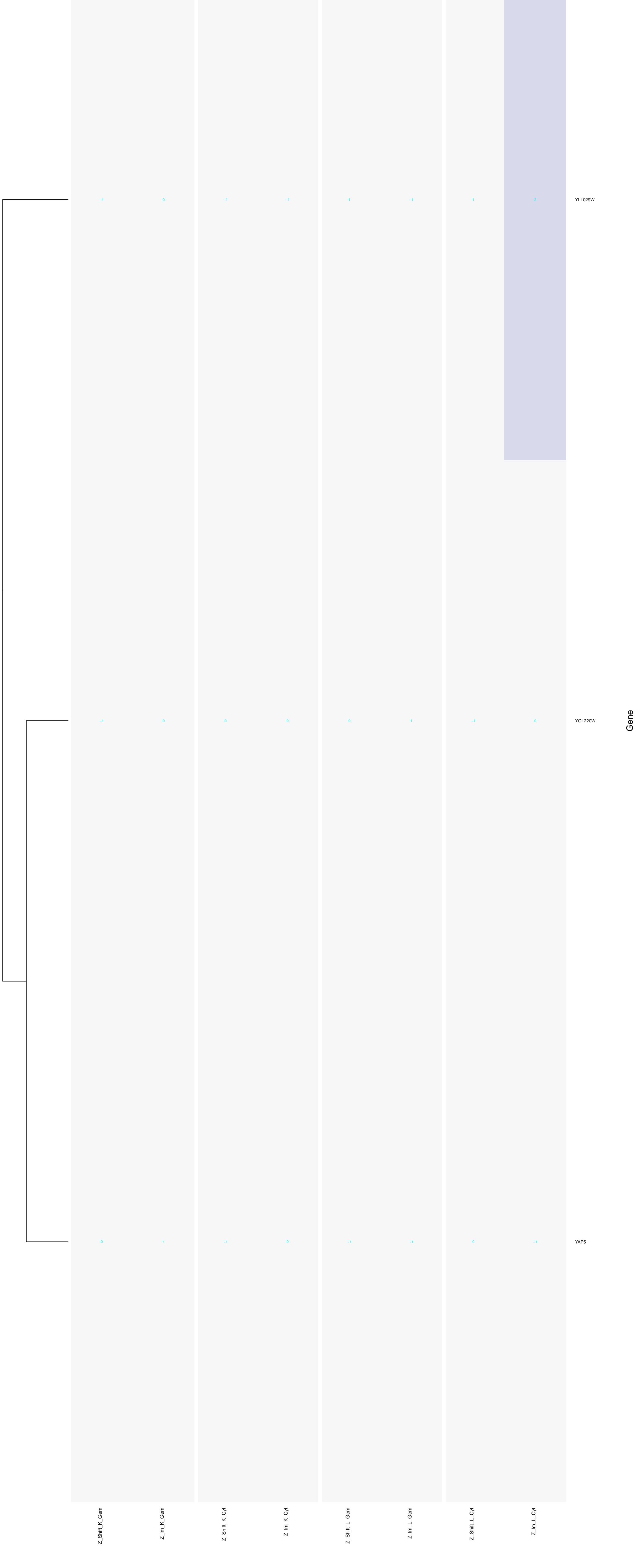
regulation of transcription elongation from RNA polymerase II promoter



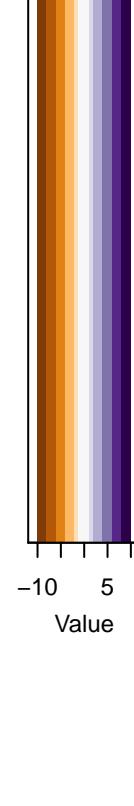
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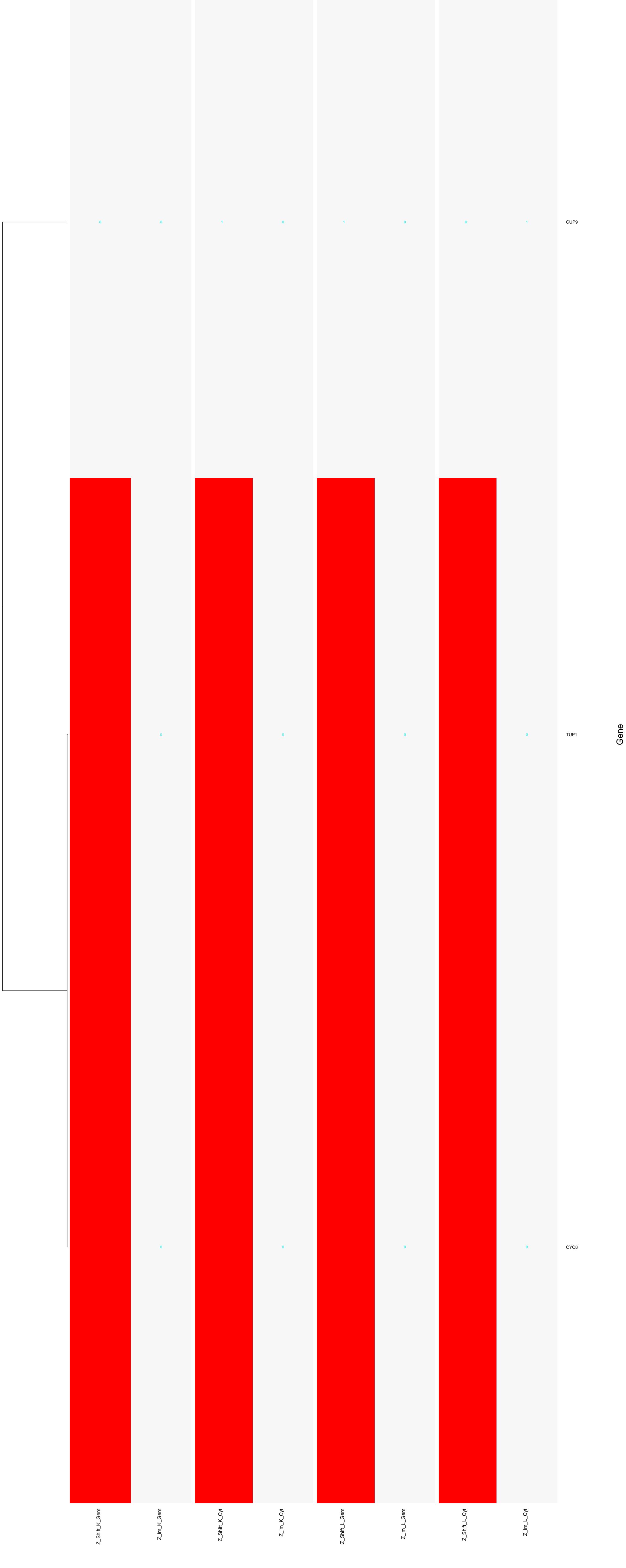
regulation of transcription from RNA polymerase II promoter in response to iron



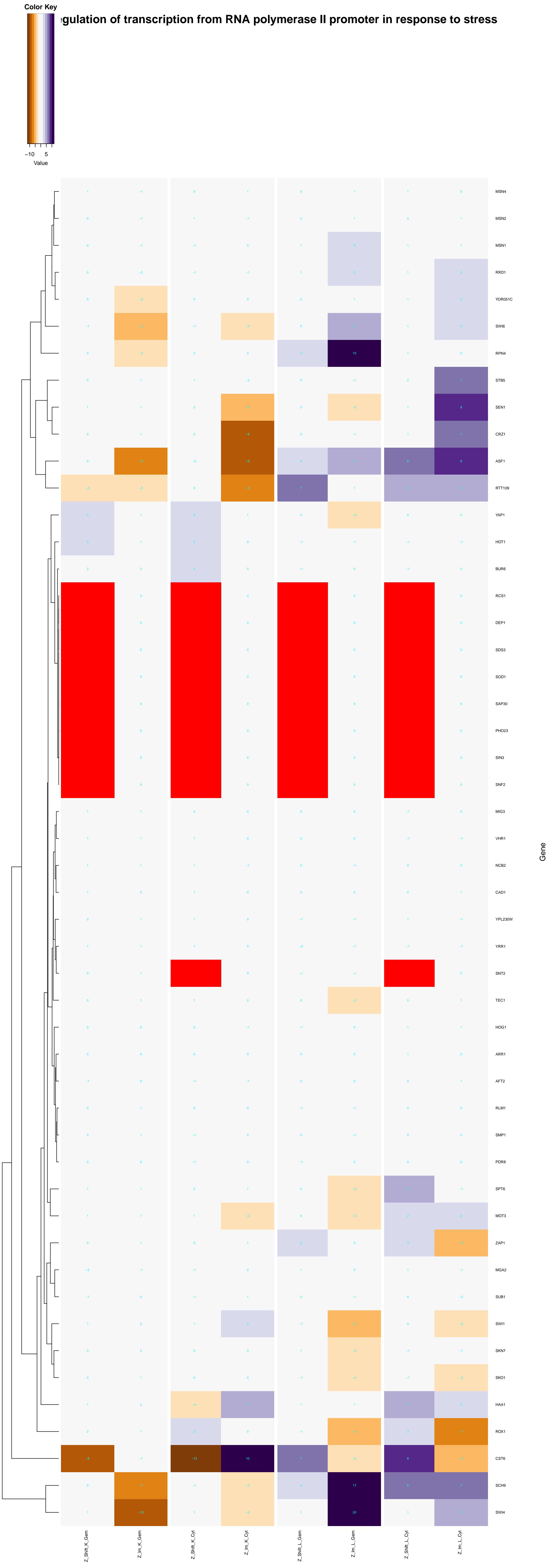
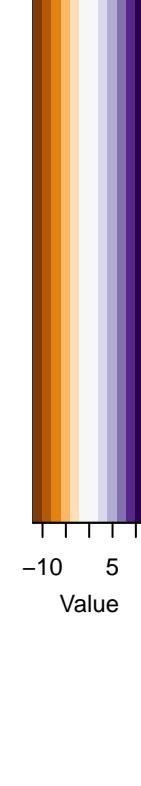
Color Key



of oligopeptide transport by regulation of transcription from RNA polymerase II promoter

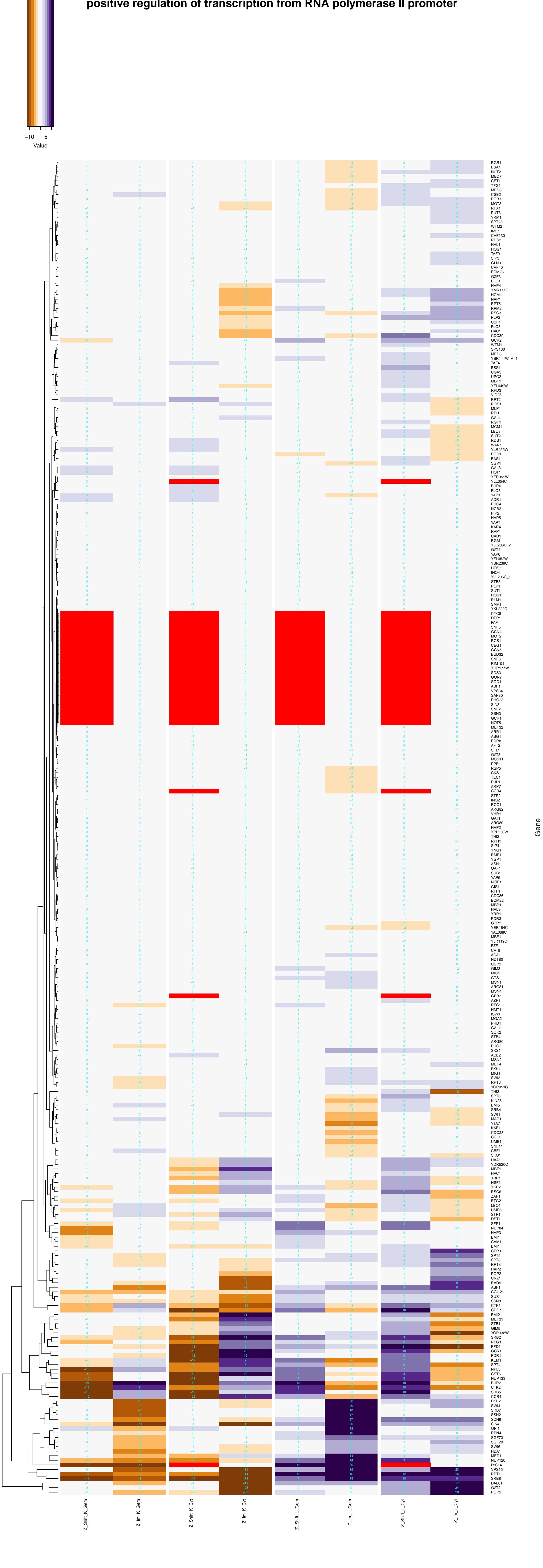


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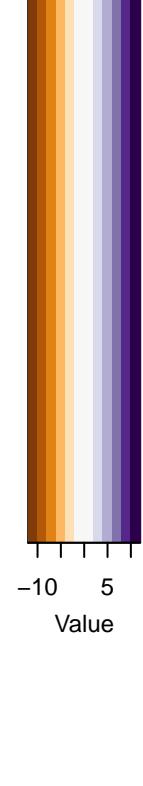


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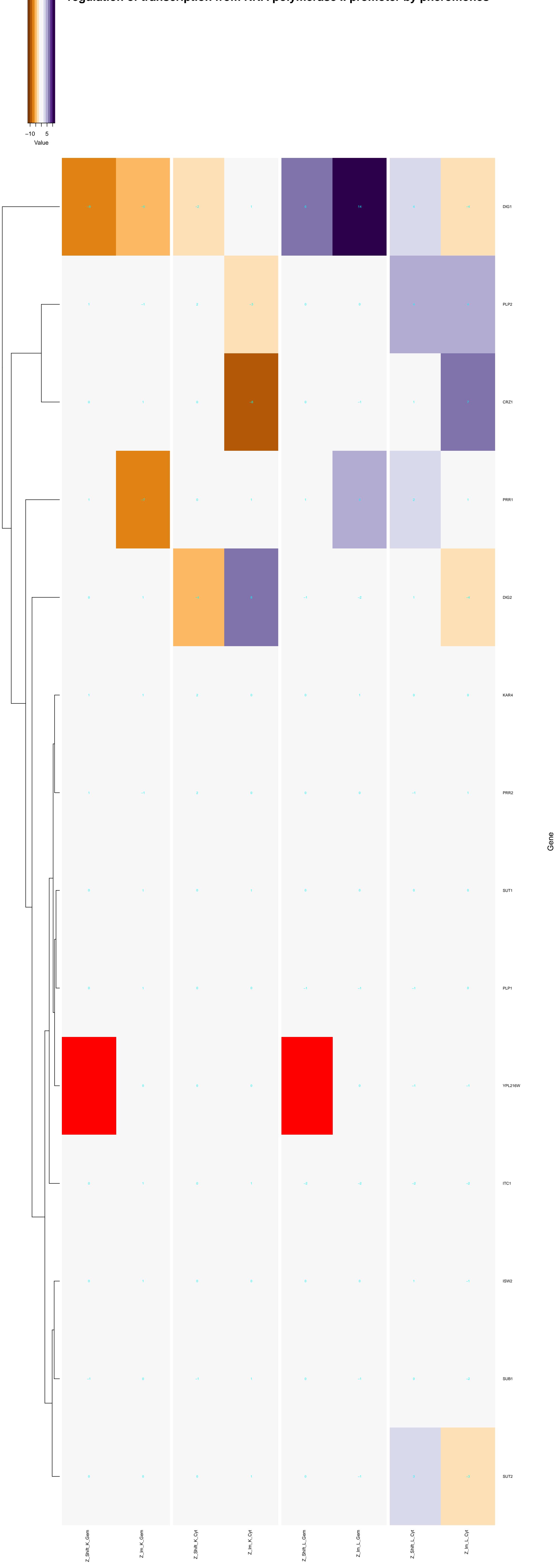
positive regulation of transcription from RNA polymerase II promoter



Color Key



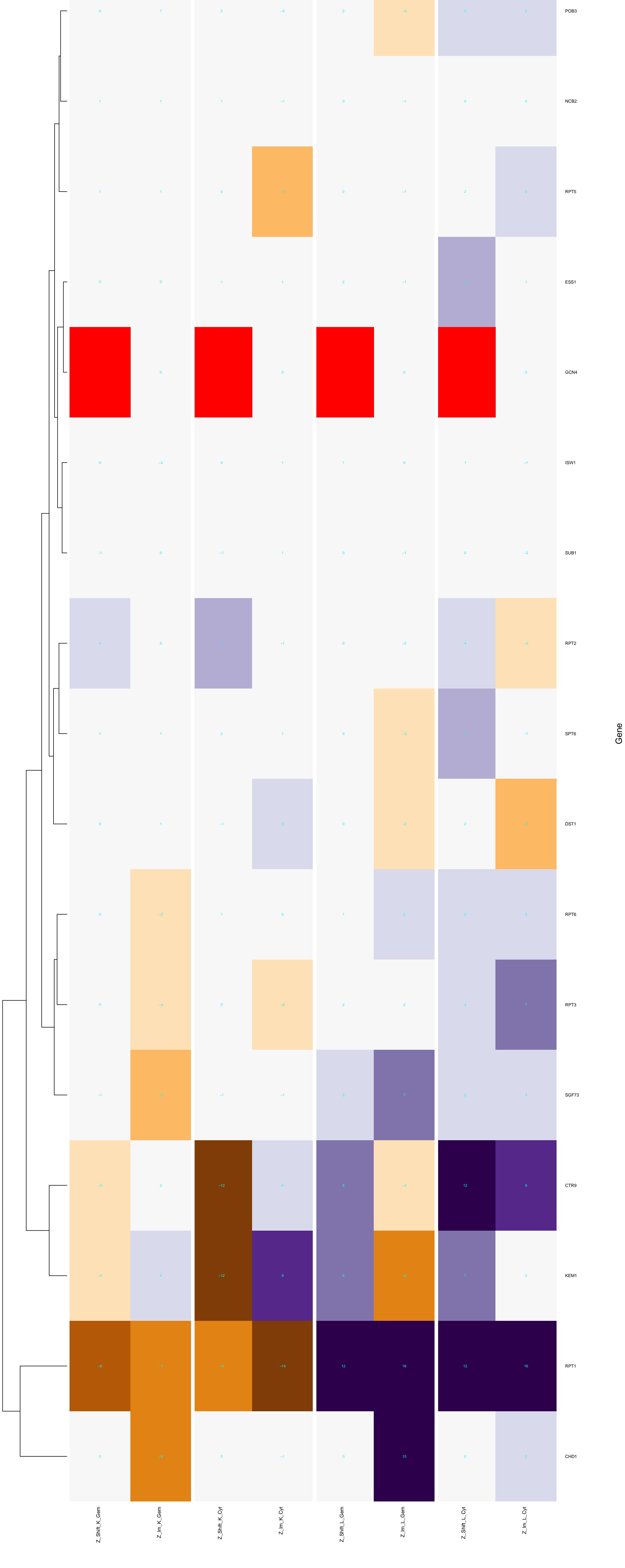
regulation of transcription from RNA polymerase II promoter by pheromones

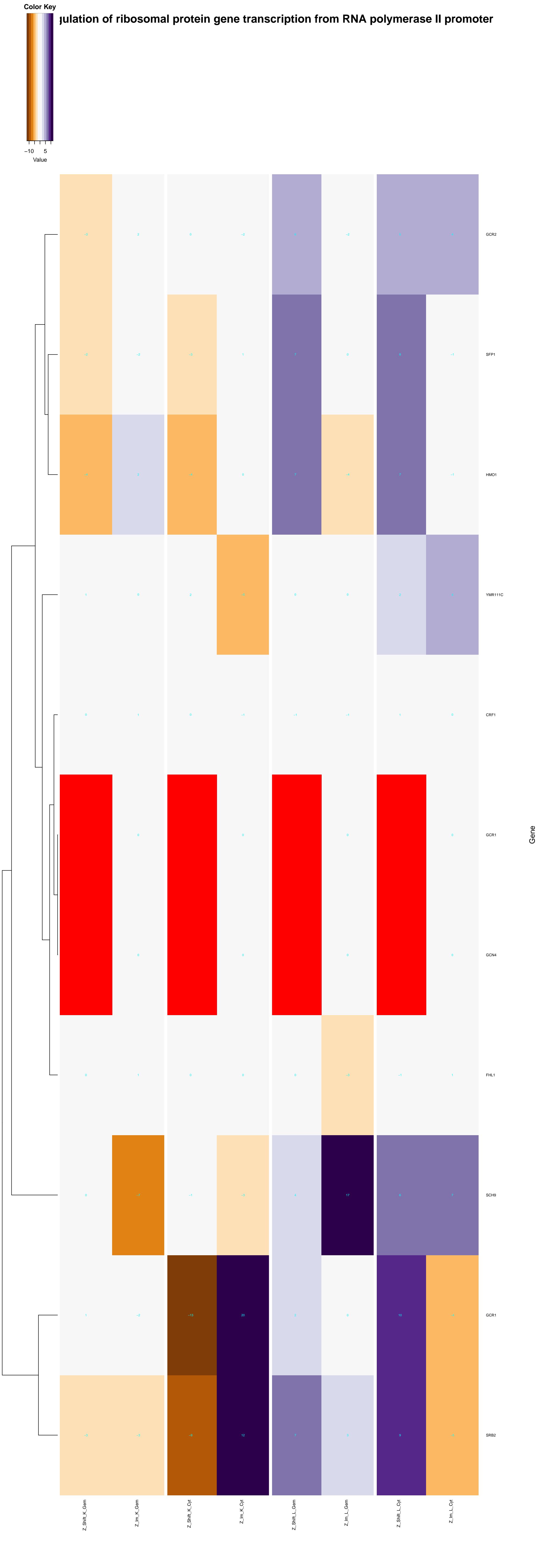
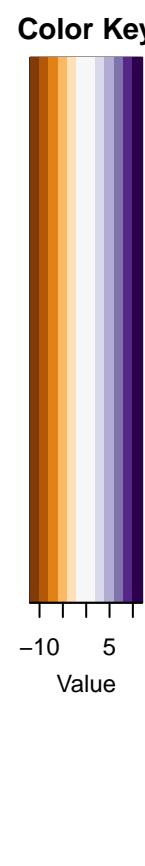


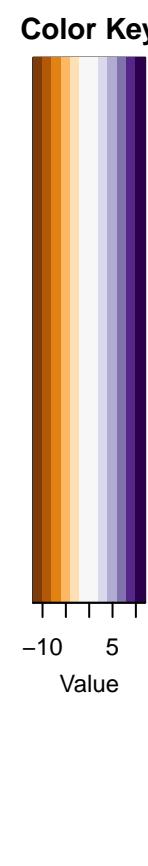
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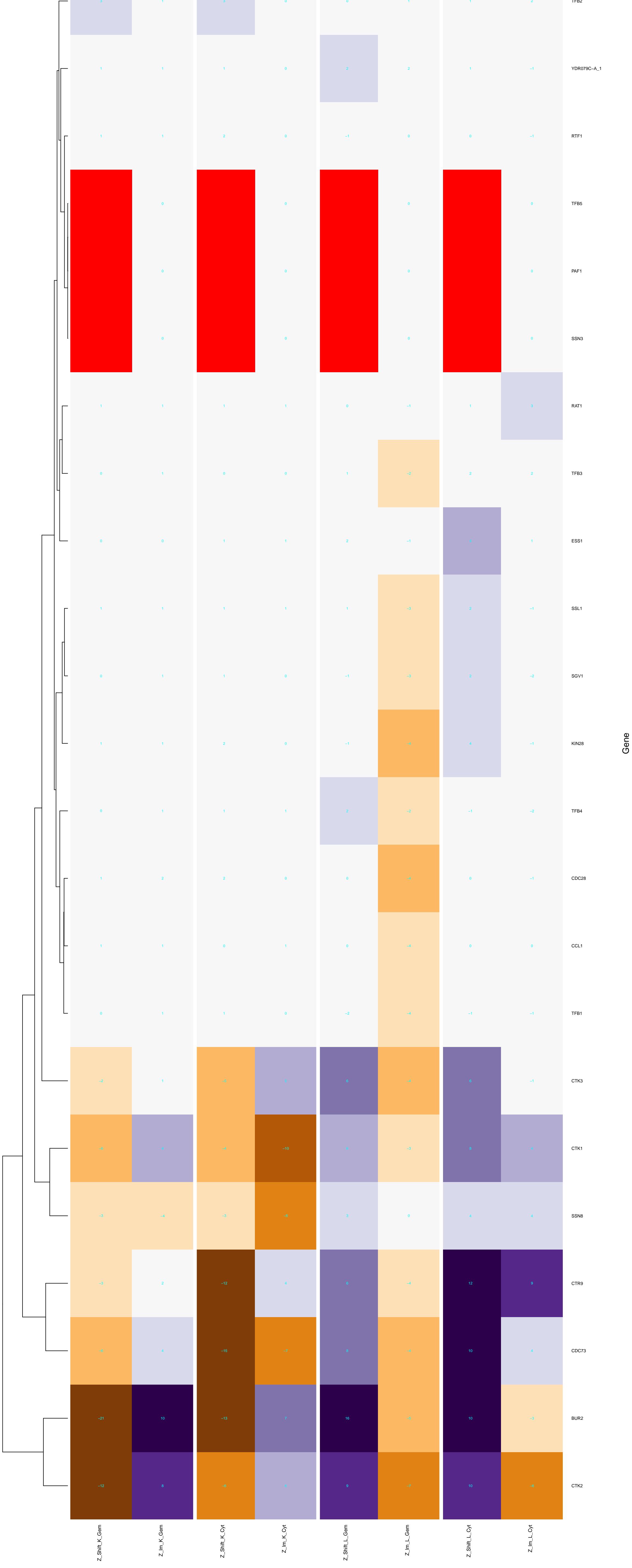
regulation of transcription initiation from RNA polymerase II promoter



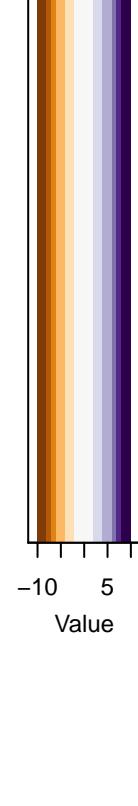




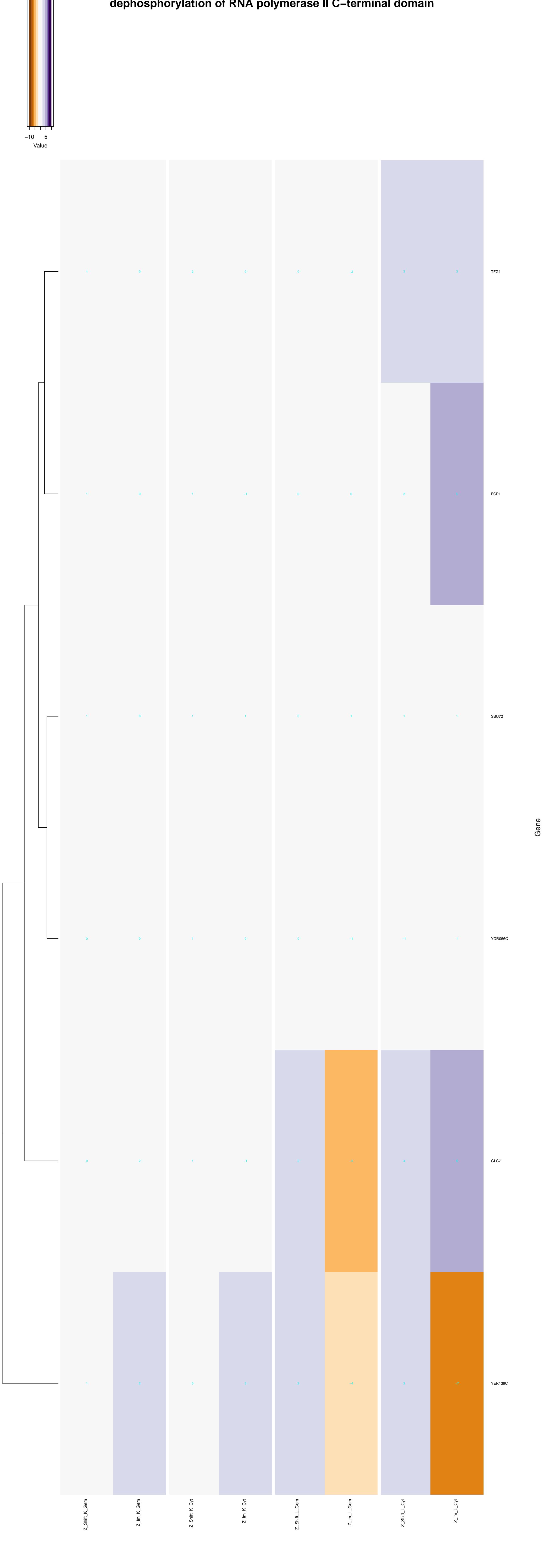
phosphorylation of RNA polymerase II C-terminal domain



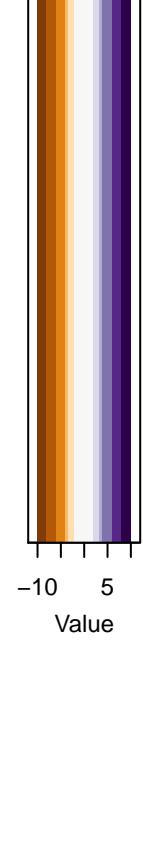
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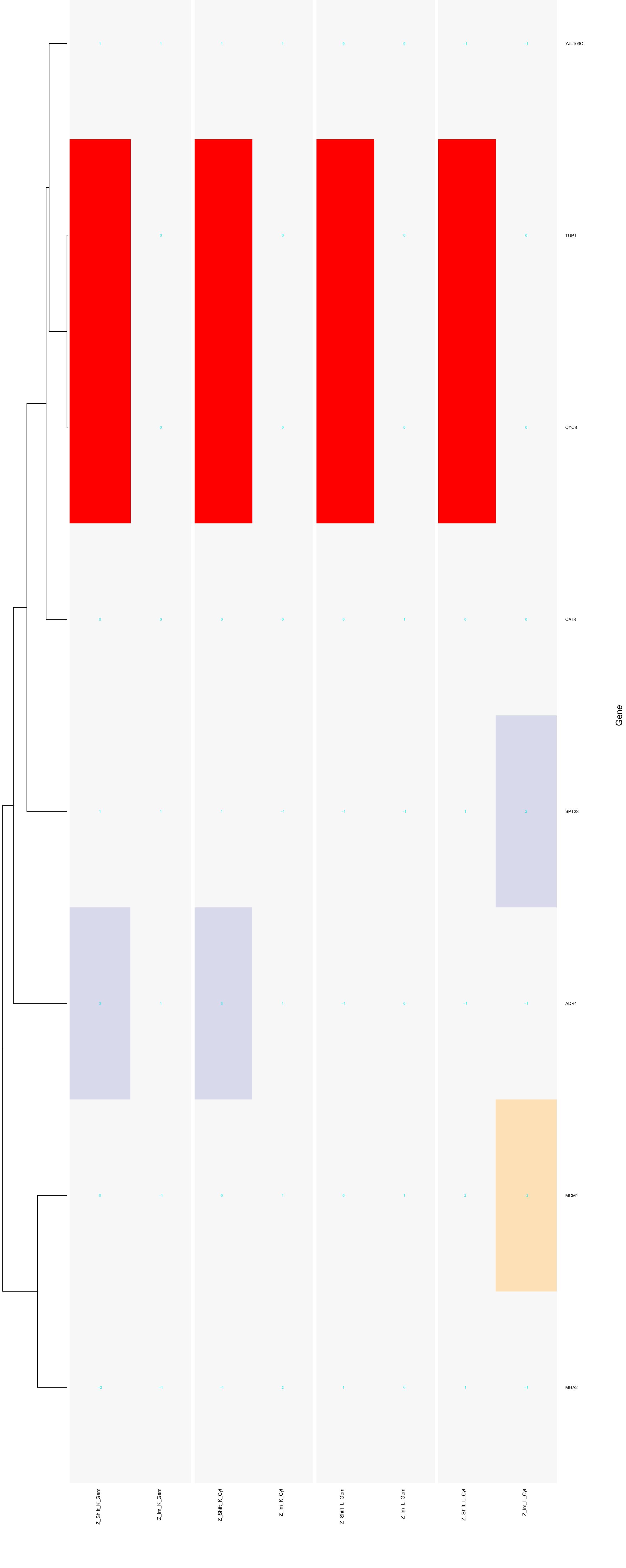
dephosphorylation of RNA polymerase II C-terminal domain



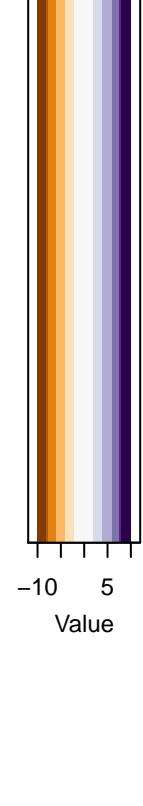
Color Key



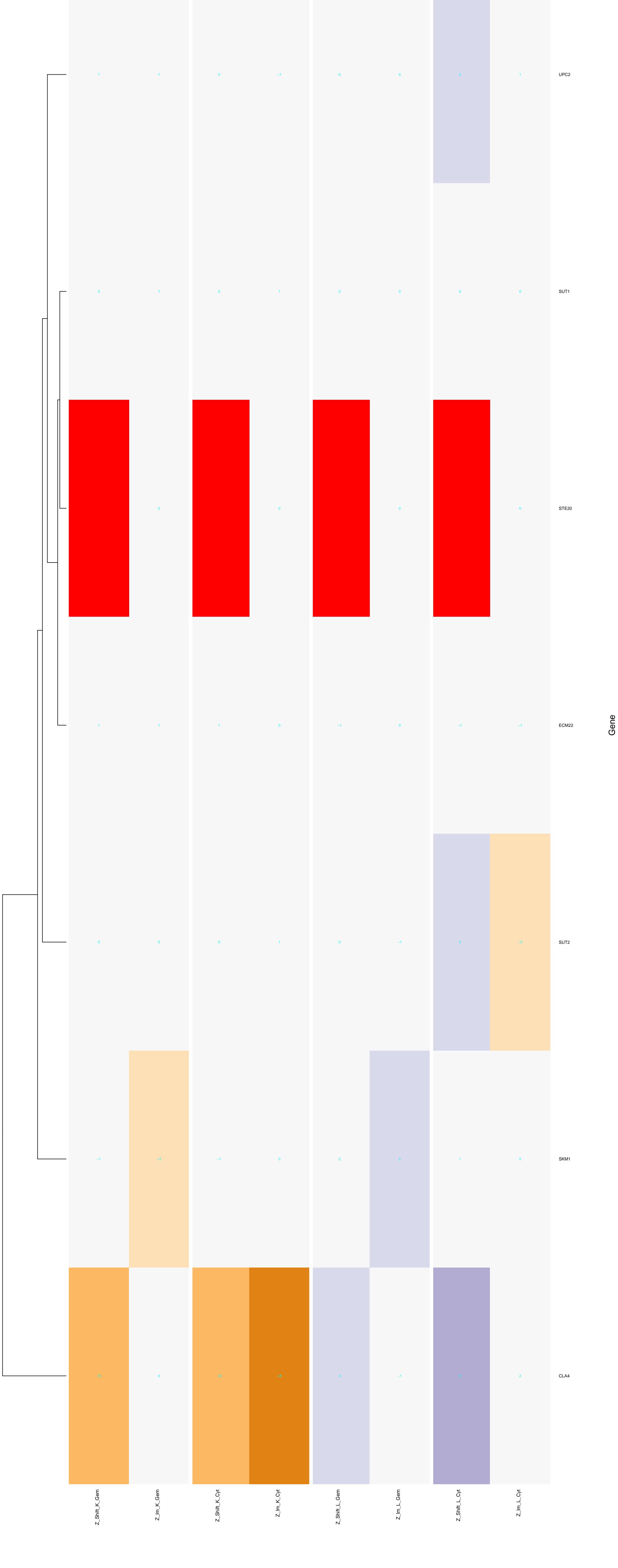
Lipid metabolism by regulation of transcription from RNA polymerase II promoter



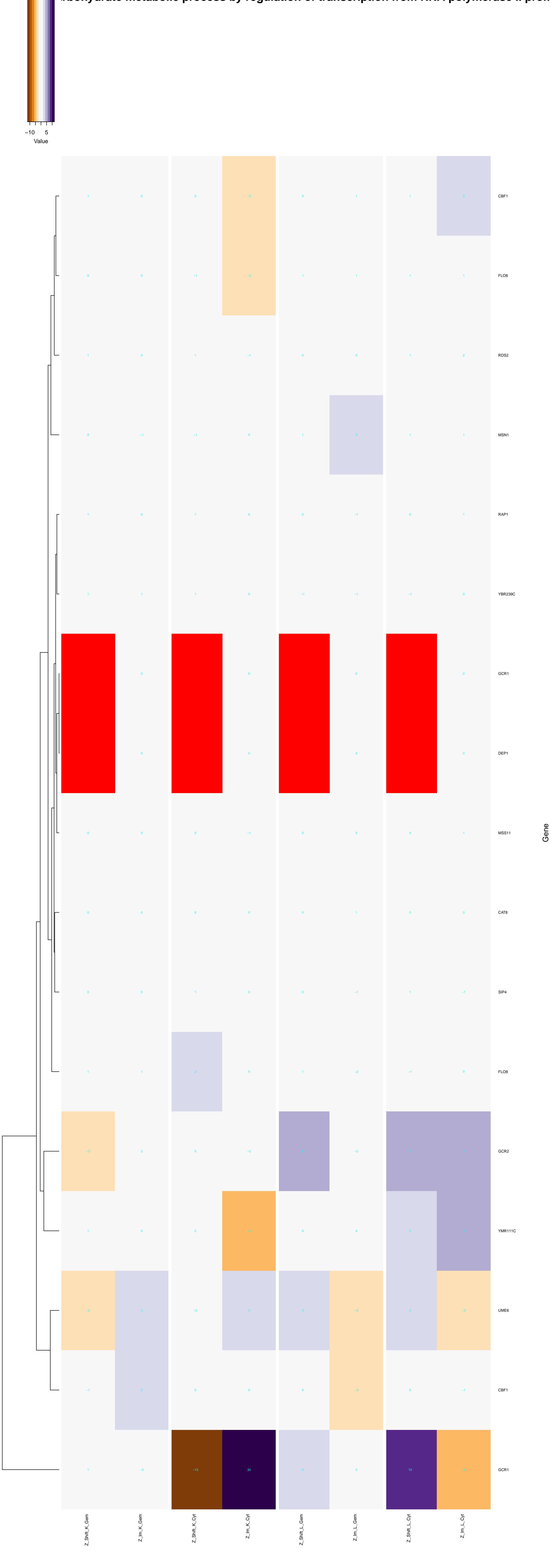
Color Key



Regulation of lipid transport by regulation of transcription from RNA polymerase II promoter



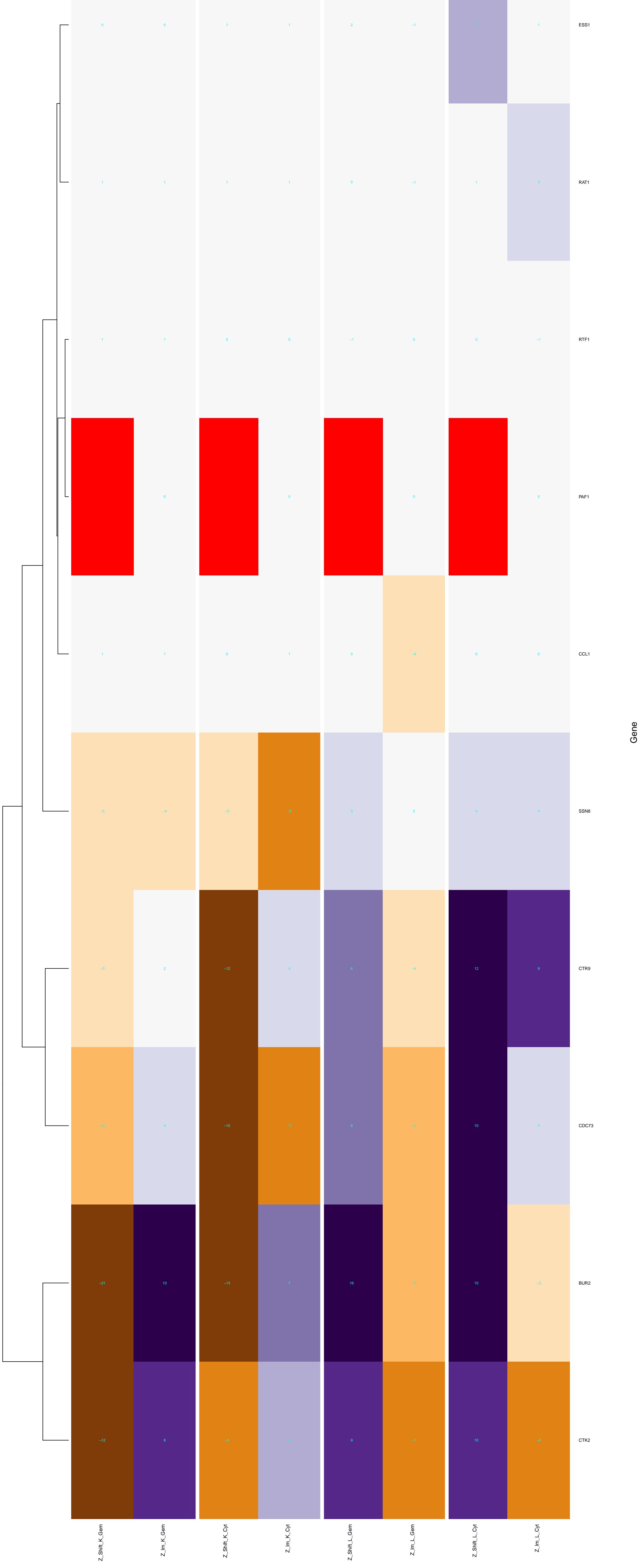
Color Key



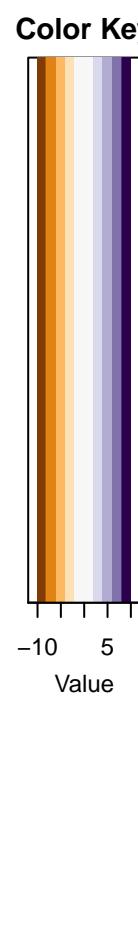
Color Key



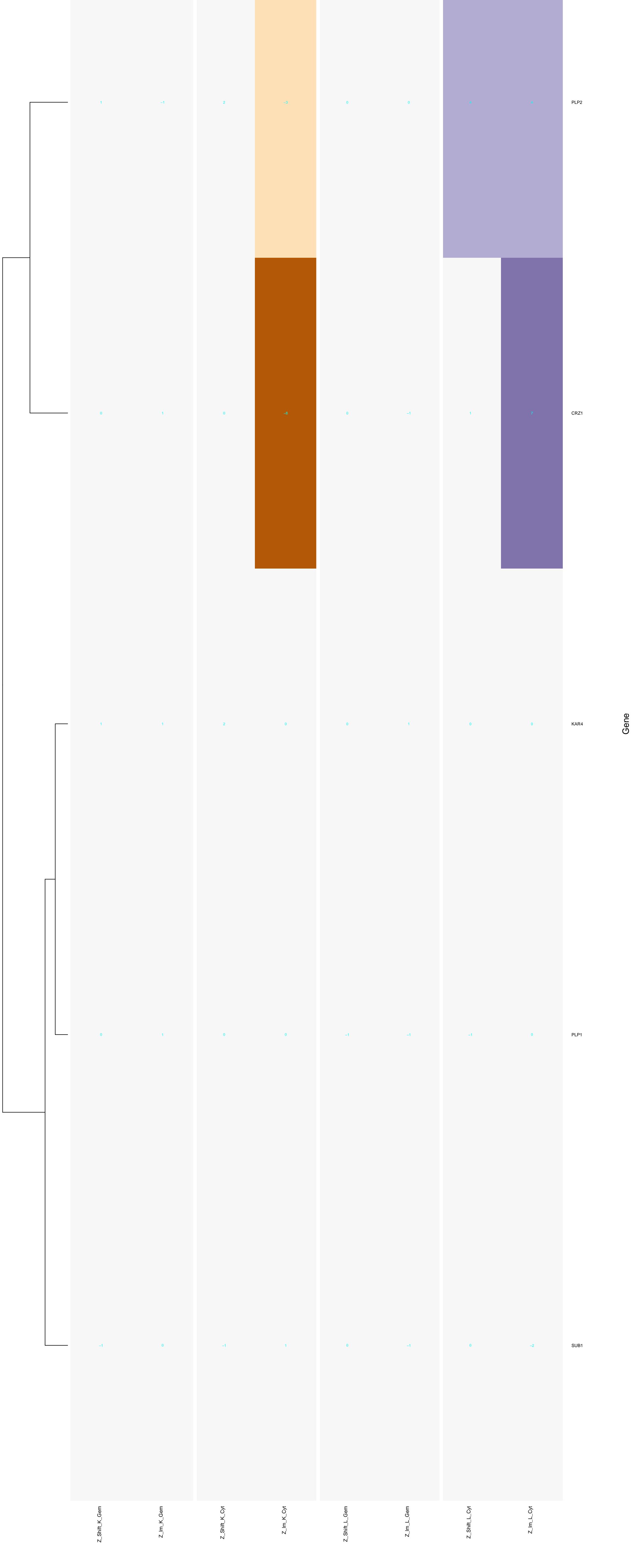
regulation of phosphorylation of RNA polymerase II C-terminal domain



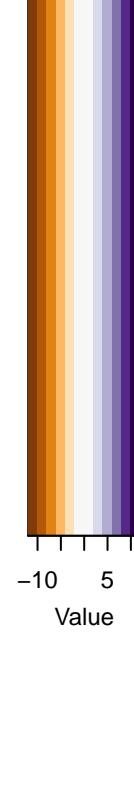
Color Key



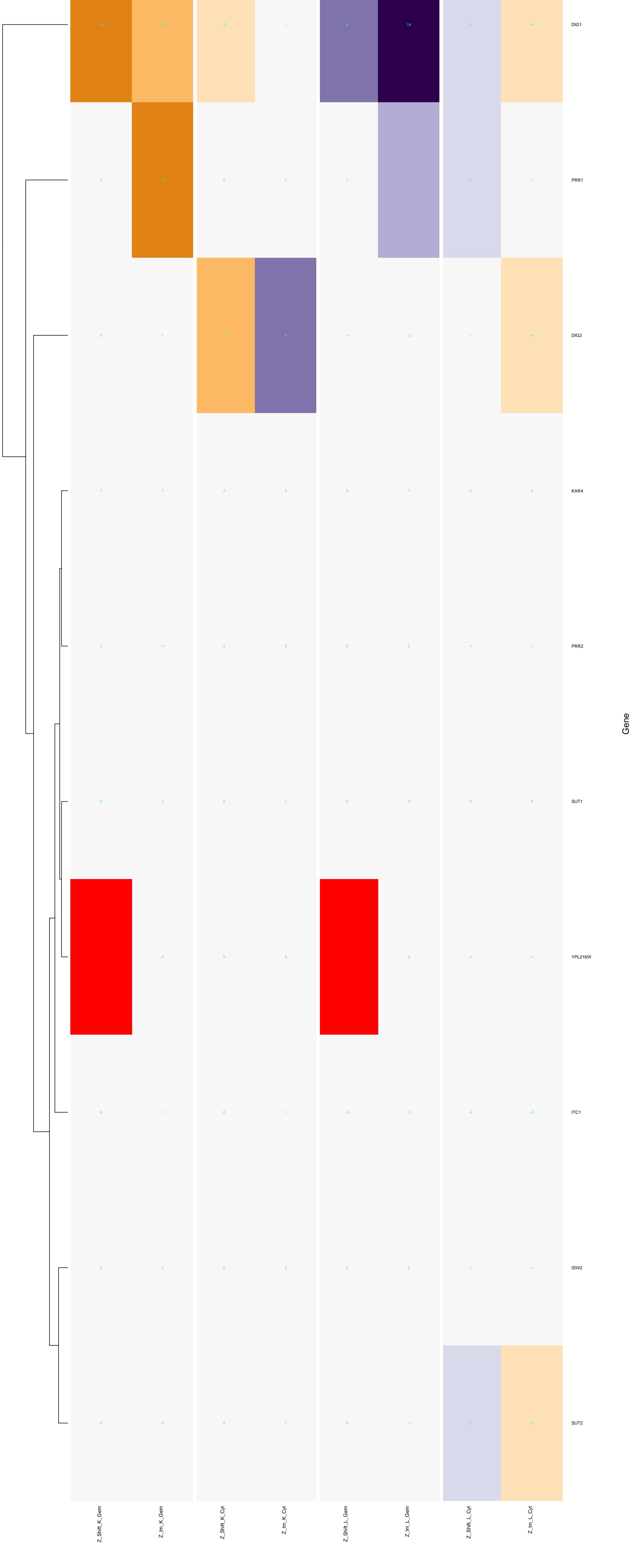
positive regulation of transcription by pheromones

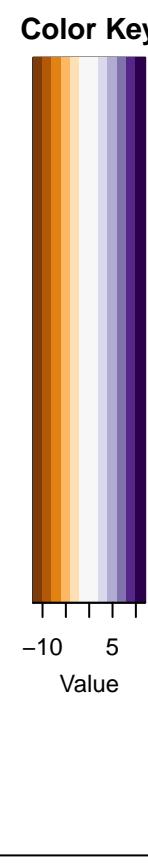


Color Key

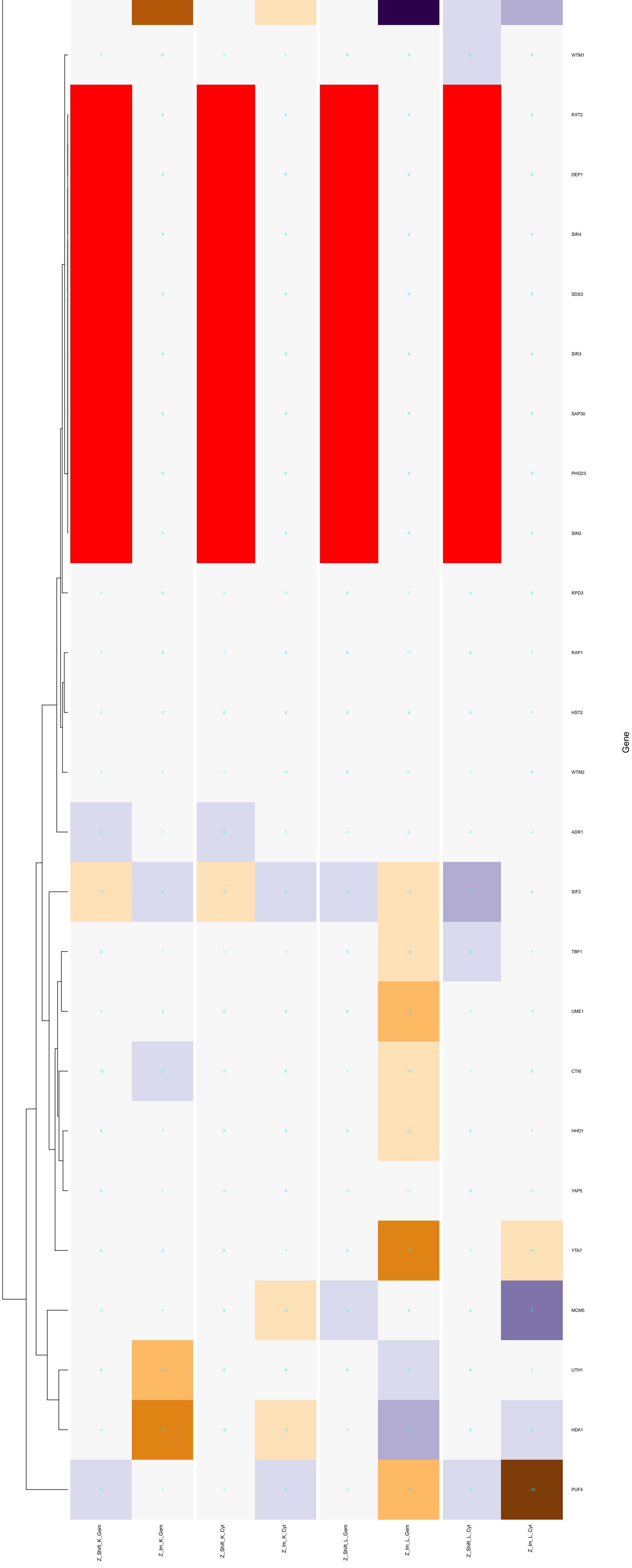


negative regulation of transcription by pheromones





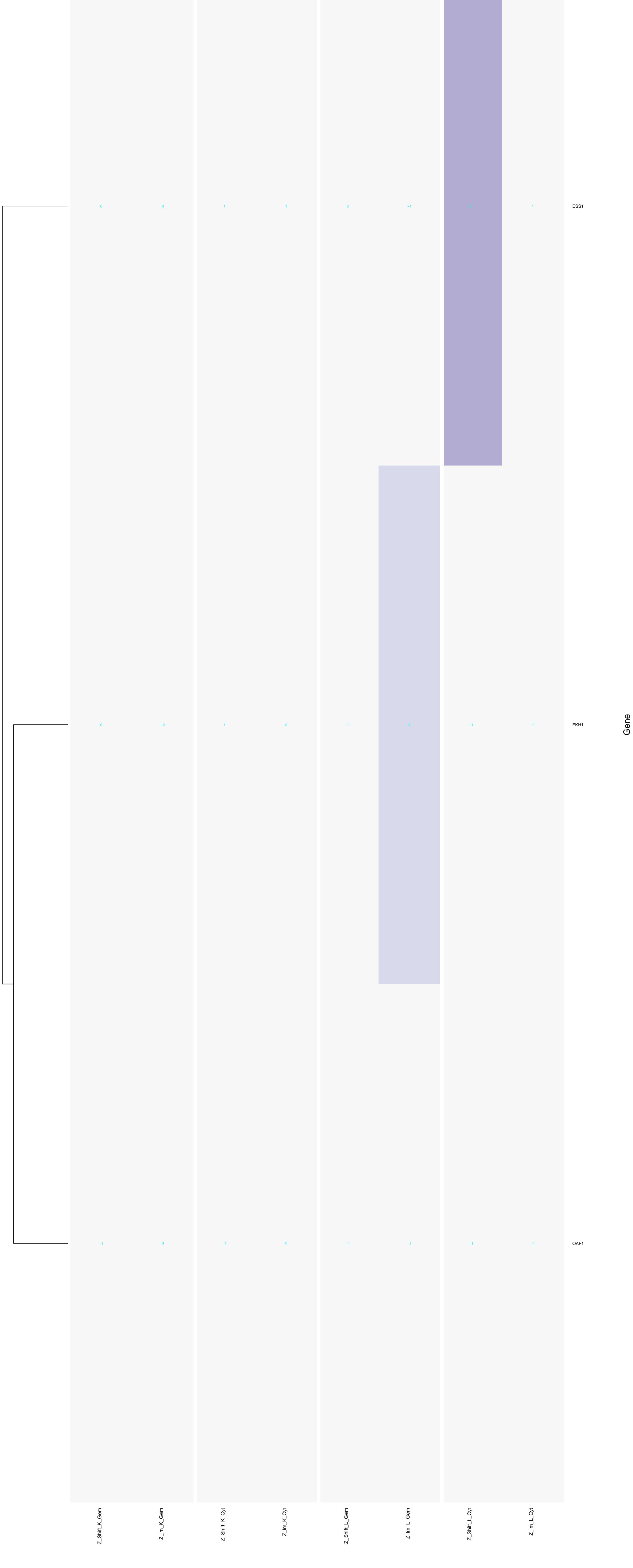
negative regulation of chromatin silencing

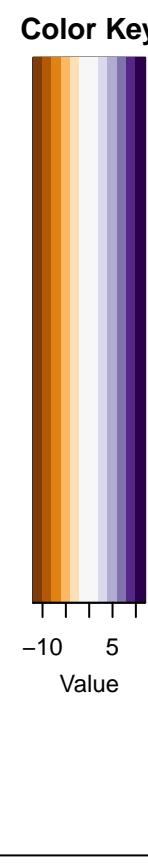


Color Key

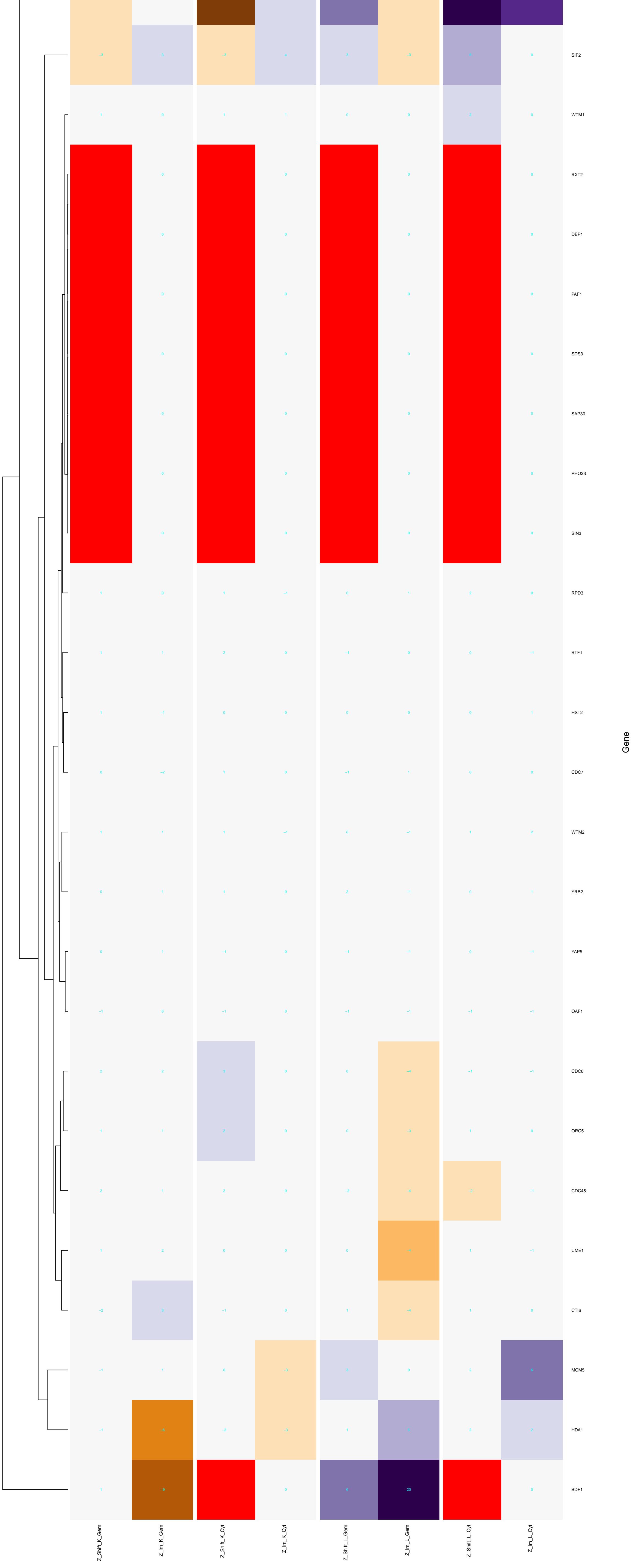


positive regulation of chromatin silencing

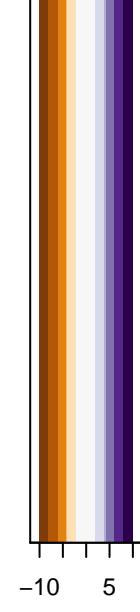




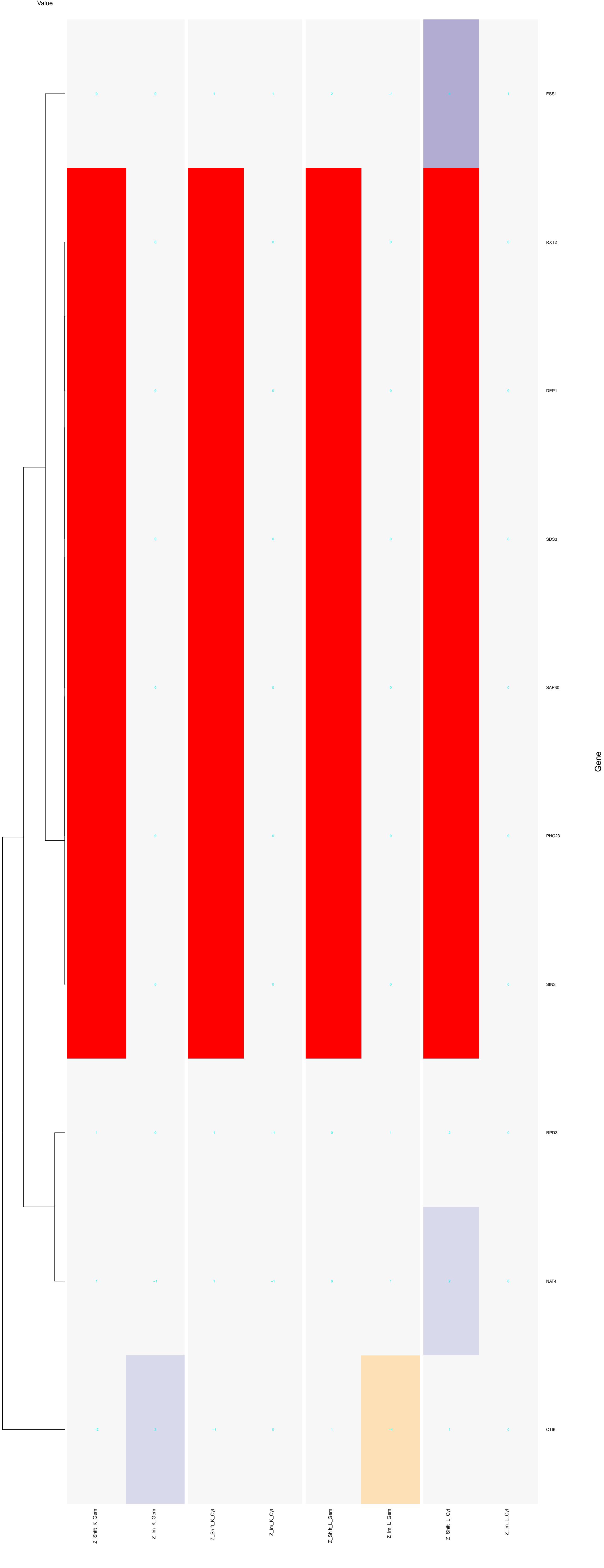
regulation of chromatin silencing at telomere



Color Key



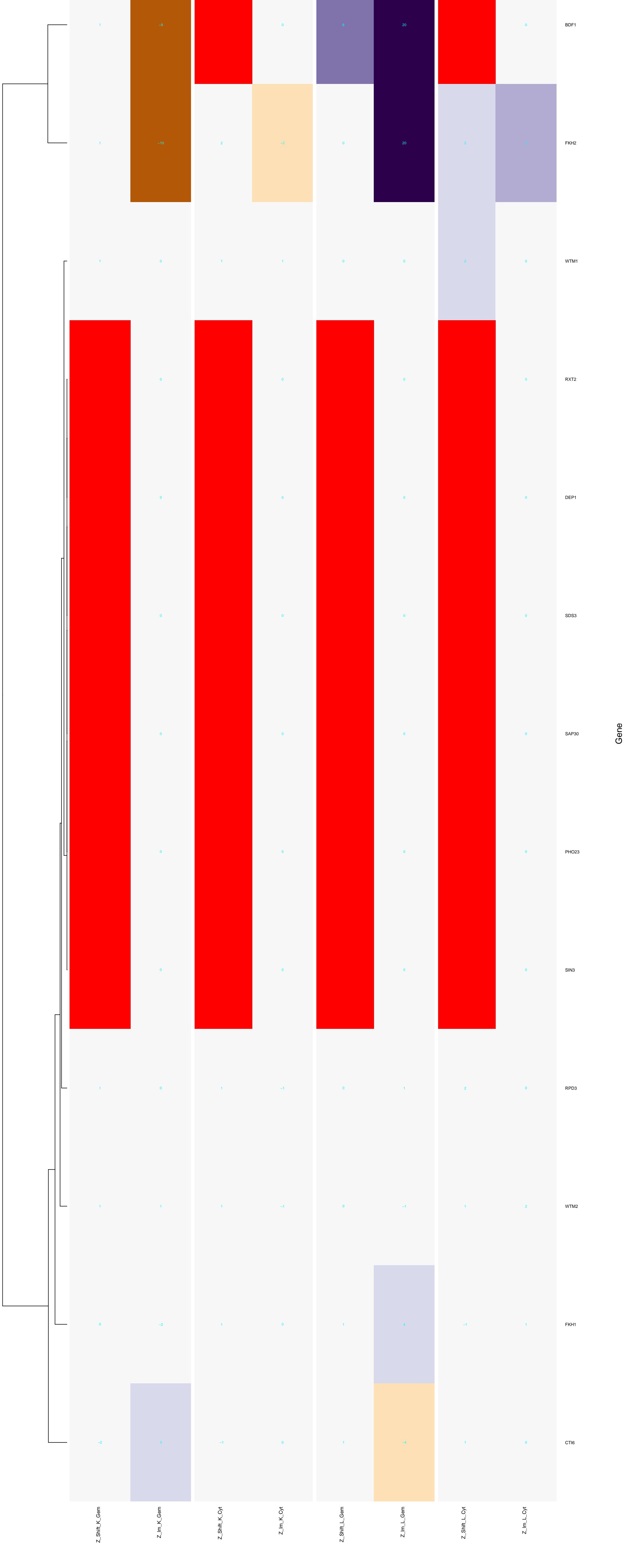
regulation of chromatin silencing at rDNA



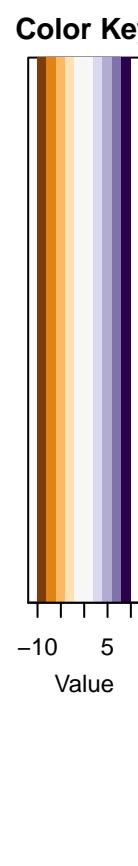
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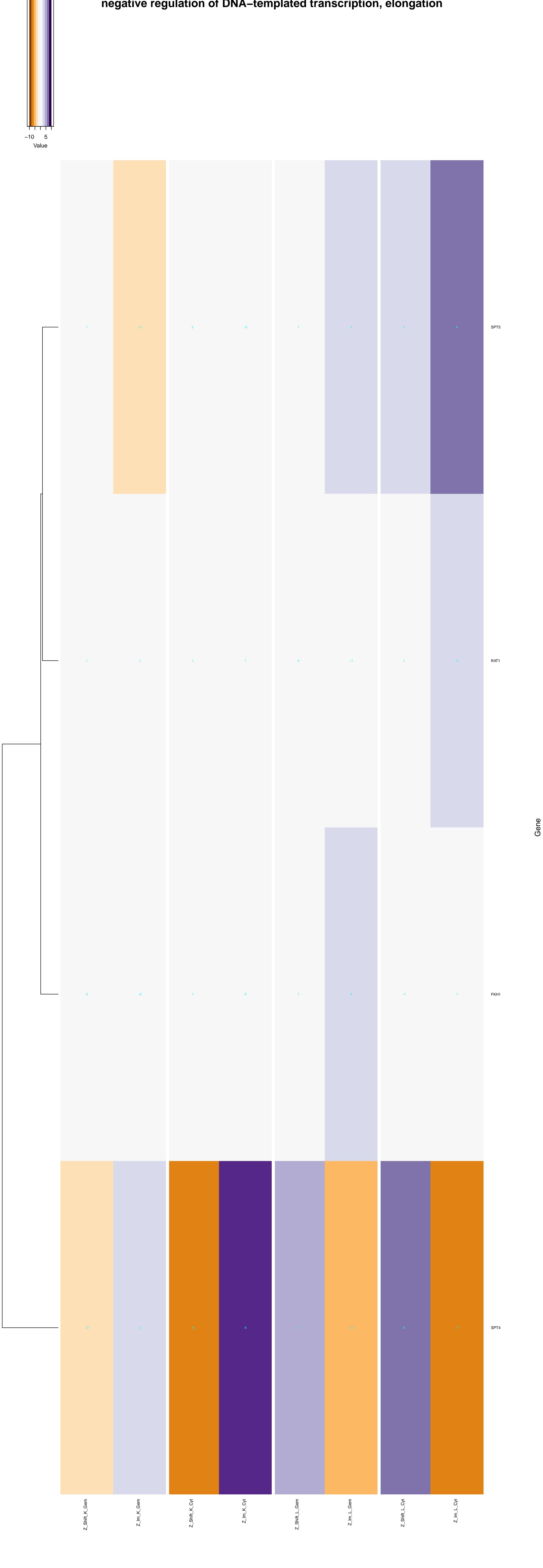
regulation of chromatin silencing at silent mating-type cassette

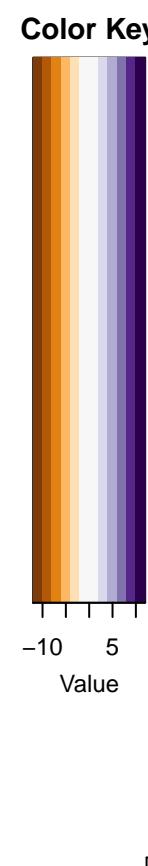


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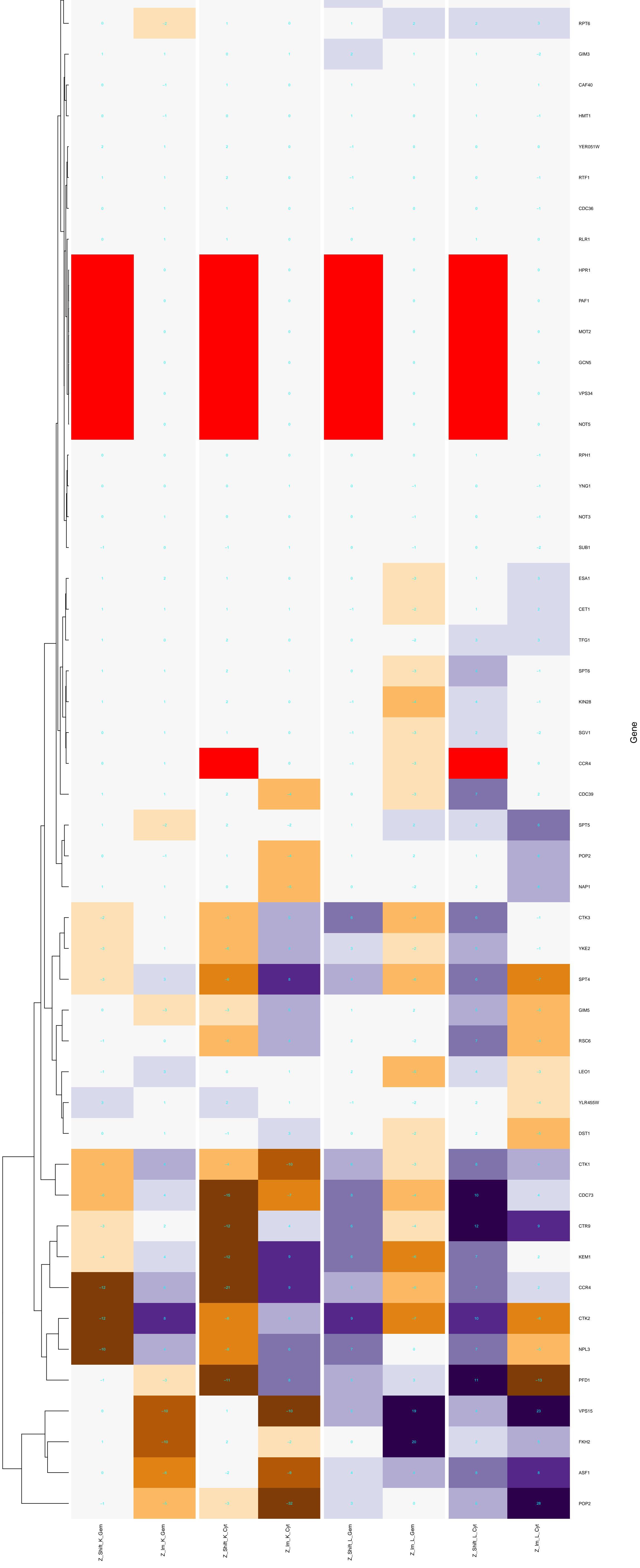


negative regulation of DNA-templated transcription, elongation





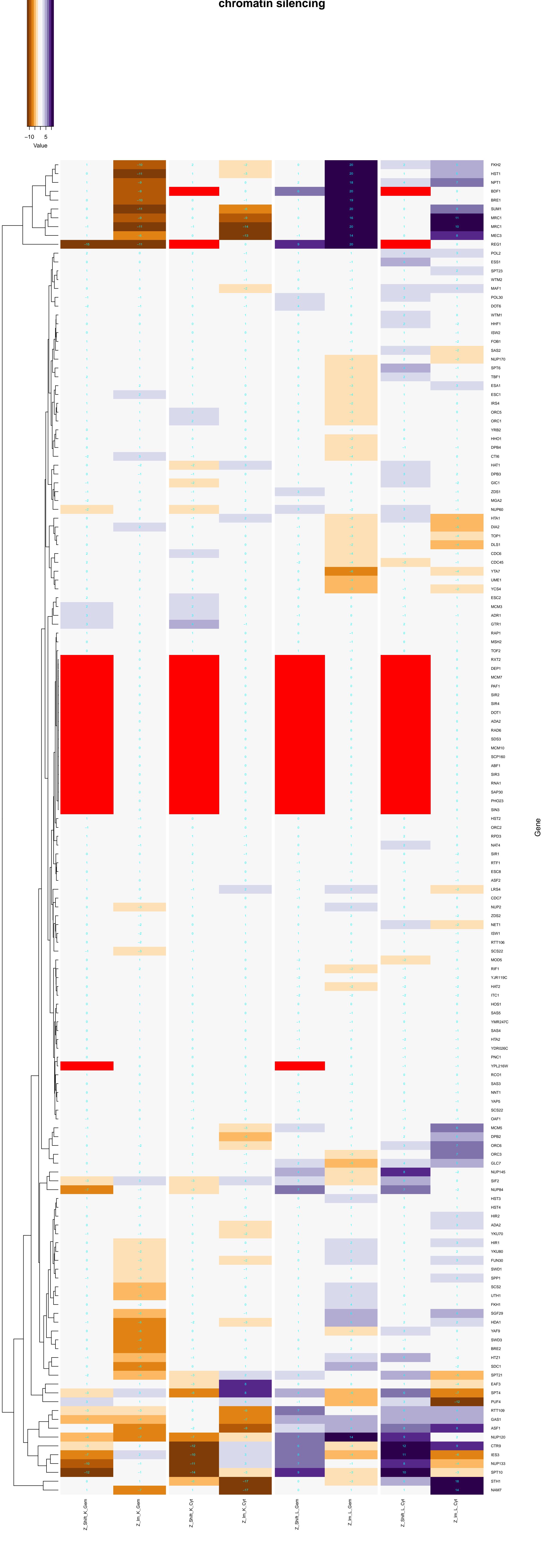
positive regulation of DNA-templated transcription, elongation



Color Key



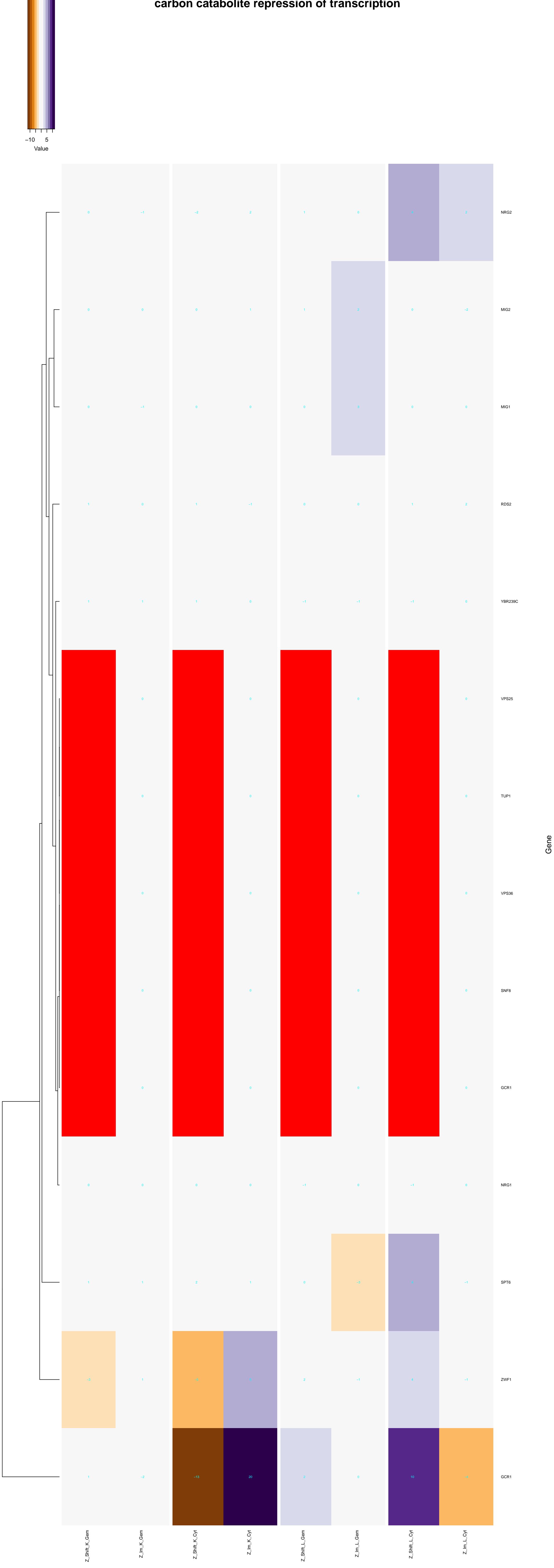
chromatin silencing



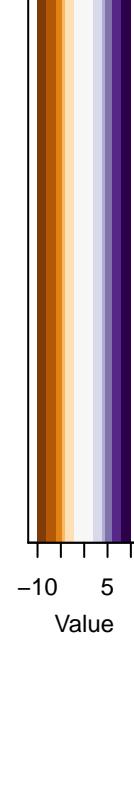
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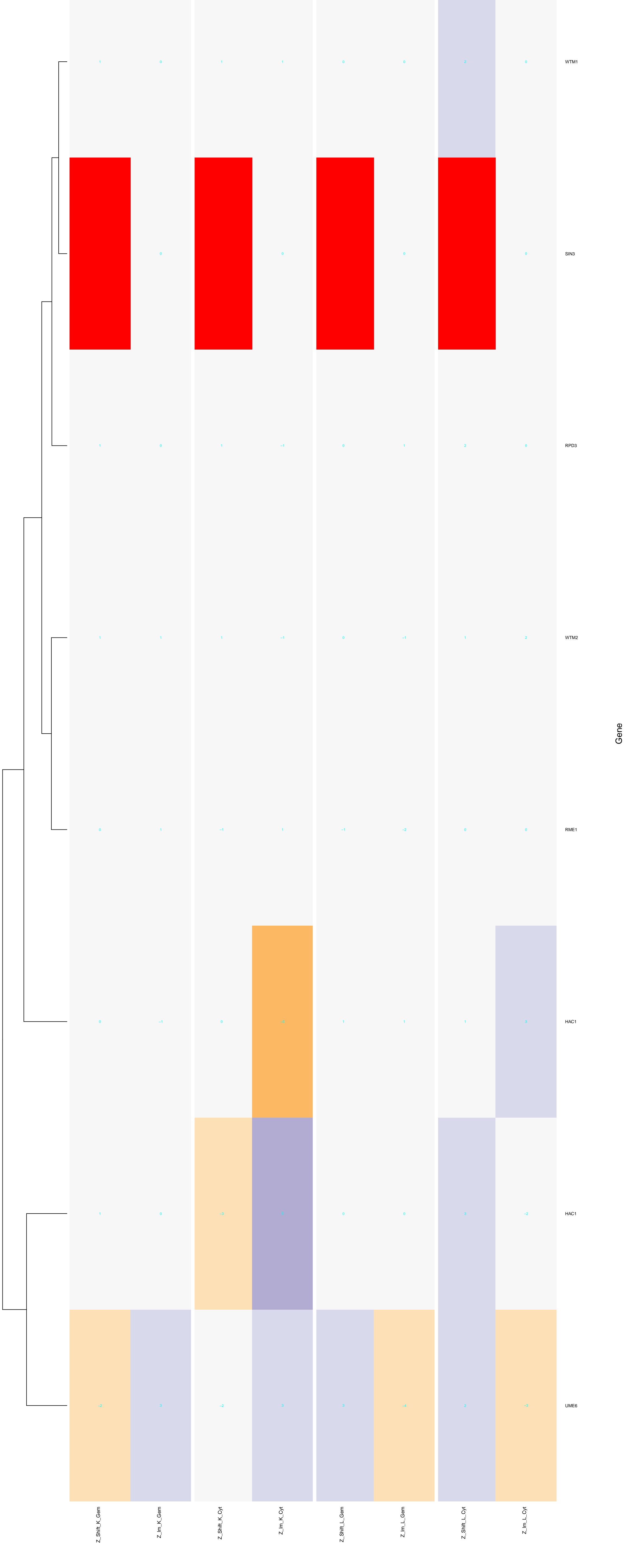
carbon catabolite repression of transcription



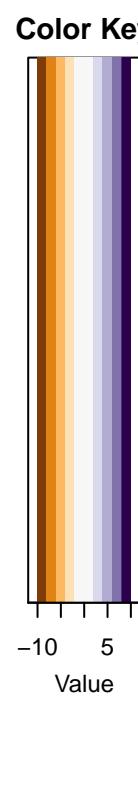
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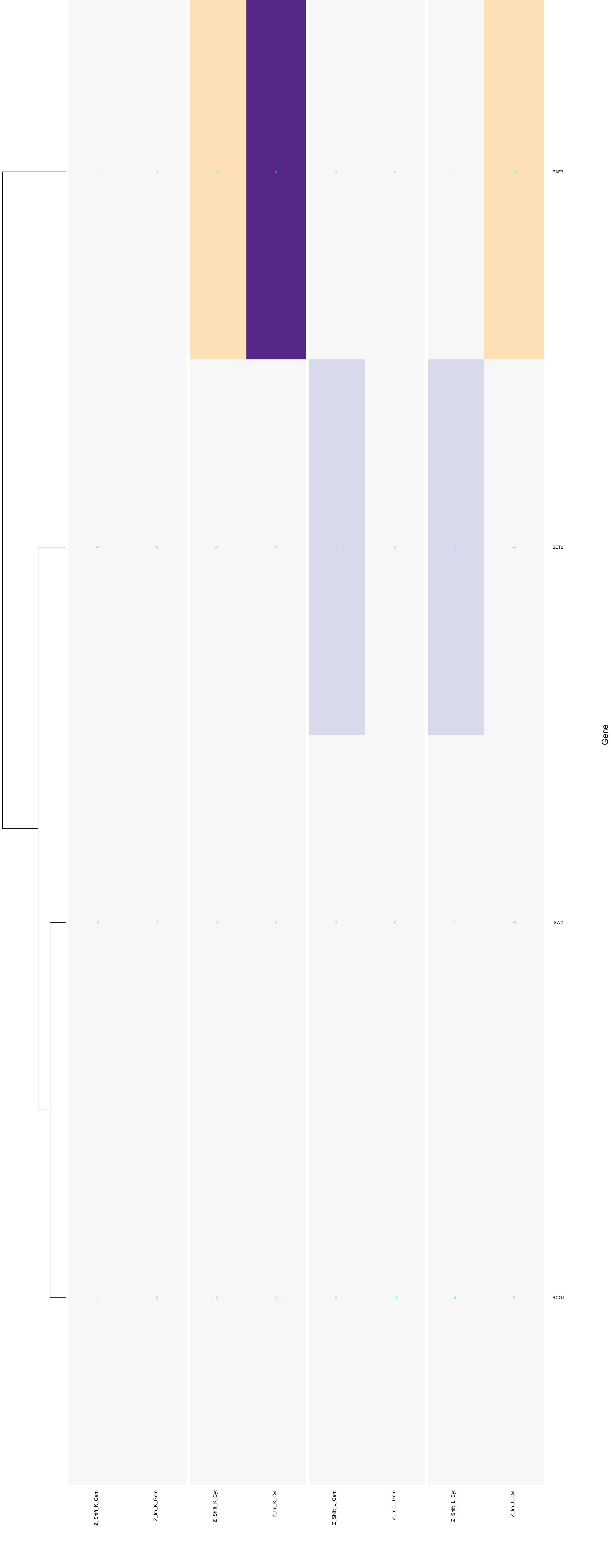
negative regulation of transcription involved in meiotic cell cycle



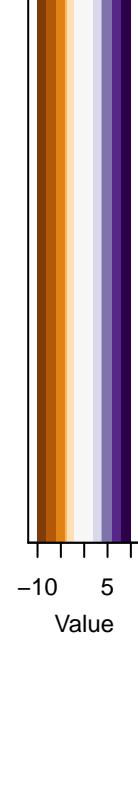
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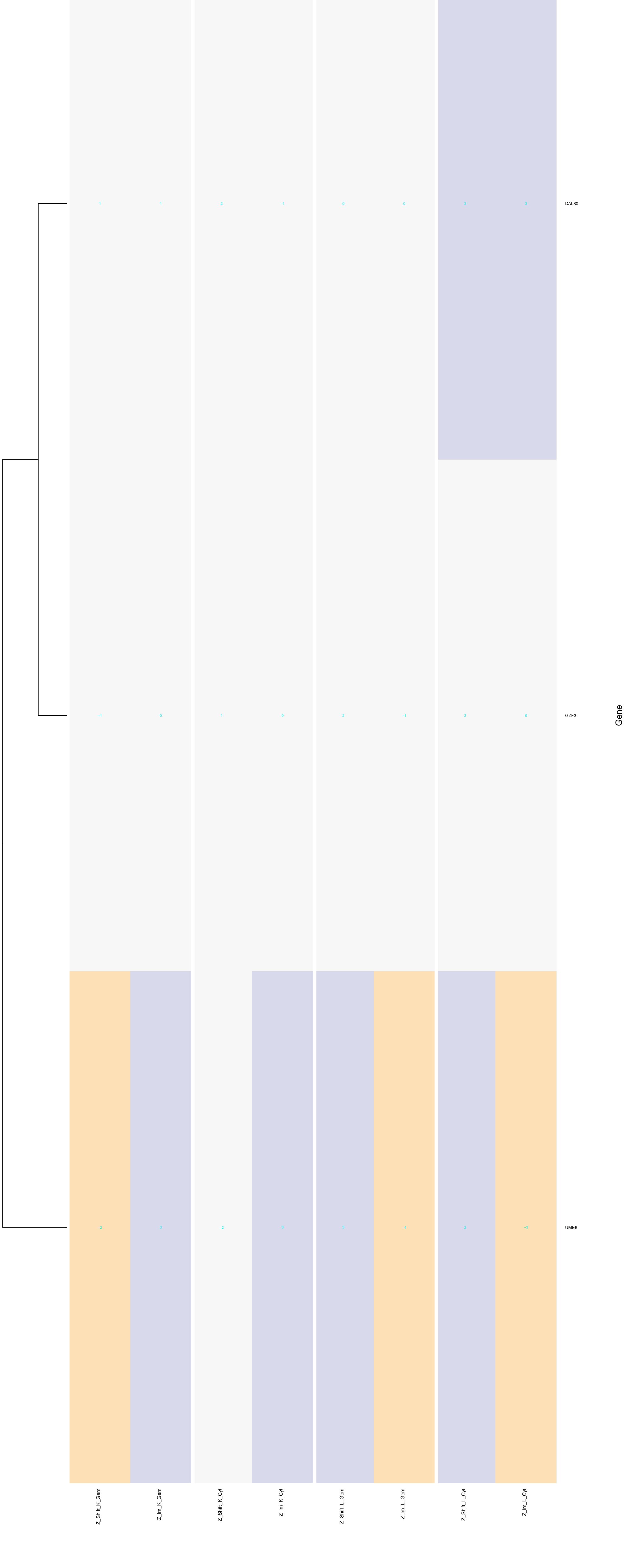
negative regulation of antisense RNA transcription



Color Key

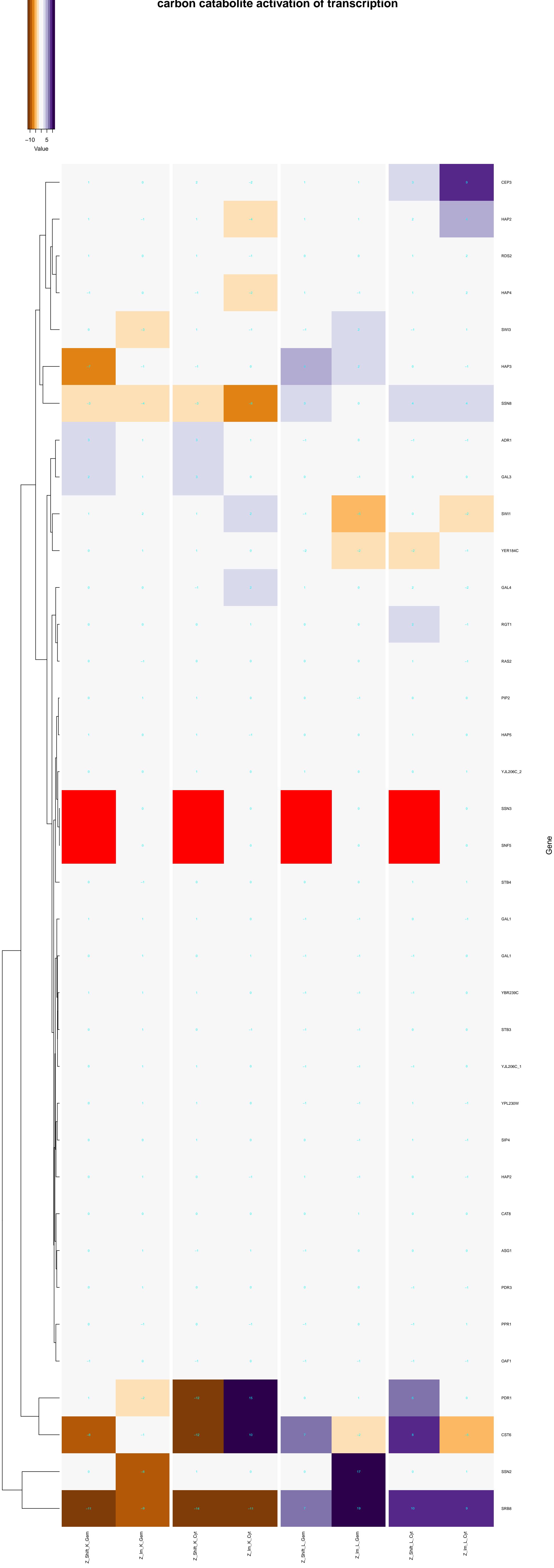


nitrogen catabolite repression of transcription

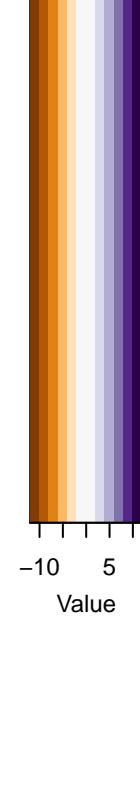


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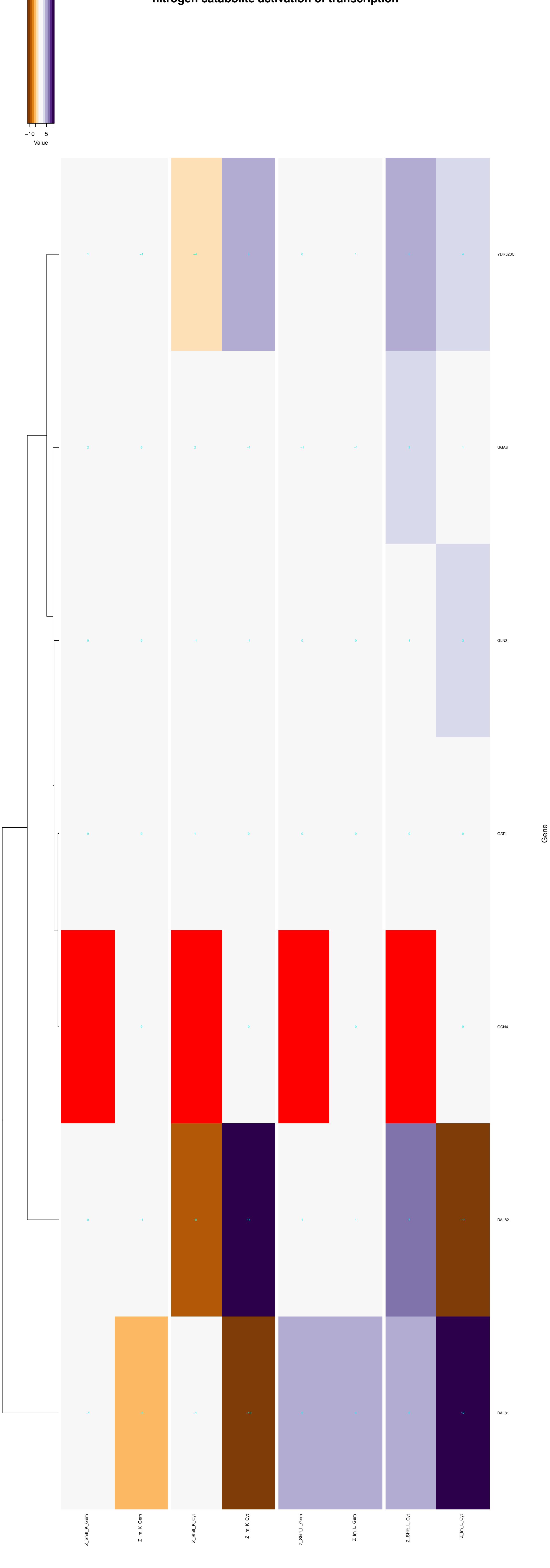
carbon catabolite activation of transcription



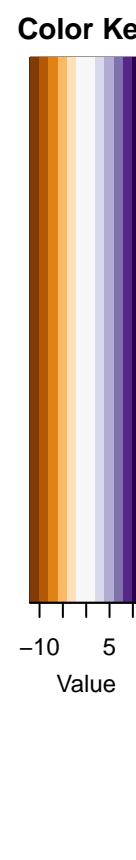
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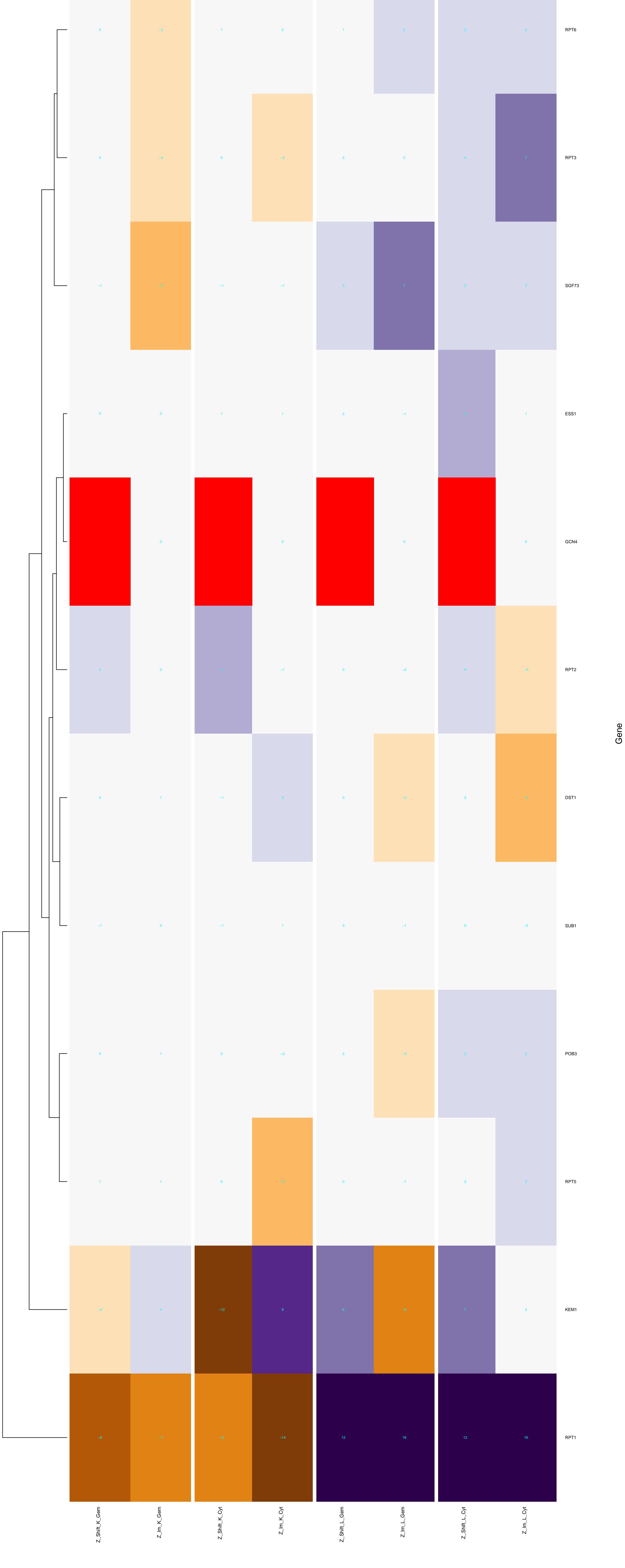
nitrogen catabolite activation of transcription



Color Key



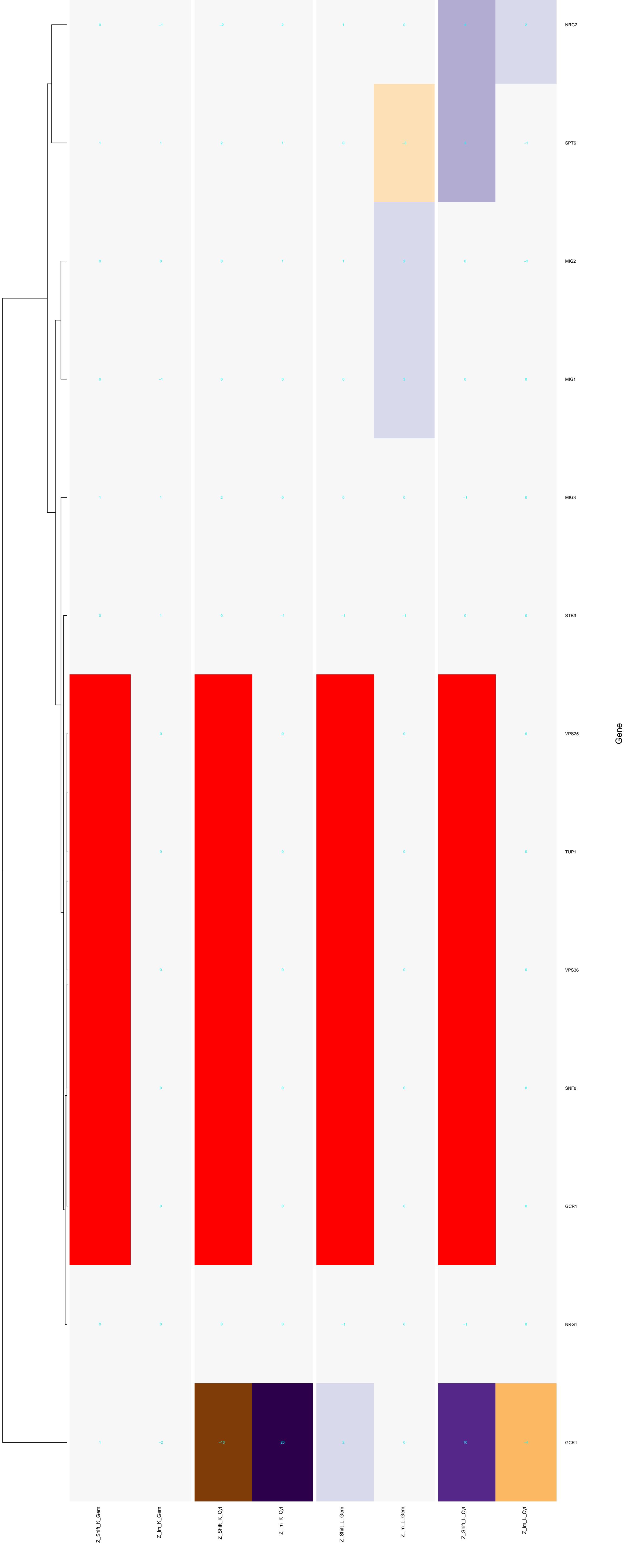
positive regulation of DNA-templated transcription, initiation



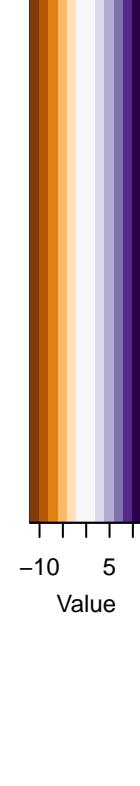
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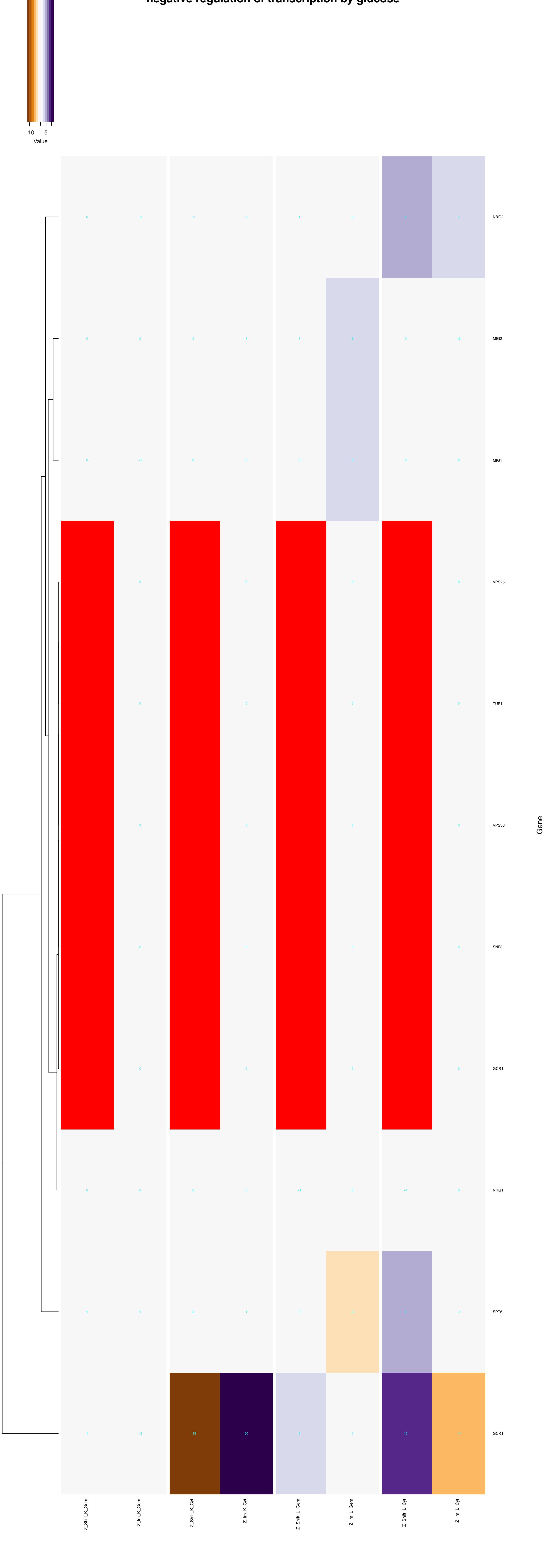
regulation of transcription from RNA polymerase II promoter by glucose



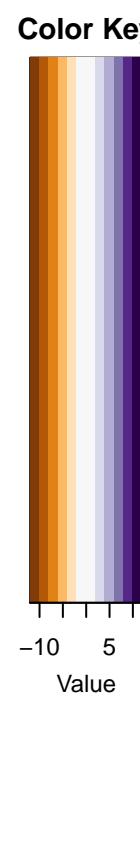
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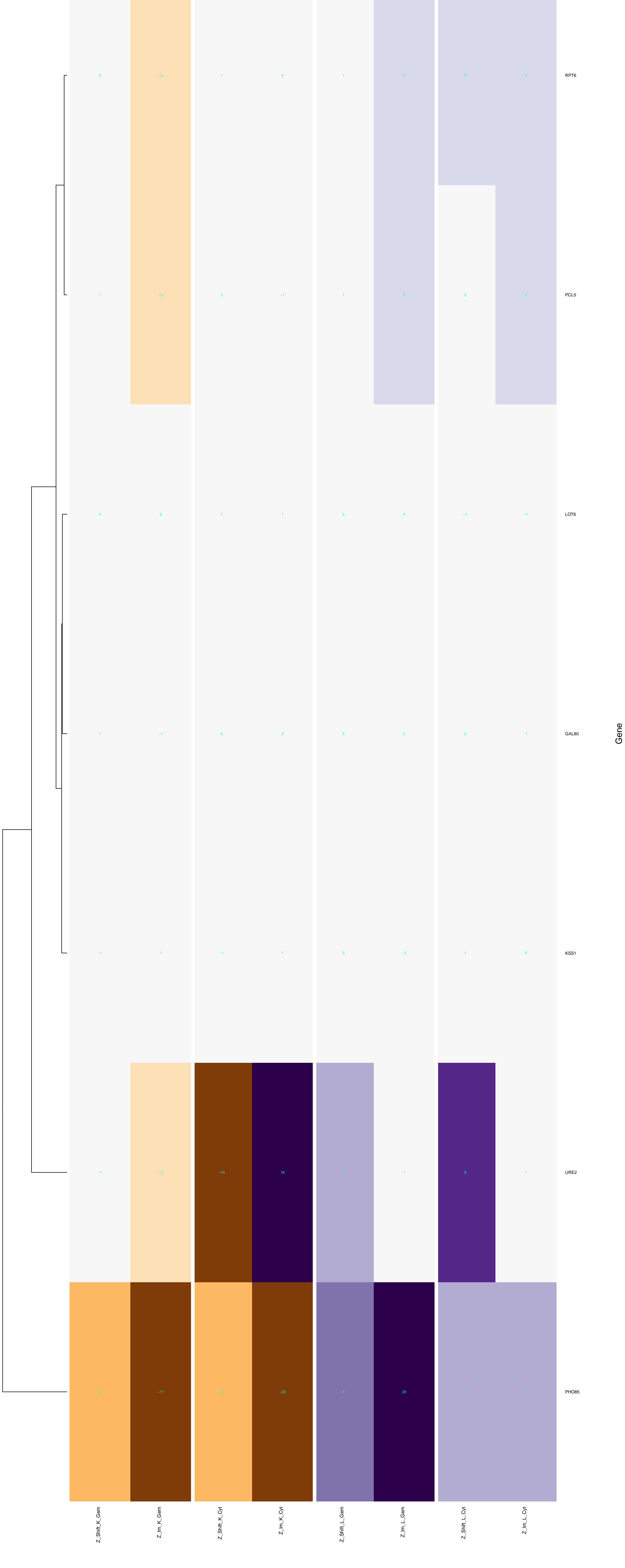
negative regulation of transcription by glucose



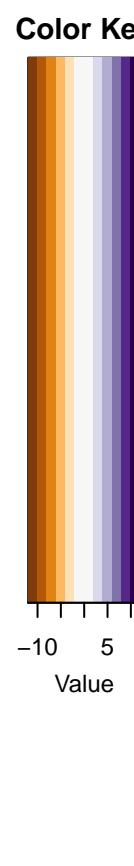
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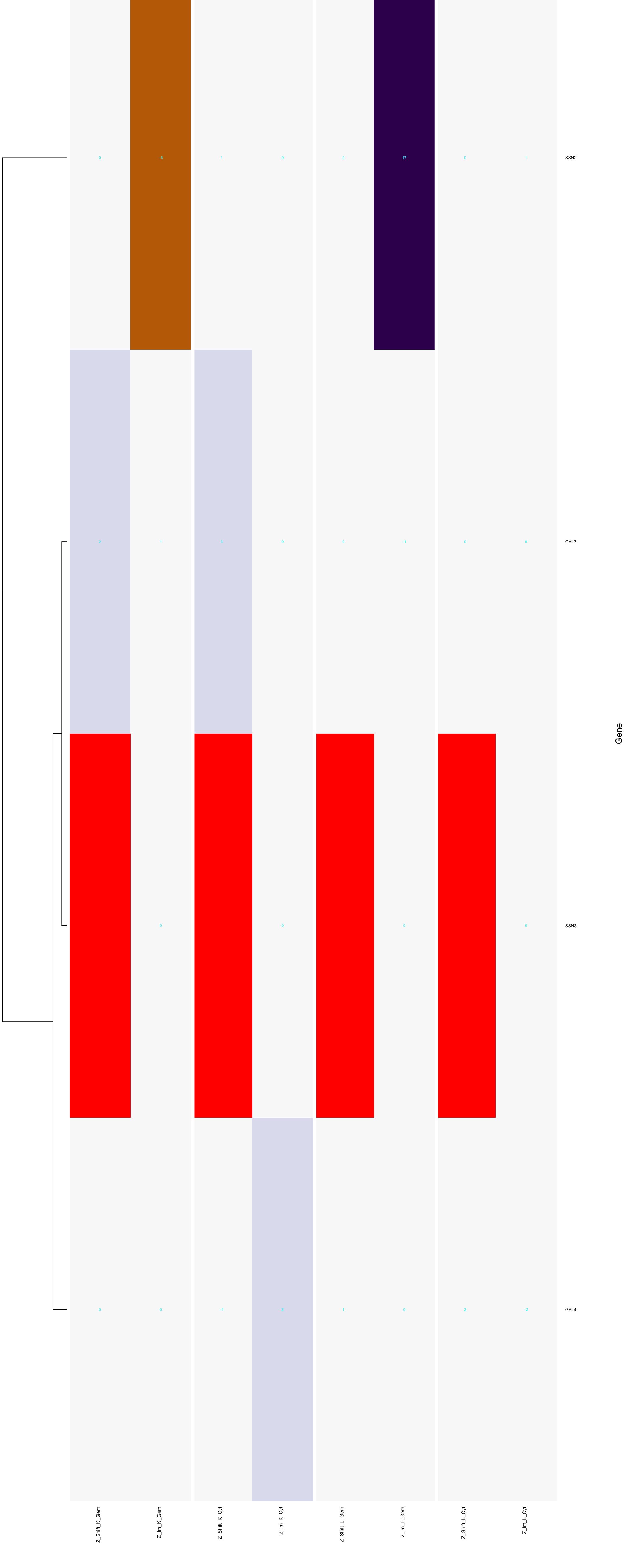
Negative regulation of sequence-specific DNA binding transcription factor activity



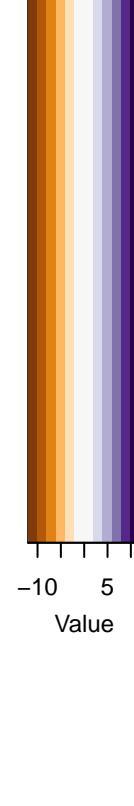
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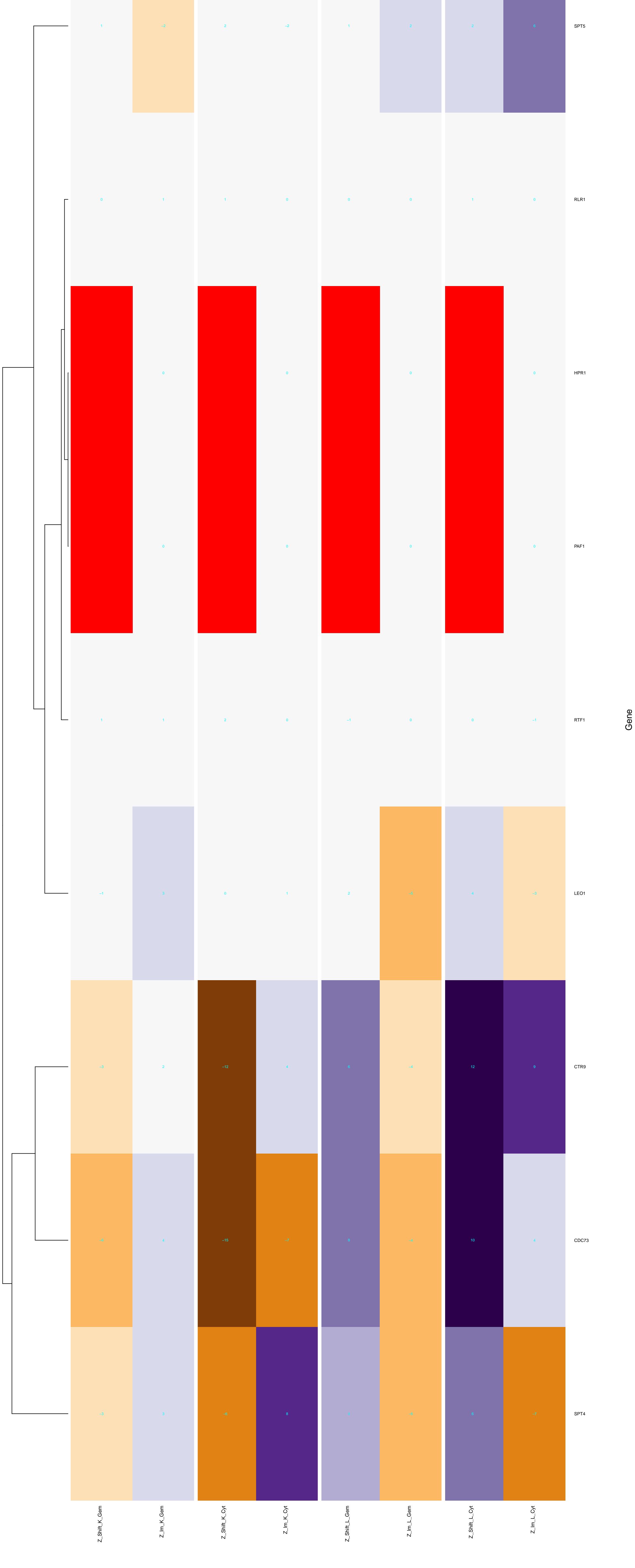
Positive regulation of transcription from RNA polymerase II promoter by galactose



Color Key



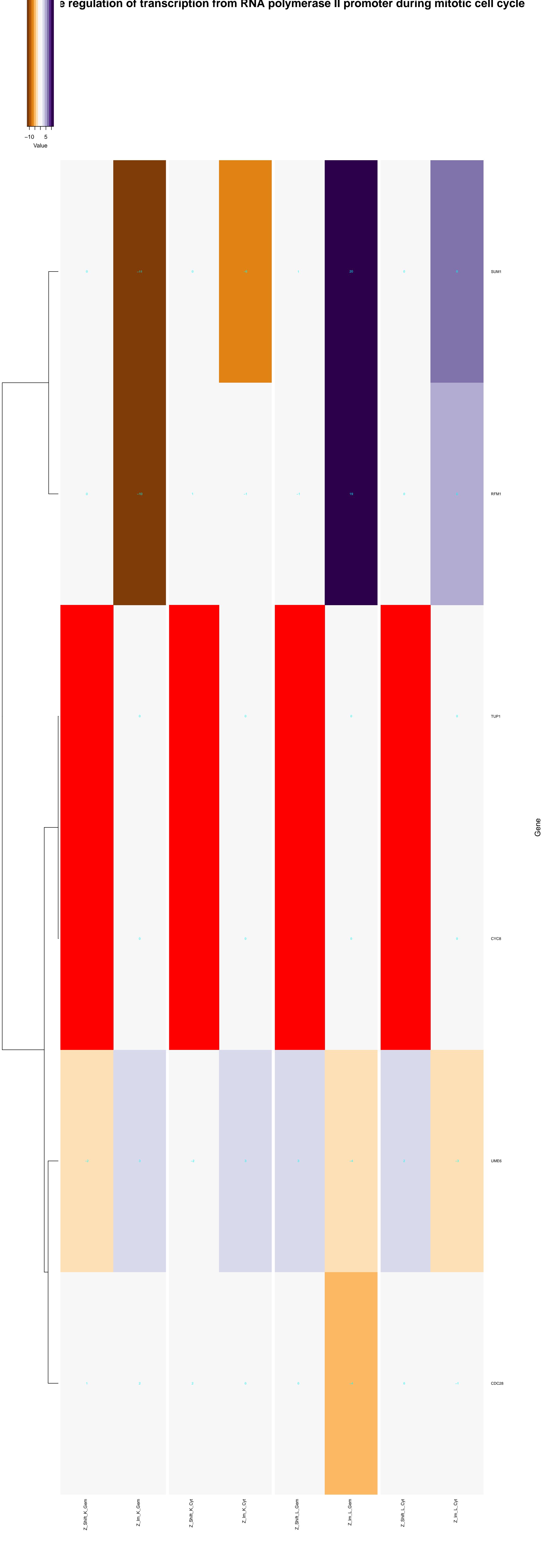
Positive regulation of transcription elongation from RNA polymerase I promoter



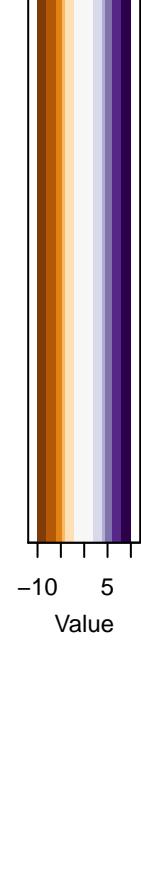
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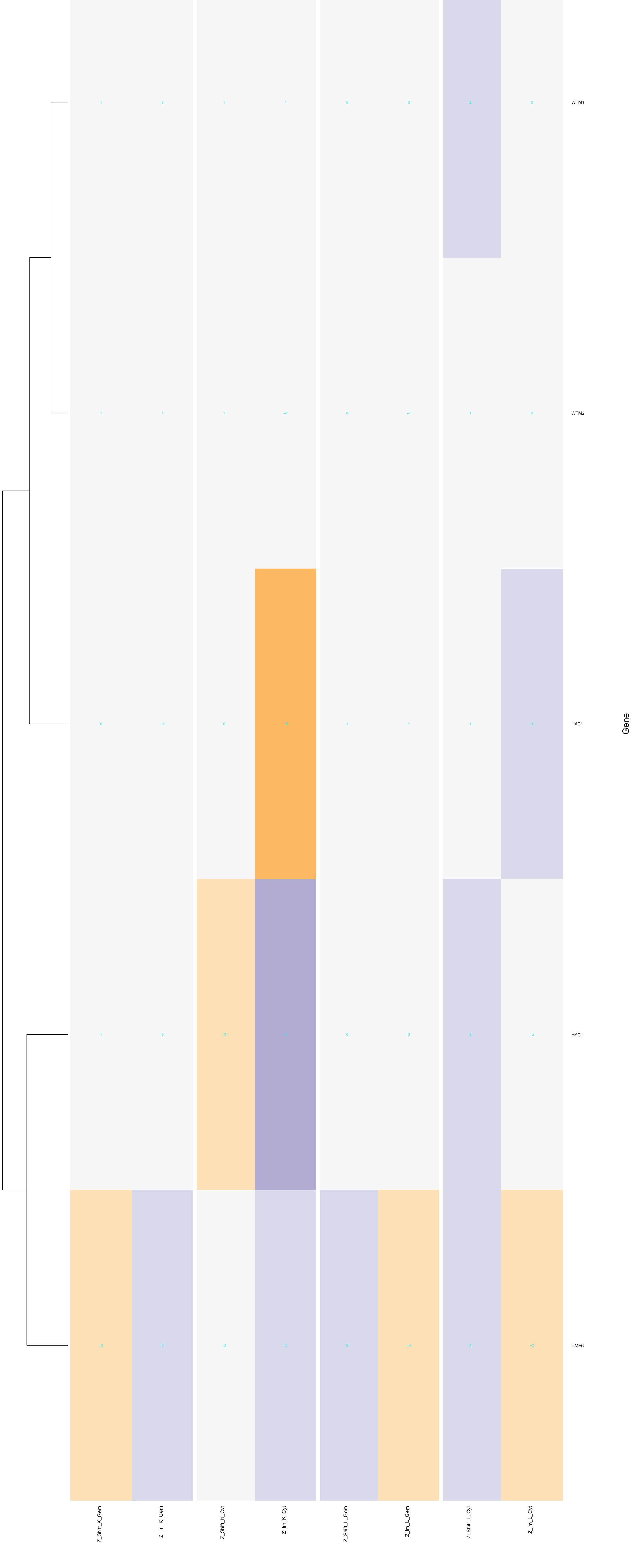
Regulation of transcription from RNA polymerase II promoter during mitotic cell cycle



Color Key



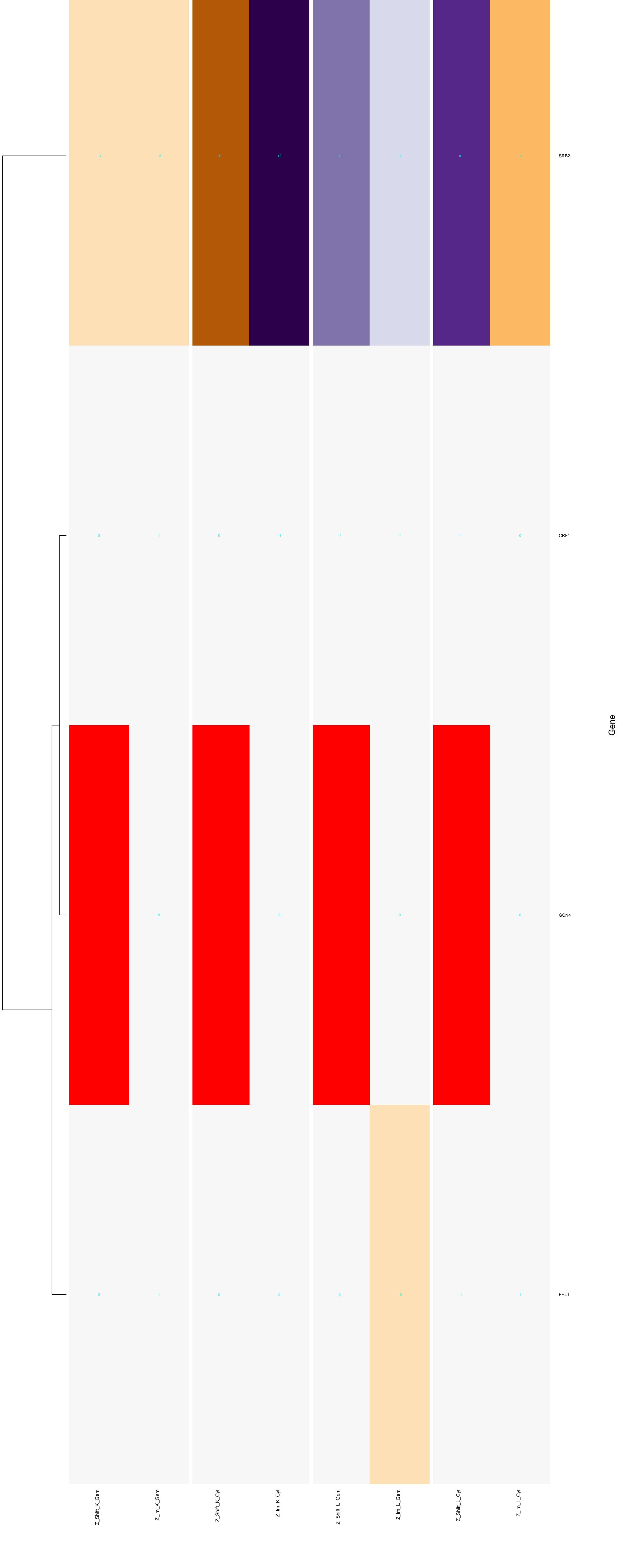
regulation of transcription from RNA polymerase II promoter involved in meiotic cell cycle

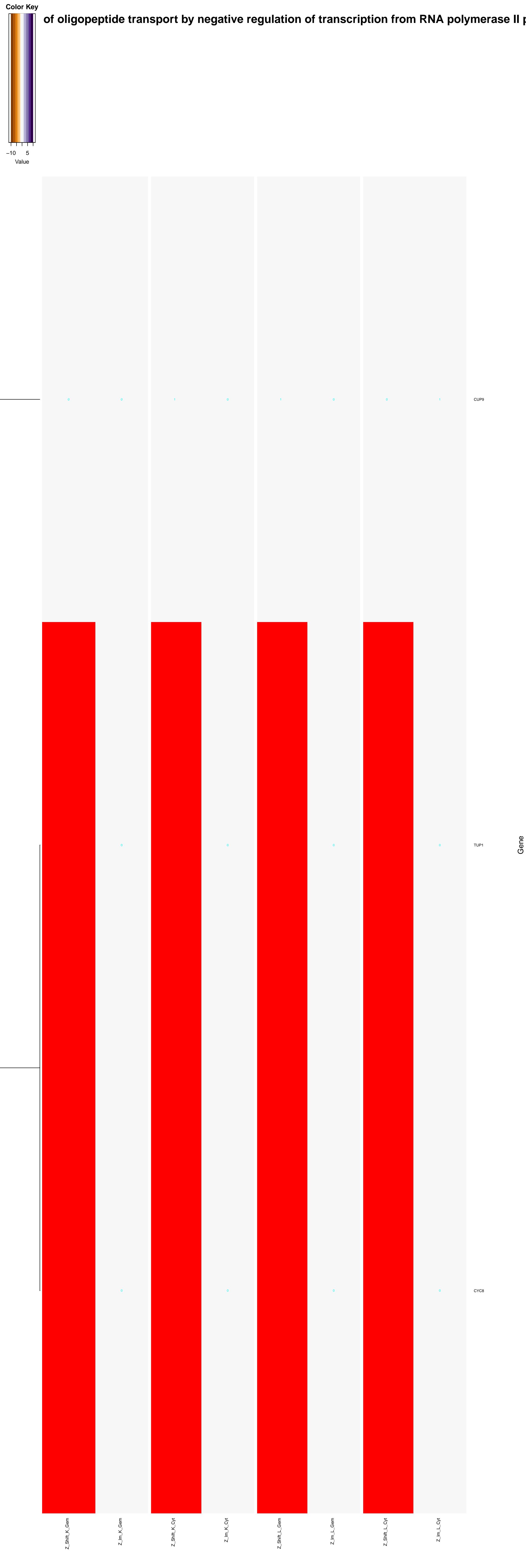


Color Key



e regulation of ribosomal protein gene transcription from RNA polymerase II promoter

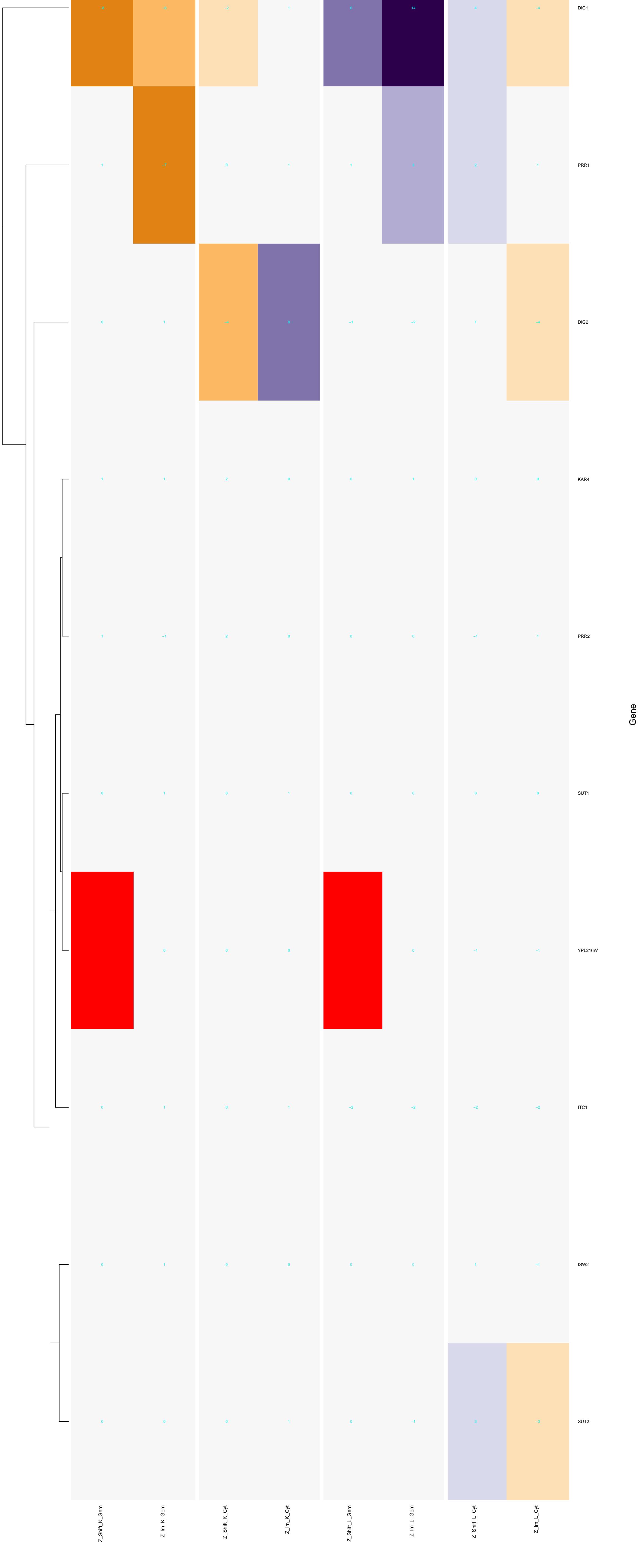




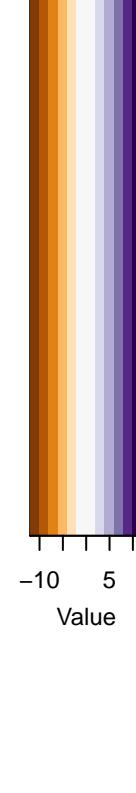
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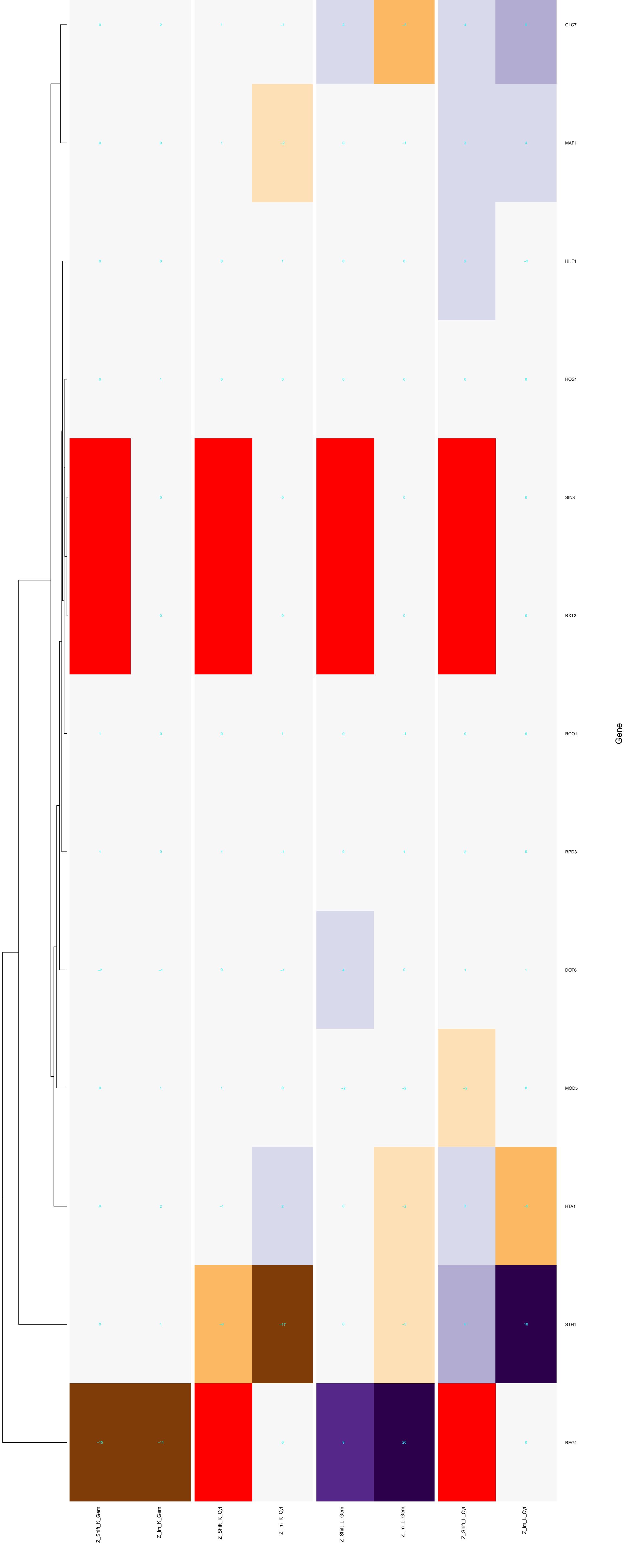
Positive regulation of transcription from RNA polymerase II promoter by pheromones



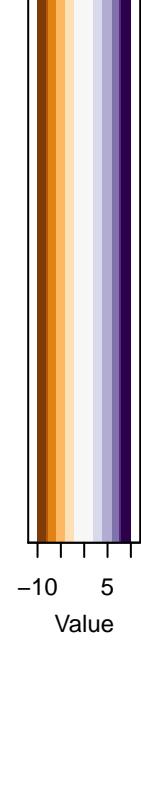
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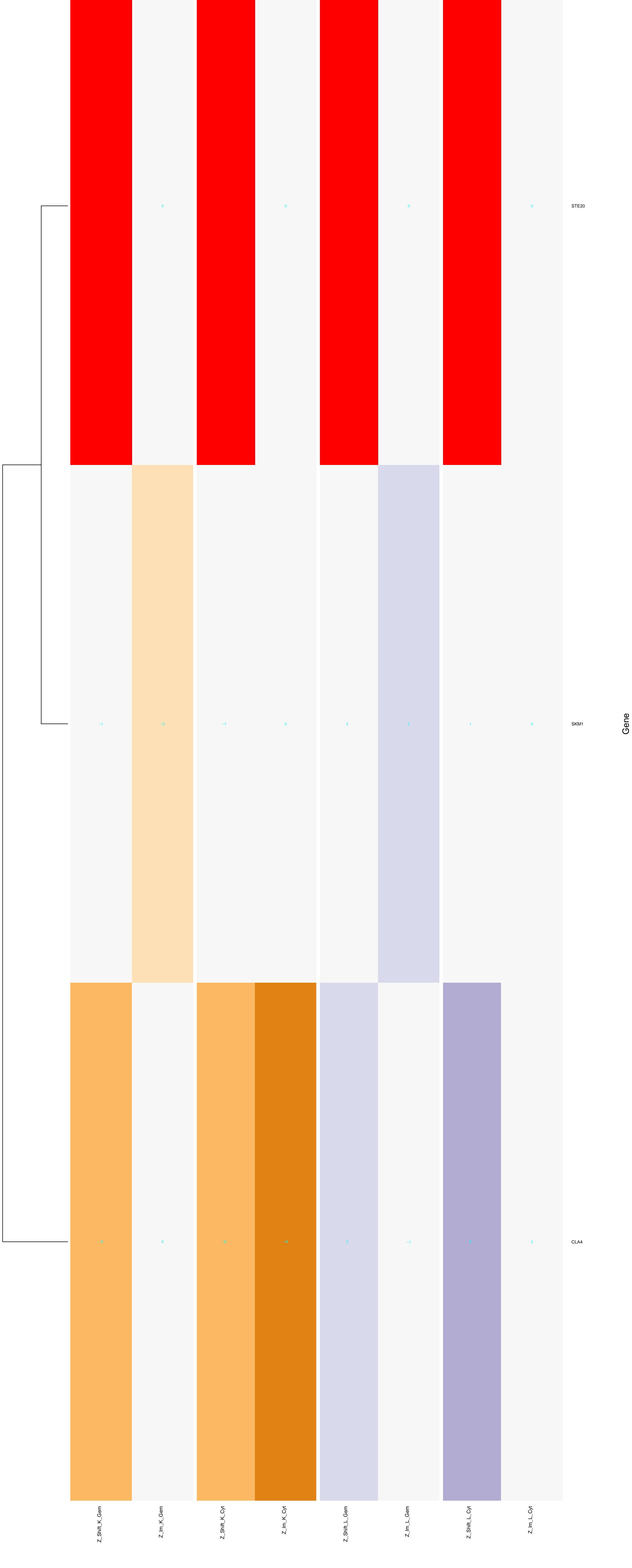
transfer RNA gene-mediated silencing



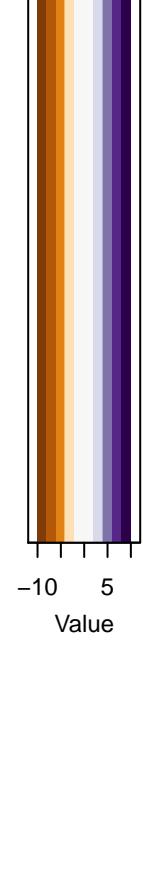
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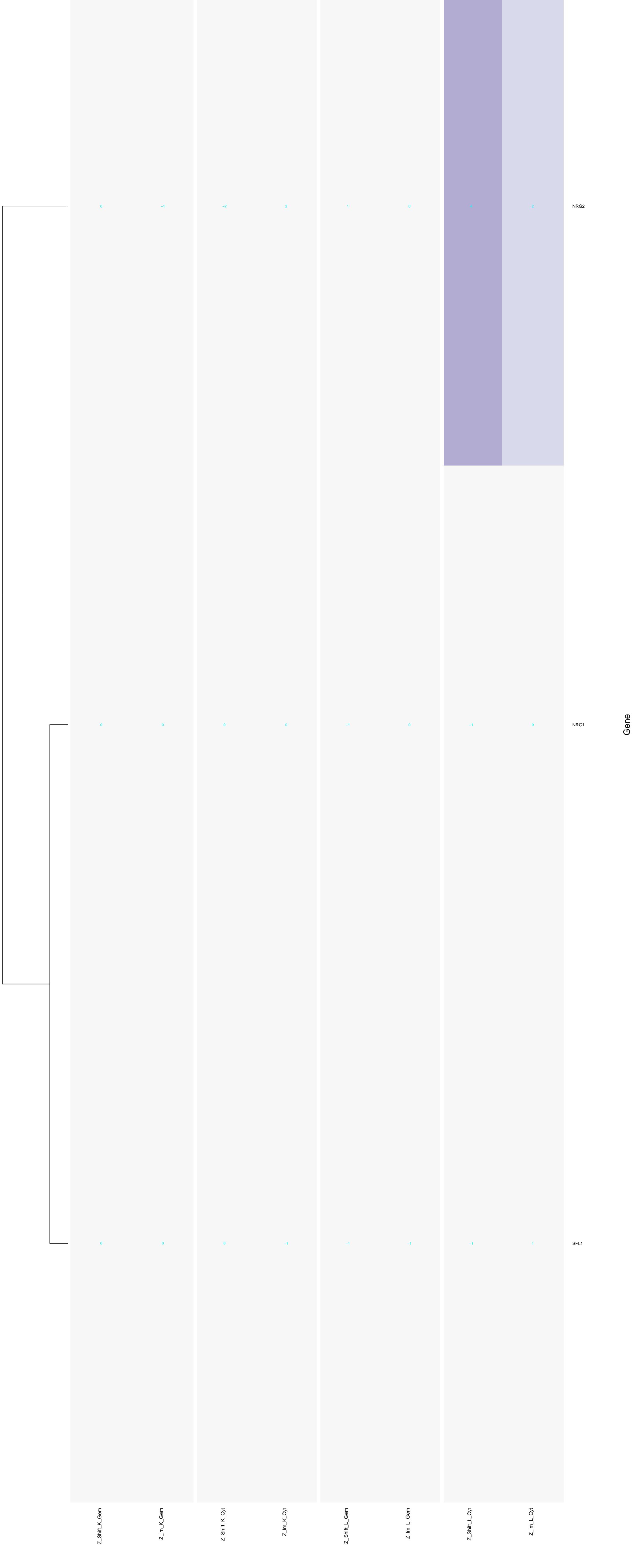
of lipid transport by negative regulation of transcription from RNA polymerase II promoter

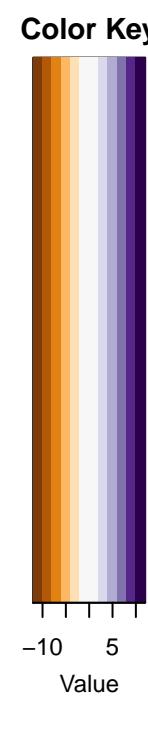


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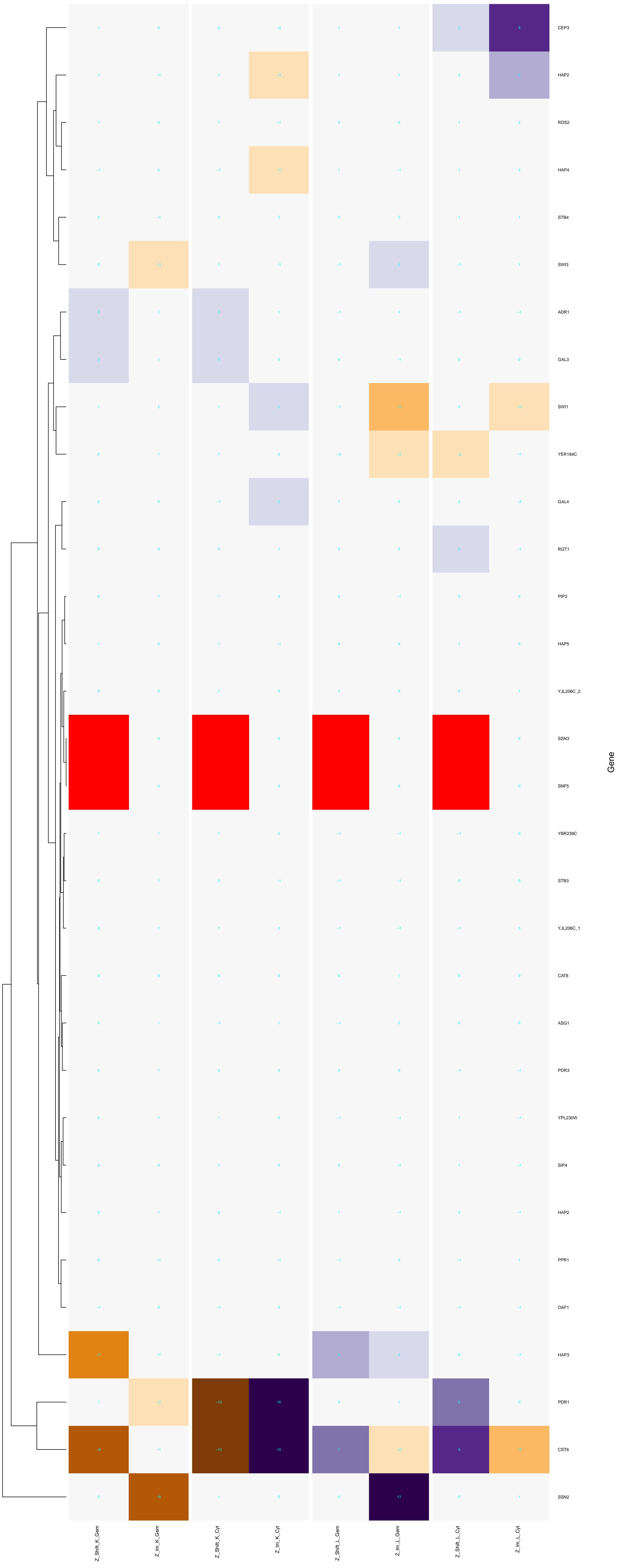


rowth in response to glucose limitation by negative regulation of transcription from RNA p





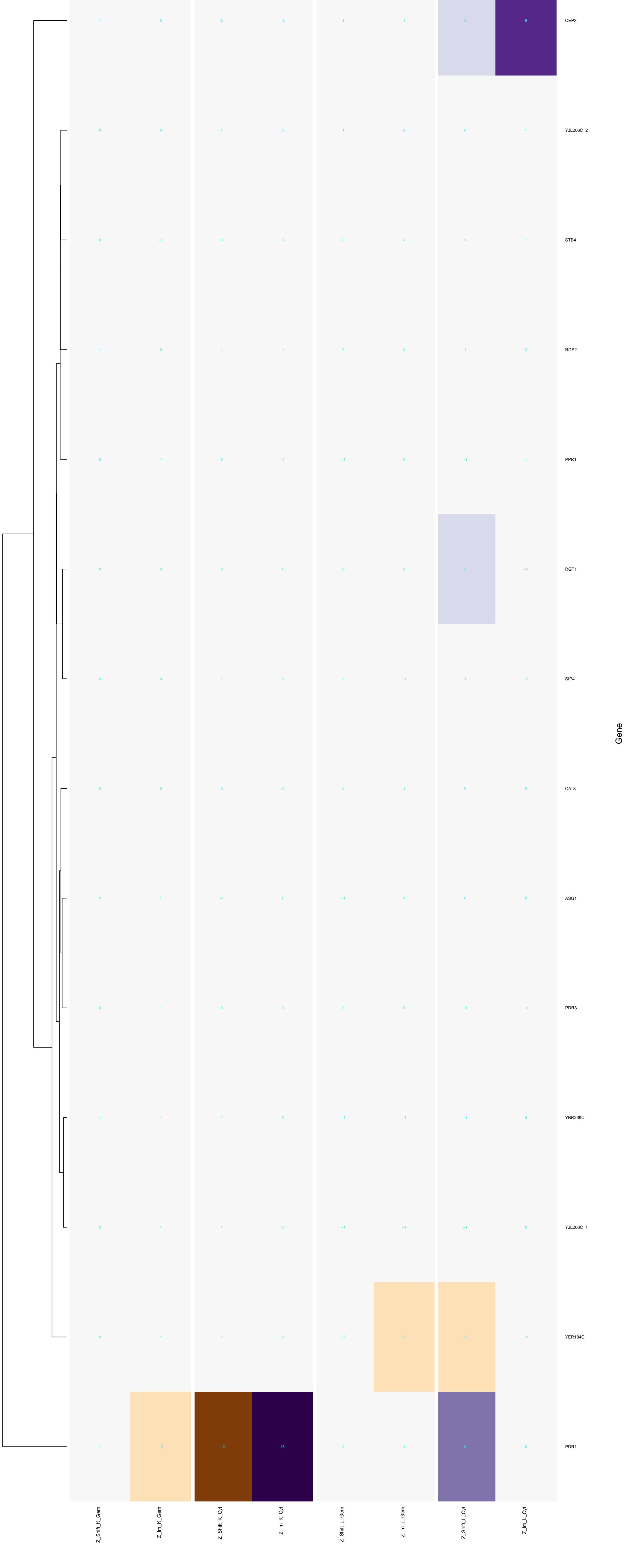
carbon catabolite activation of transcription from RNA polymerase II promoter



Color Key



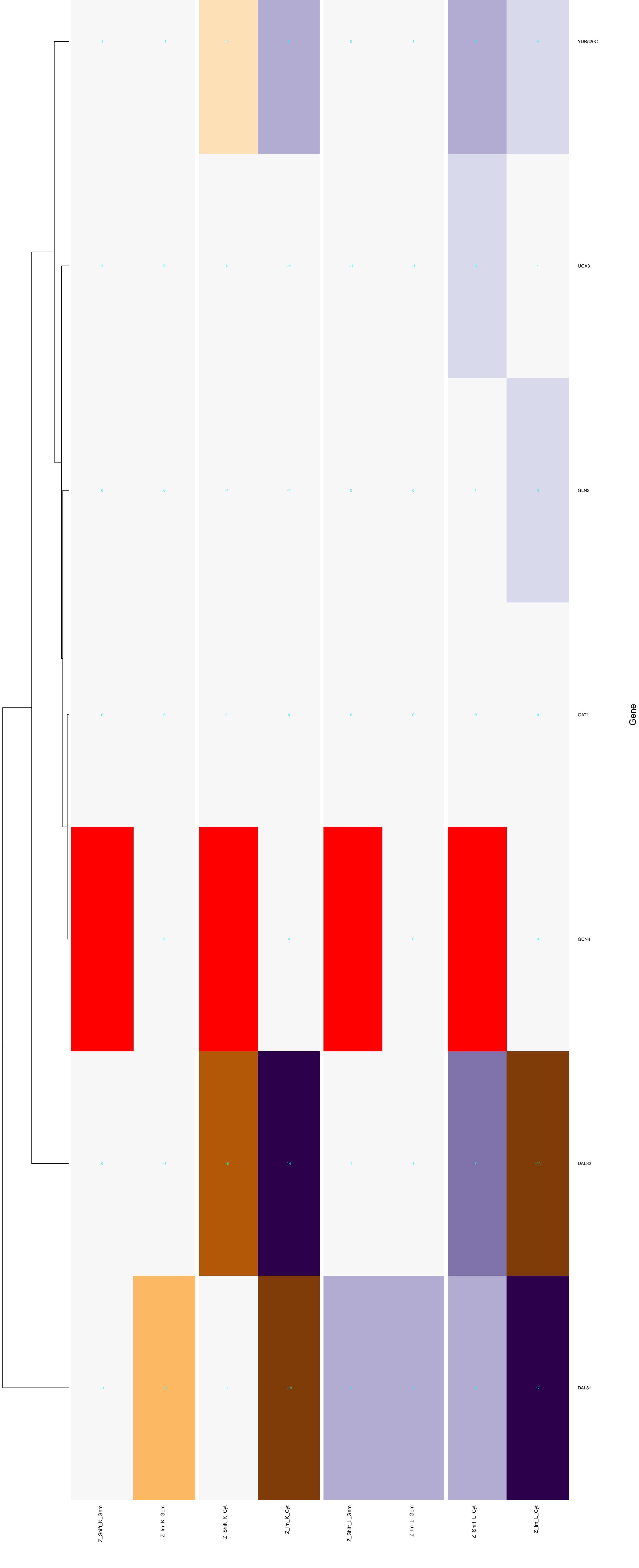
n of transcription from RNA polymerase II promoter by a nonfermentable carbon source



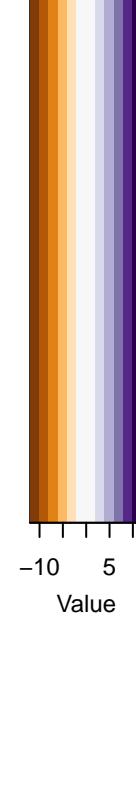
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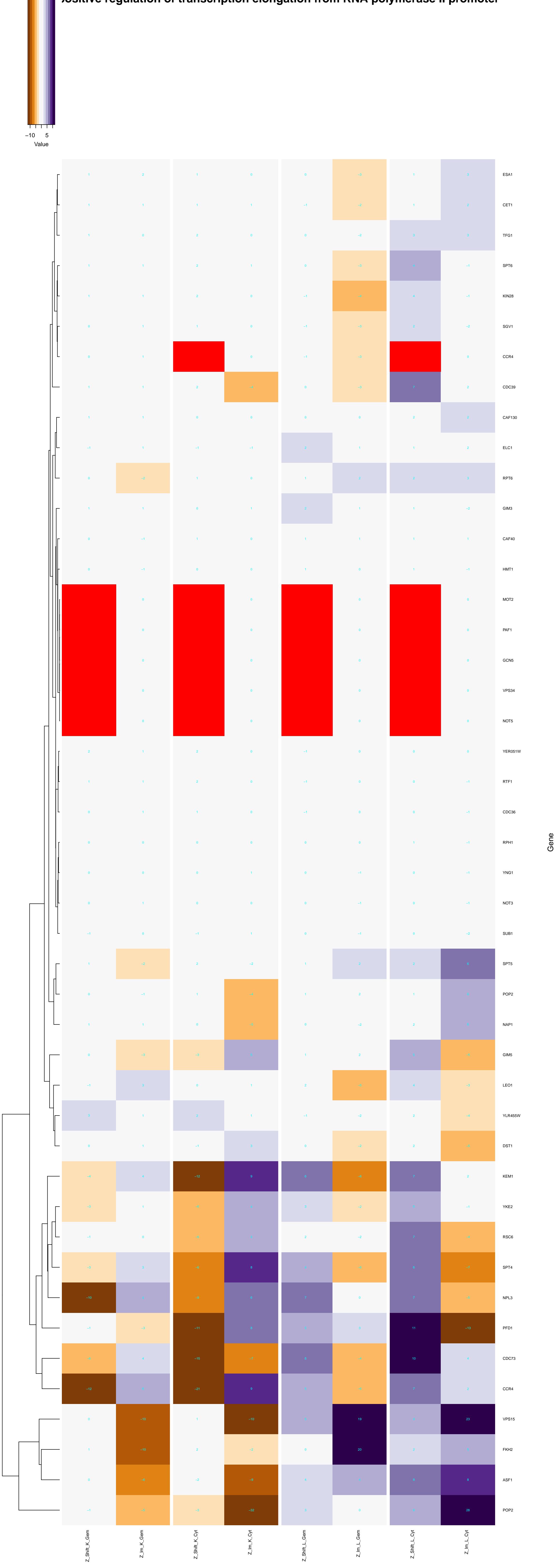
nitrogen catabolite activation of transcription from RNA polymerase II promoter



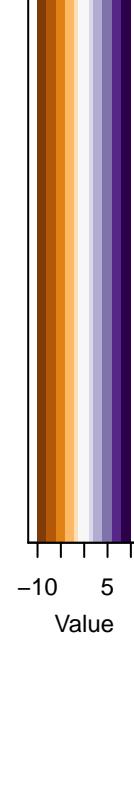
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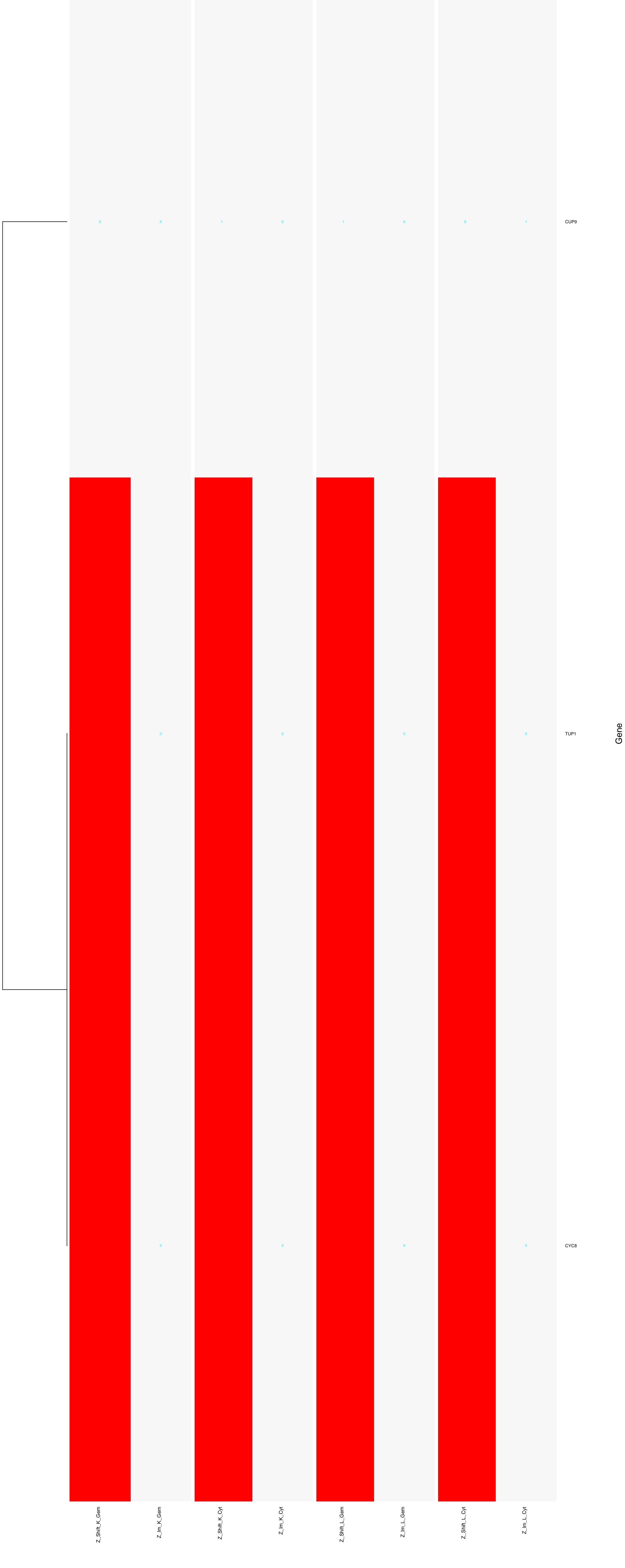
Positive regulation of transcription elongation from RNA polymerase II promoter



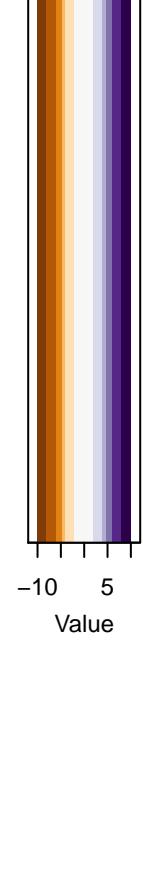
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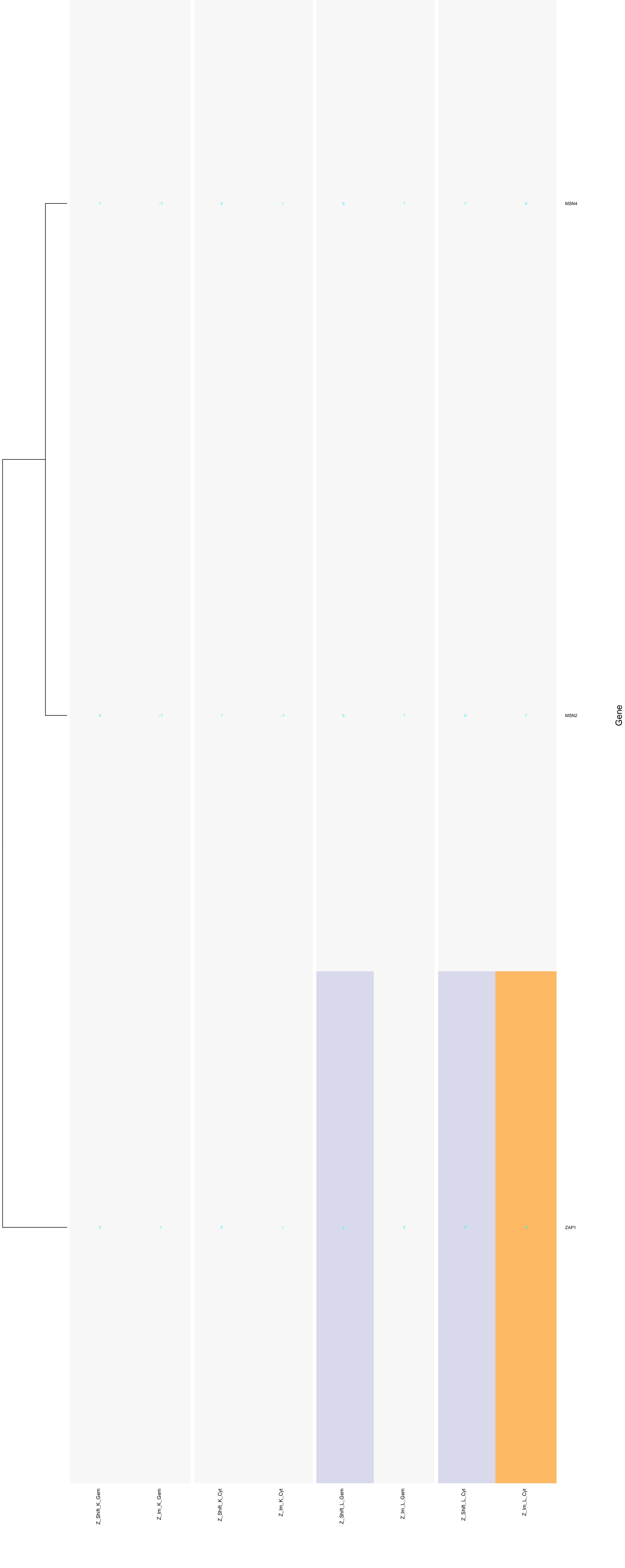
n of dipeptide transport by regulation of transcription from RNA polymerase II promoter



Color Key

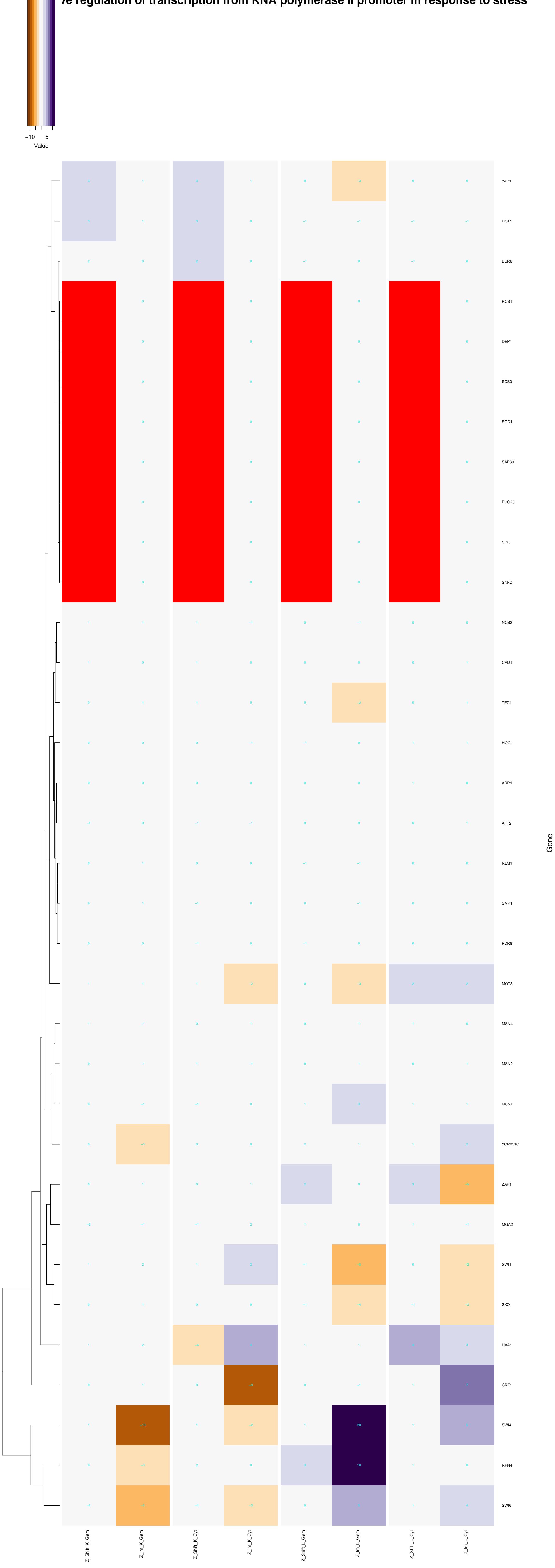


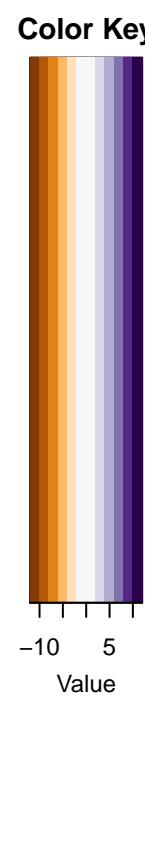
on of transcription from RNA polymerase II promoter in response to zinc ion starvation



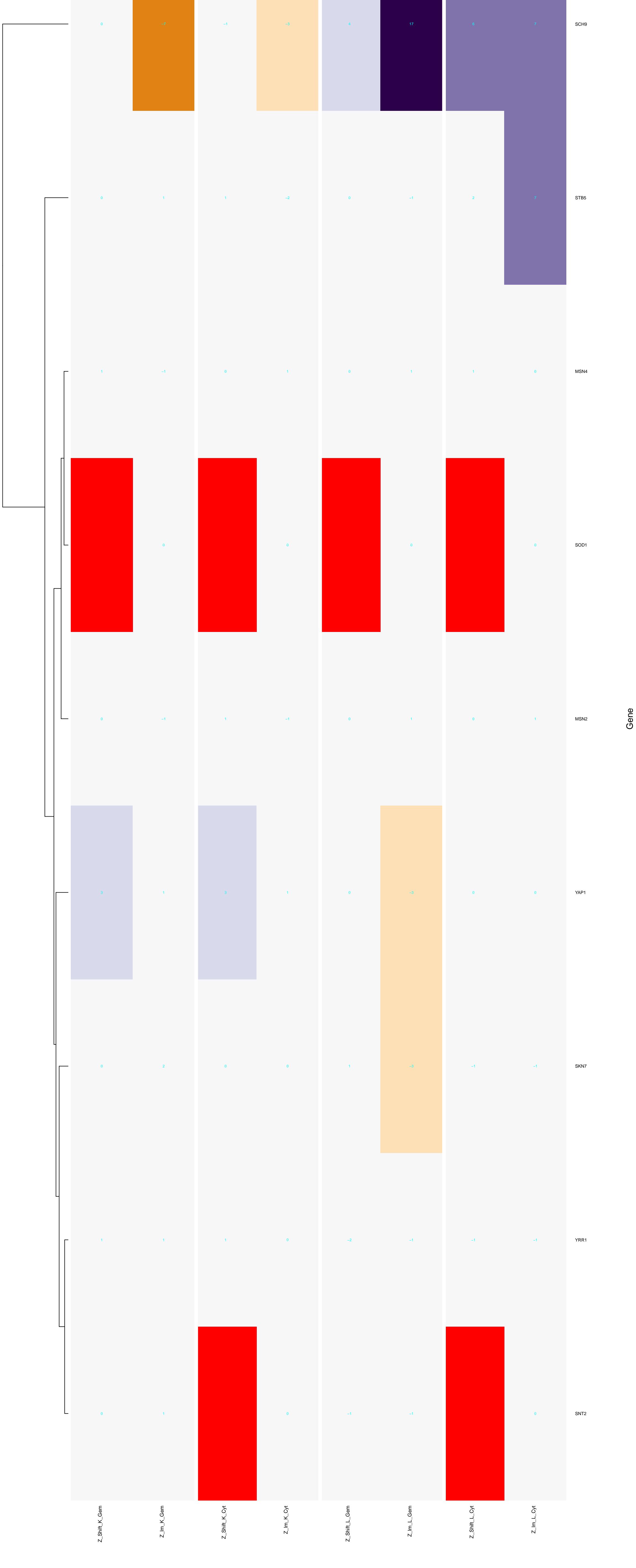
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Positive regulation of transcription from RNA polymerase II promoter in response to stress

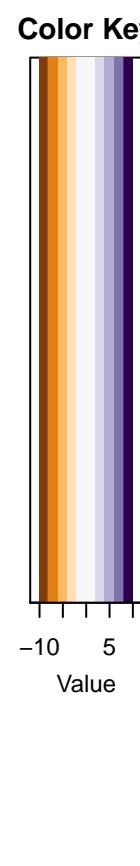




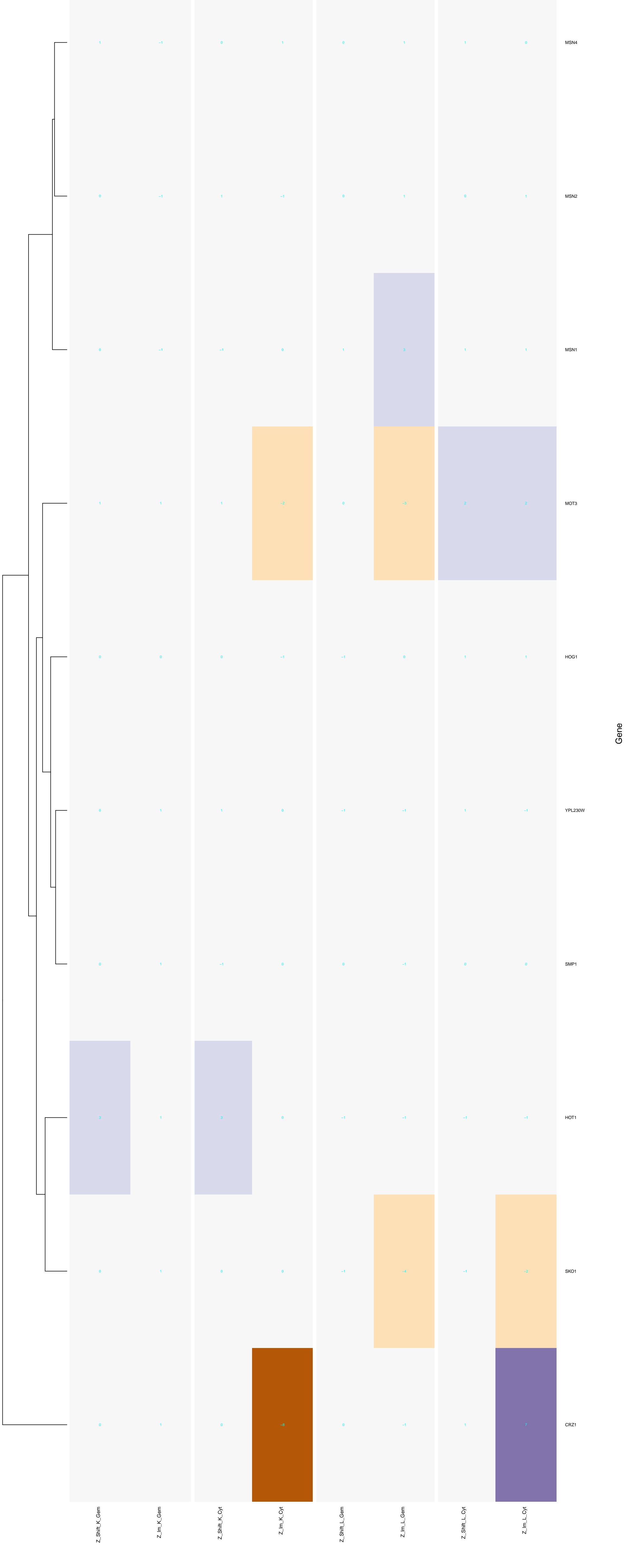
Induction of transcription from RNA polymerase II promoter in response to oxidative stress



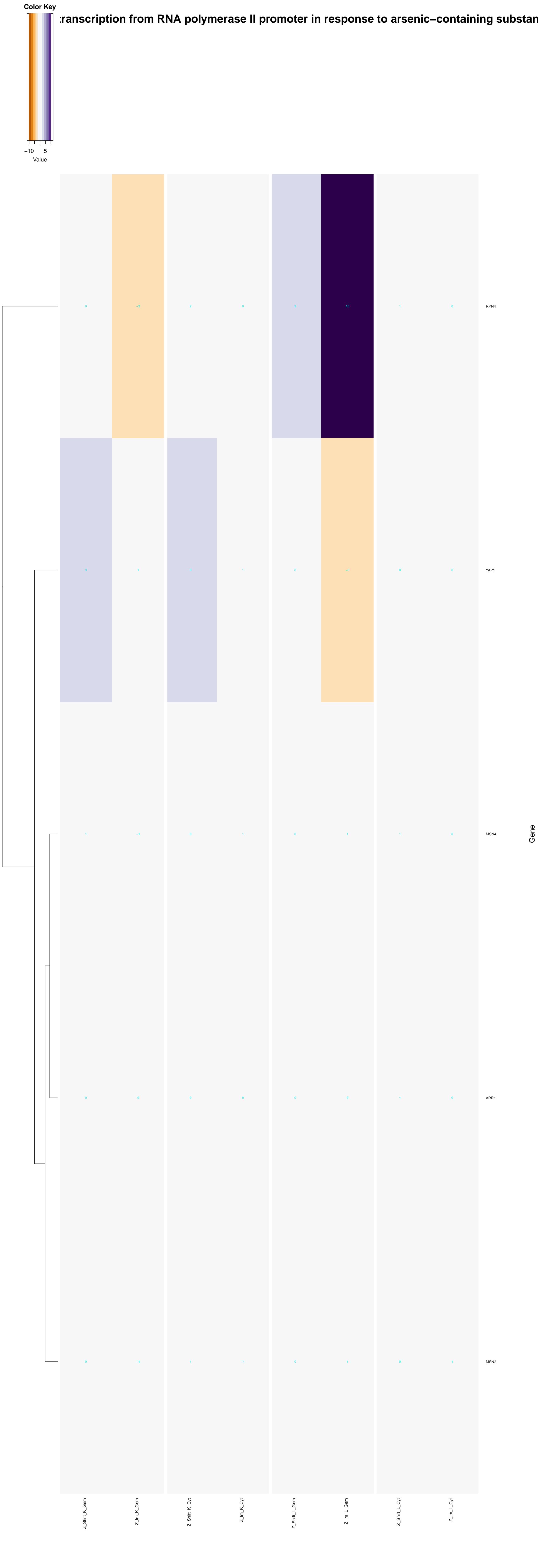
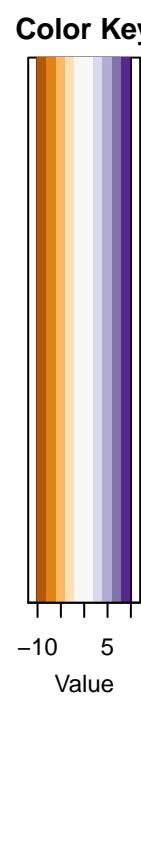
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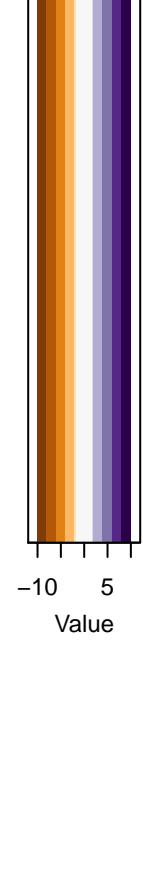
Activation of transcription from RNA polymerase II promoter in response to osmotic stress



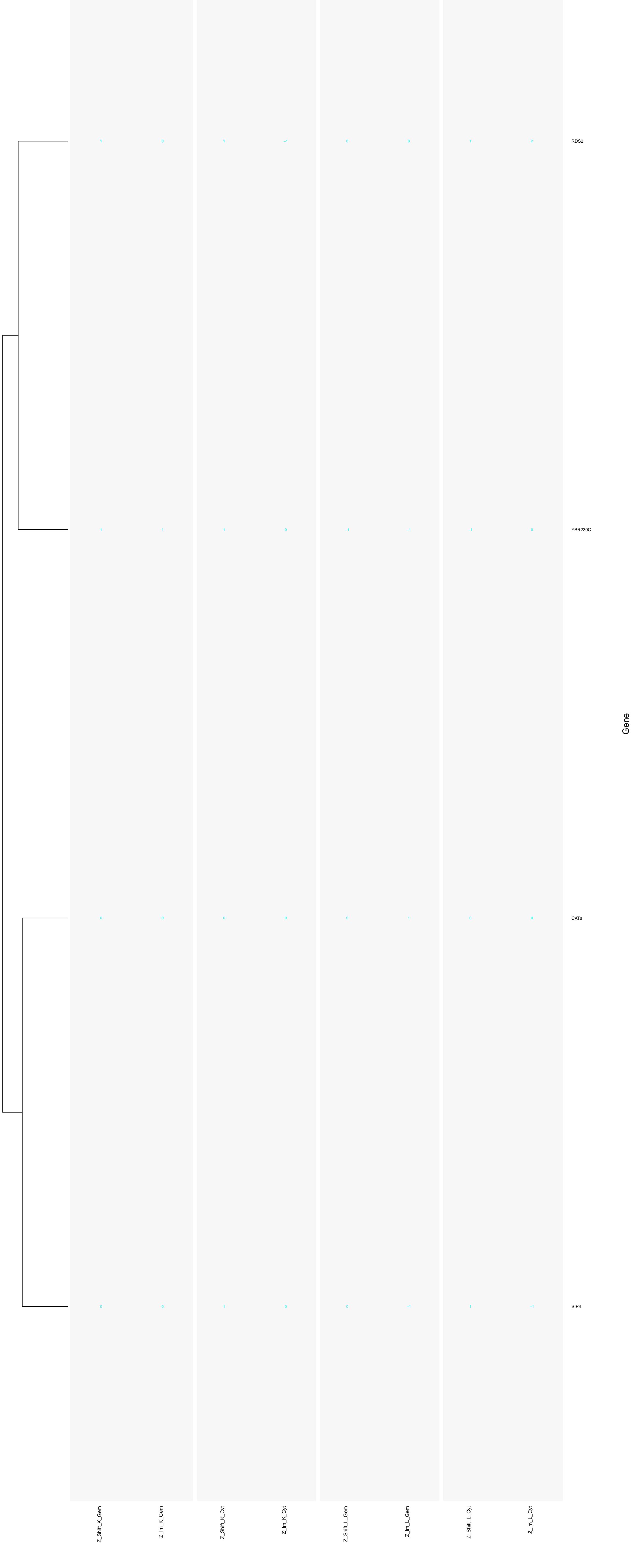
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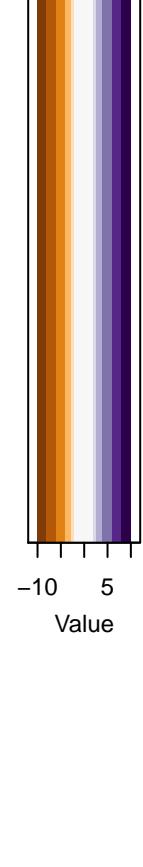
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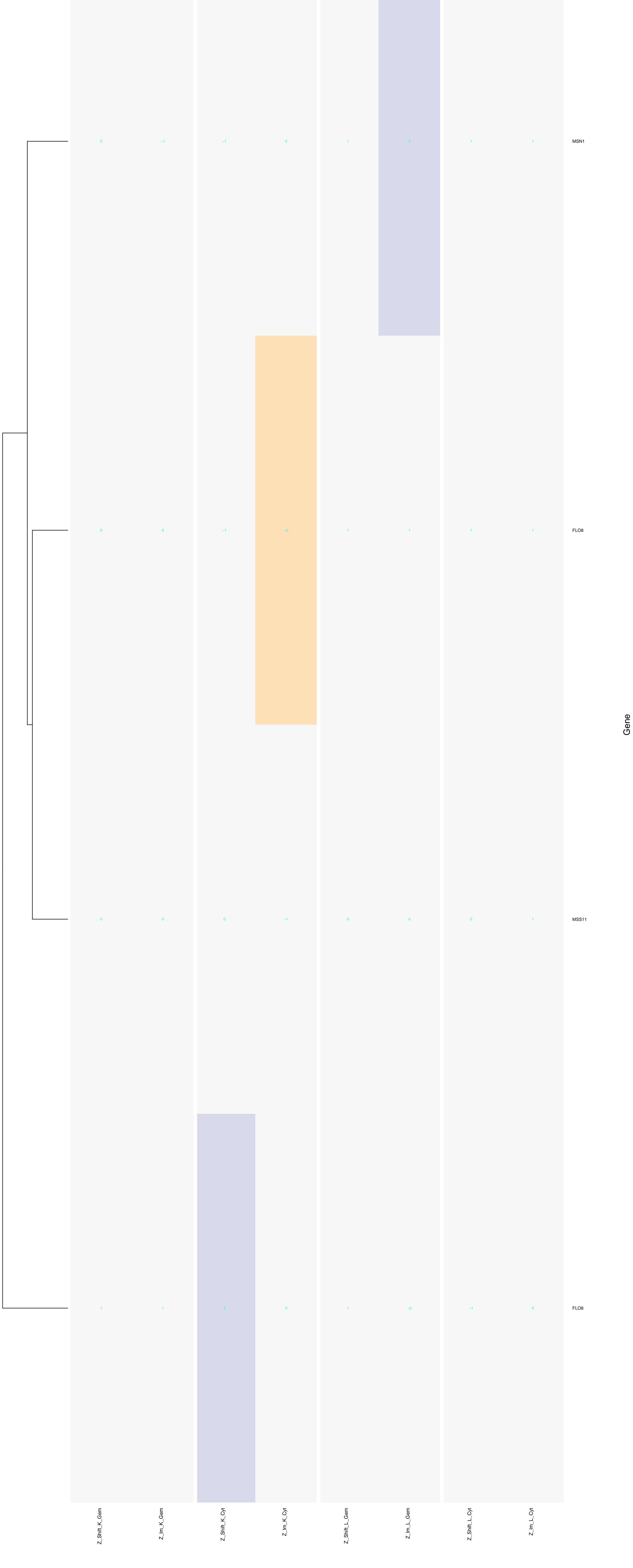
on of gluconeogenesis by positive regulation of transcription from RNA polymerase II pro-



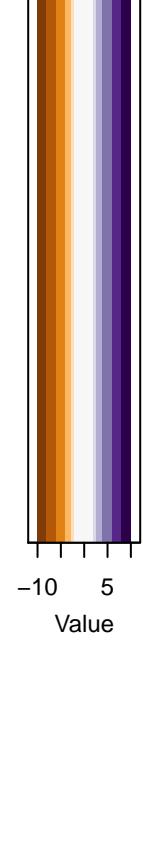
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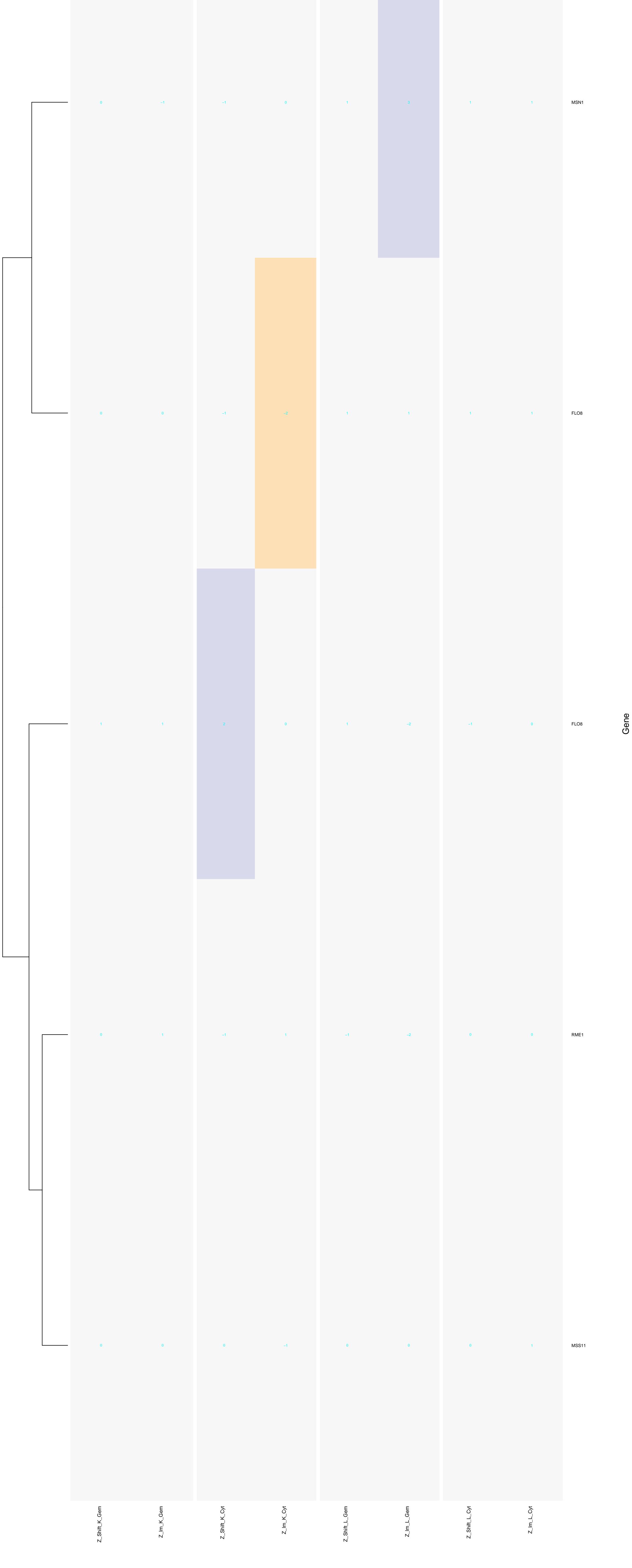
of starch catabolic process by positive regulation of transcription from RNA polymerase II



Color Key



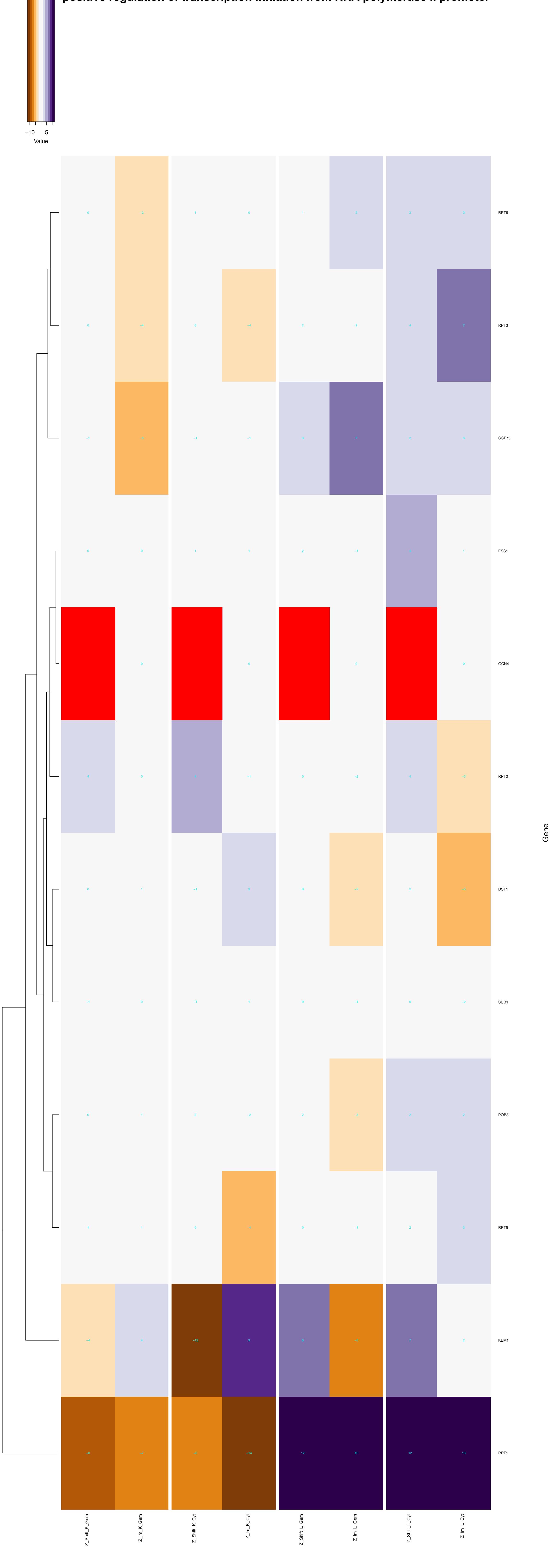
Growth in response to glucose limitation by positive regulation of transcription from RNA pol



Color Key



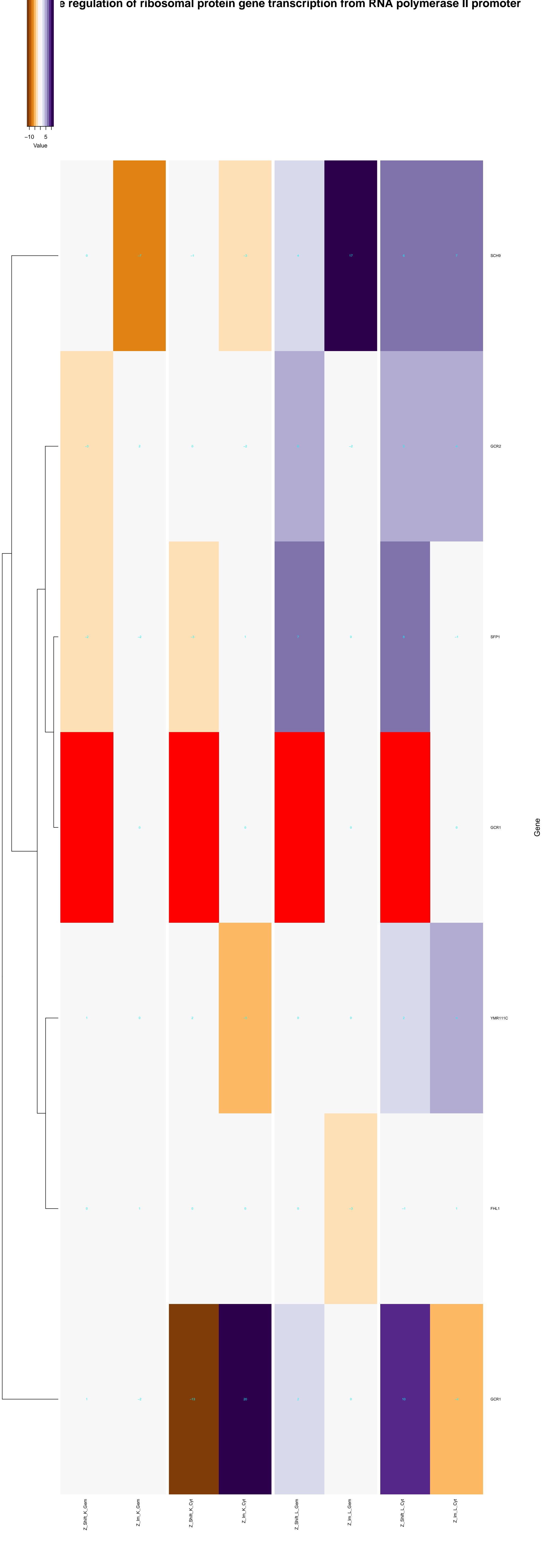
positive regulation of transcription initiation from RNA polymerase II promoter



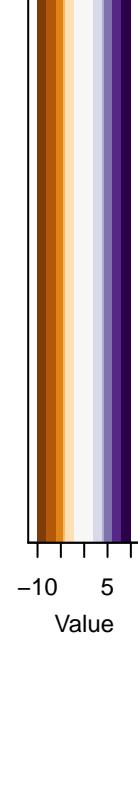
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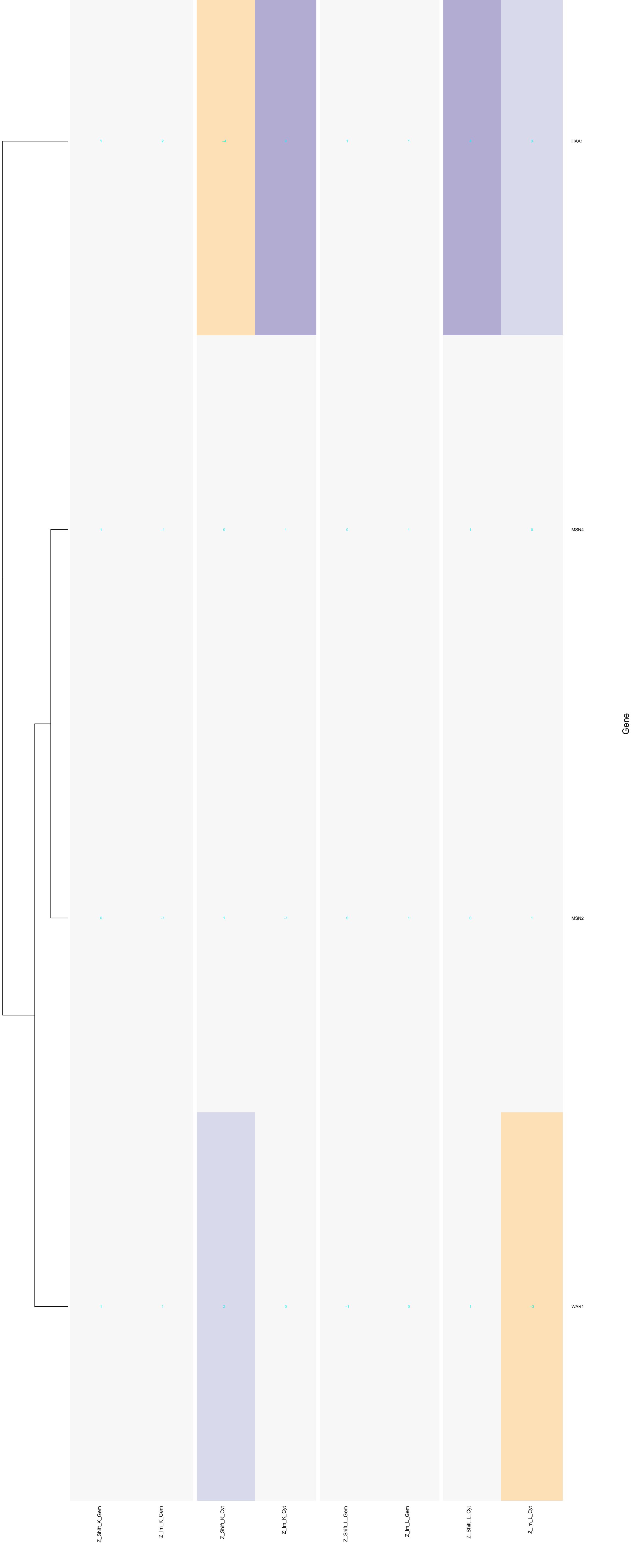
e regulation of ribosomal protein gene transcription from RNA polymerase II promoter



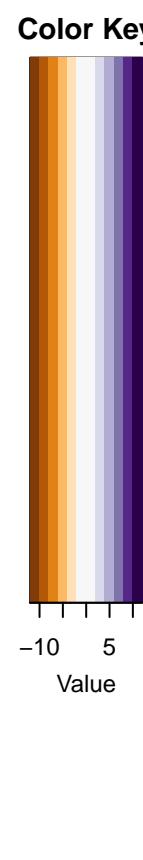
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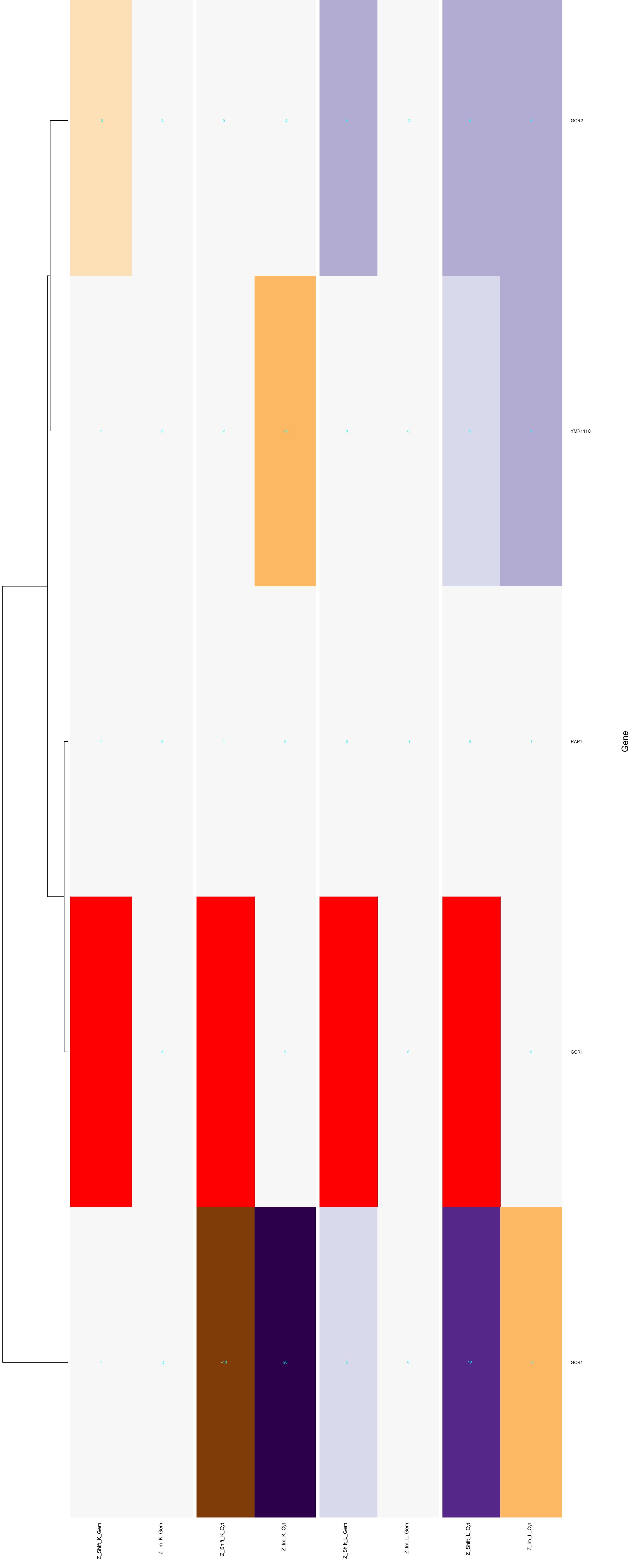
regulation of transcription from RNA polymerase II promoter in response to acidic pH



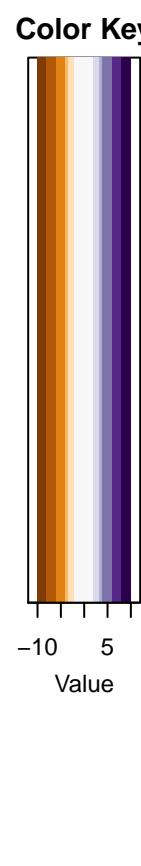
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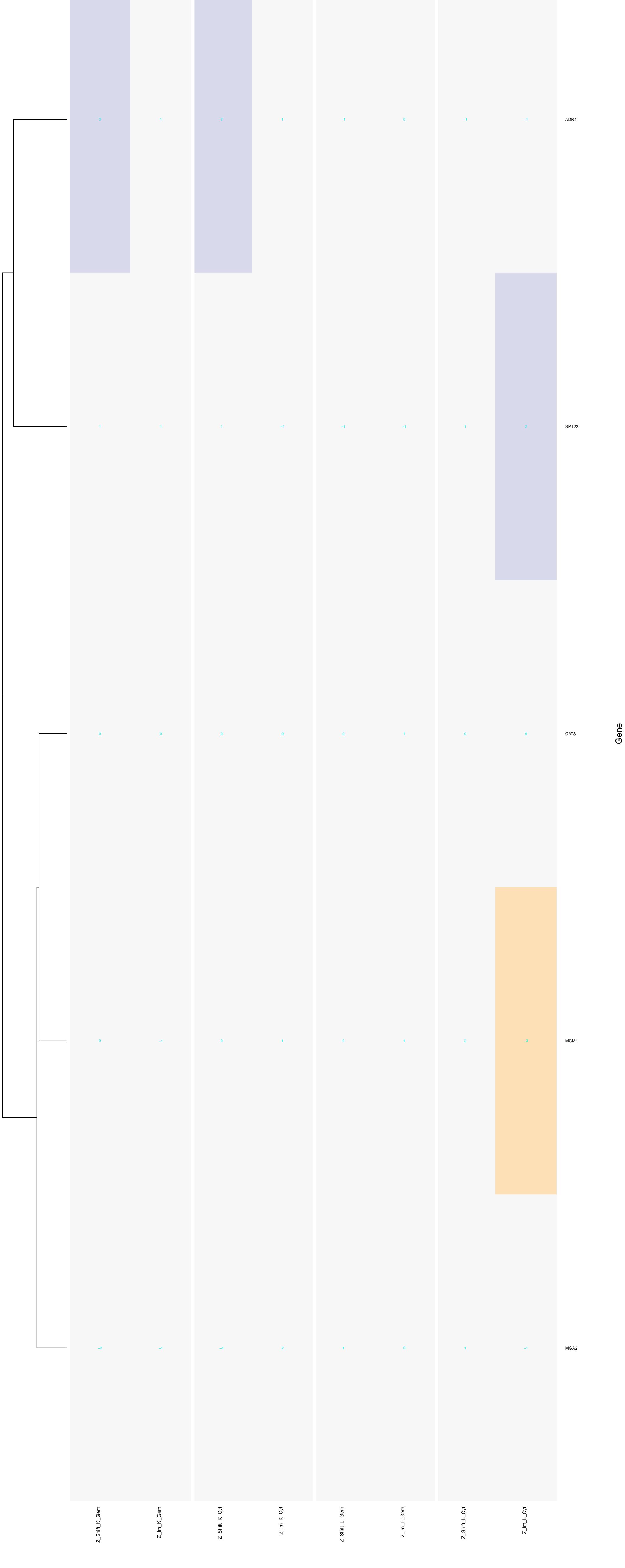
glycolytic process by positive regulation of transcription from RNA polymerase II promoter



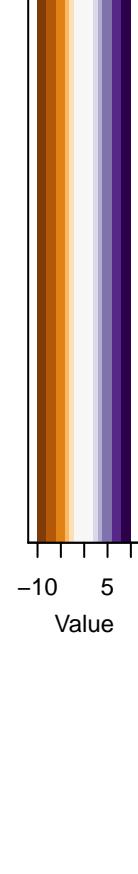
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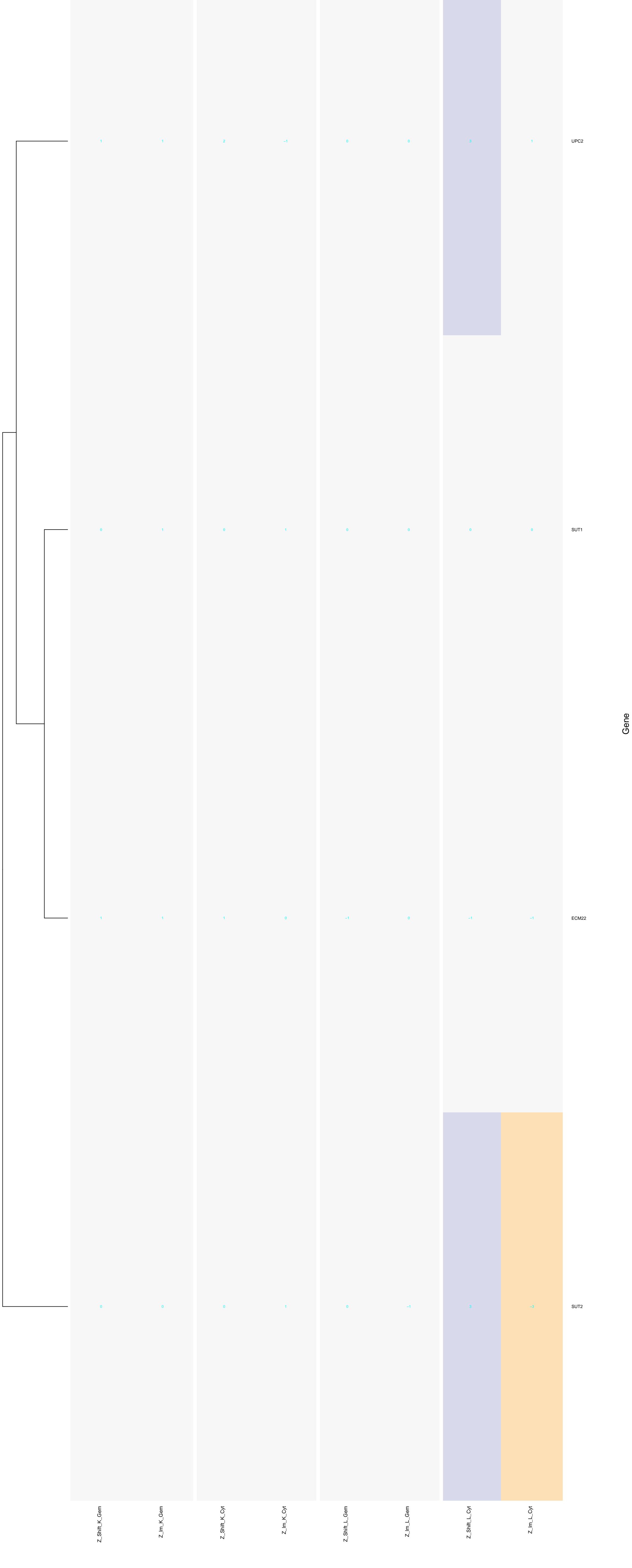
ketone metabolic process by positive regulation of transcription from RNA polymerase II

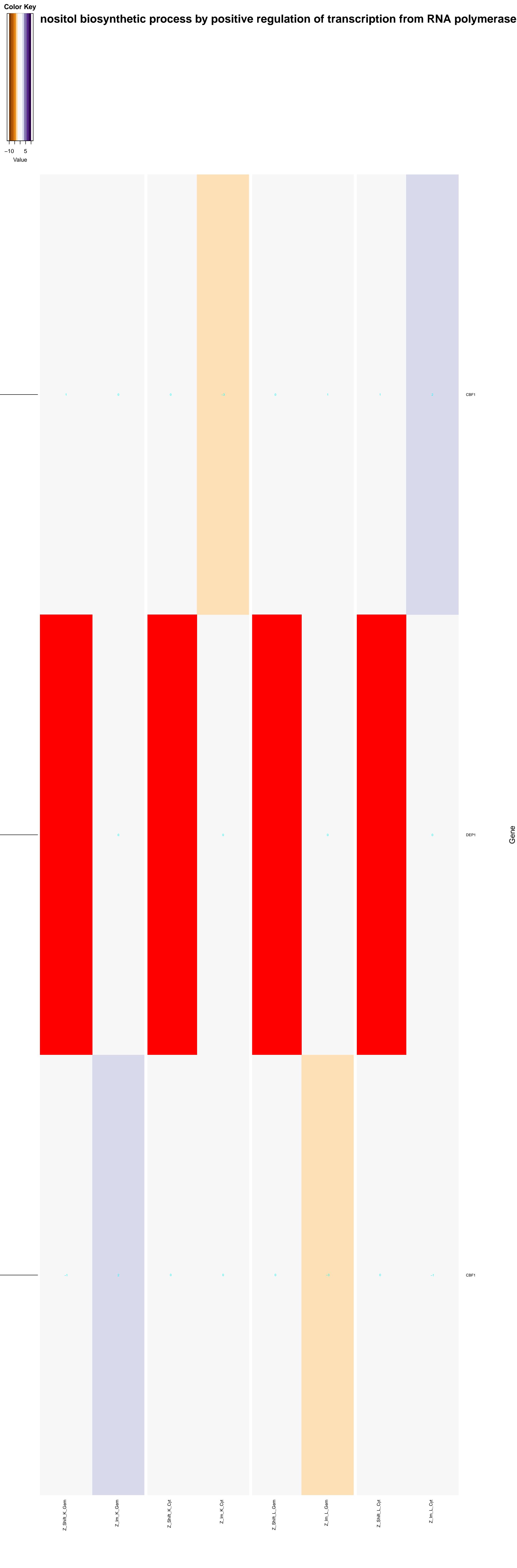


Color Key

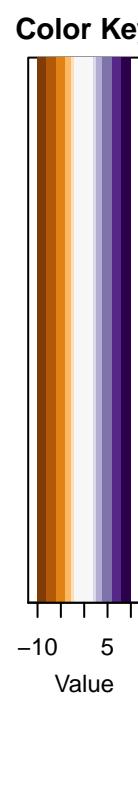


of lipid transport by positive regulation of transcription from RNA polymerase II promoter

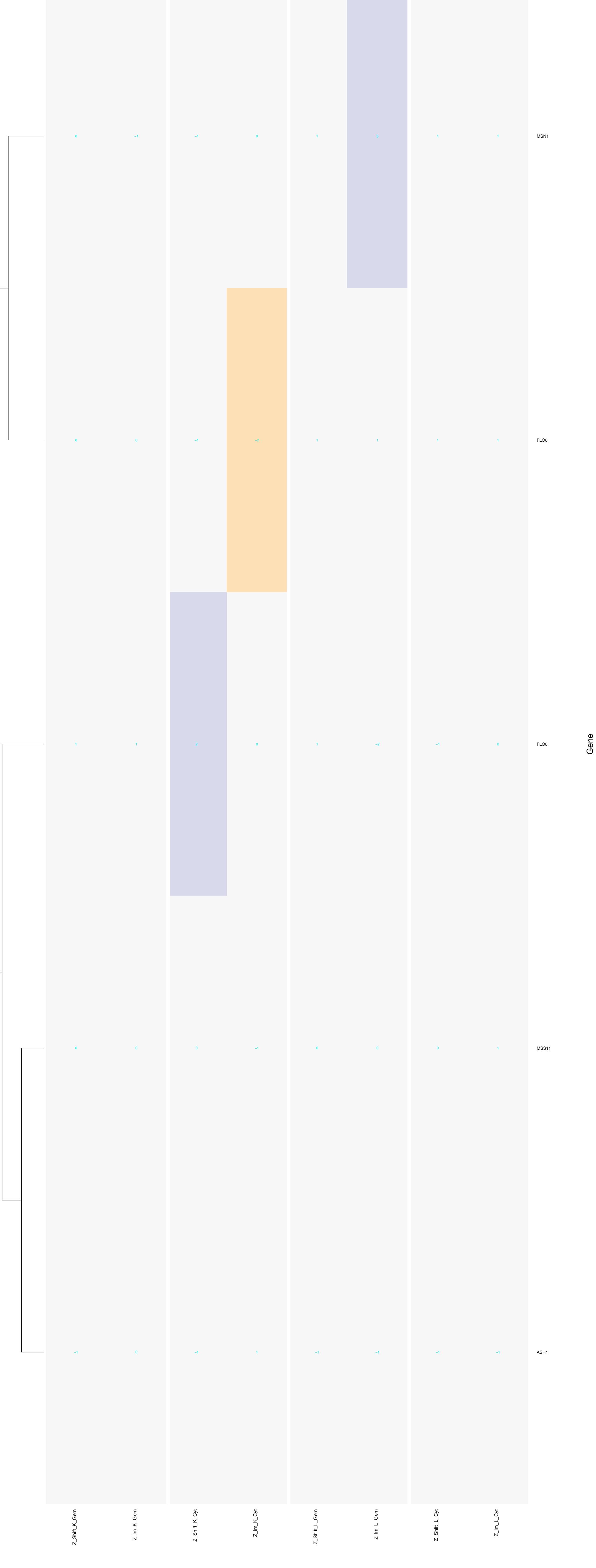




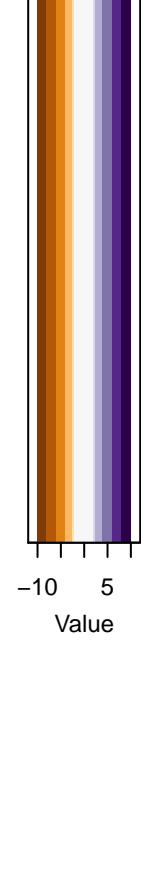
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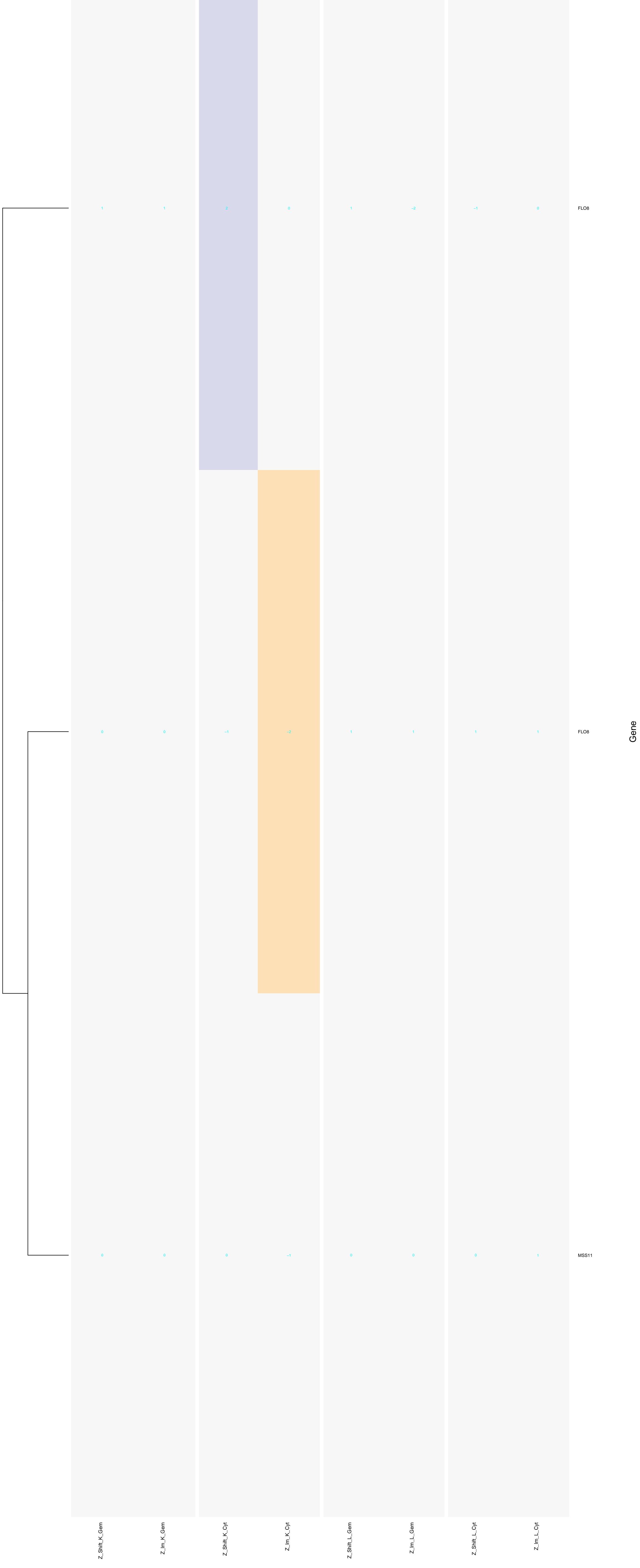
of pseudohyphal growth by positive regulation of transcription from RNA polymerase II pr



Color Key

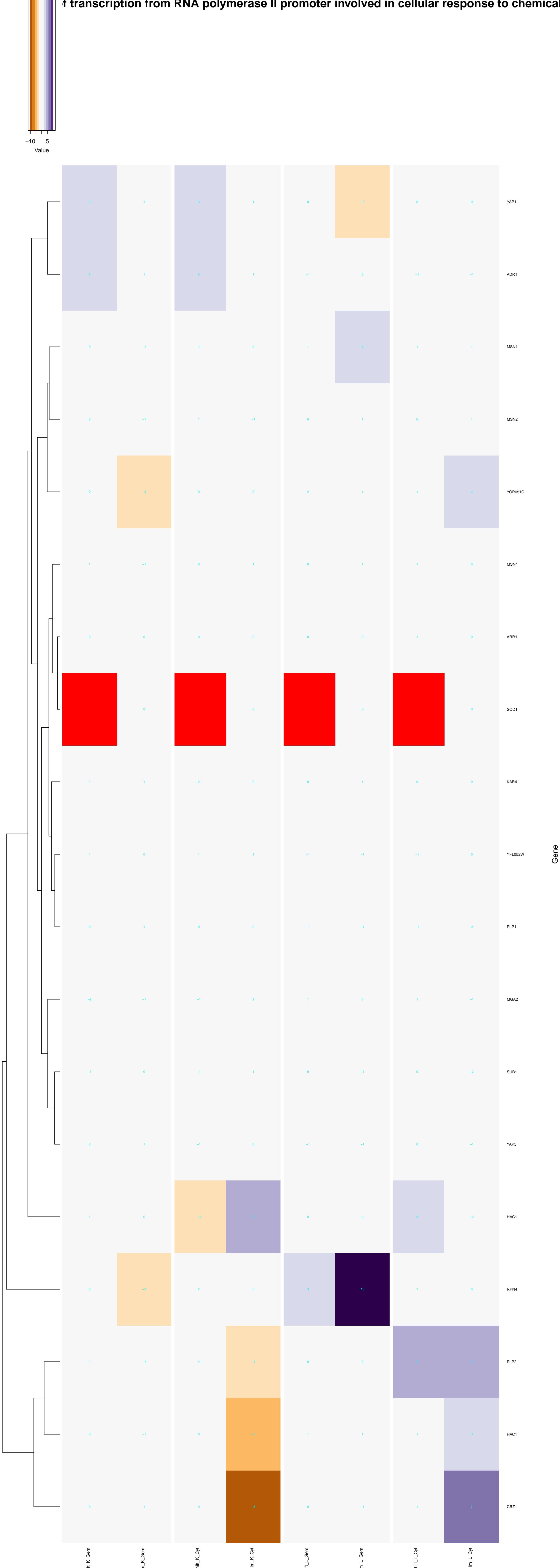


cell wall protein–carbohydrate interaction by positive regulation of transcription from RN

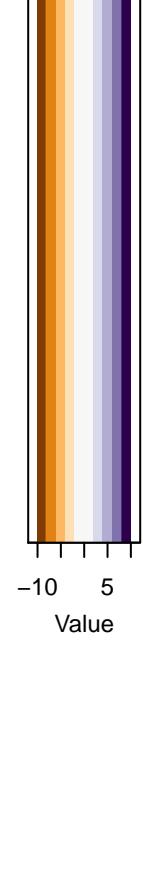


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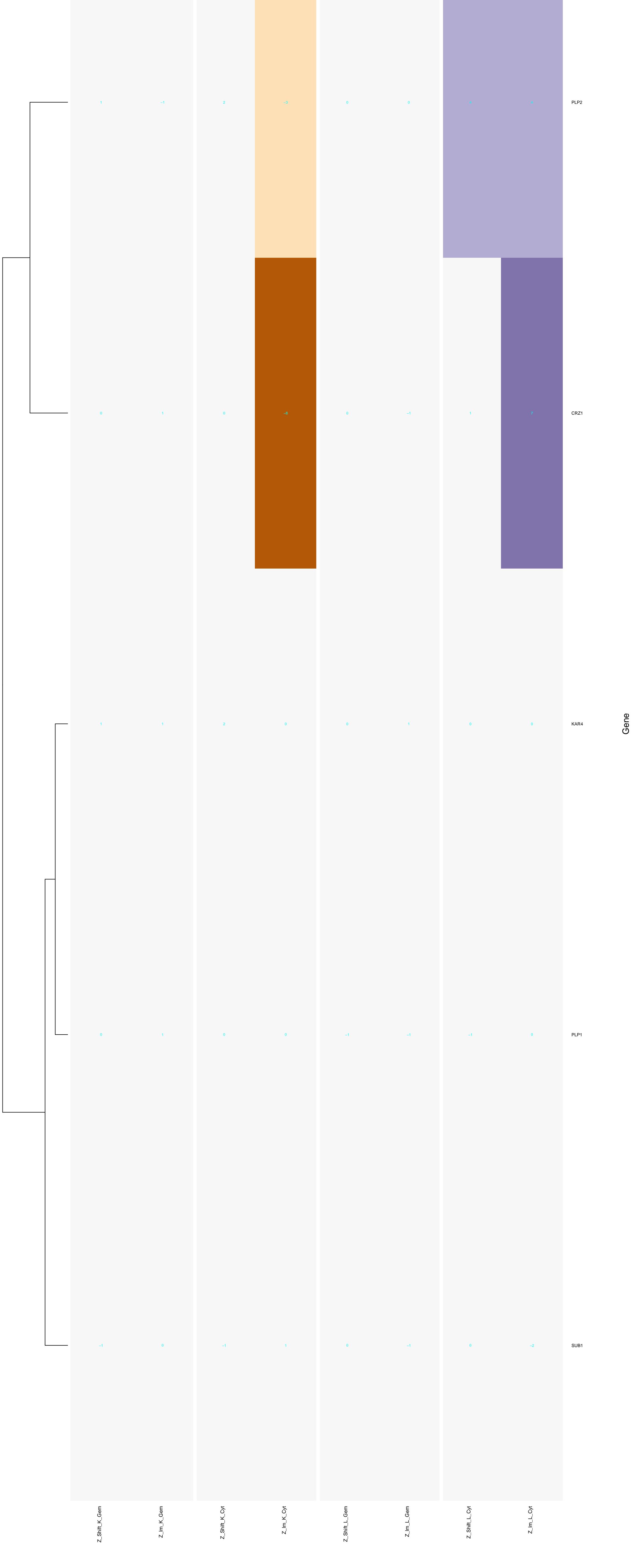
f transcription from RNA polymerase II promoter involved in cellular response to chemical



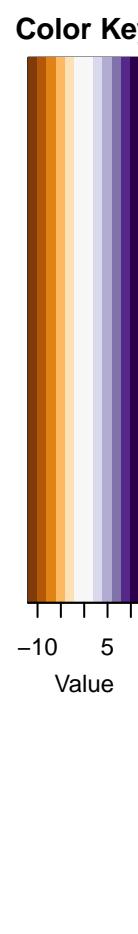
Color Key



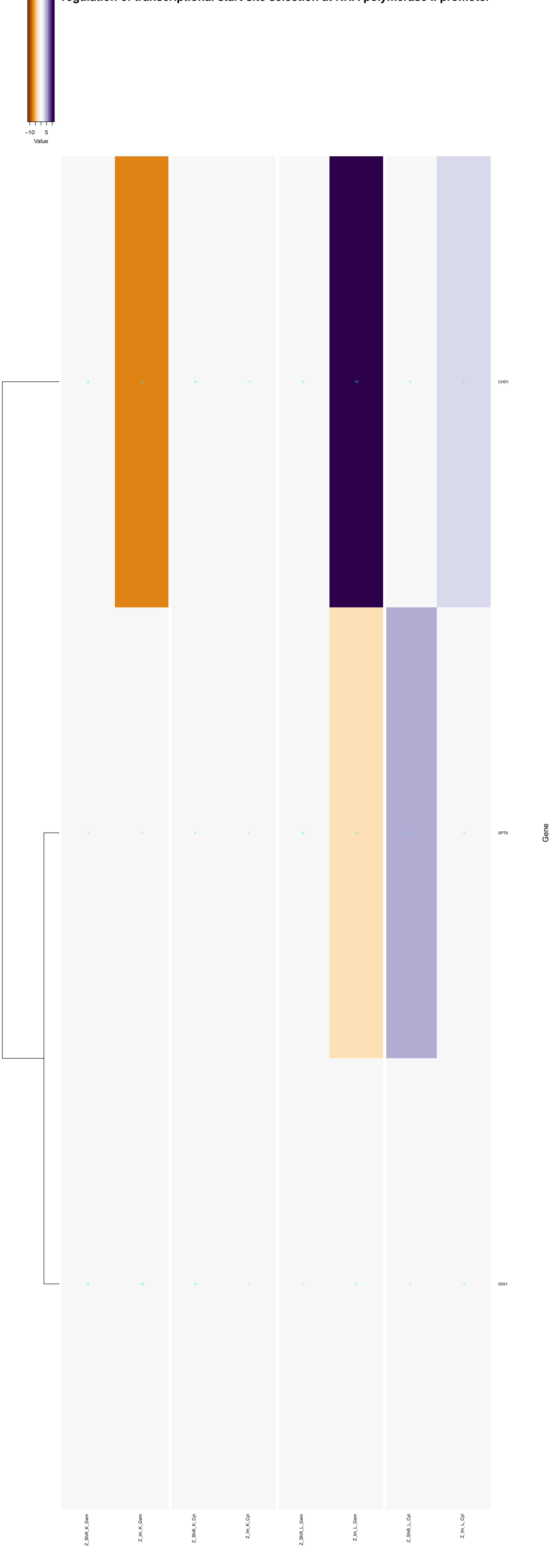
Positive regulation of transcription from RNA polymerase II promoter by pheromones



Color Key



regulation of transcriptional start site selection at RNA polymerase II promoter



Color Key



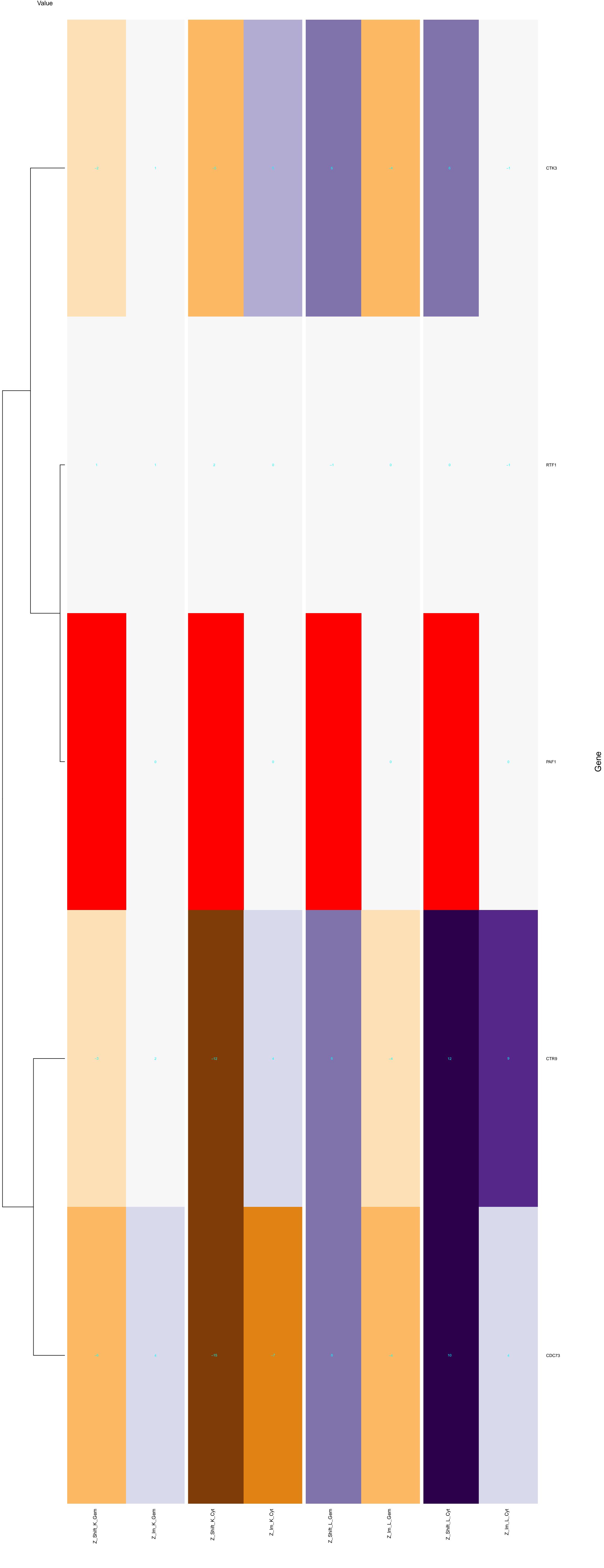
regulation of RNA polymerase II transcriptional preinitiation complex assembly



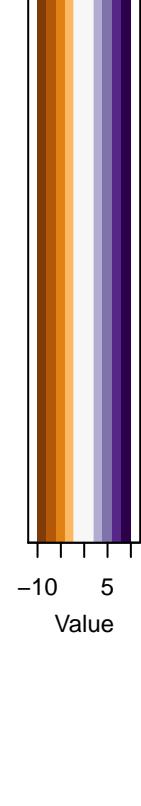
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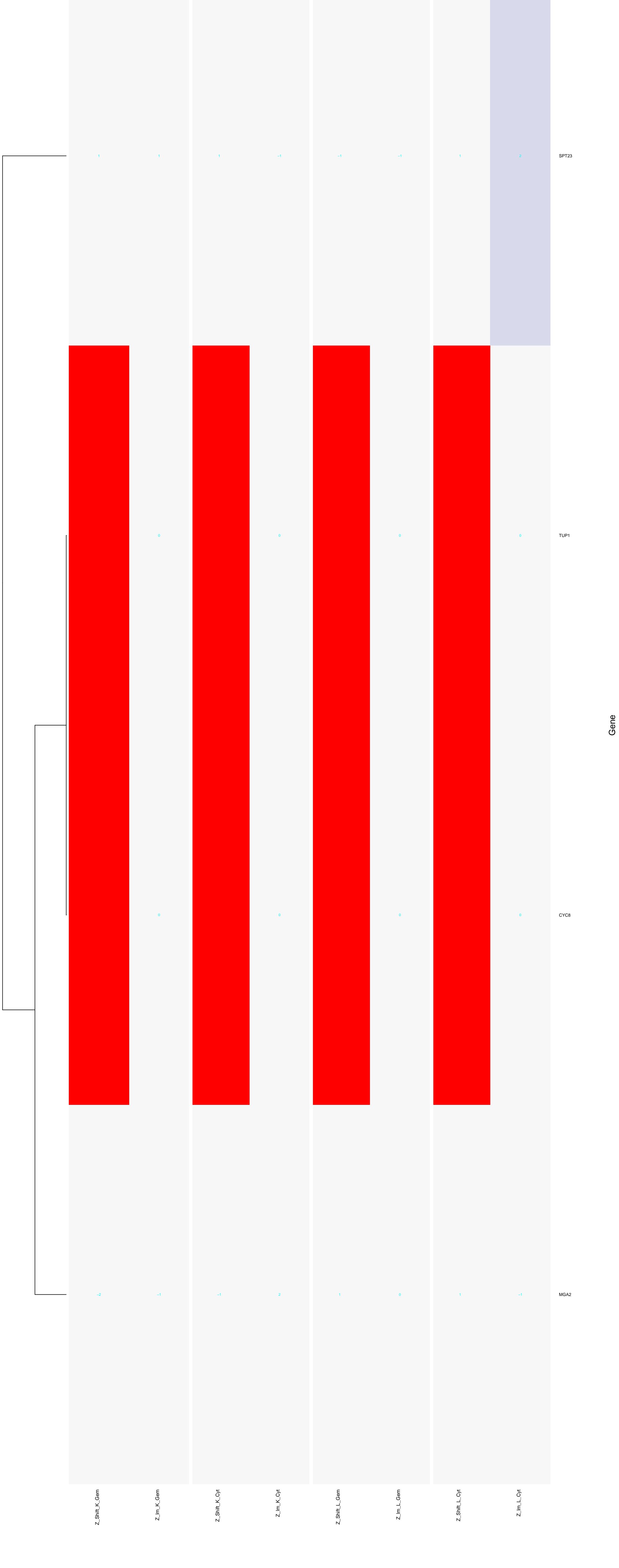
phosphorylation of RNA polymerase II C-terminal domain serine 2 residues

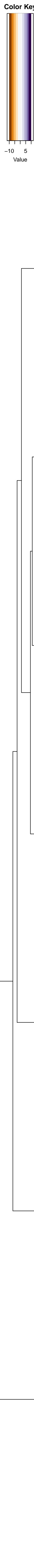


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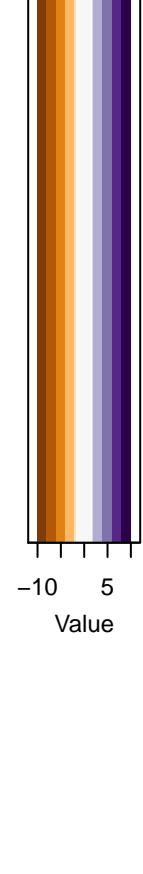


atty acid biosynthetic process by regulation of transcription from RNA polymerase II promoter

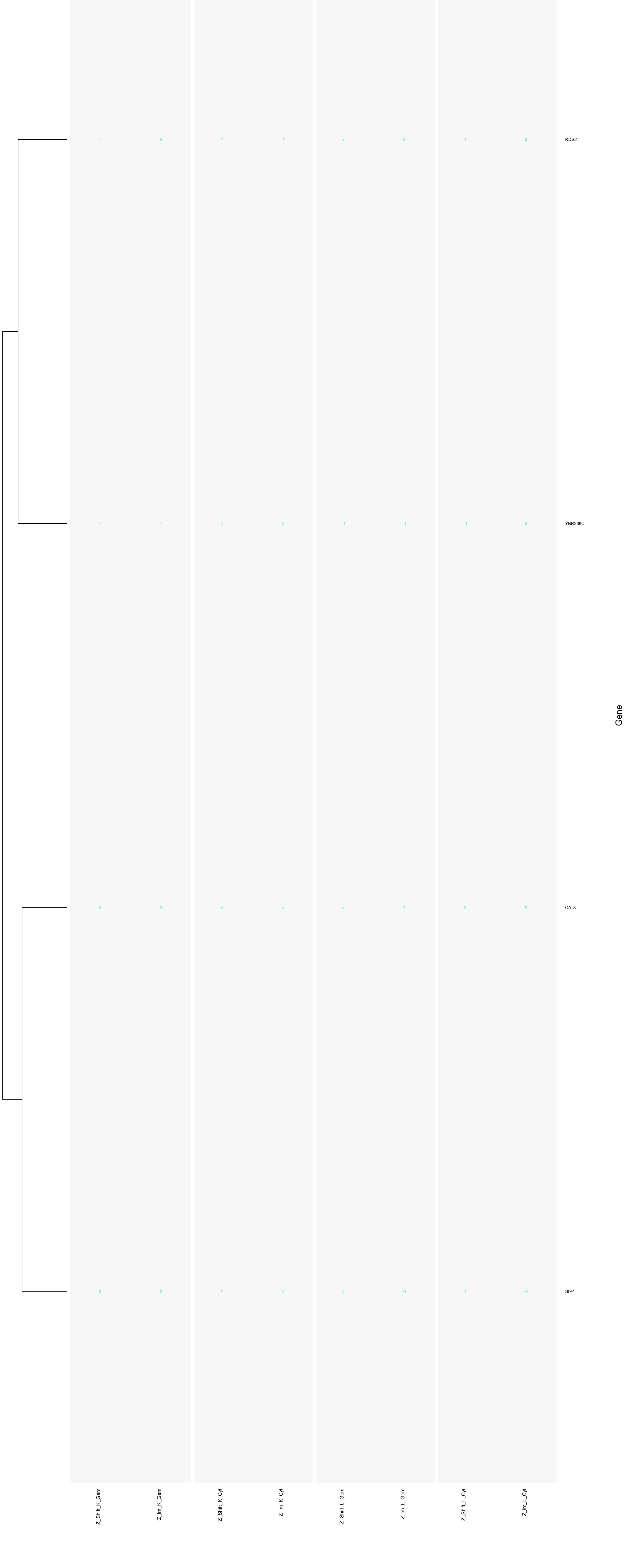




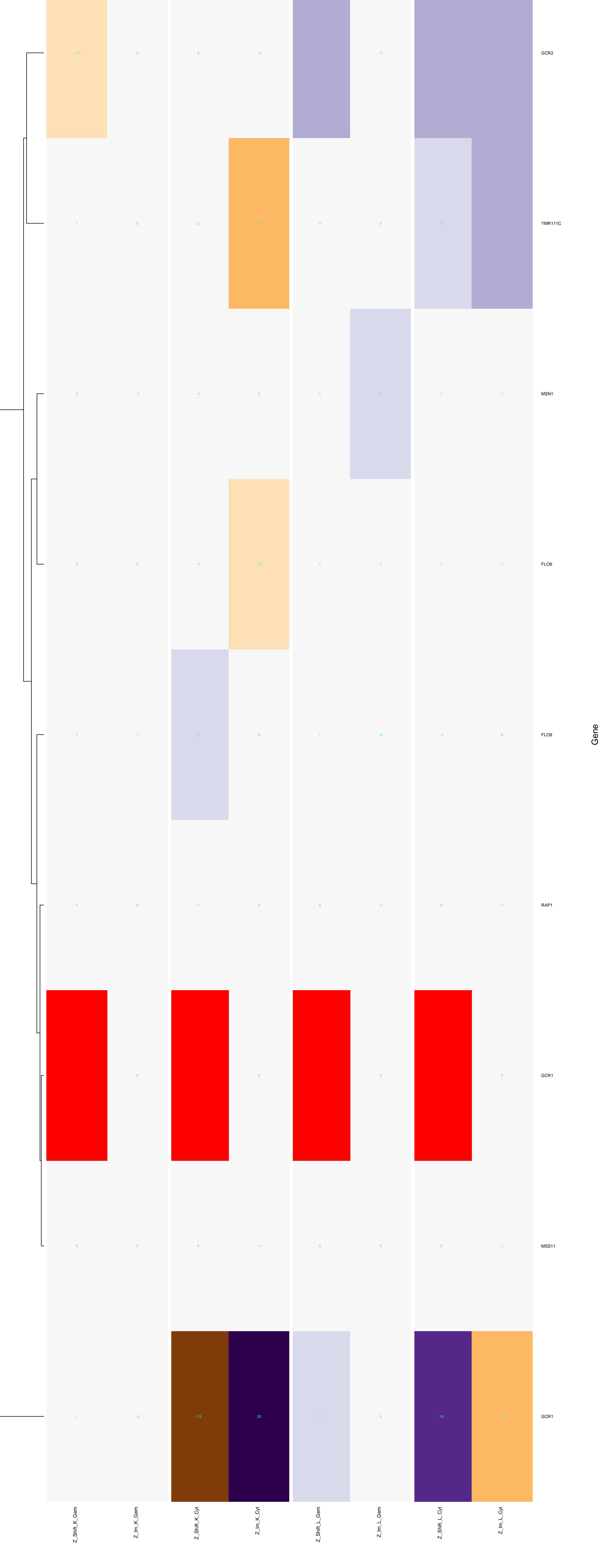
Color Key



Regulation of gluconeogenesis by regulation of transcription from RNA polymerase II promoter



Color Key

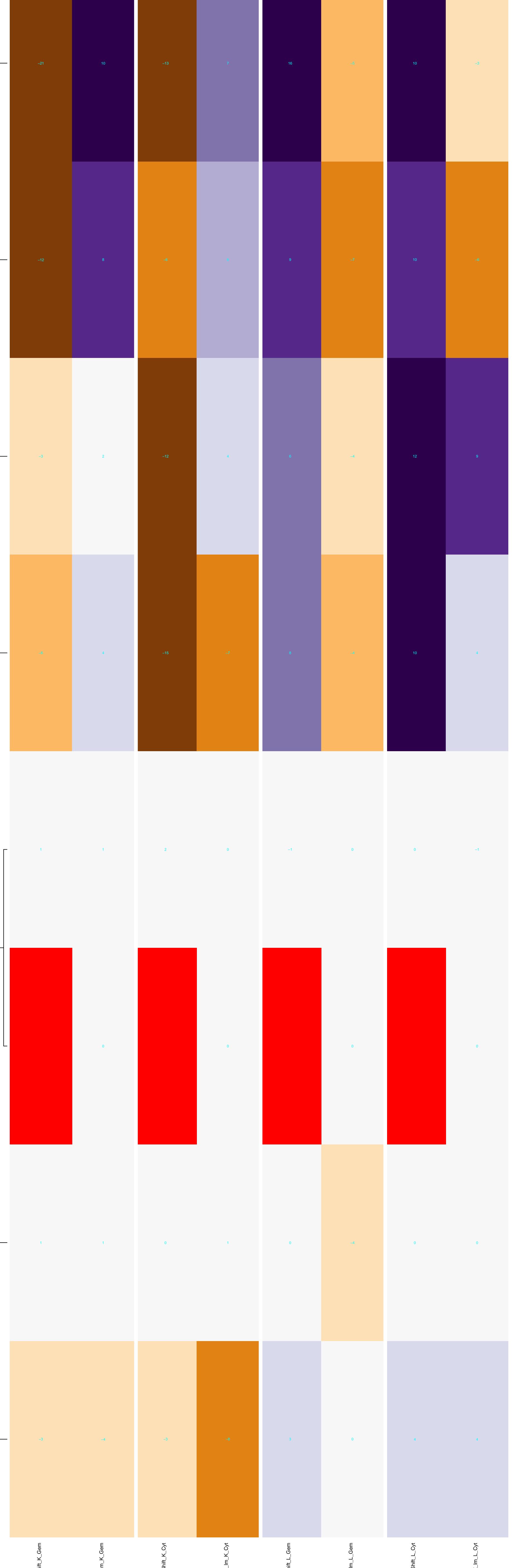


Color Key



Positive regulation of phosphorylation of RNA polymerase II C-terminal domain

-10 5
Value

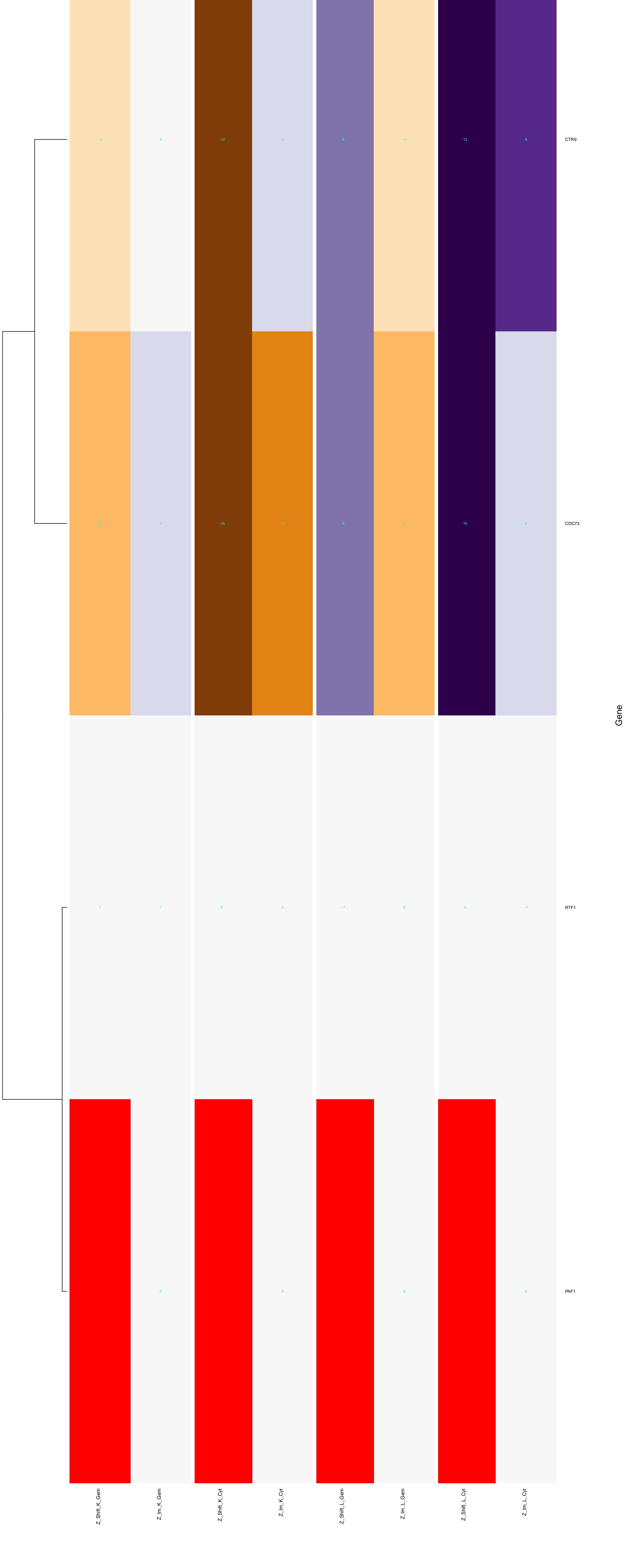


Gene

Color Key



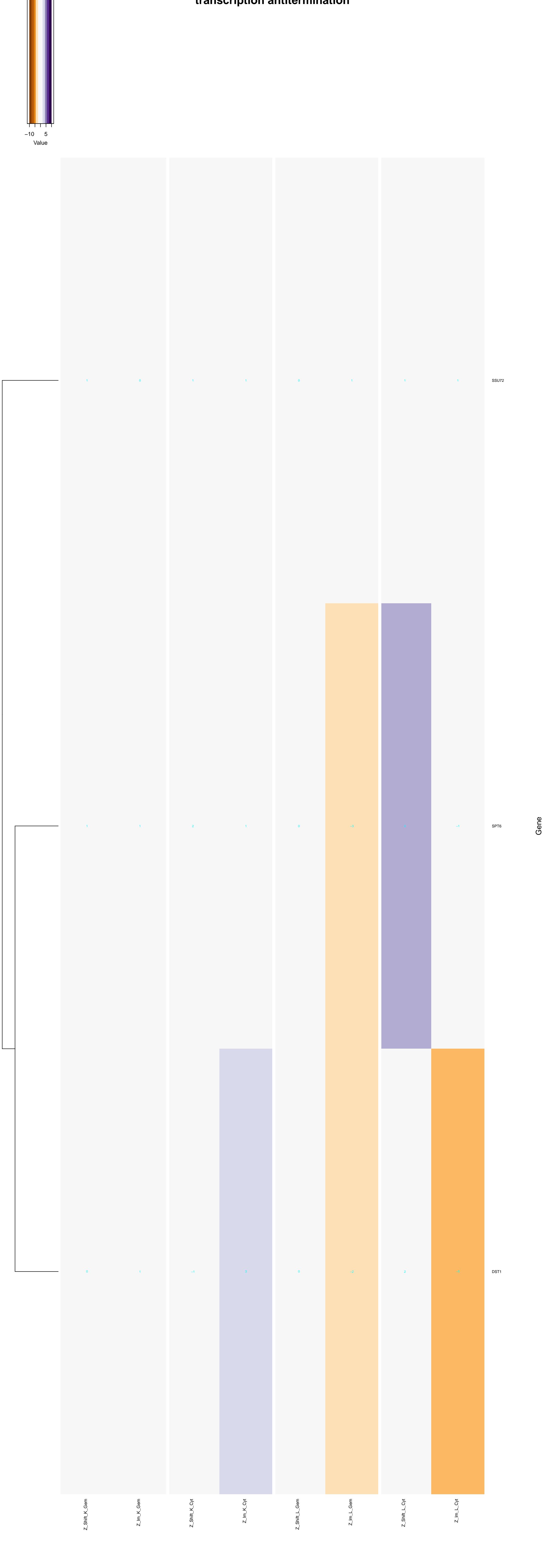
Activation of phosphorylation of RNA polymerase II C-terminal domain serine 2 residues



Color Key



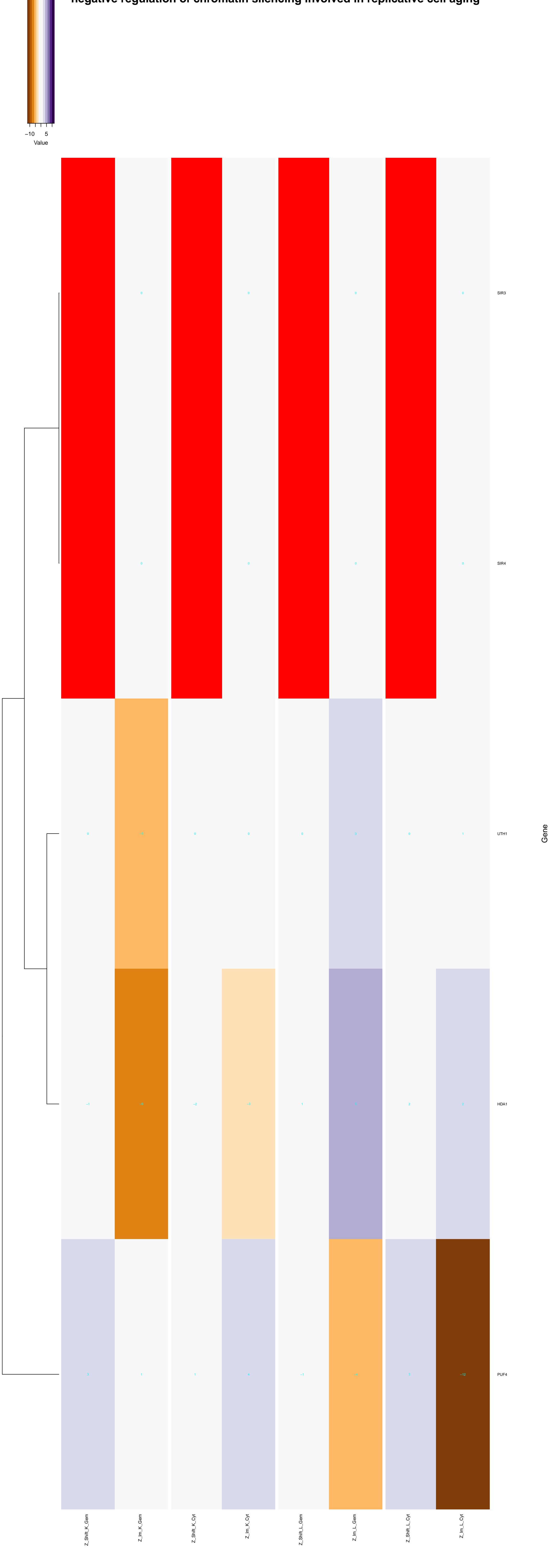
transcription antitermination



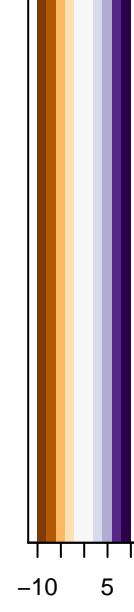
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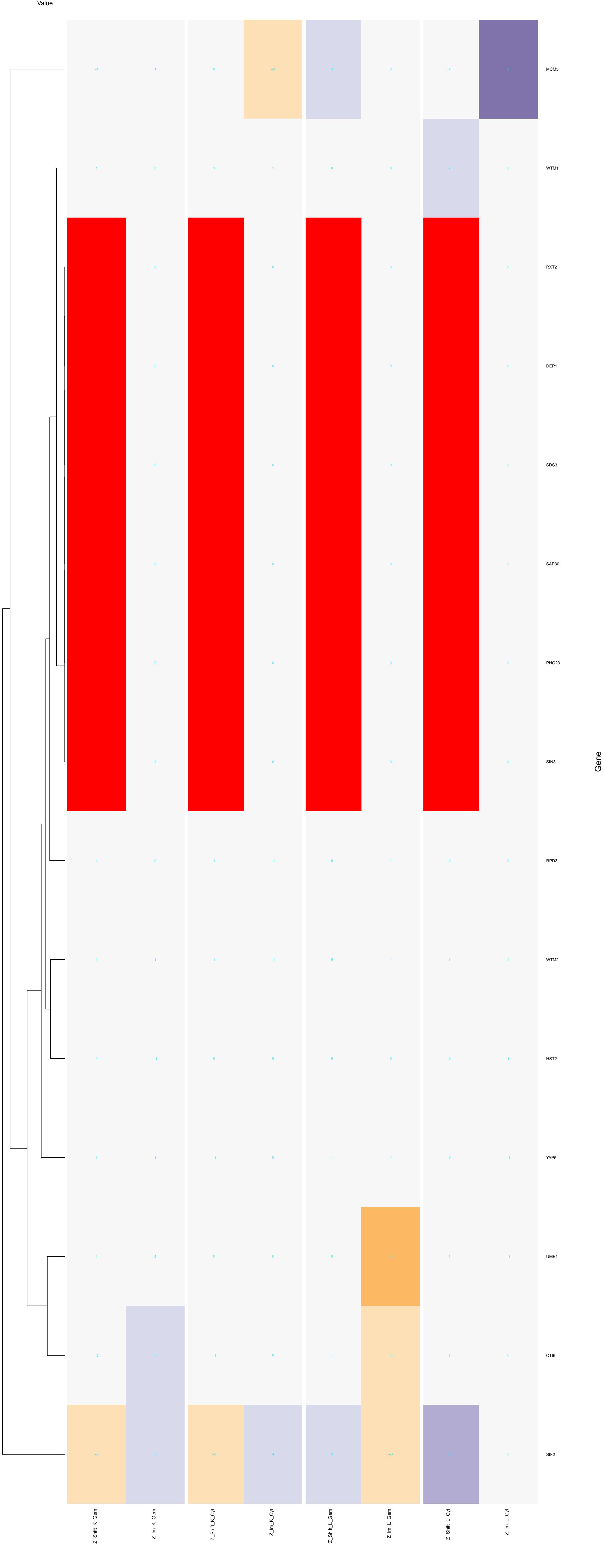
negative regulation of chromatin silencing involved in replicative cell aging



Color Key



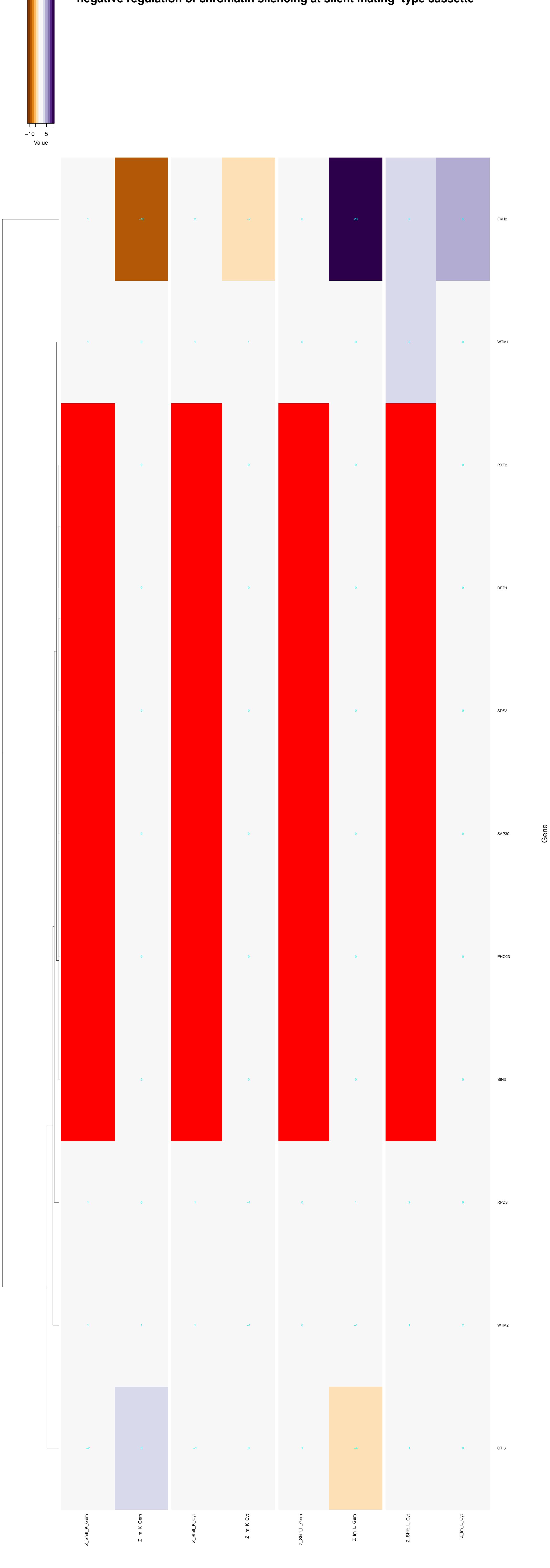
negative regulation of chromatin silencing at telomere

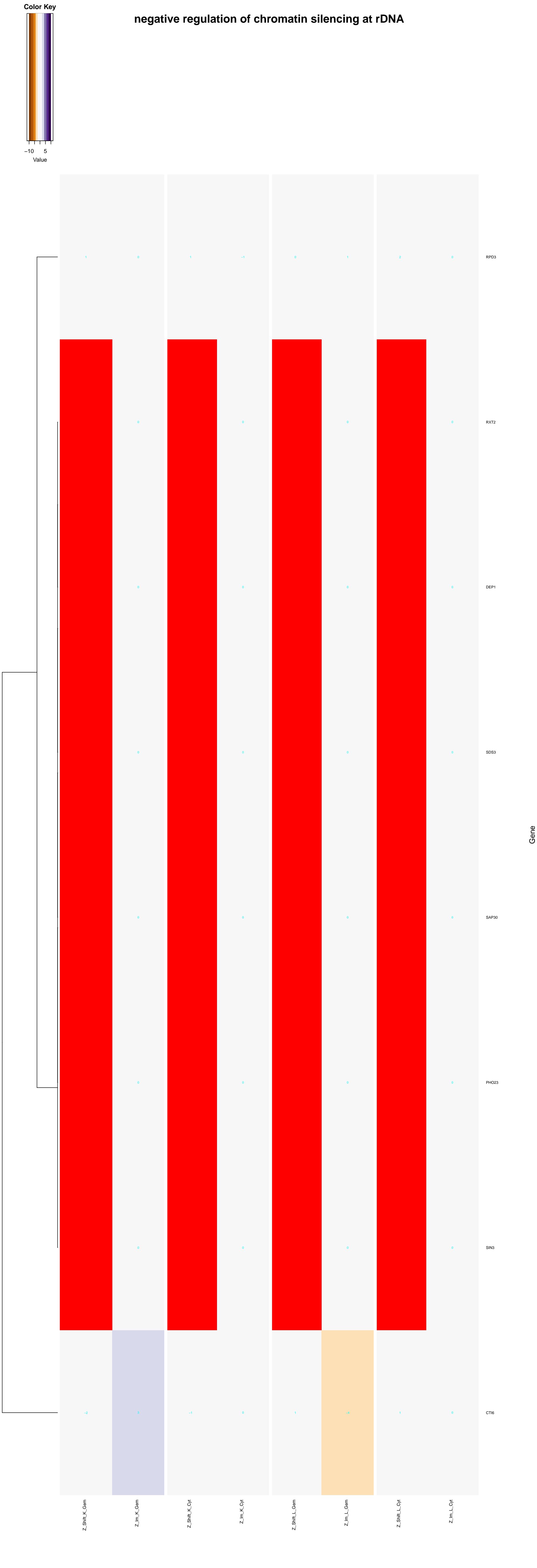


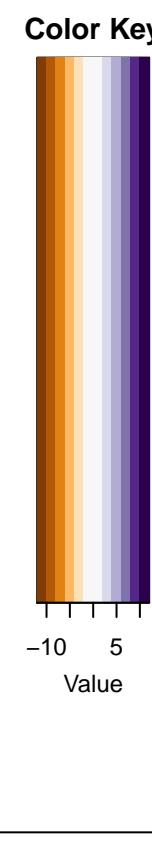
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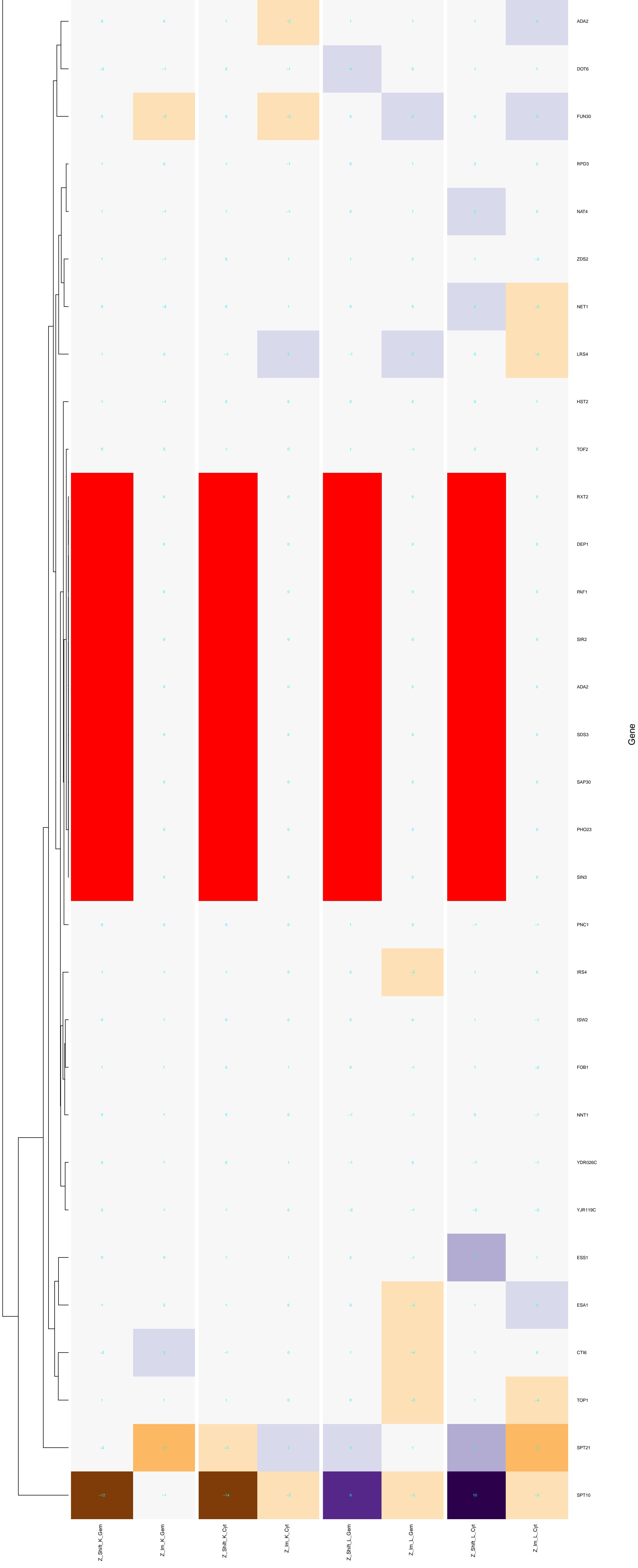
negative regulation of chromatin silencing at silent mating-type cassette

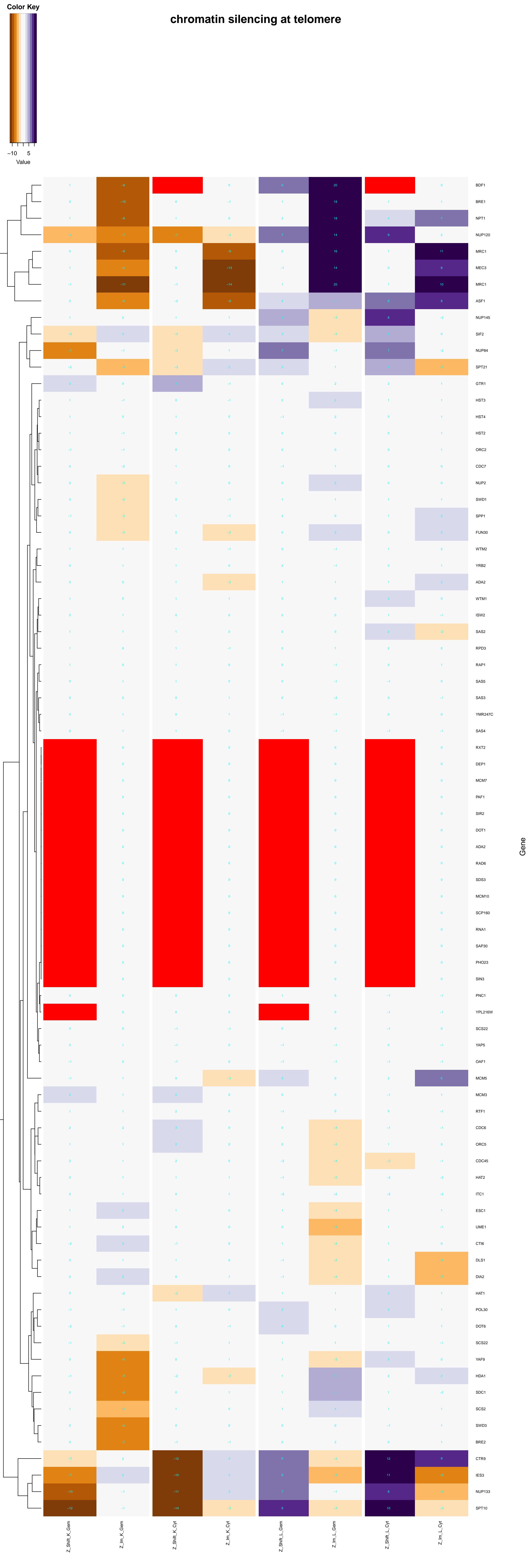


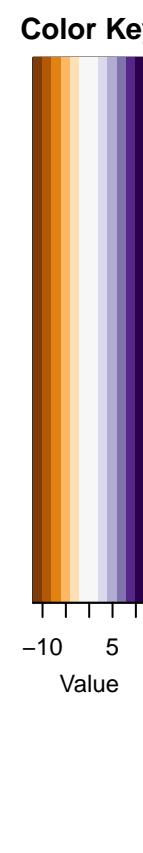




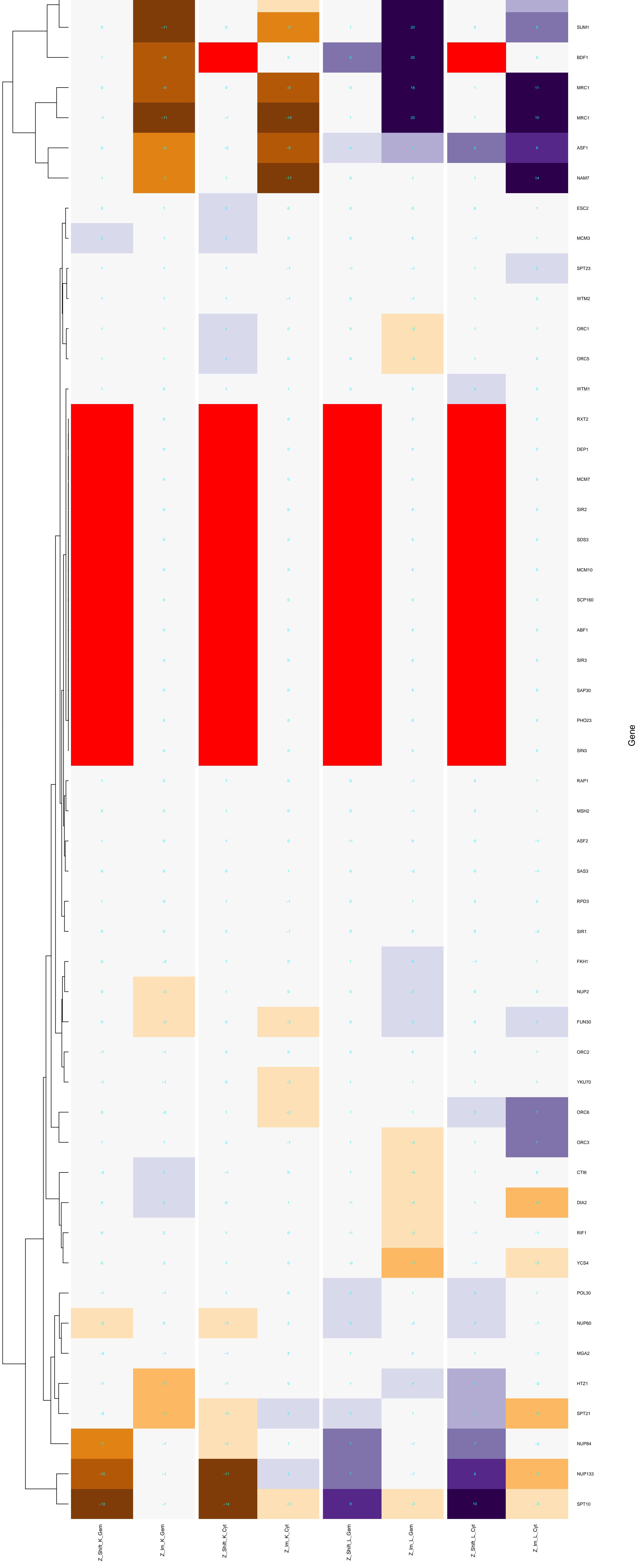
chromatin silencing at rDNA



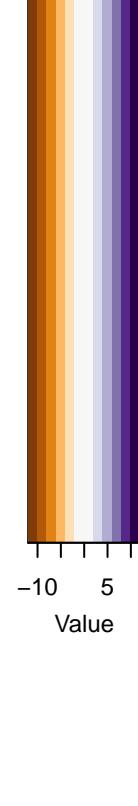




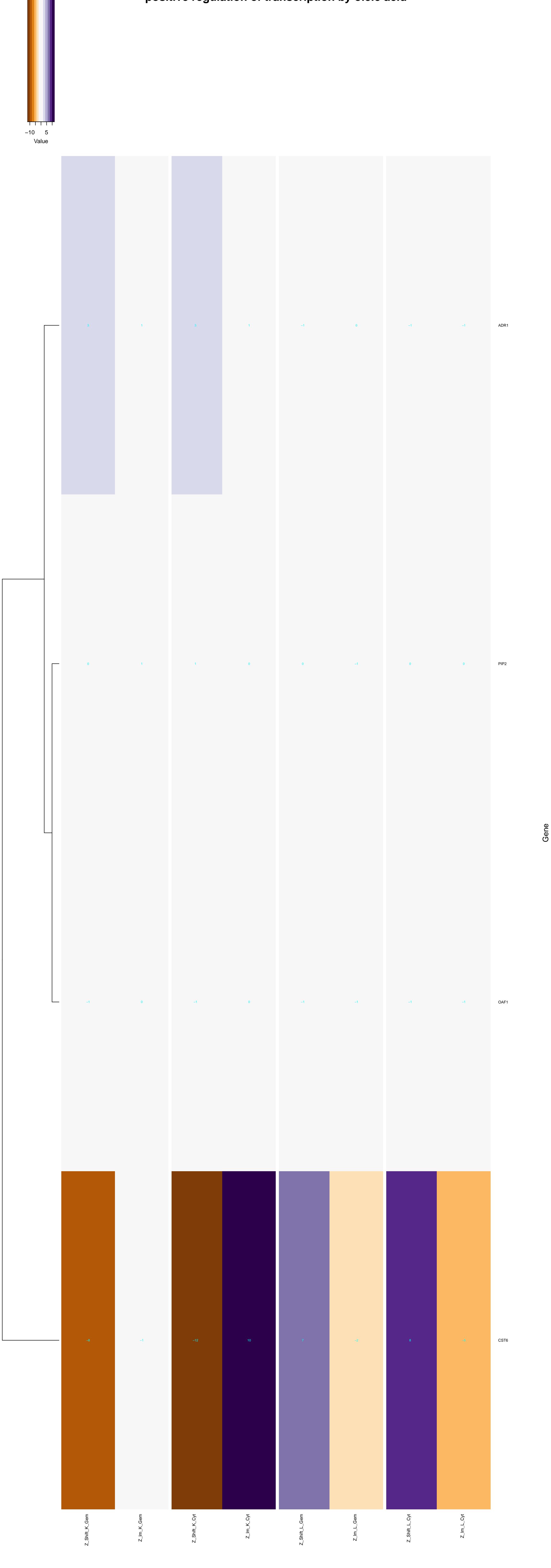
chromatin silencing at silent mating-type cassette



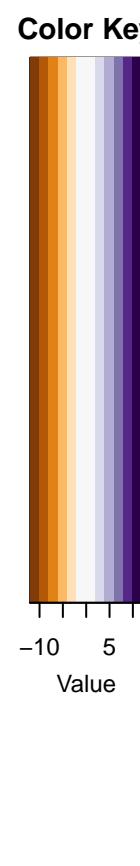
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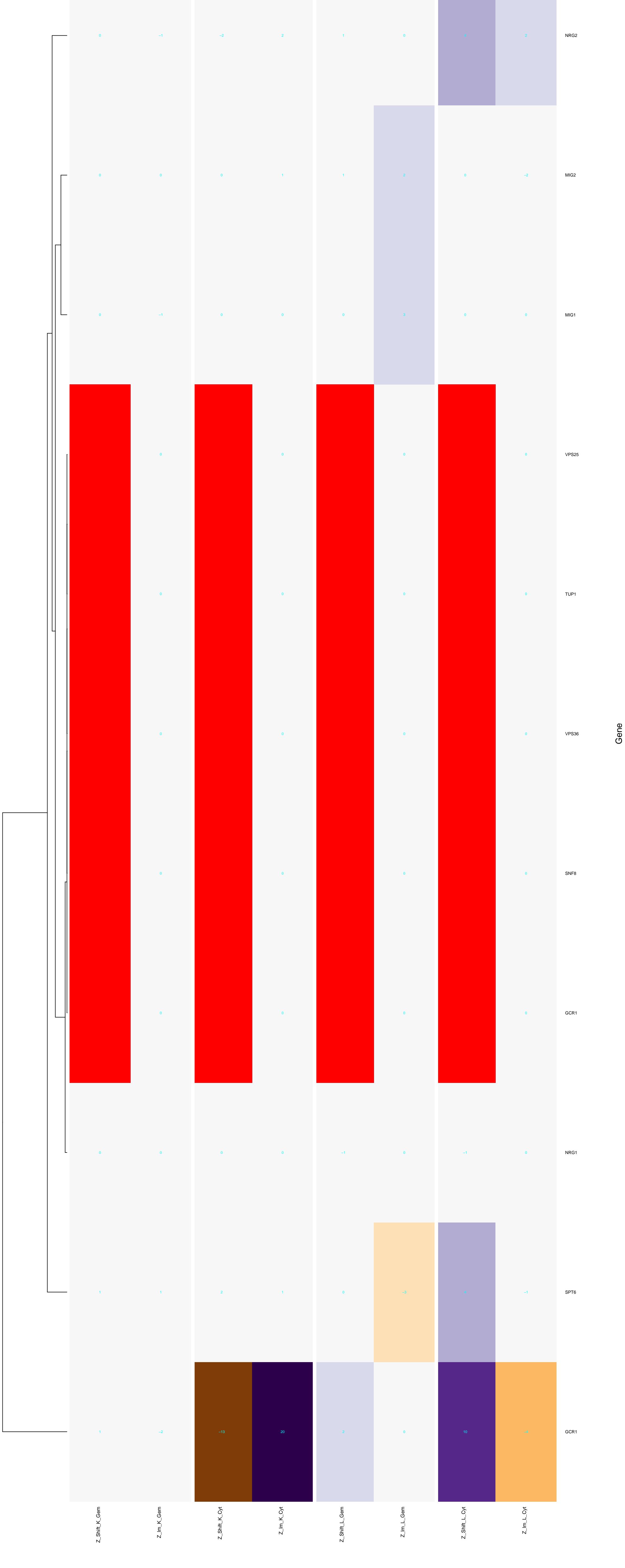
positive regulation of transcription by oleic acid



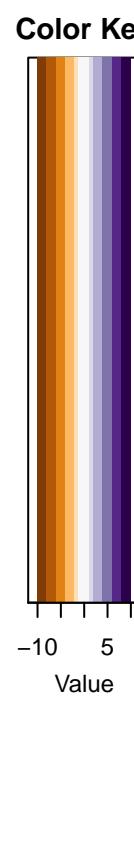
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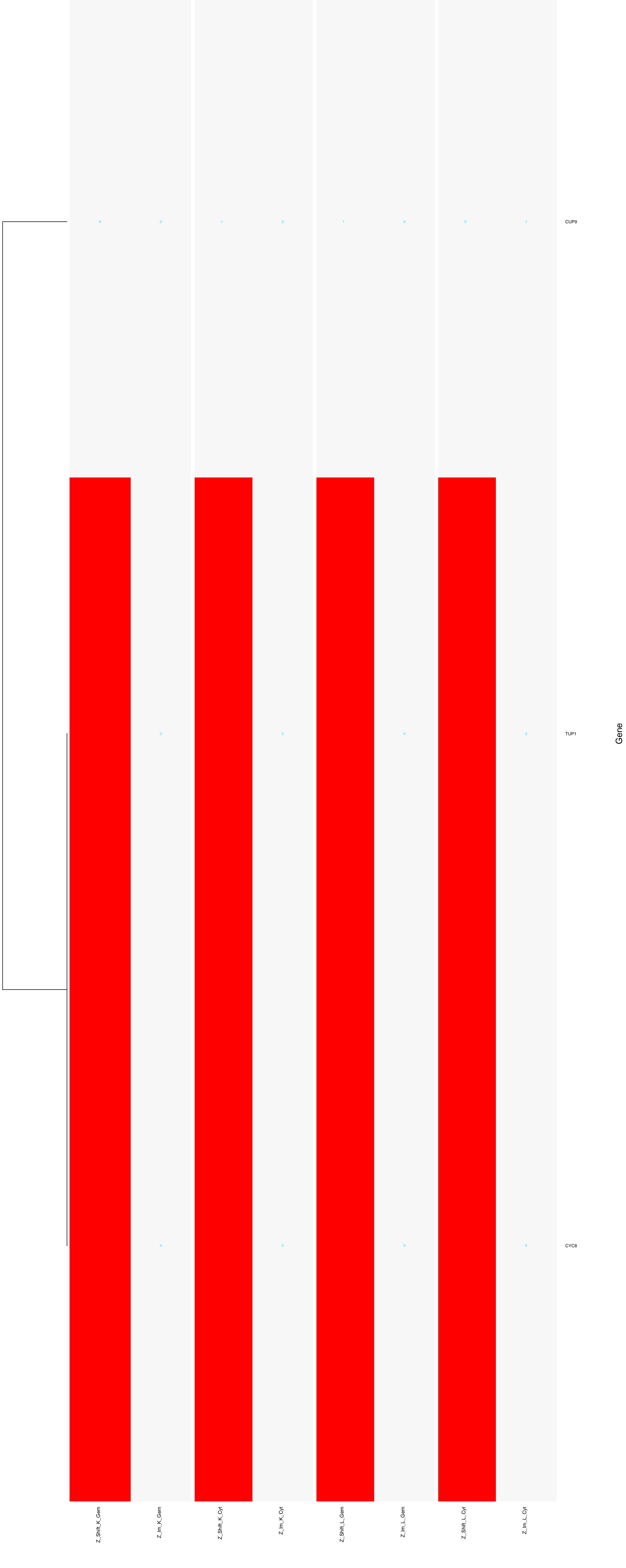
Negative regulation of transcription from RNA polymerase II promoter by glucose



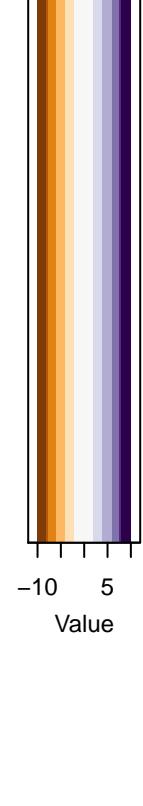
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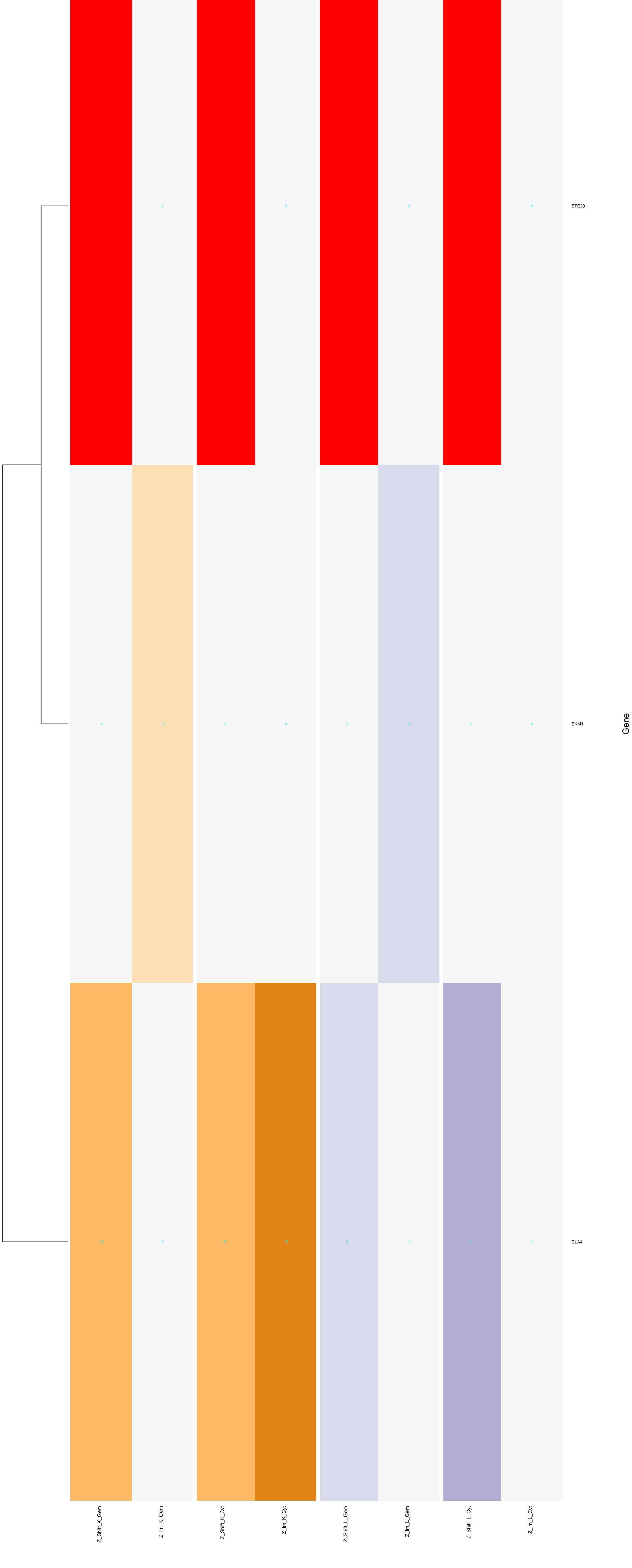
n of dipeptide transport by negative regulation of transcription from RNA polymerase II pr



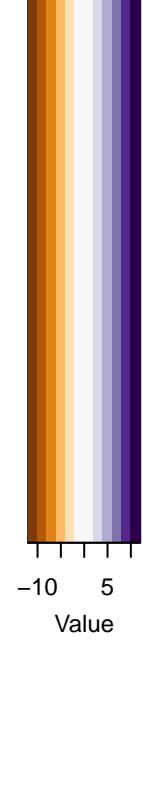
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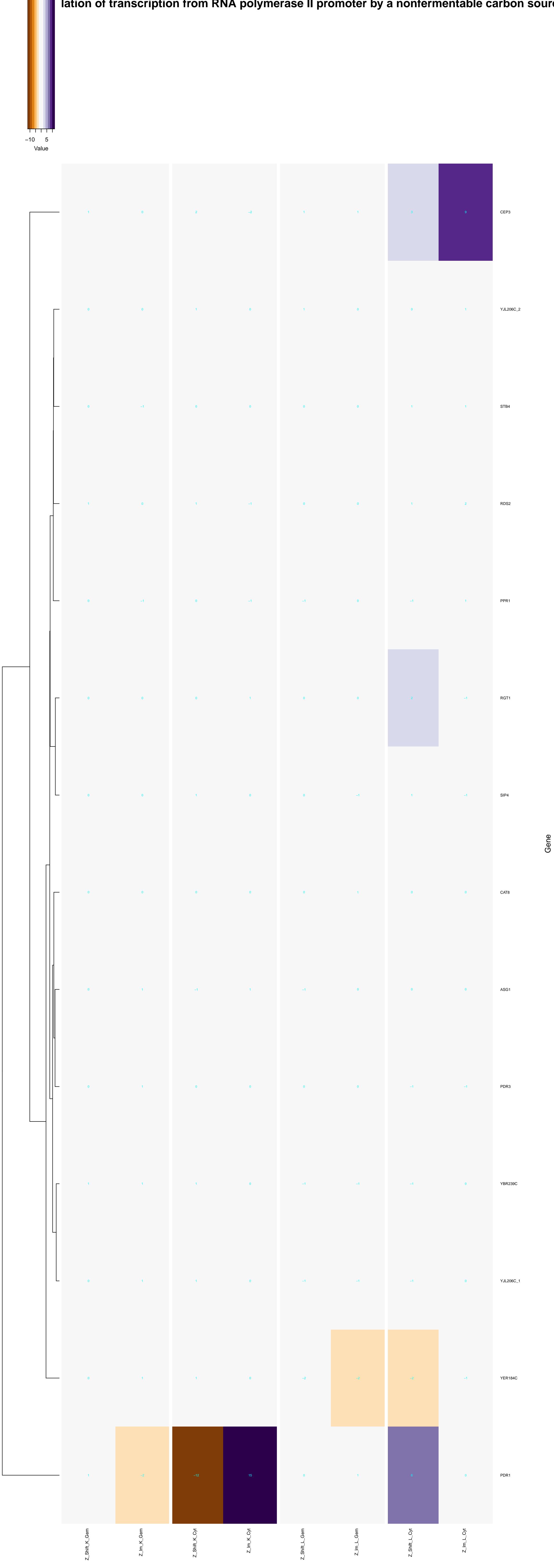
Regulation of sterol import by negative regulation of transcription from RNA polymerase II promoter



Color Key



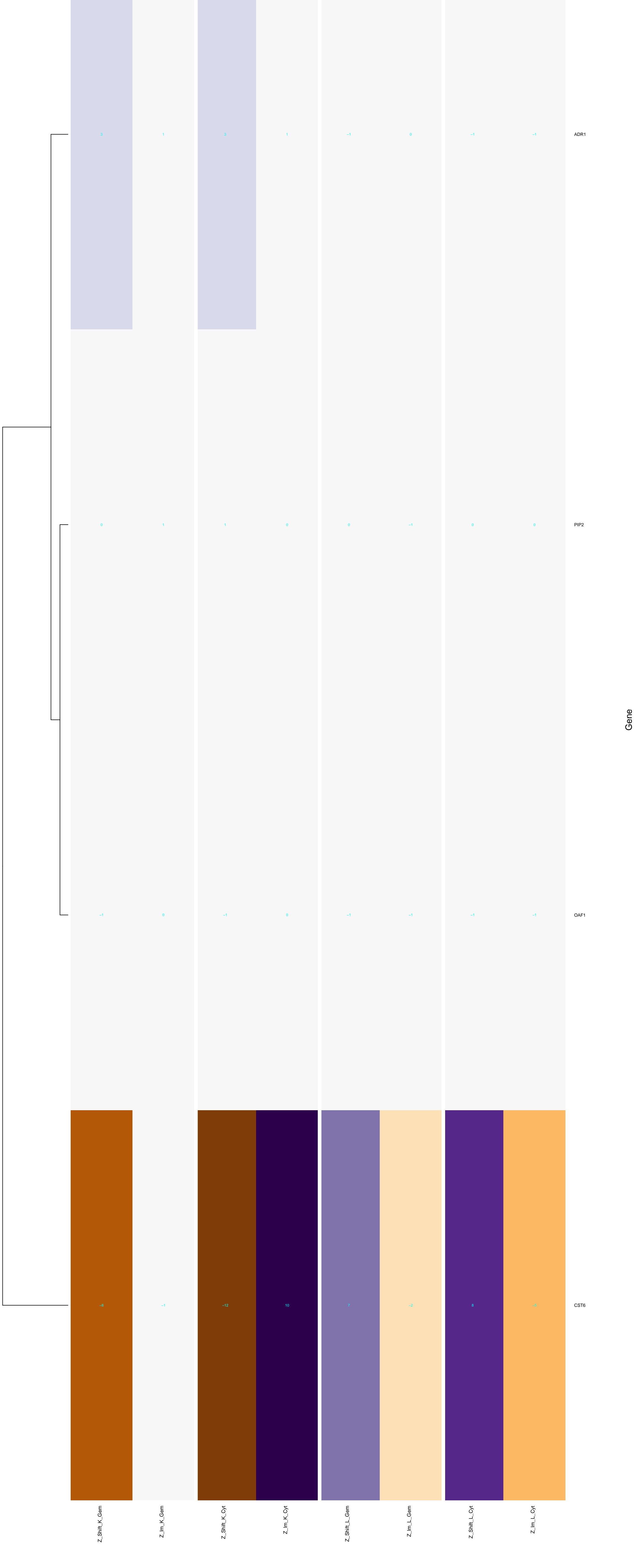
Inhibition of transcription from RNA polymerase II promoter by a nonfermentable carbon source



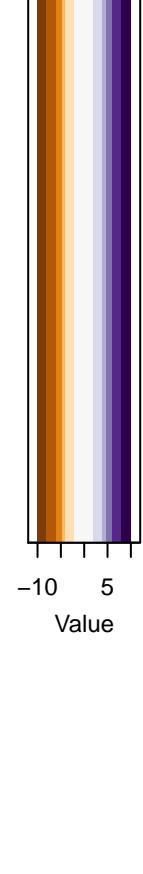
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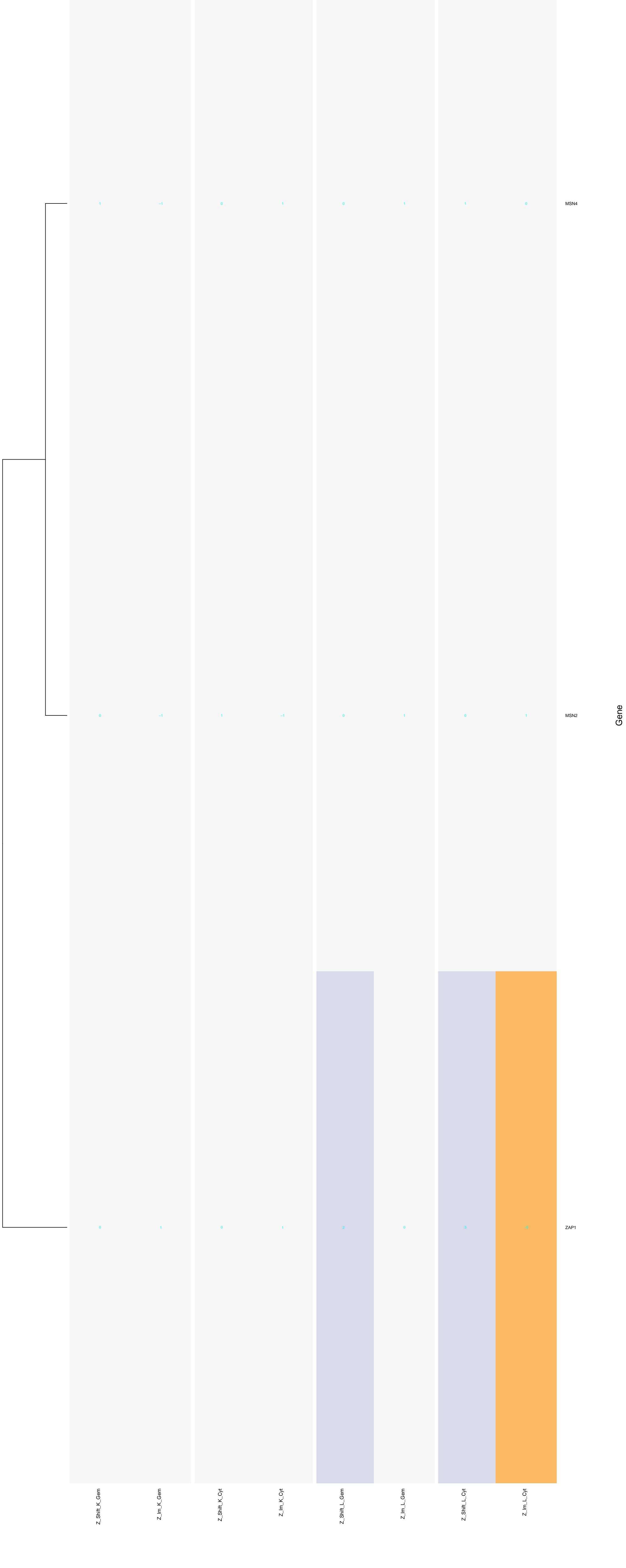
Positive regulation of transcription from RNA polymerase II promoter by oleic acid



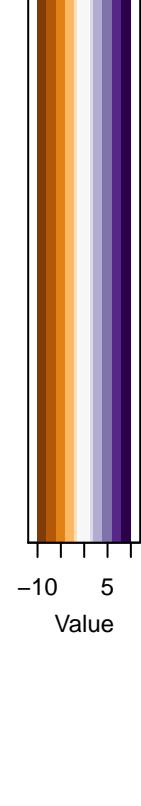
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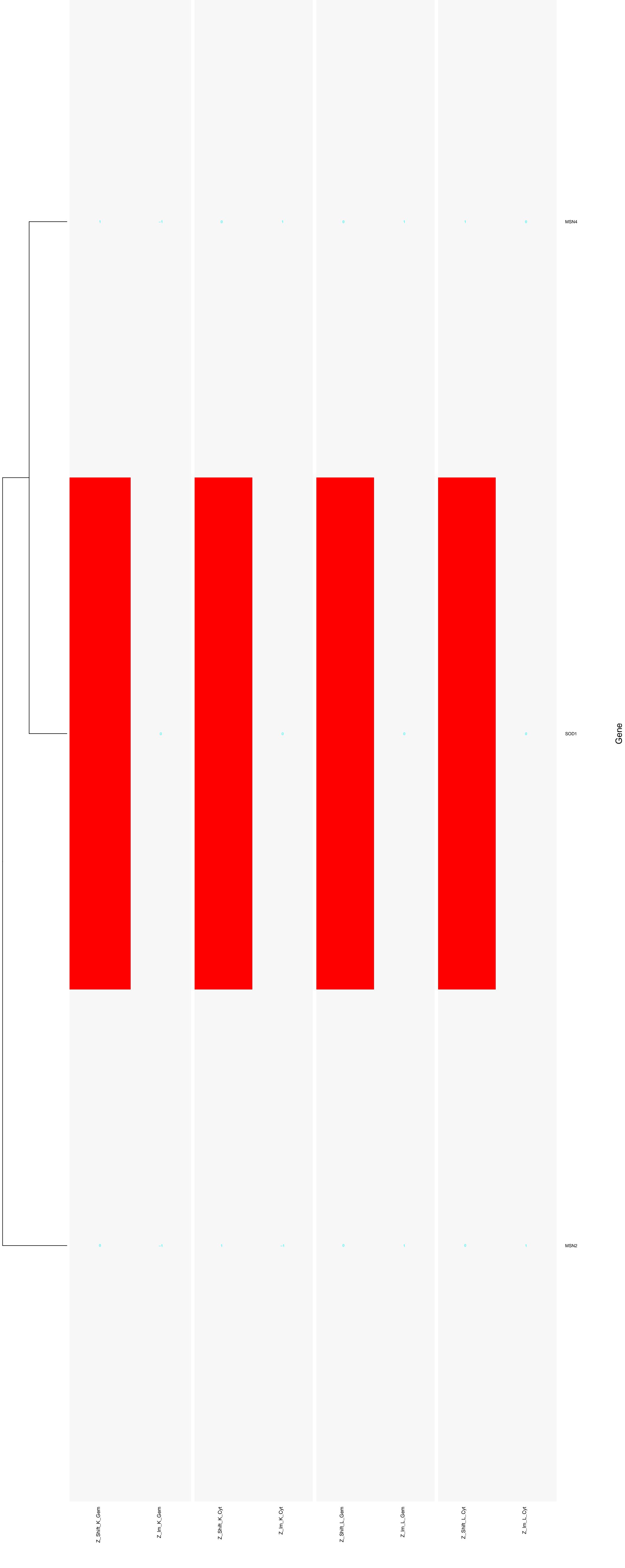
ulation of transcription from RNA polymerase II promoter in response to zinc ion starvation



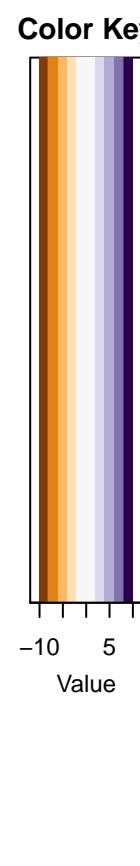
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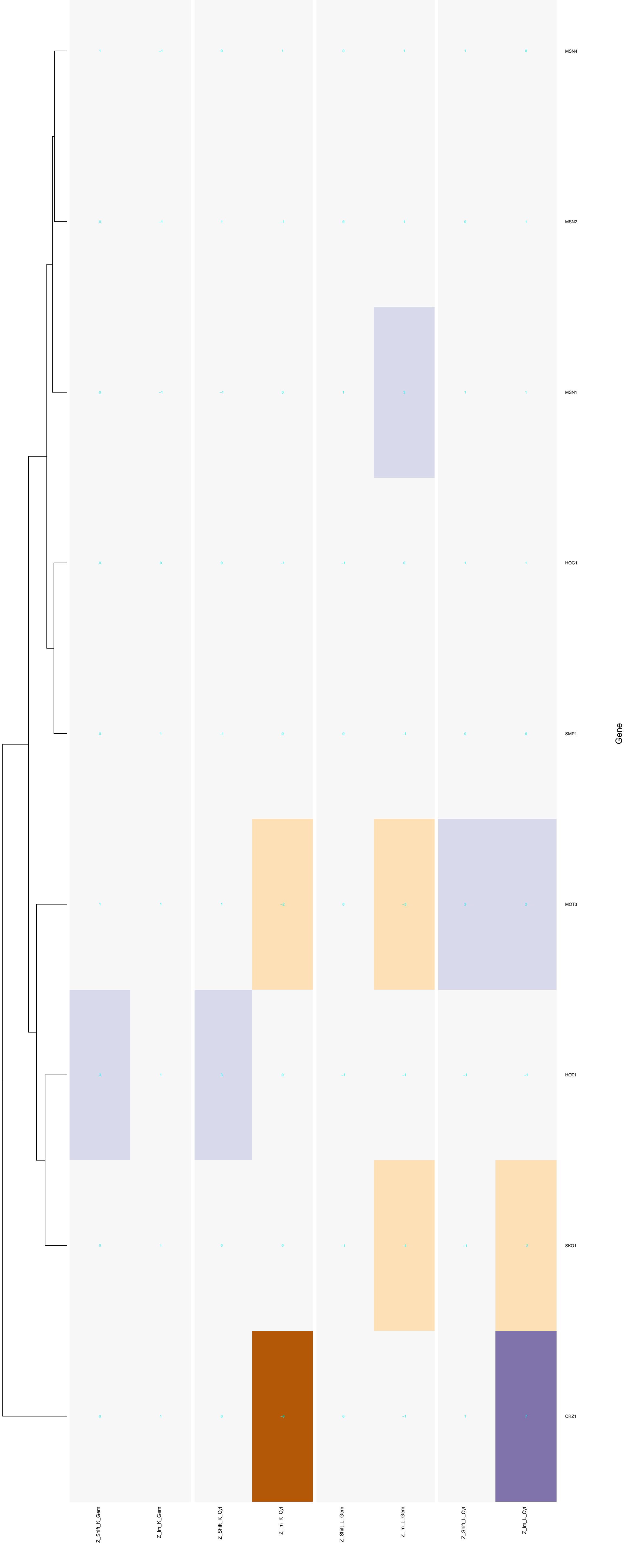
gulation of transcription from RNA polymerase II promoter in response to oxidative stress



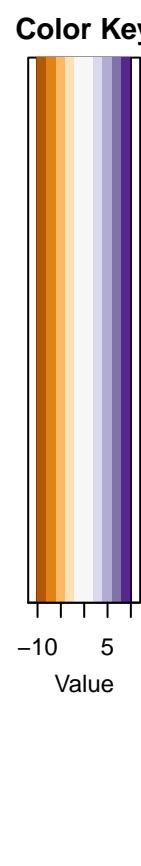
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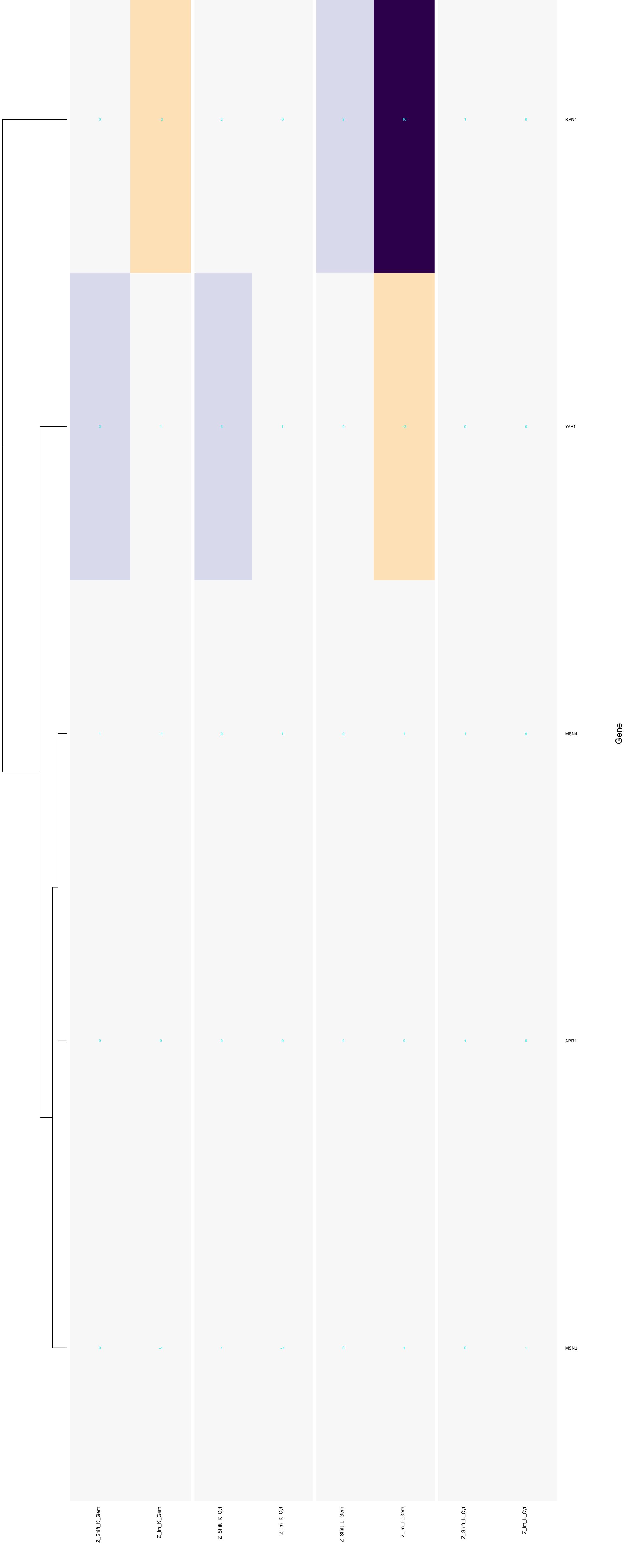
Regulation of transcription from RNA polymerase II promoter in response to osmotic stress



Color Key



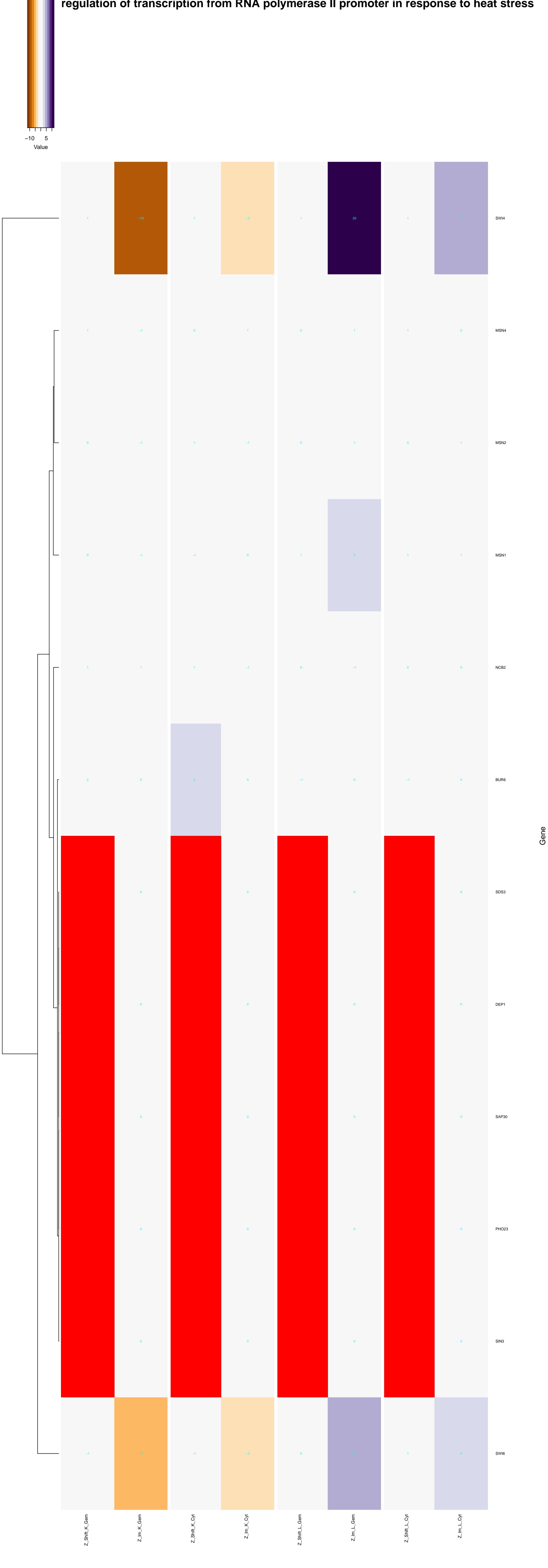
ChIP of transcription from RNA polymerase II promoter in response to arsenic-containing sub



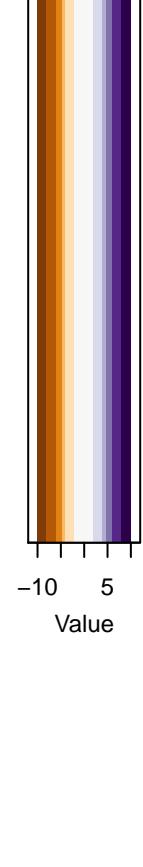
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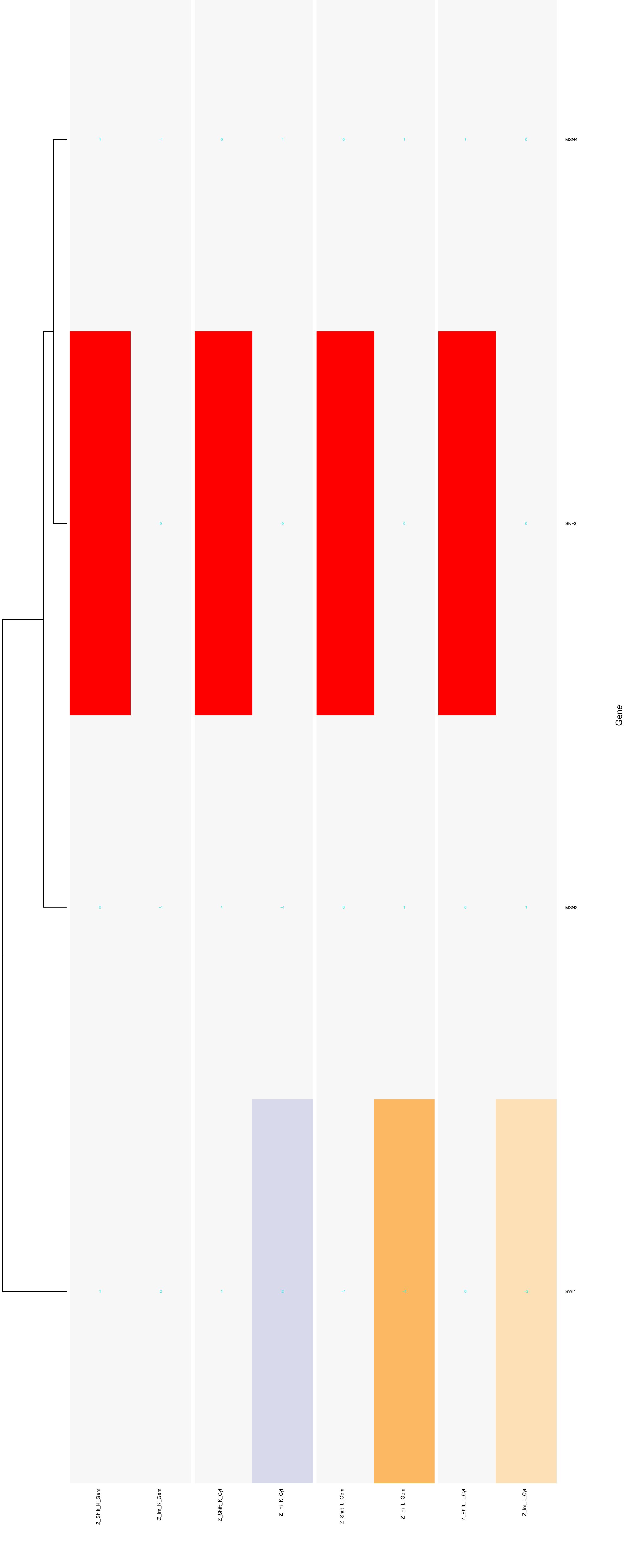
regulation of transcription from RNA polymerase II promoter in response to heat stress



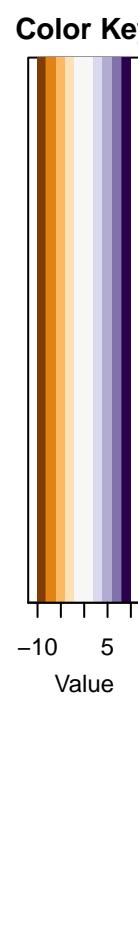
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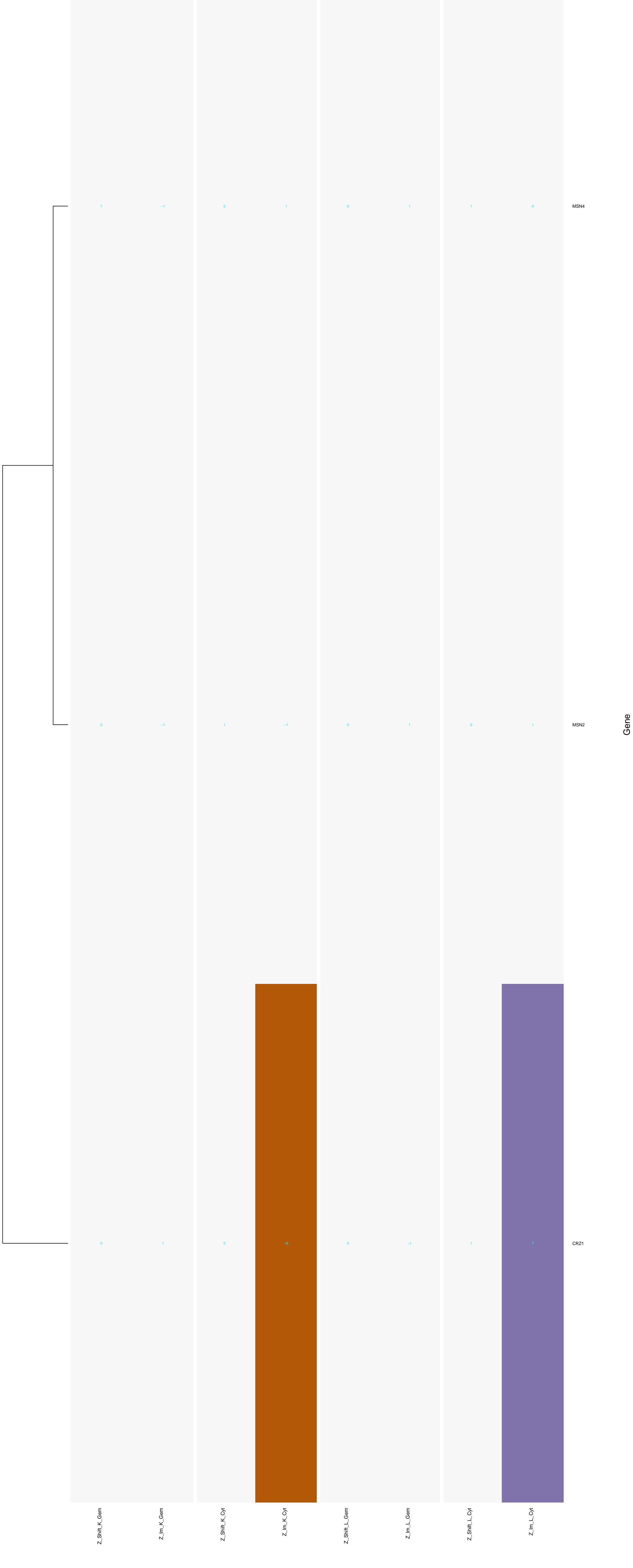
ation of transcription from RNA polymerase II promoter in response to amino acid starvati



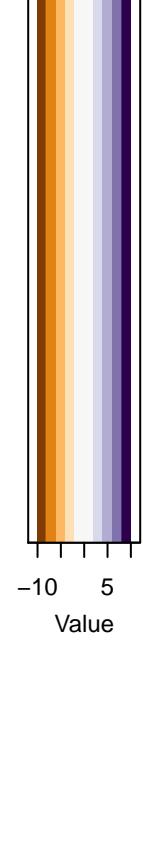
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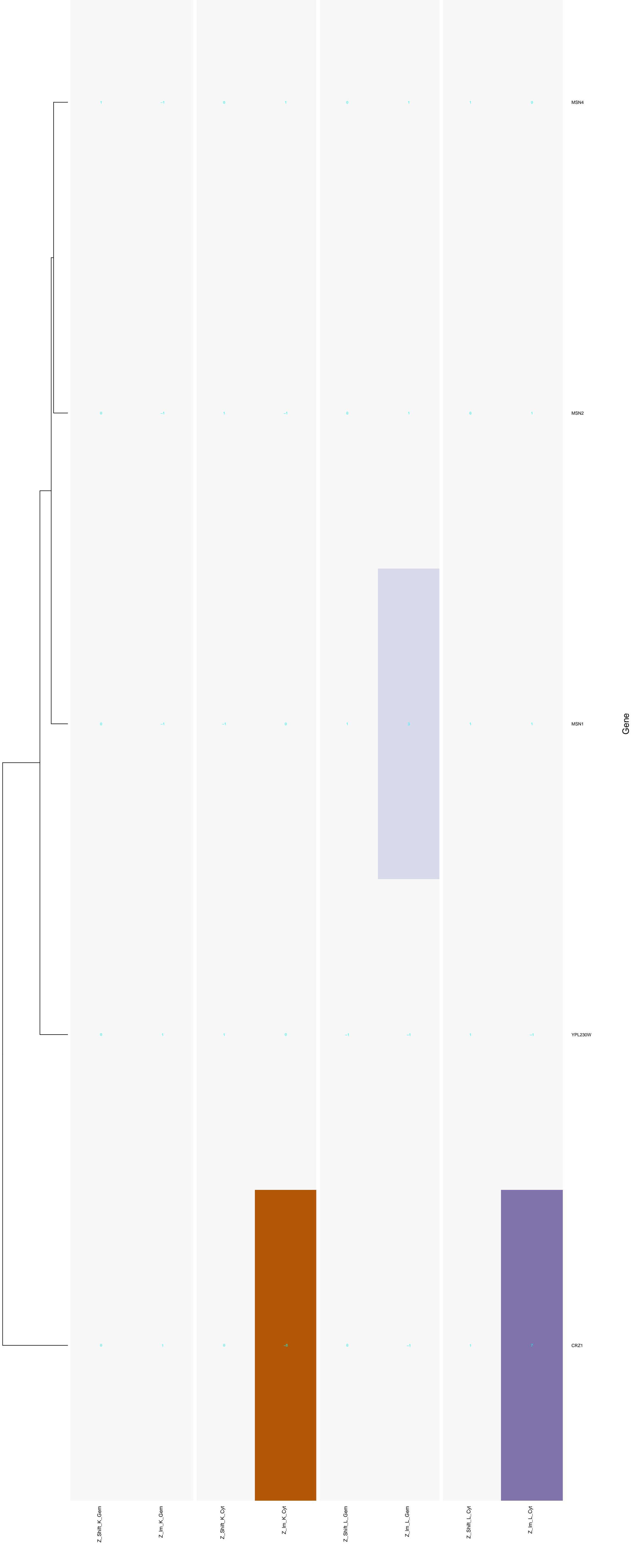
regulation of transcription from RNA polymerase II promoter in response to alkaline pH

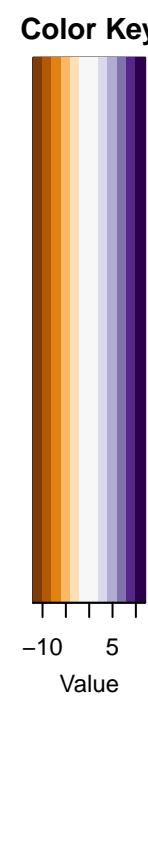


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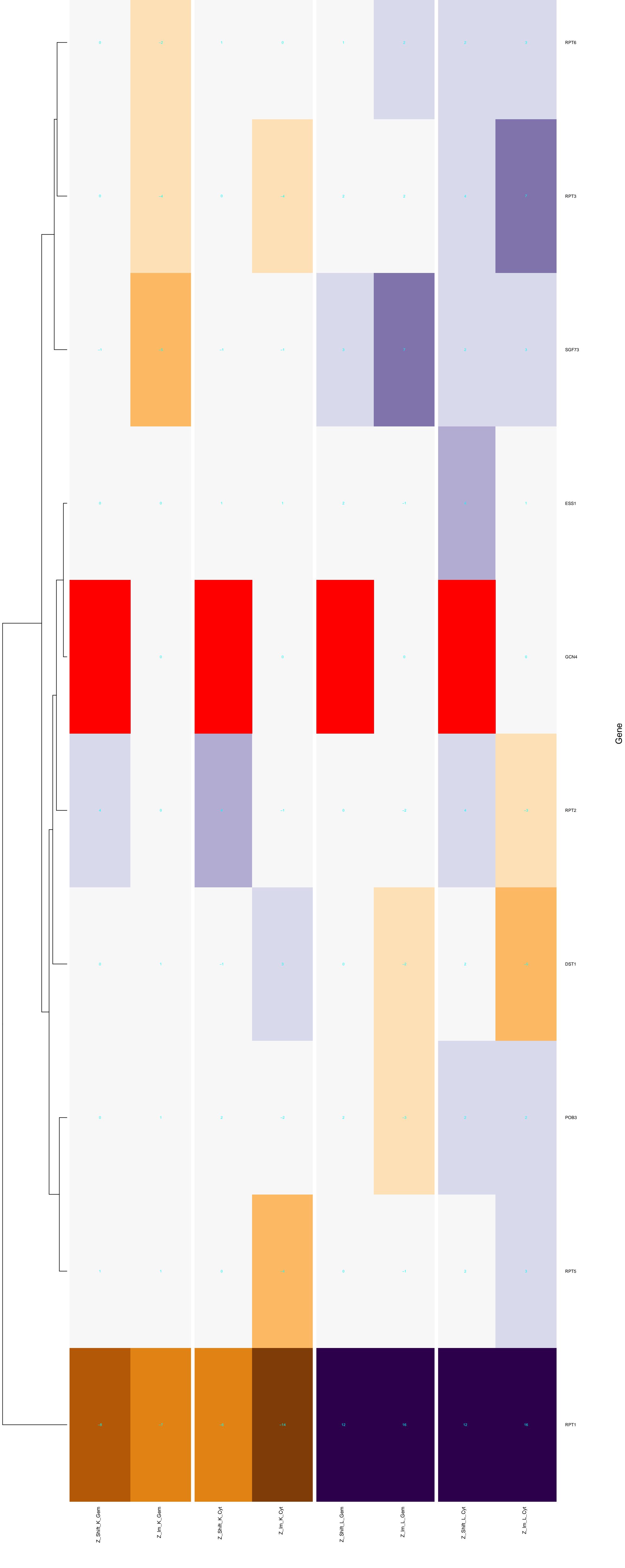


Evaluation of transcription from RNA polymerase II promoter in response to salt stress

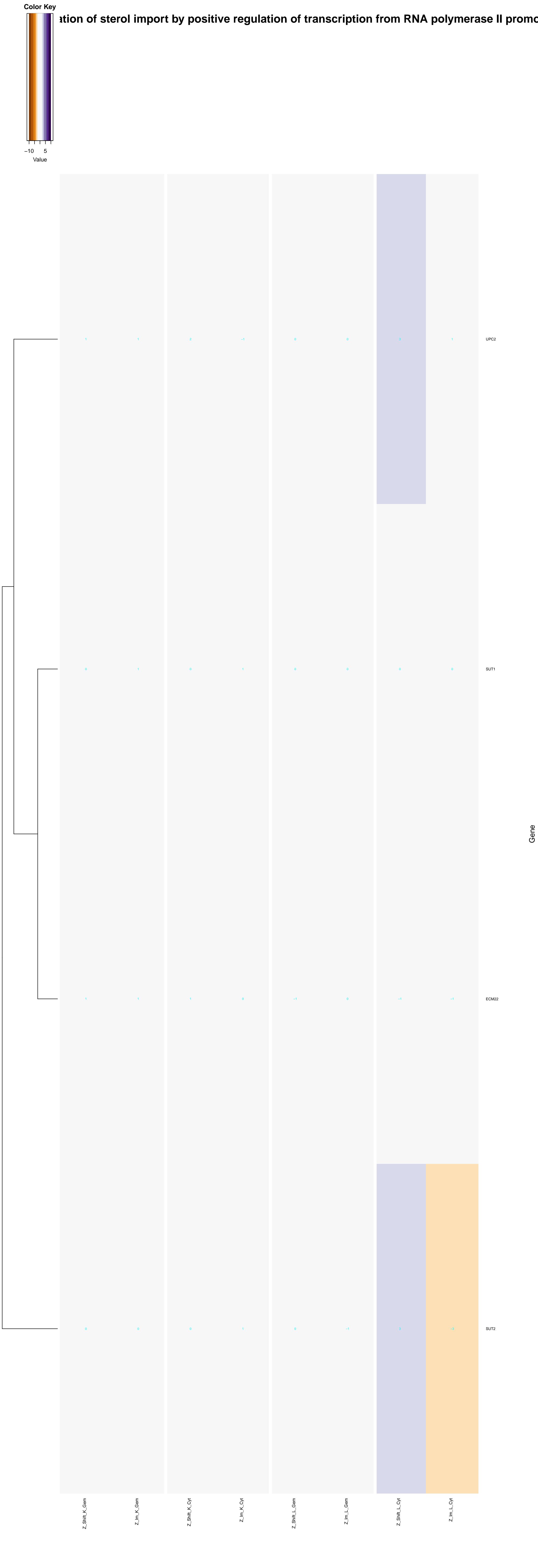
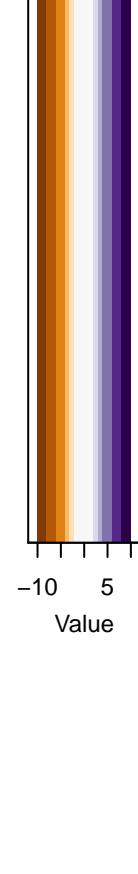




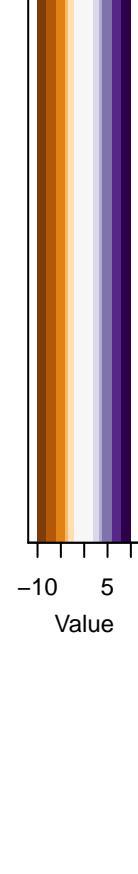
Relative regulation of RNA polymerase II transcriptional preinitiation complex assembly



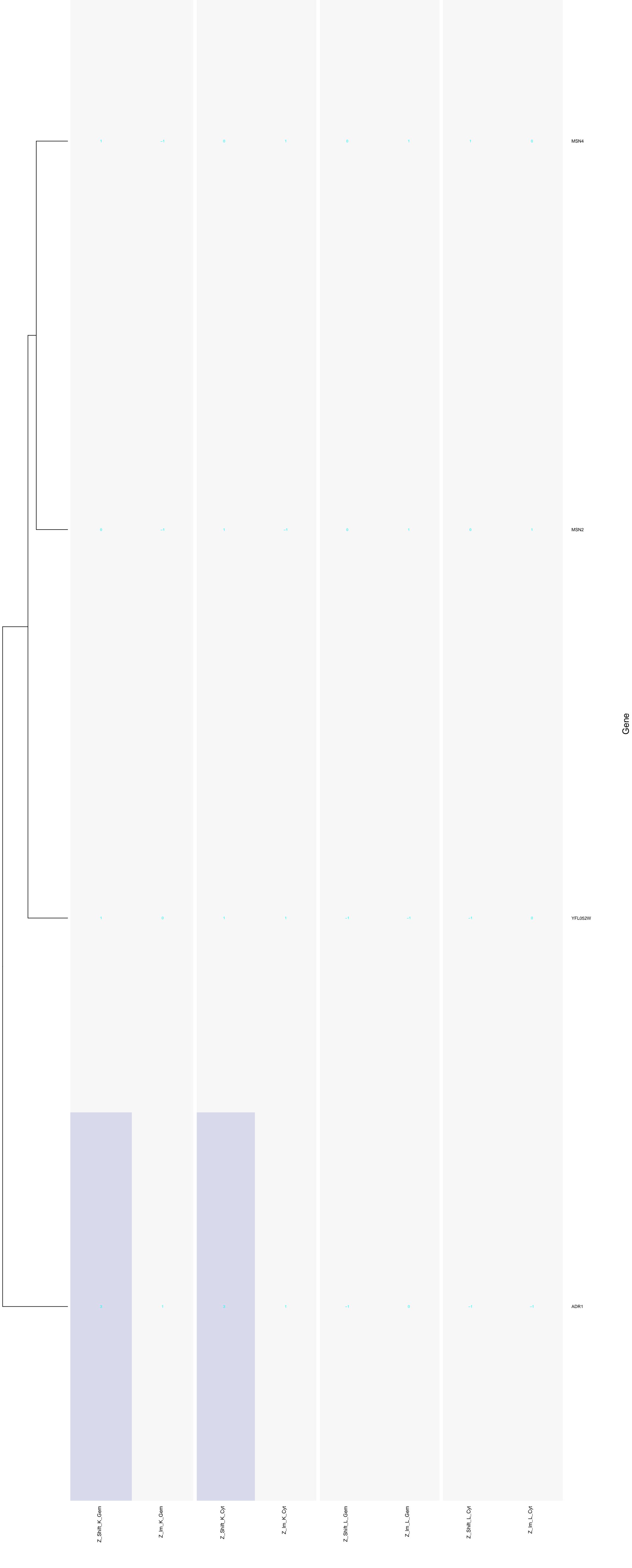
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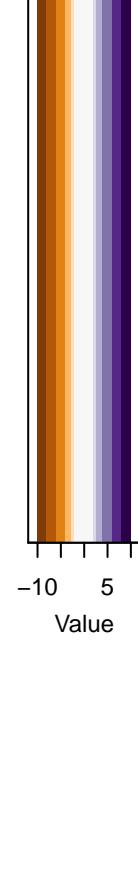
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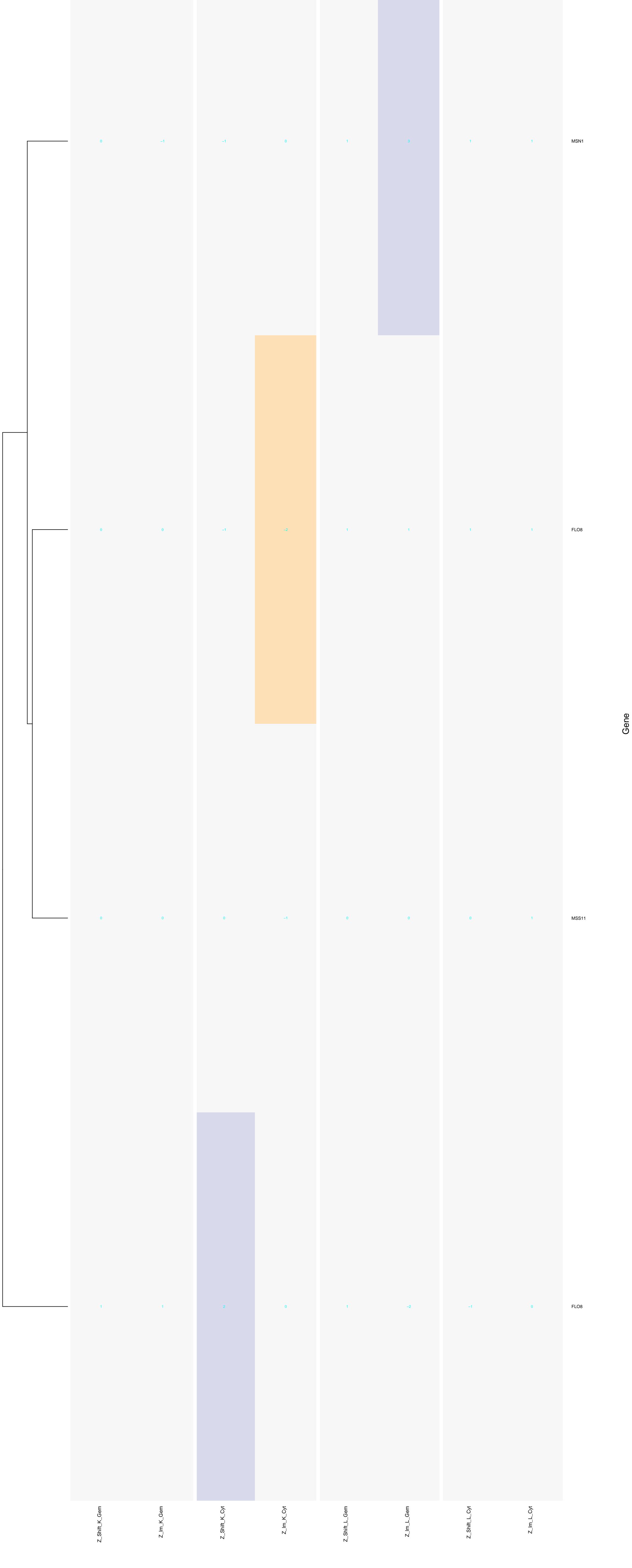
The regulation of transcription from RNA polymerase II promoter in response to ethanol



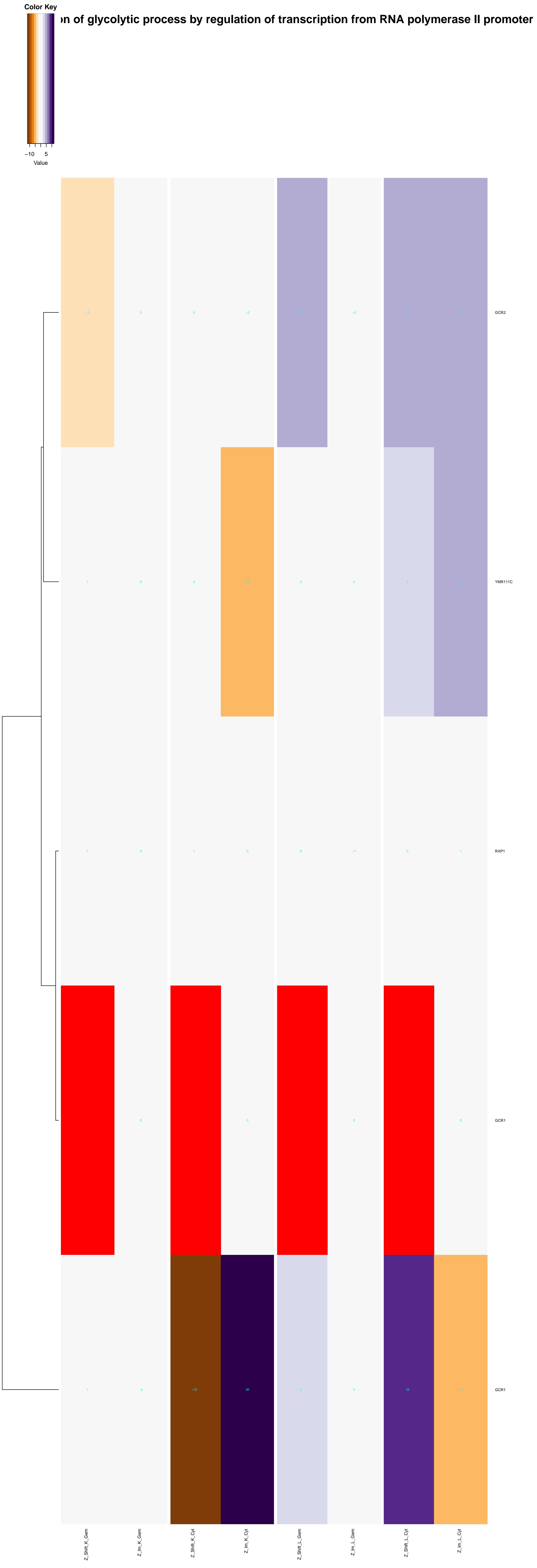
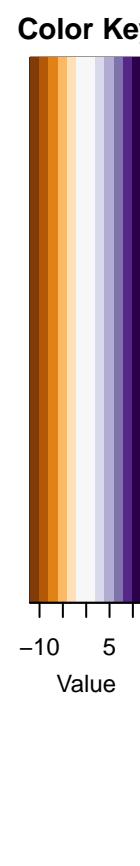
Color Key



of starch catabolic process by regulation of transcription from RNA polymerase II promoter



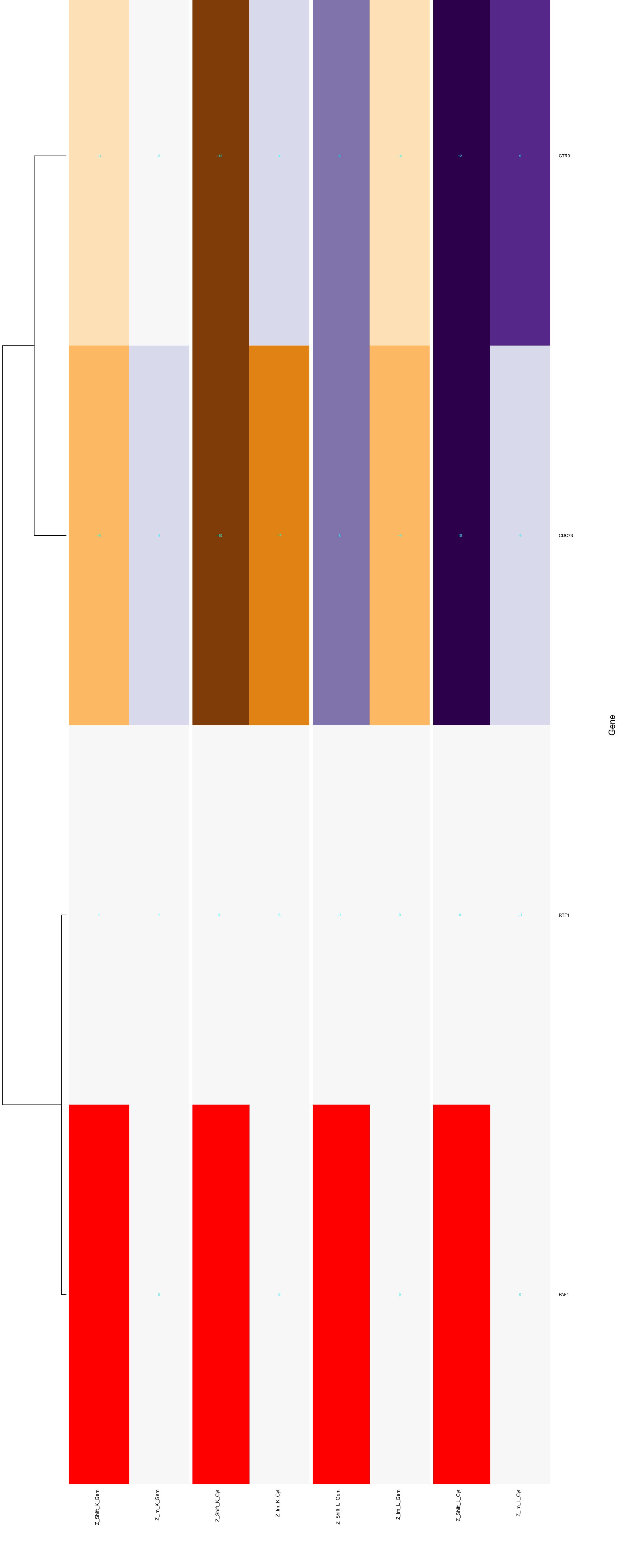
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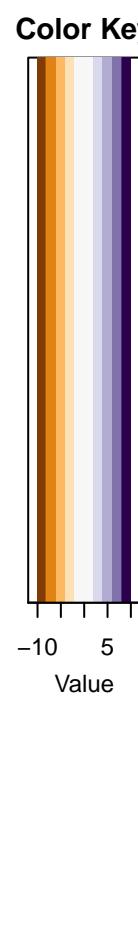
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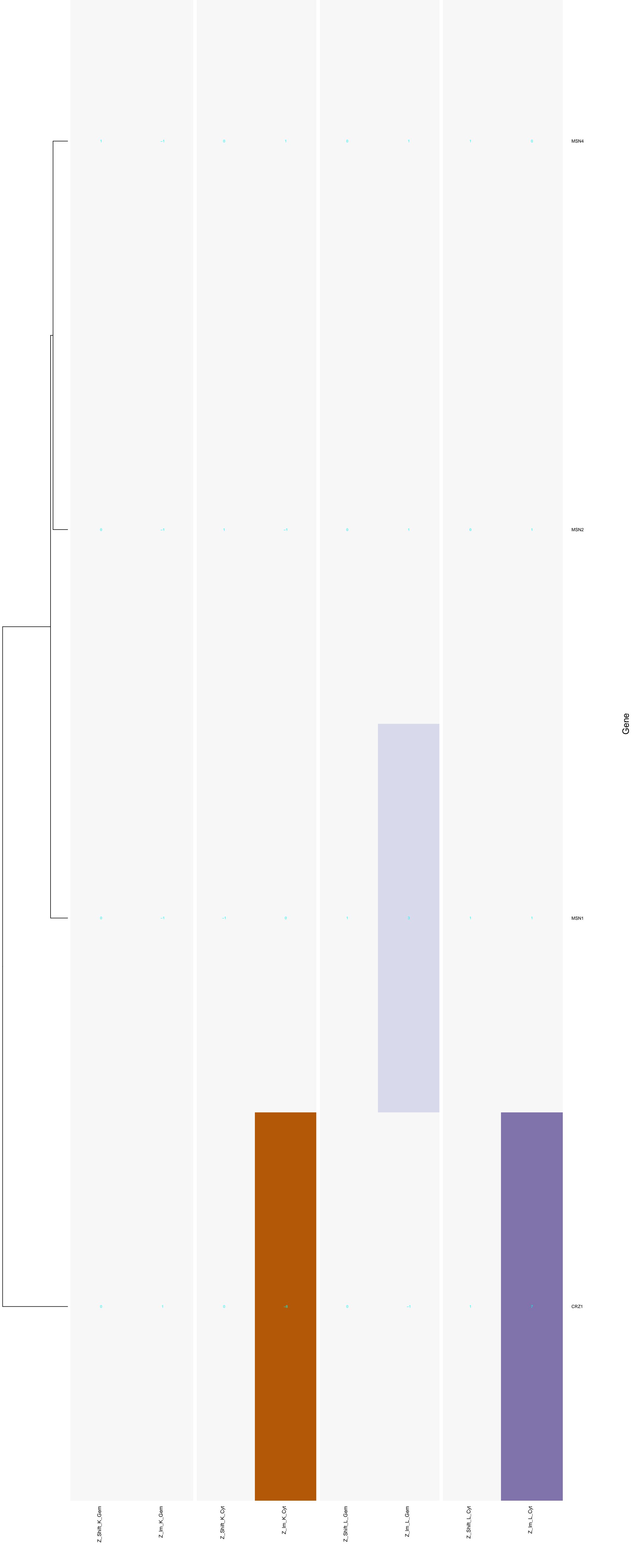
Regulation of phosphorylation of RNA polymerase II C-terminal domain serine 2 residues



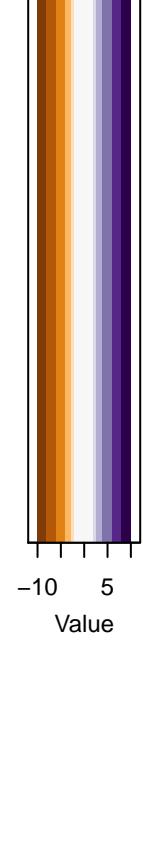
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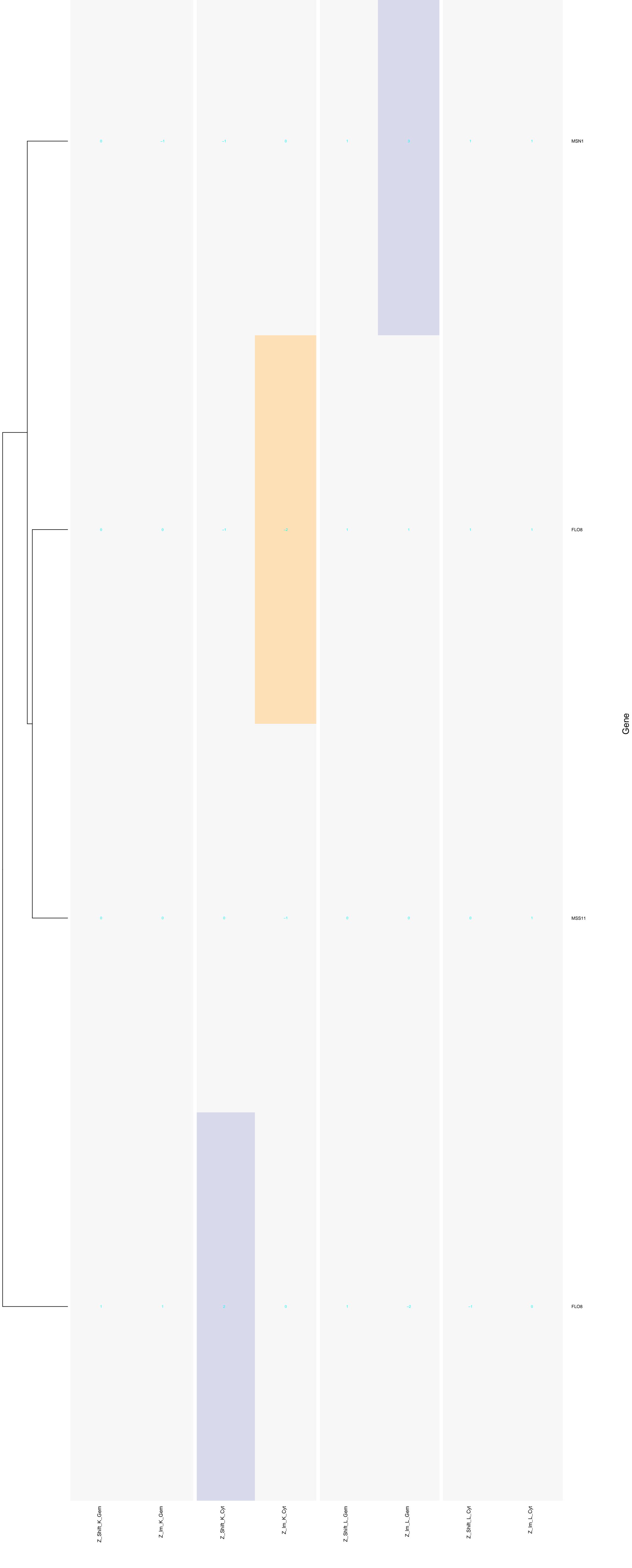
regulation of transcription from RNA polymerase II promoter in response to salt stress



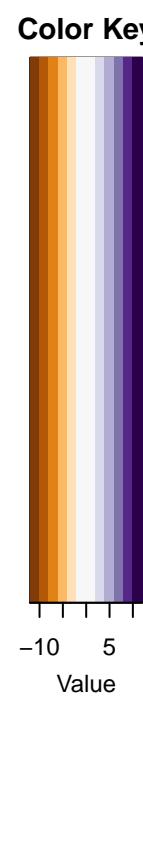
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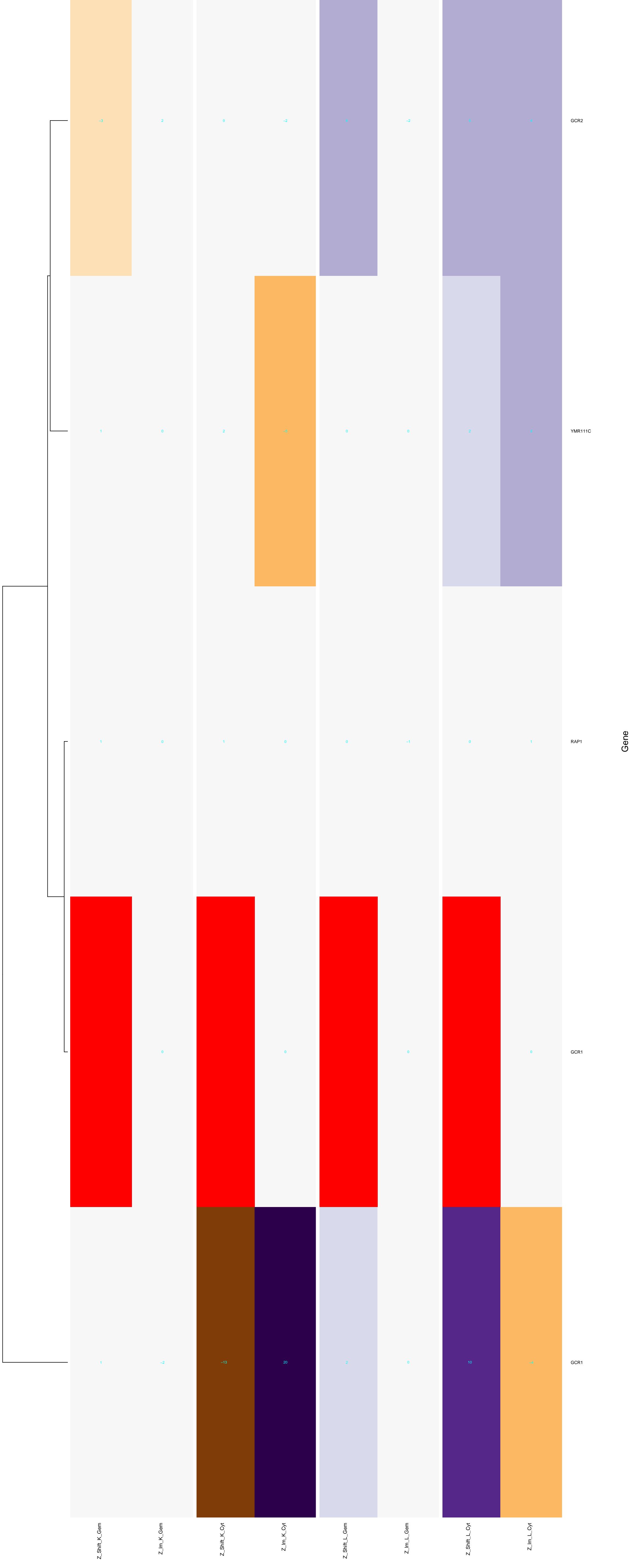
of starch catabolic process by positive regulation of transcription from RNA polymerase II



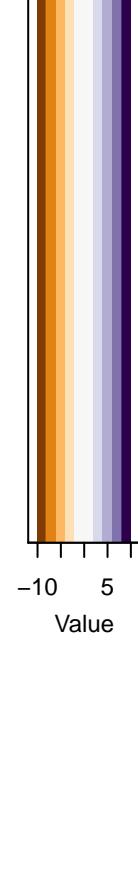
Color Key



glycolytic process by positive regulation of transcription from RNA polymerase II promoter



Color Key



Regulation of transcription from RNA polymerase II promoter in response to increased salt

