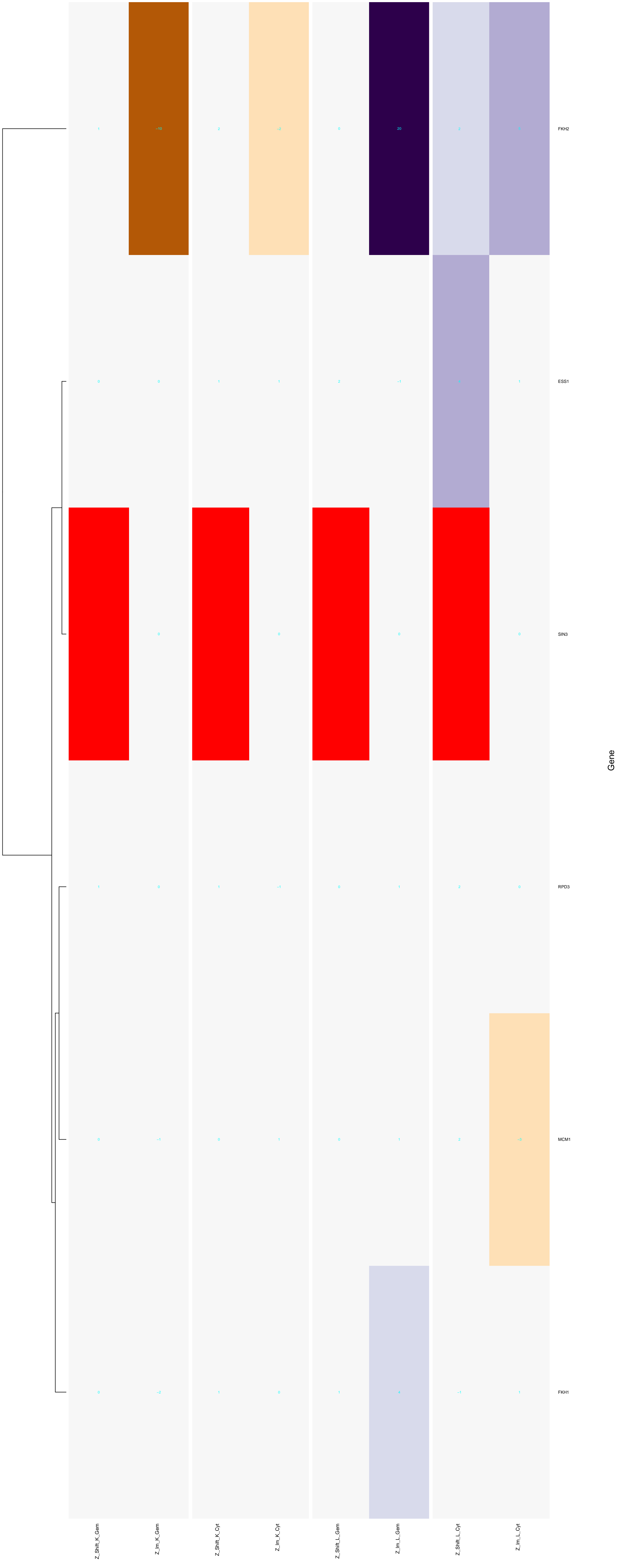
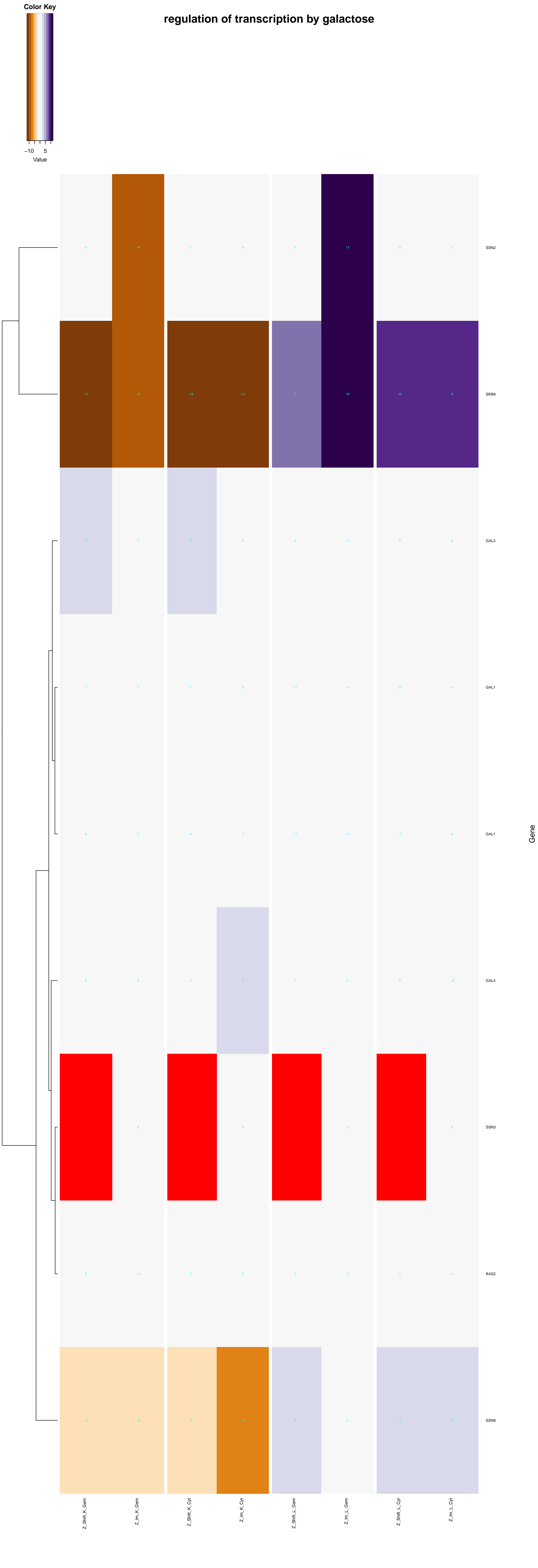
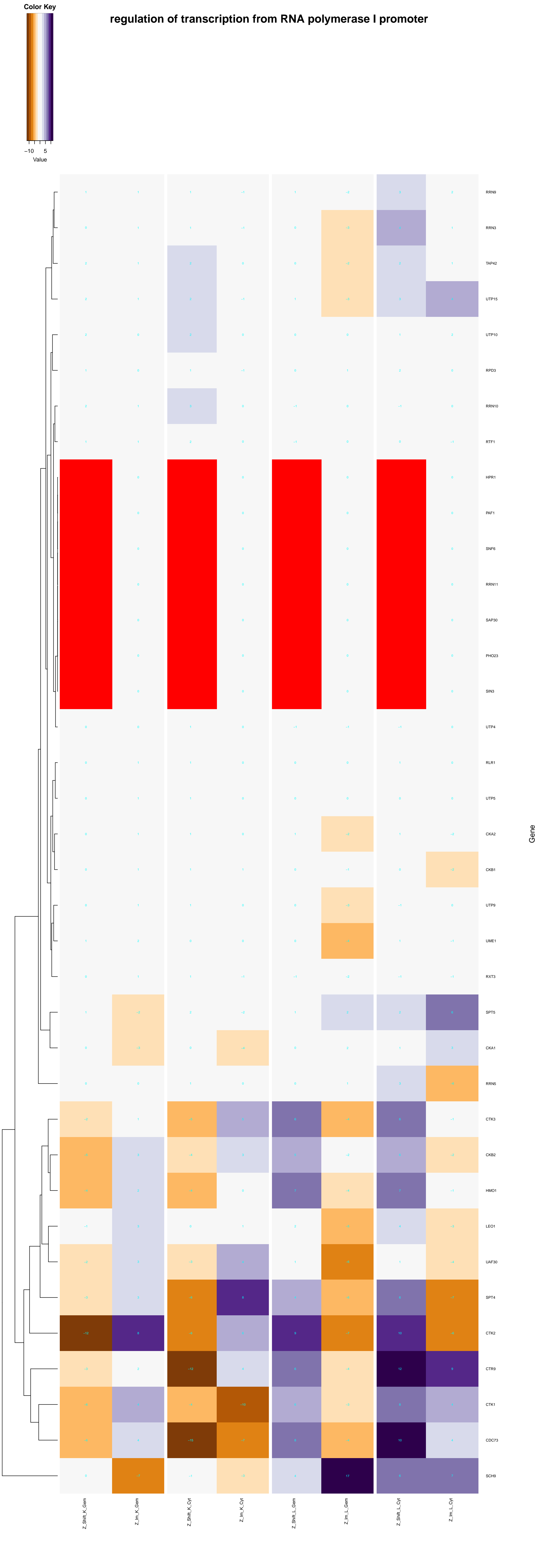


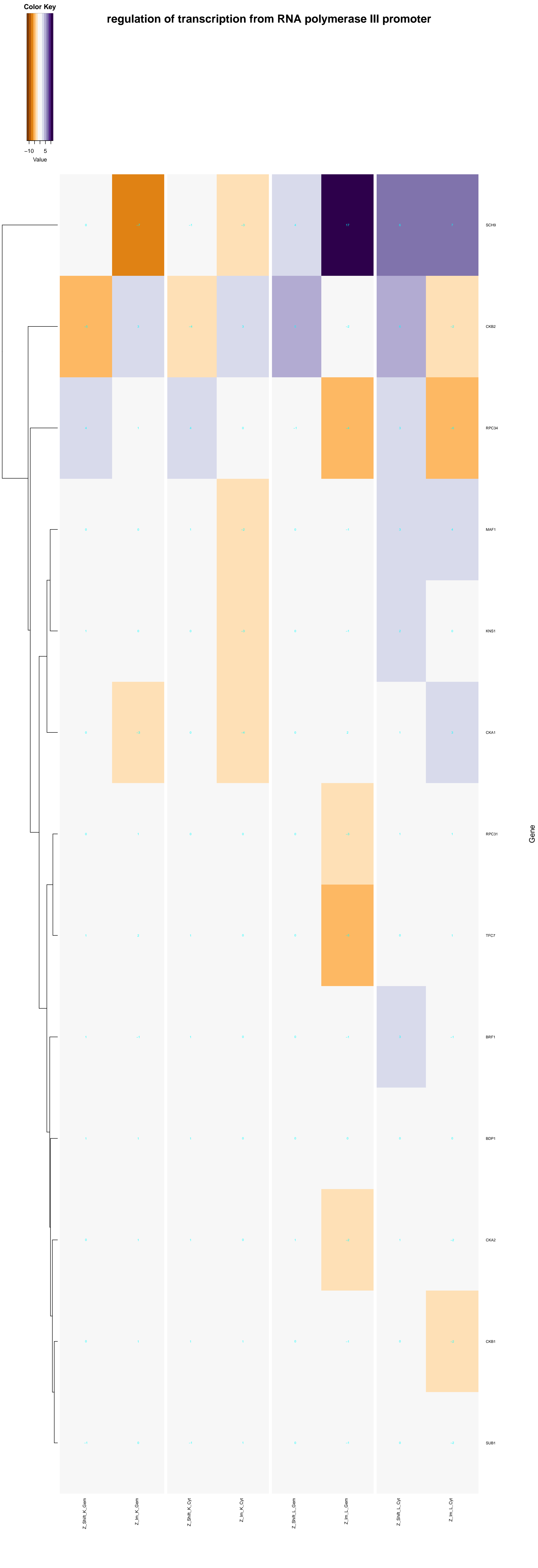
regulation of transcription involved in G2/M transition of mitotic cell cycle

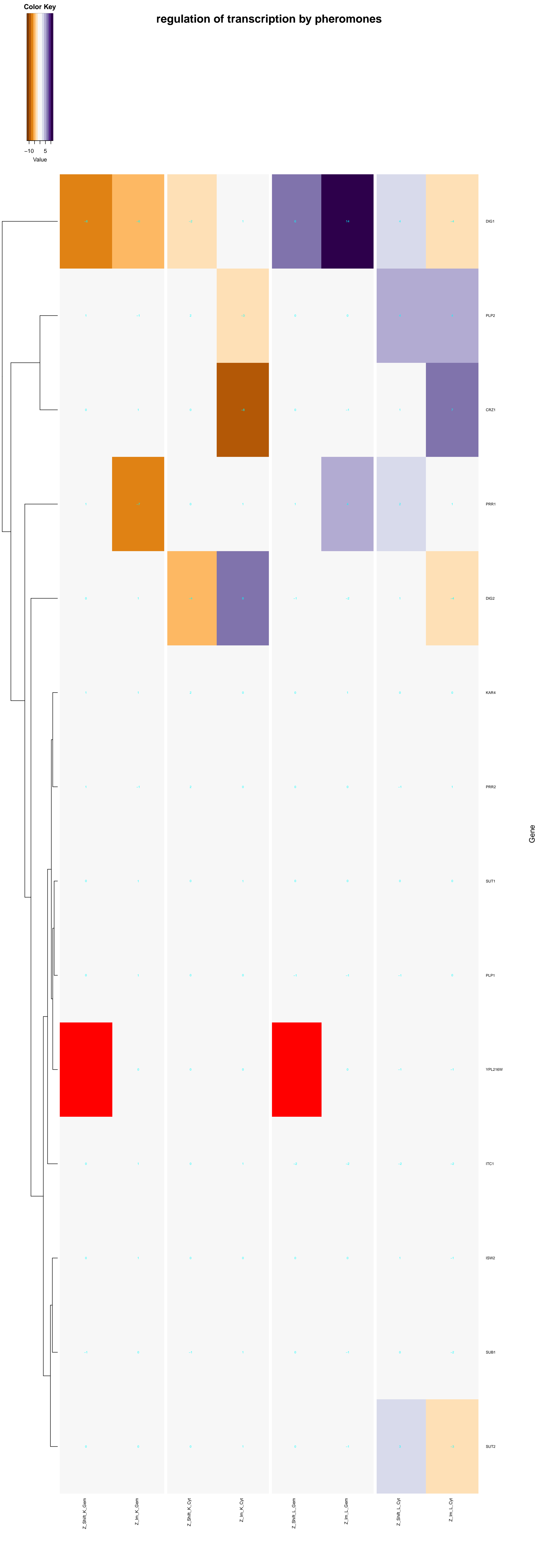


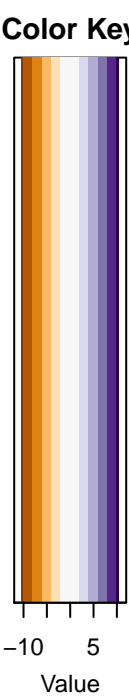




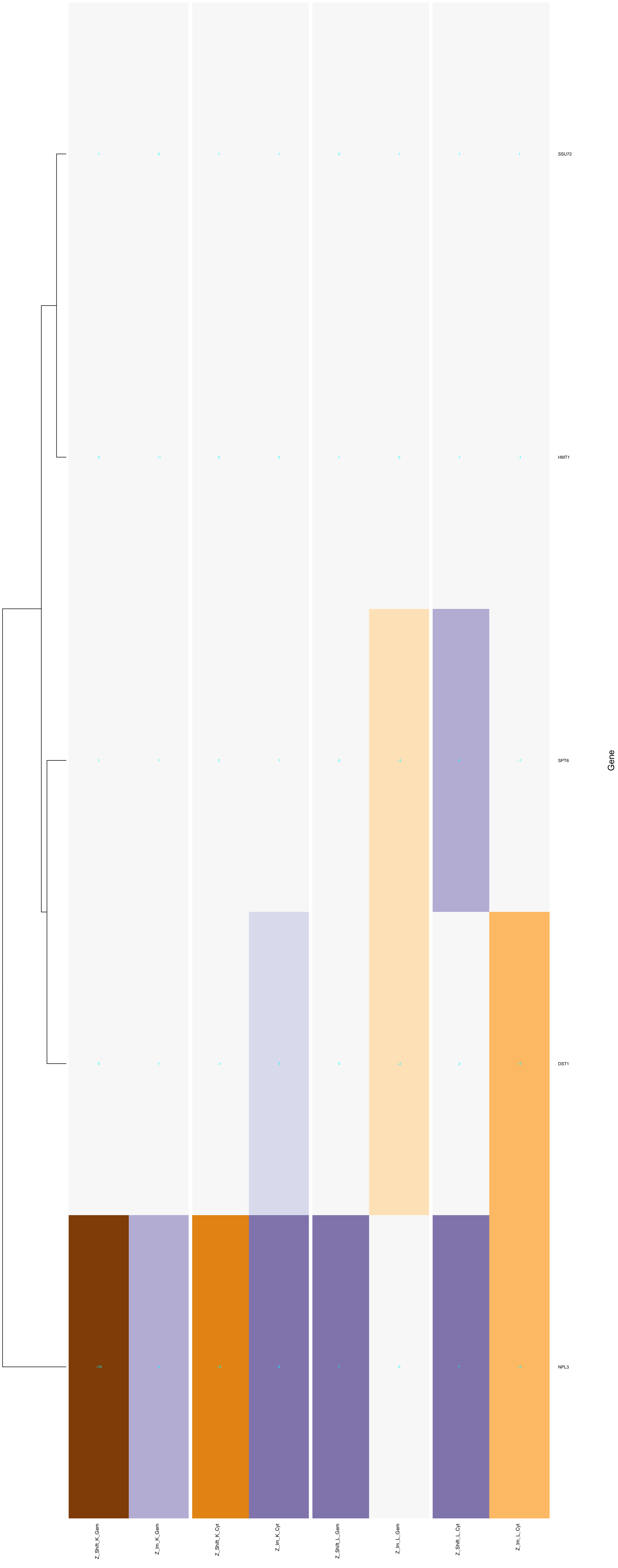


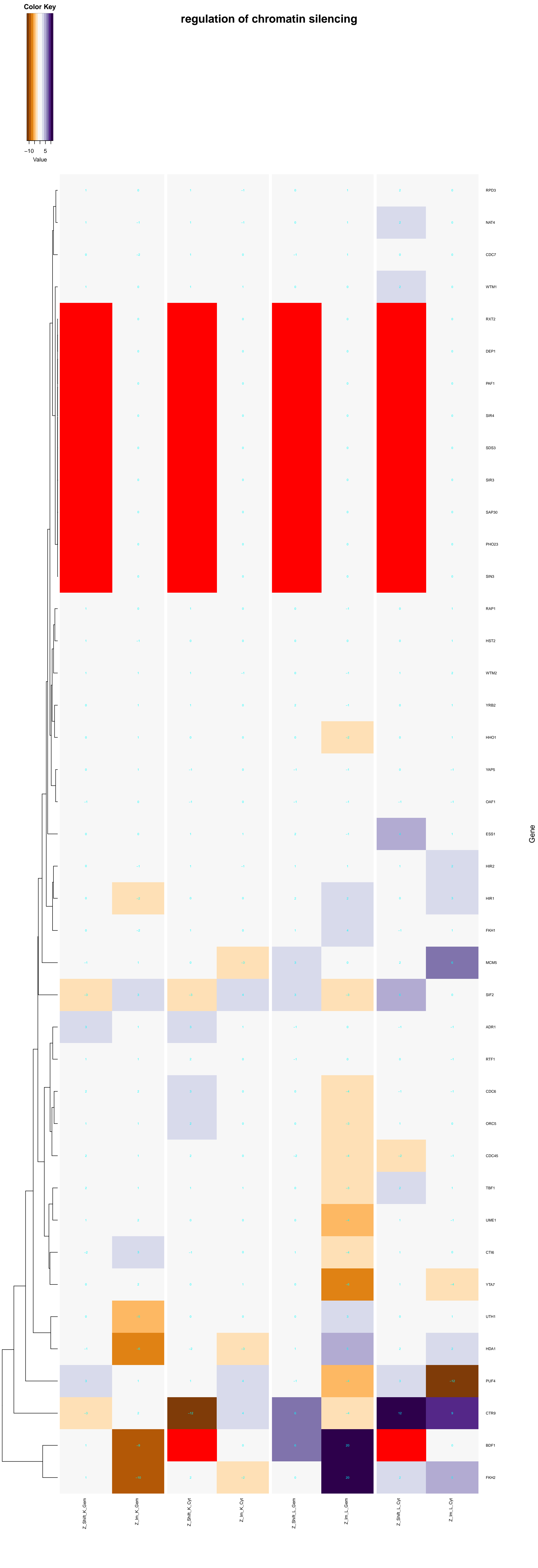


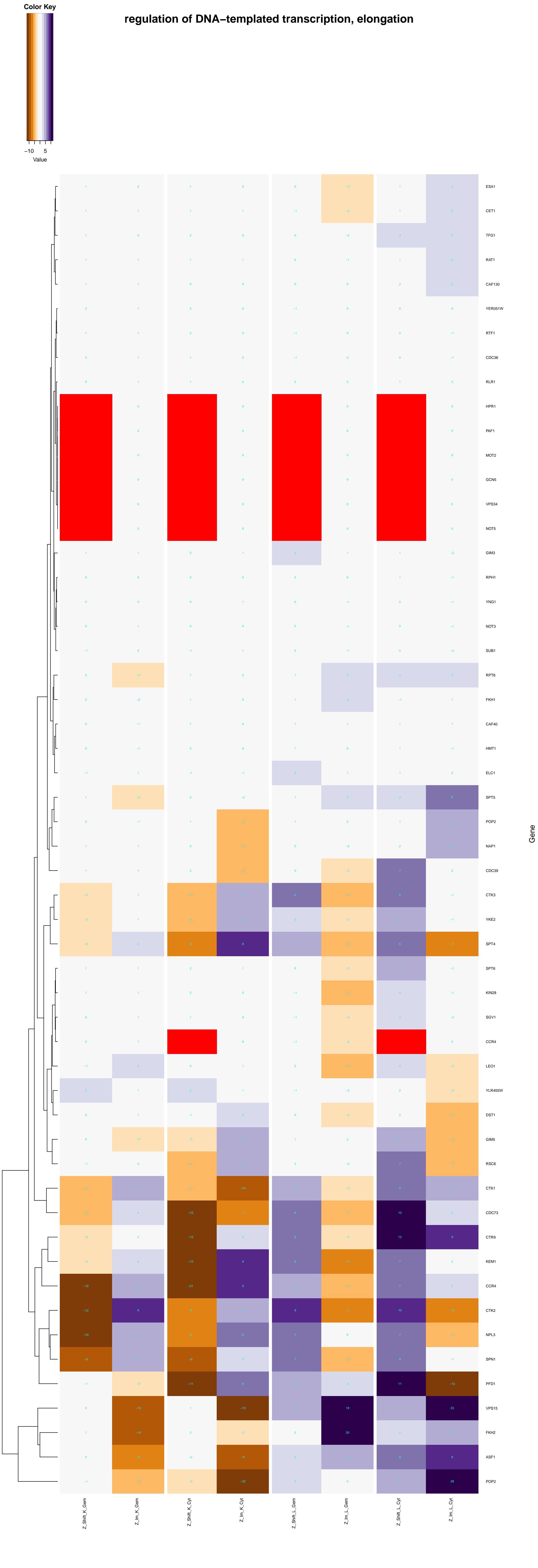


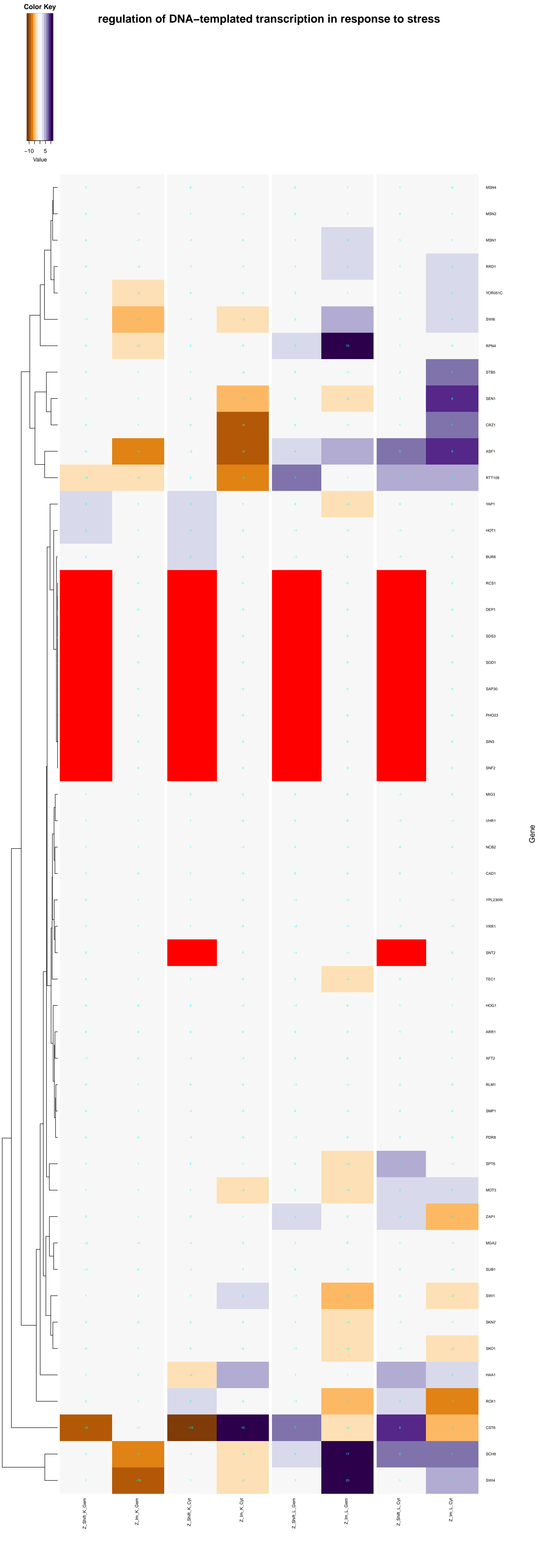


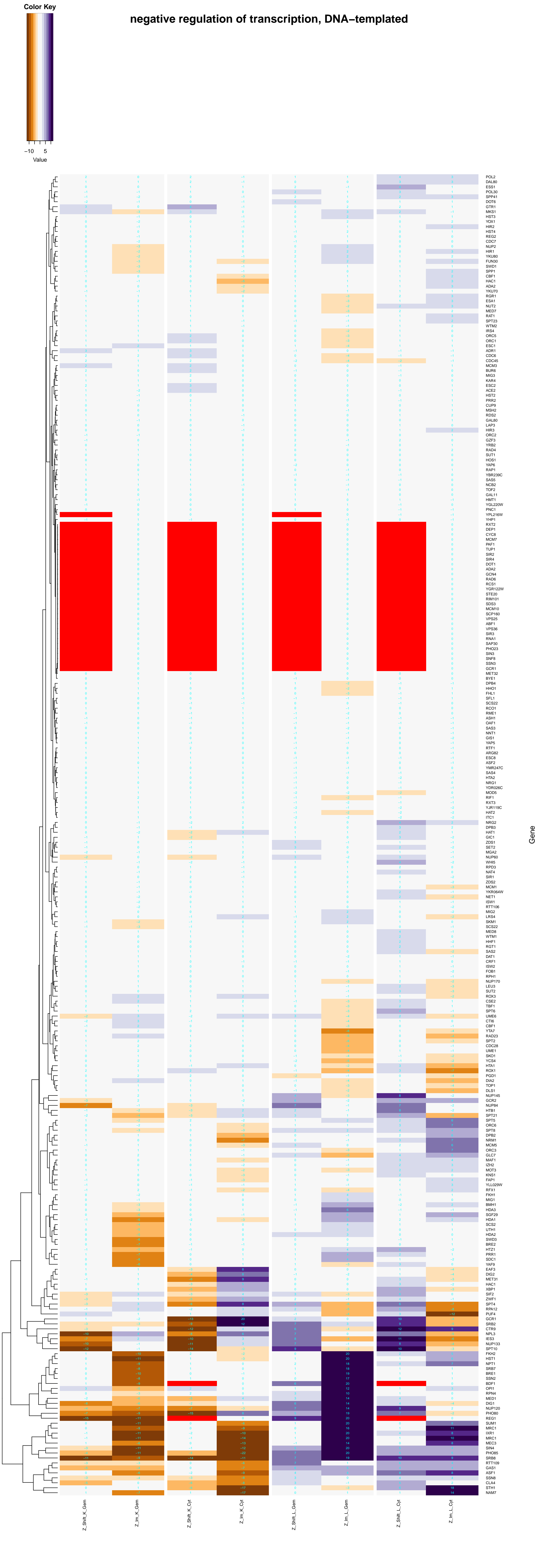
regulation of DNA-templated transcription, termination

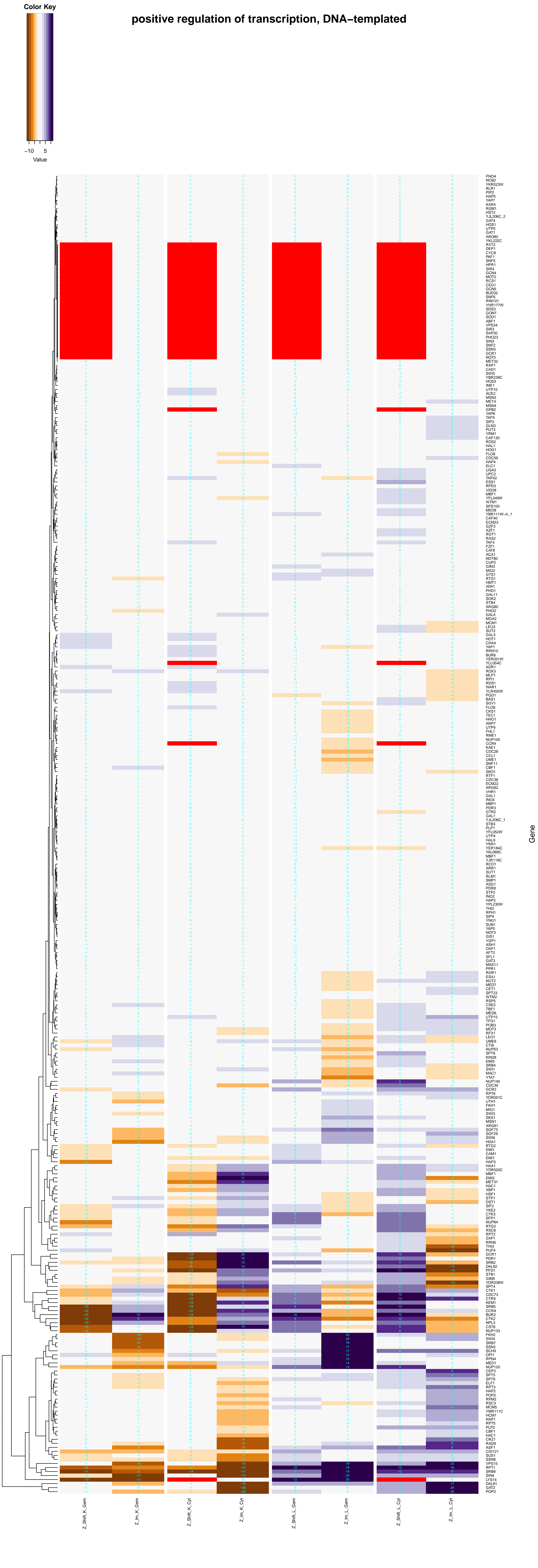


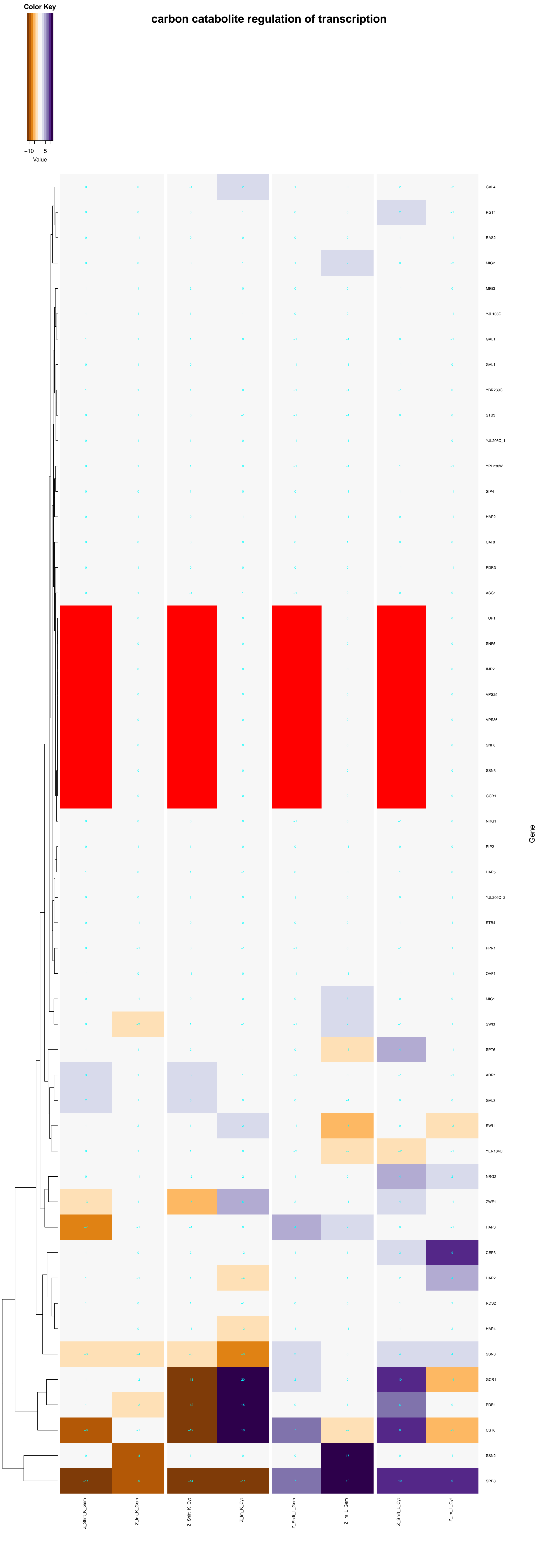


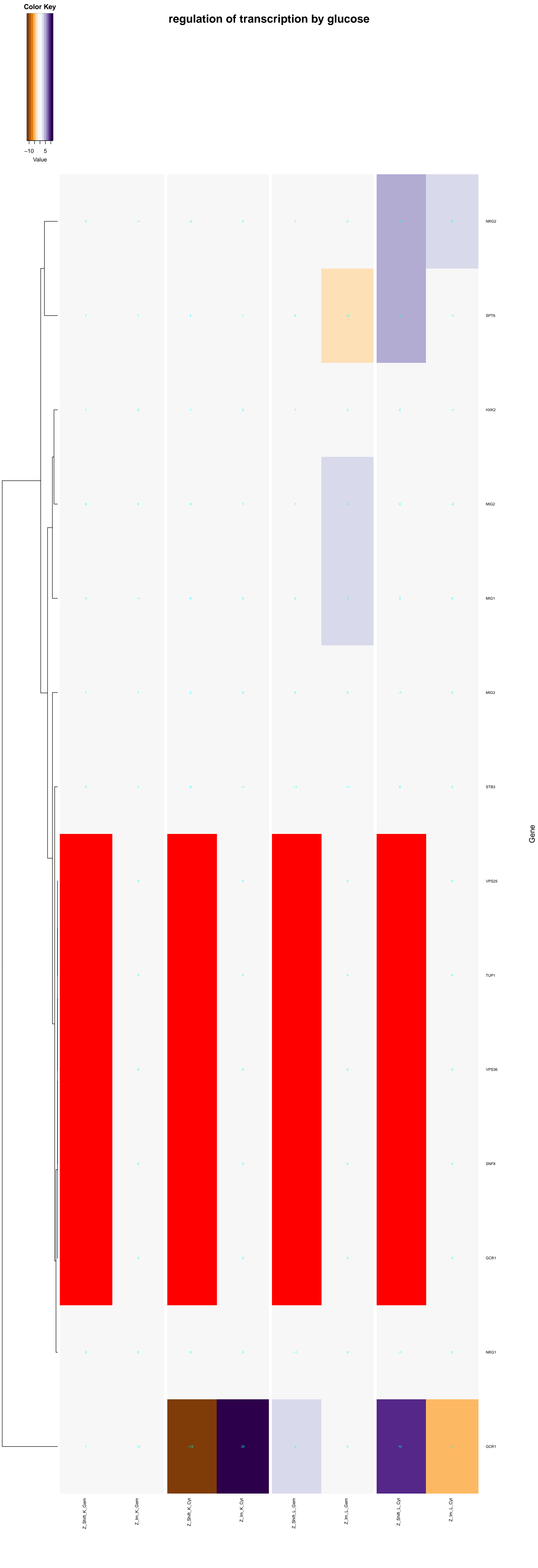


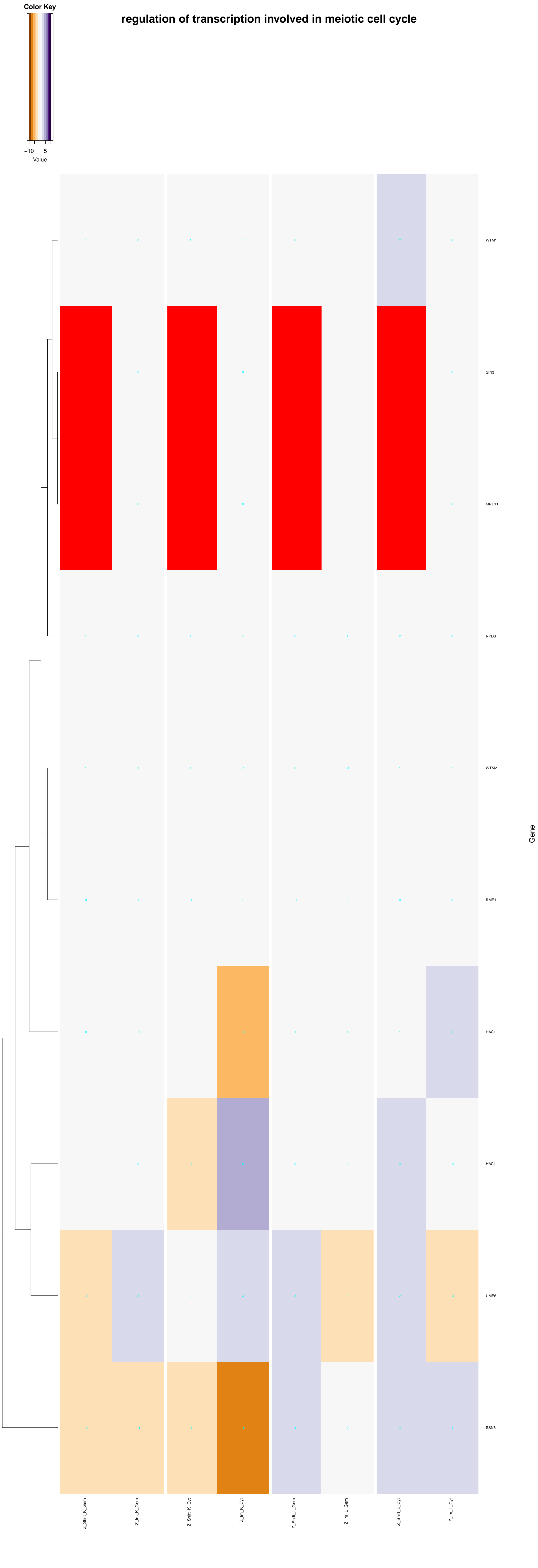


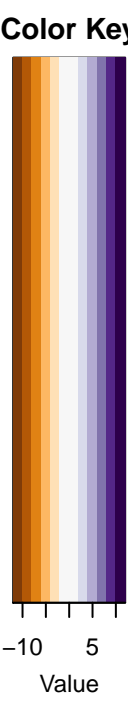




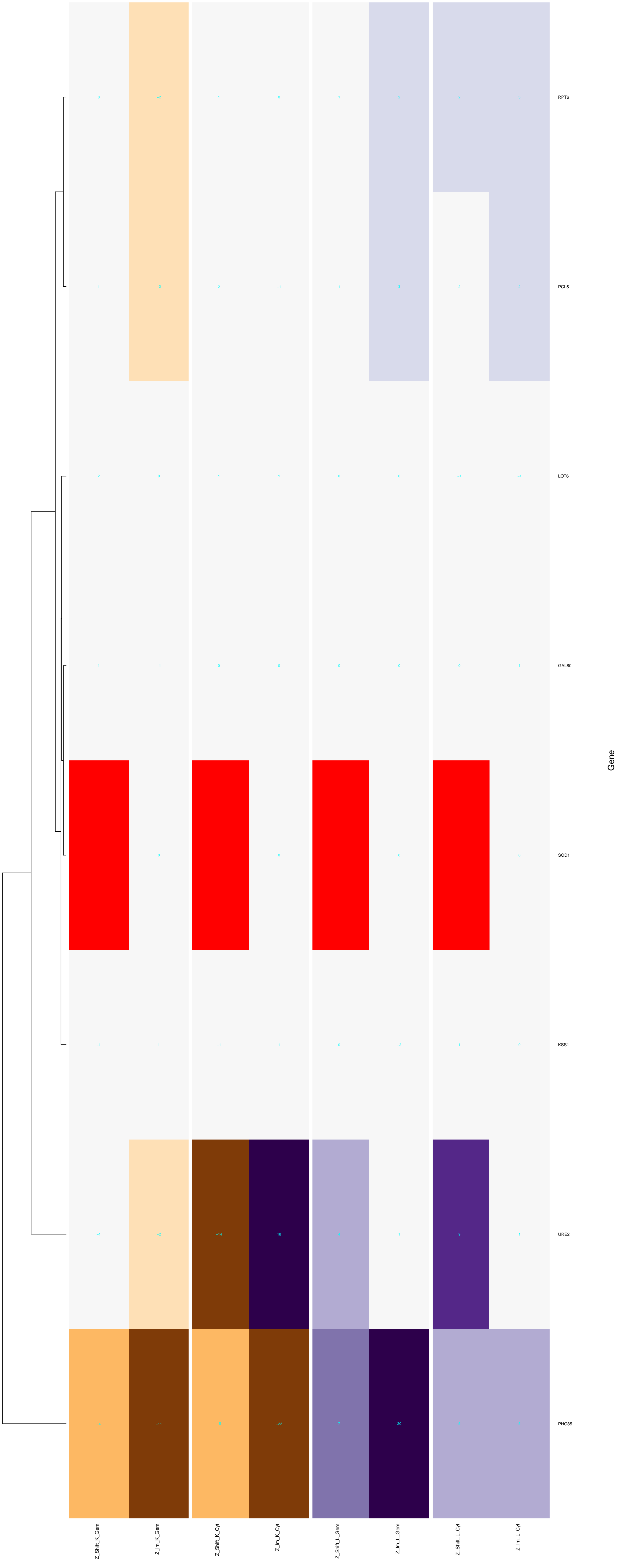




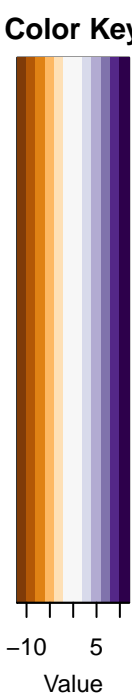




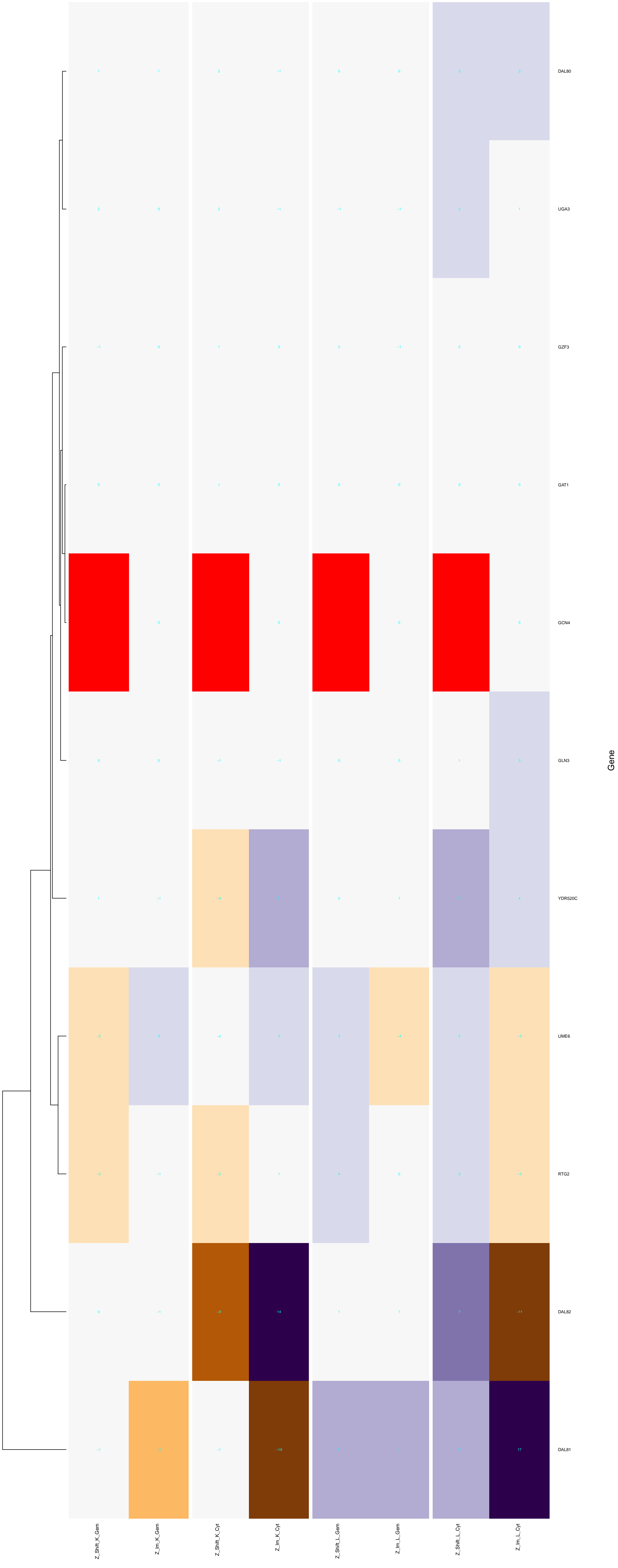
regulation of sequence-specific DNA binding transcription factor activity

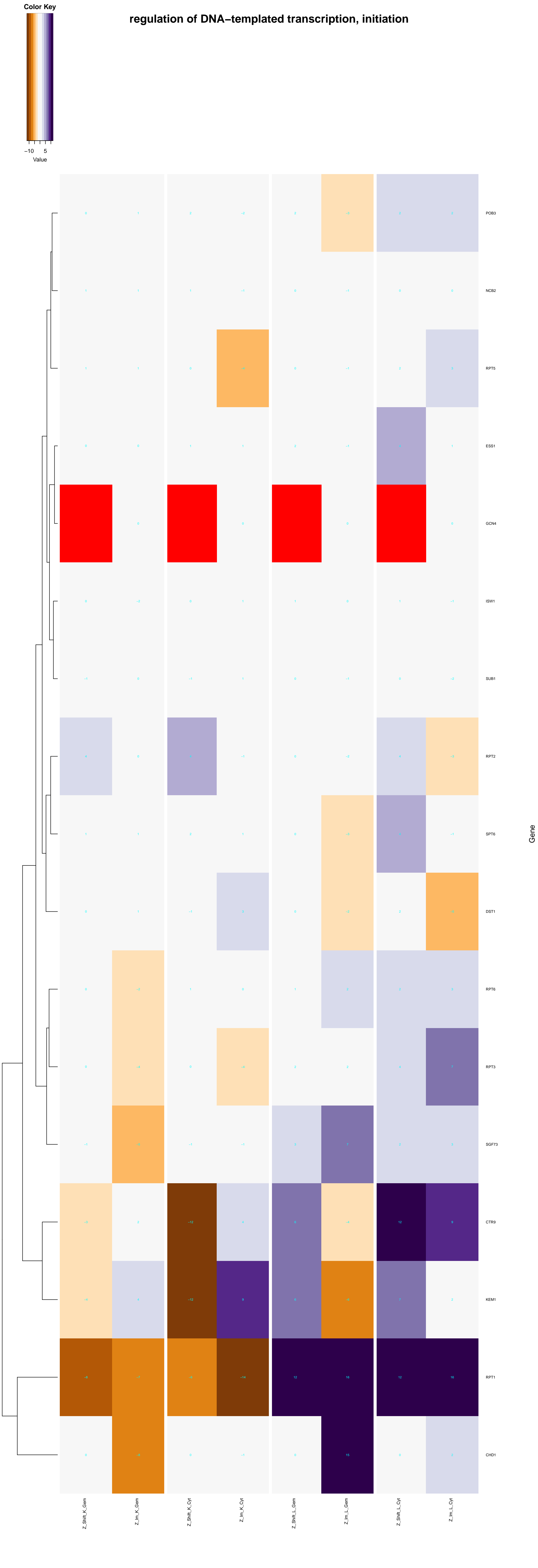






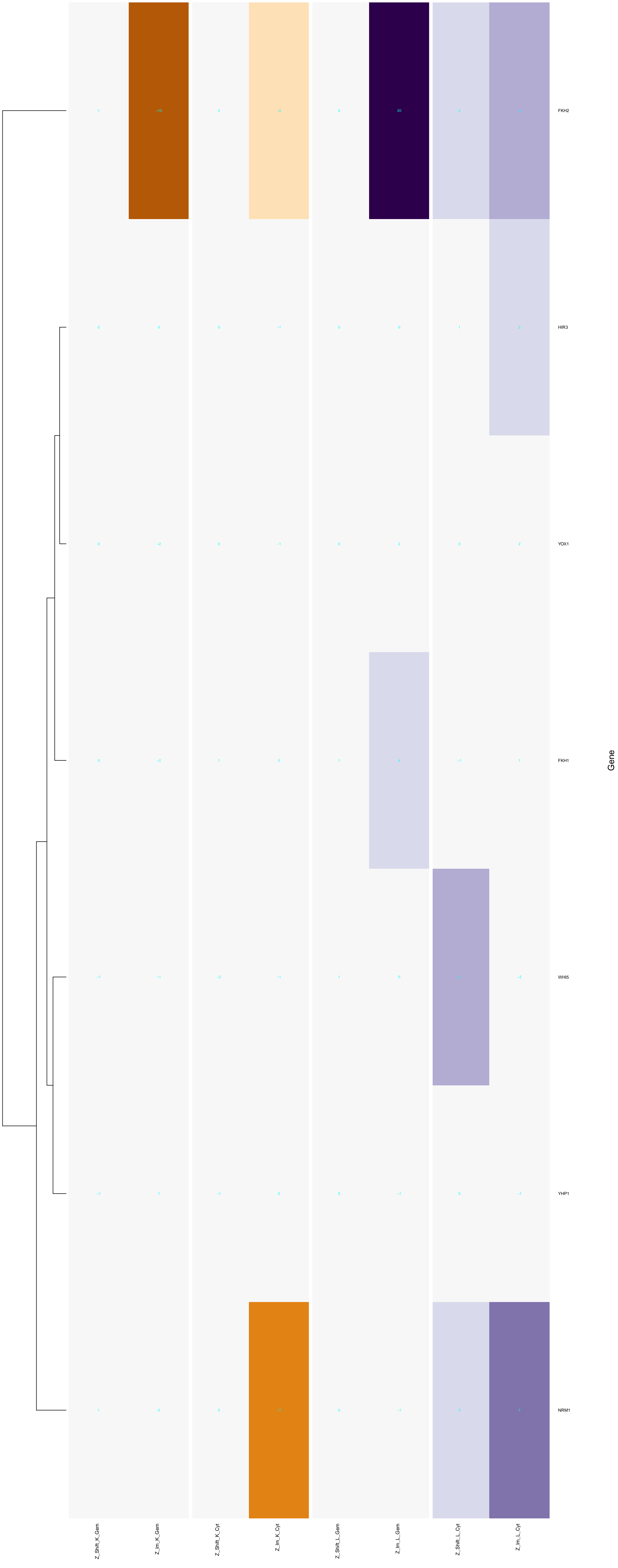
nitrogen catabolite regulation of transcription

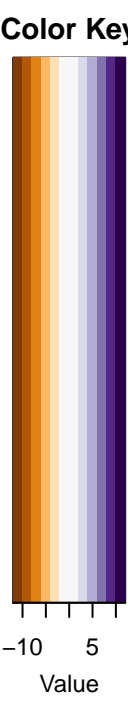




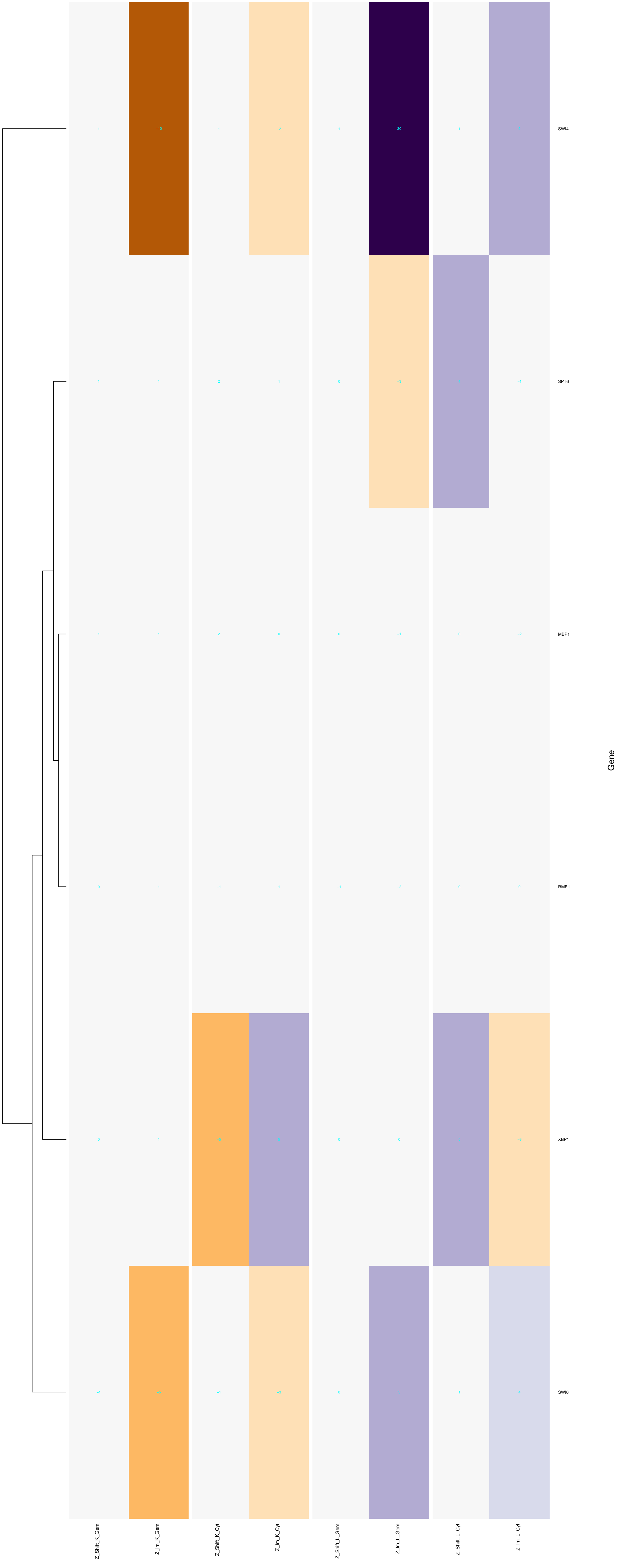


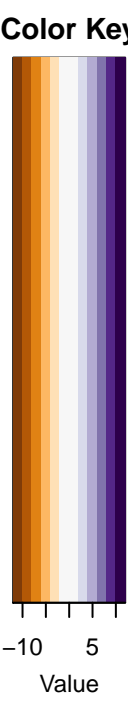
egative regulation of transcription involved in G1/S transition of mitotic cell cycle



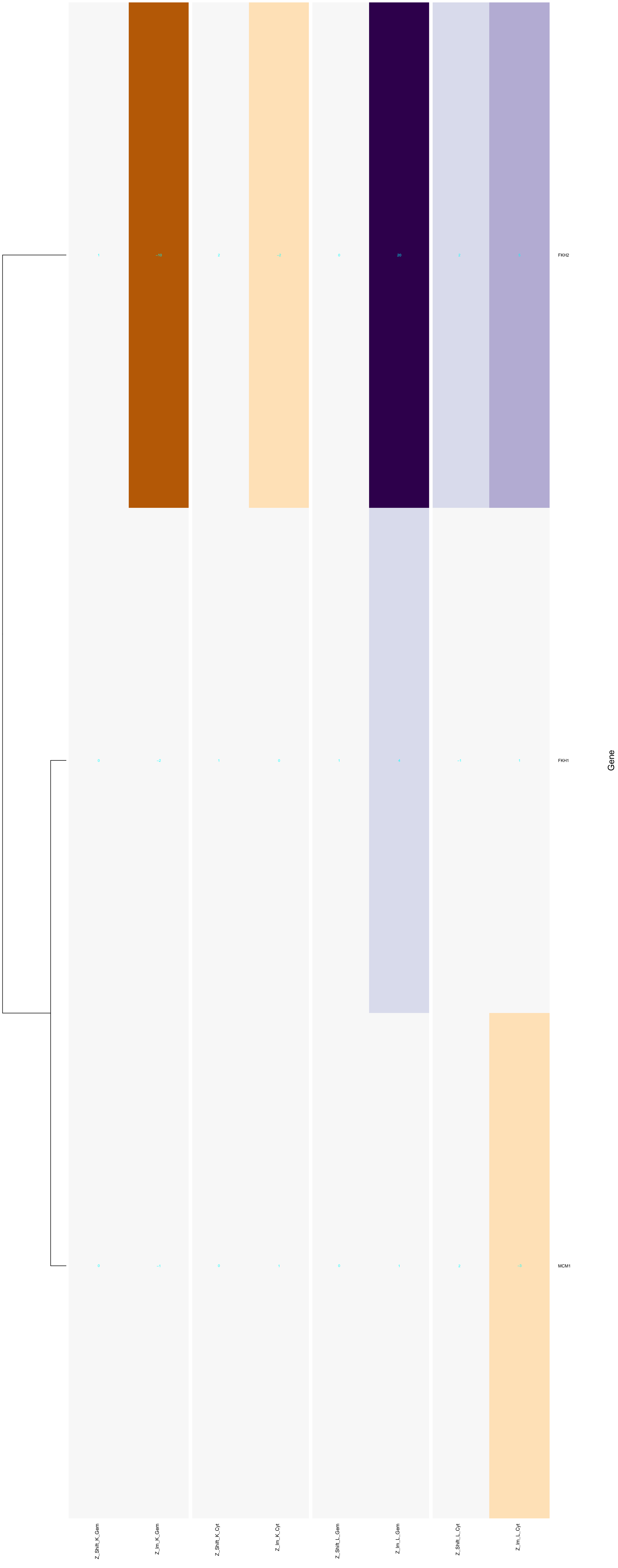


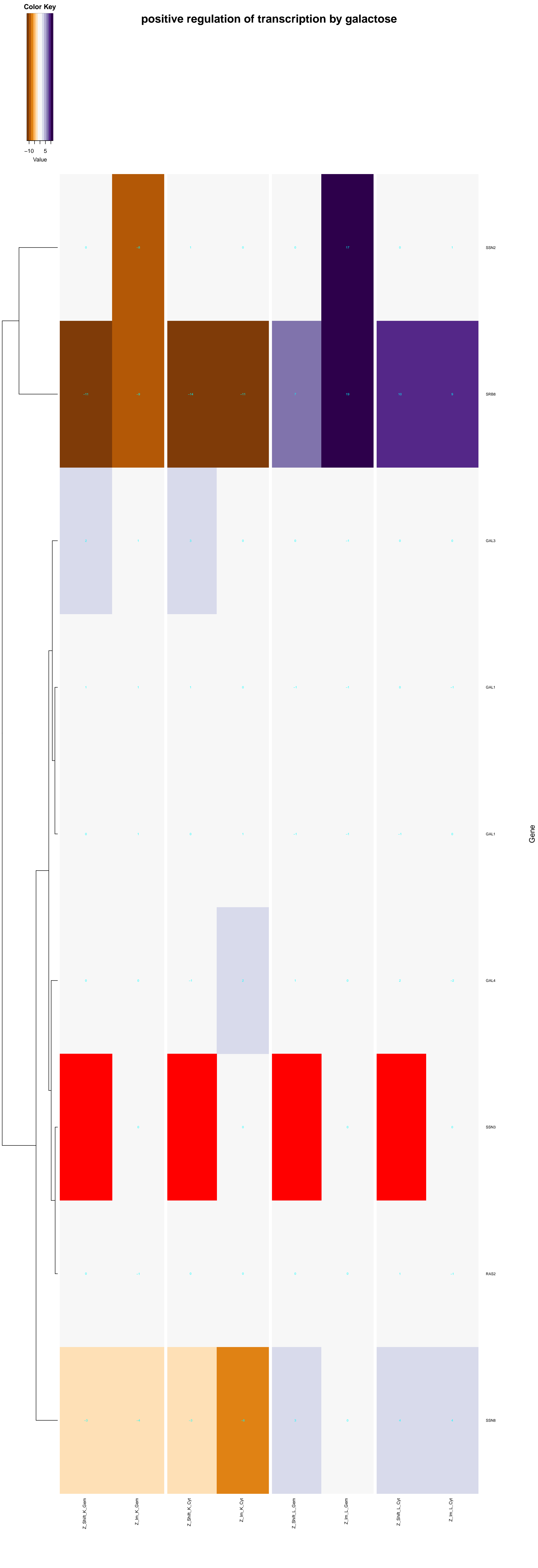
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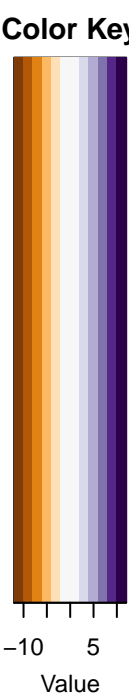




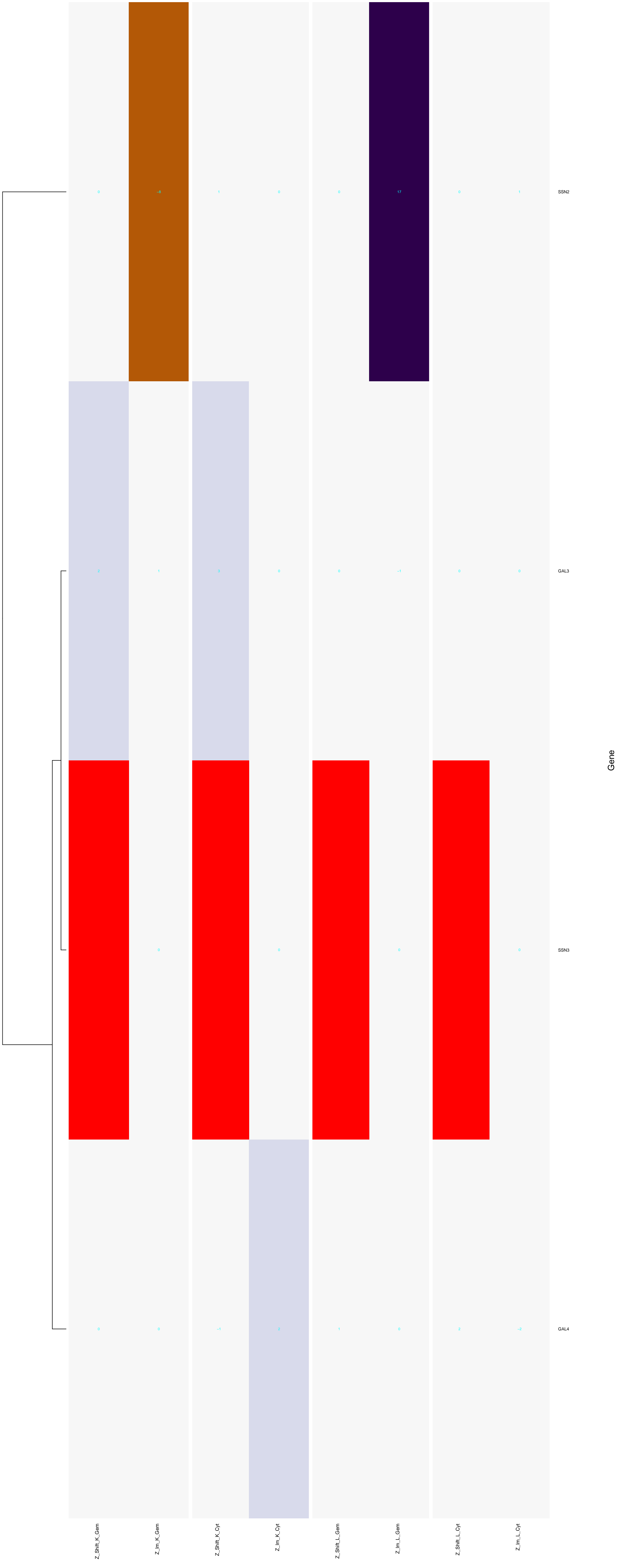
Positive regulation of transcription involved in G2/M transition of mitotic cell cycle

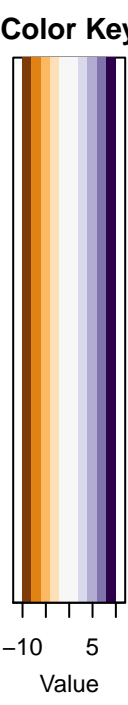




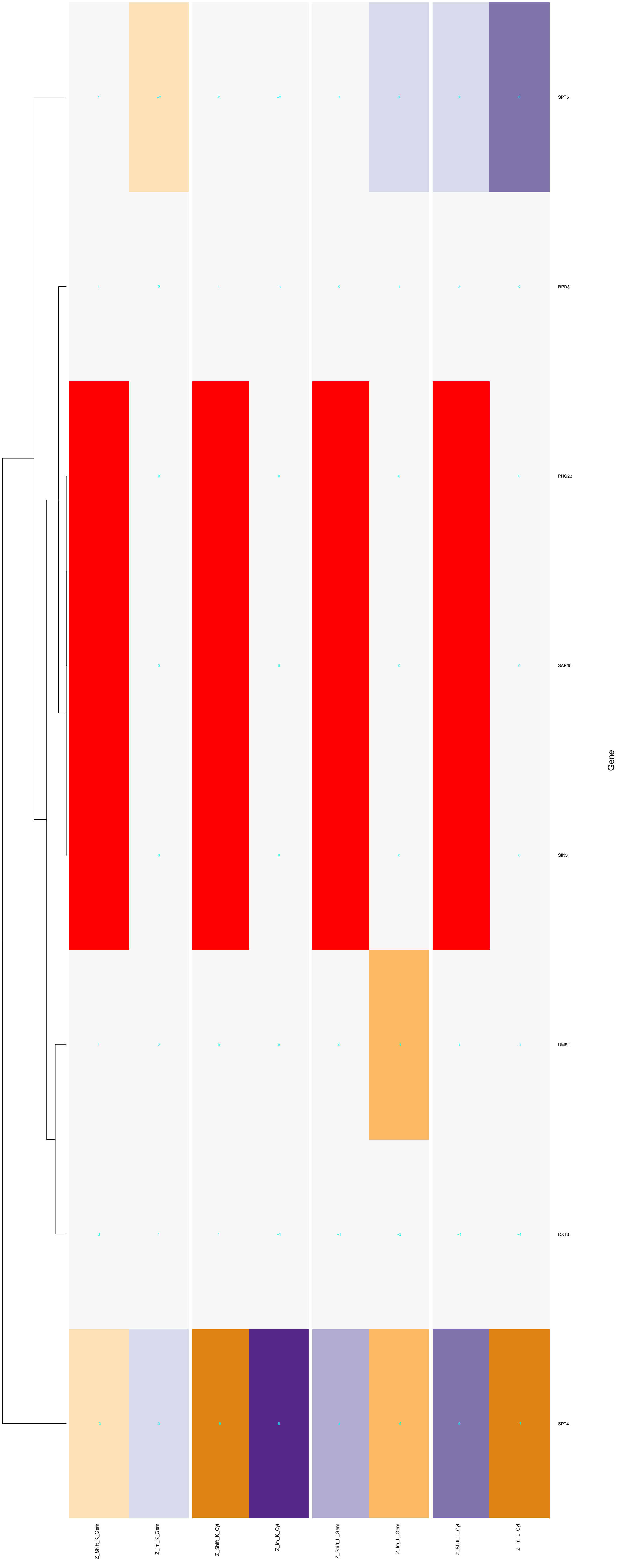


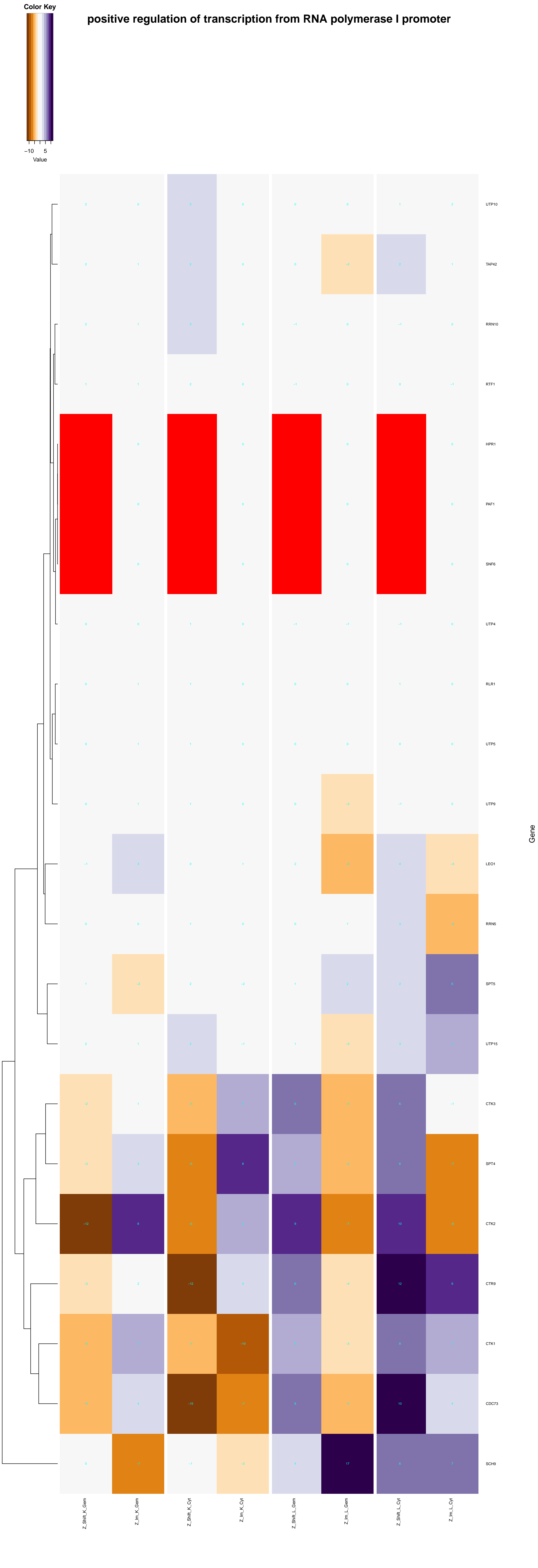
regulation of transcription from RNA polymerase II promoter by galactose





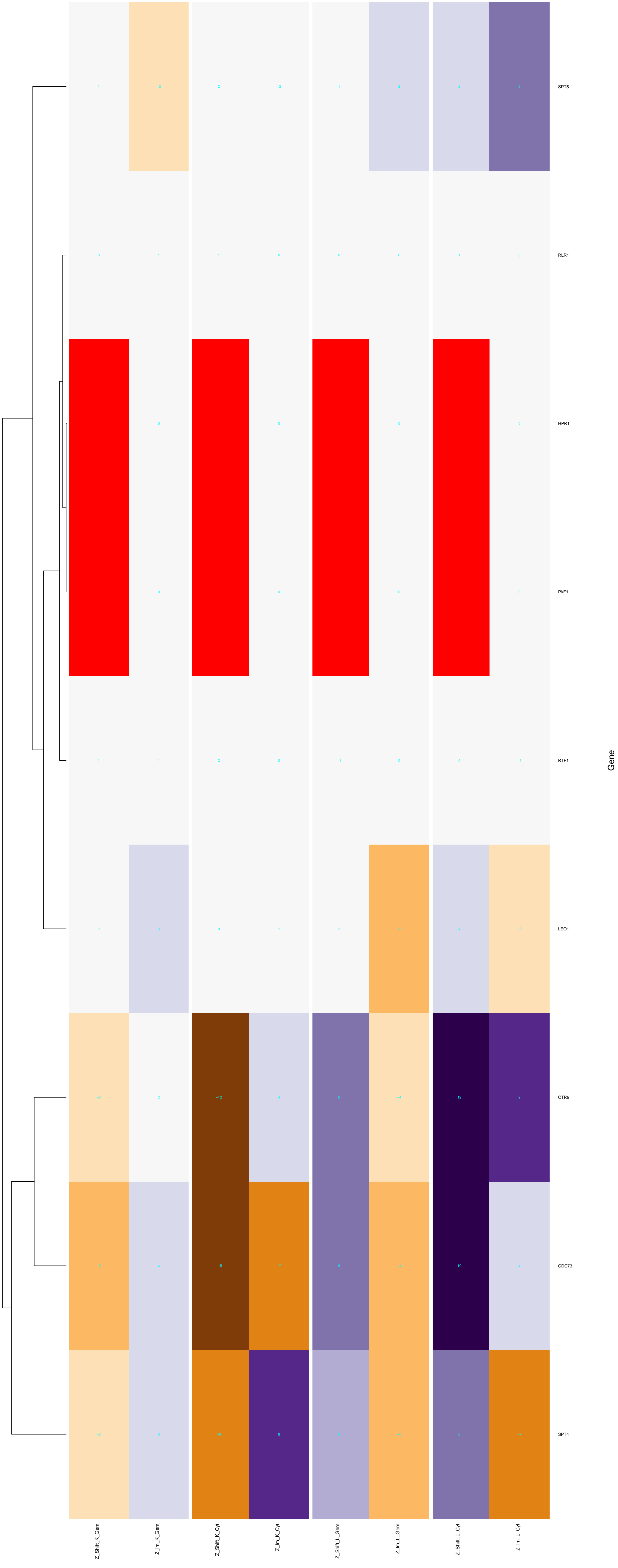
negative regulation of transcription from RNA polymerase I promoter

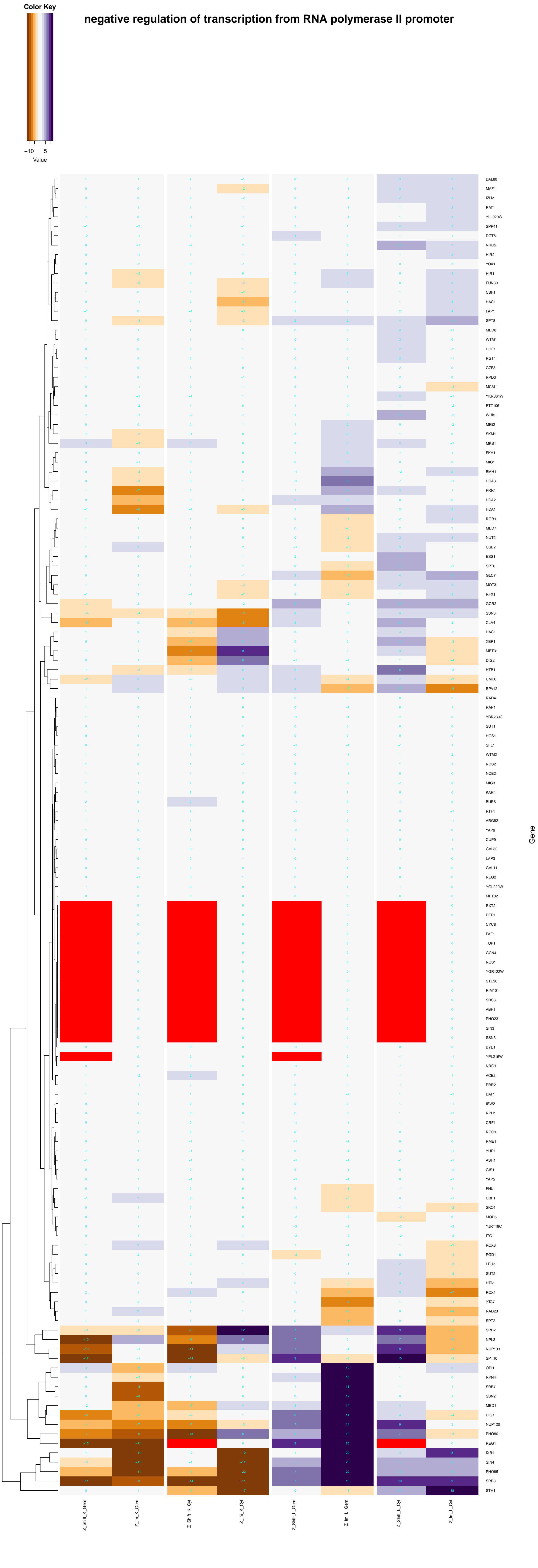


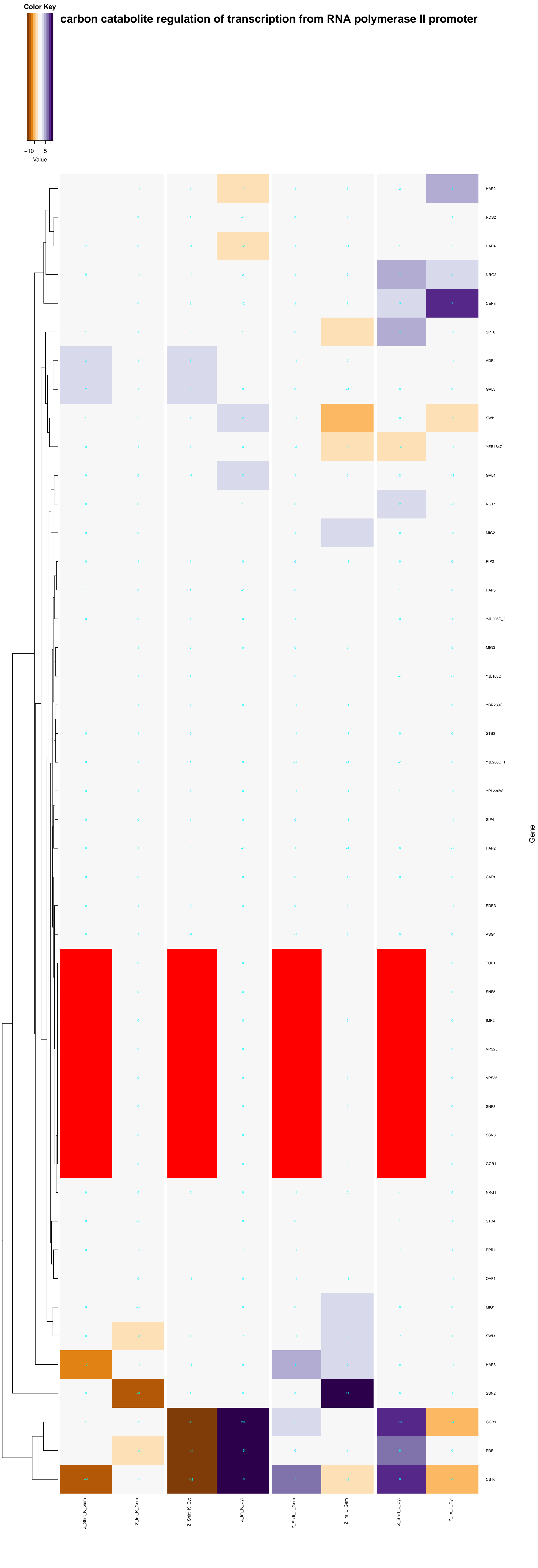




regulation of transcription elongation from RNA polymerase I promoter

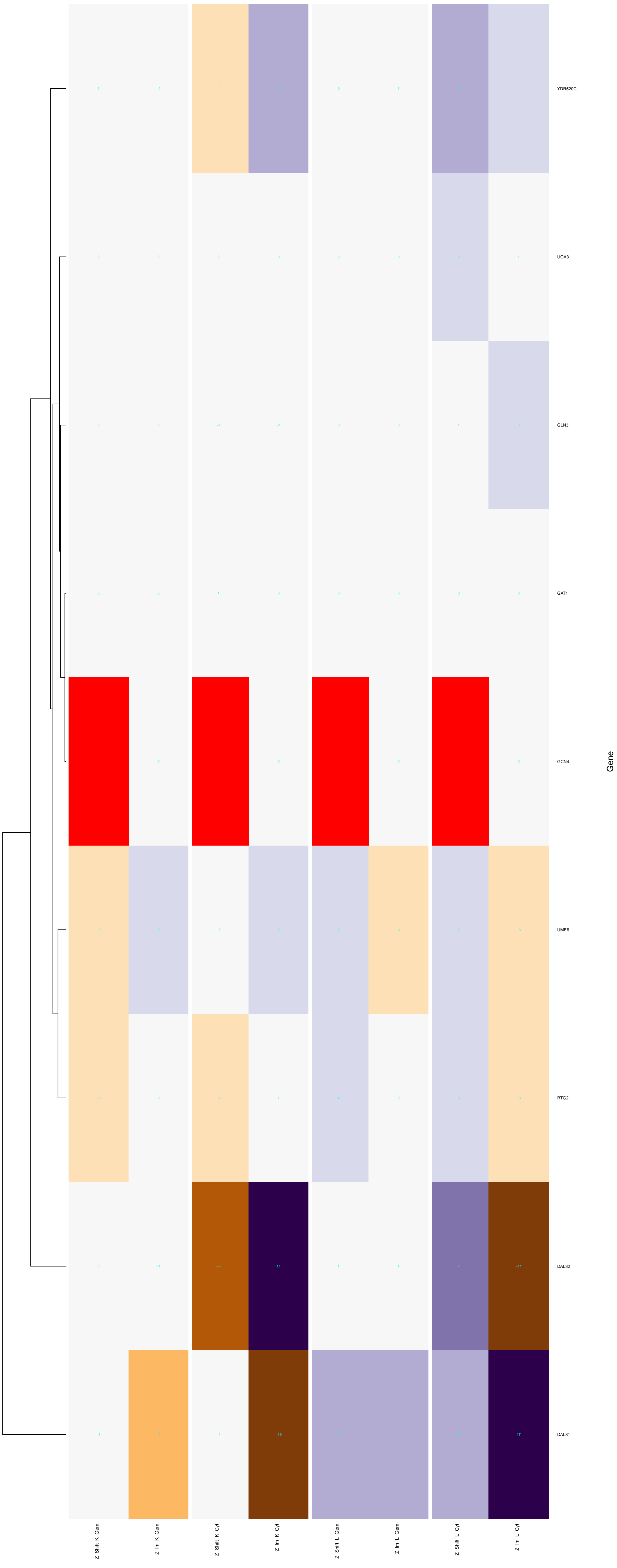








nitrogen catabolite regulation of transcription from RNA polymerase II promoter

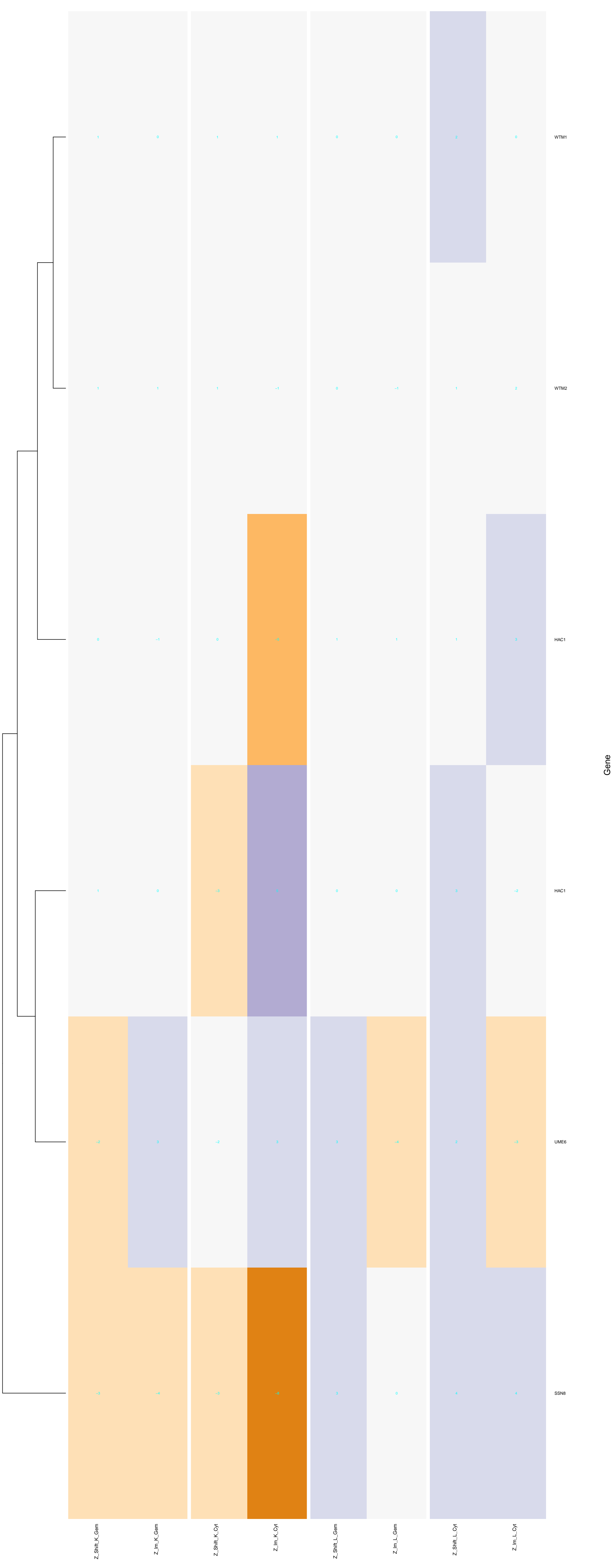


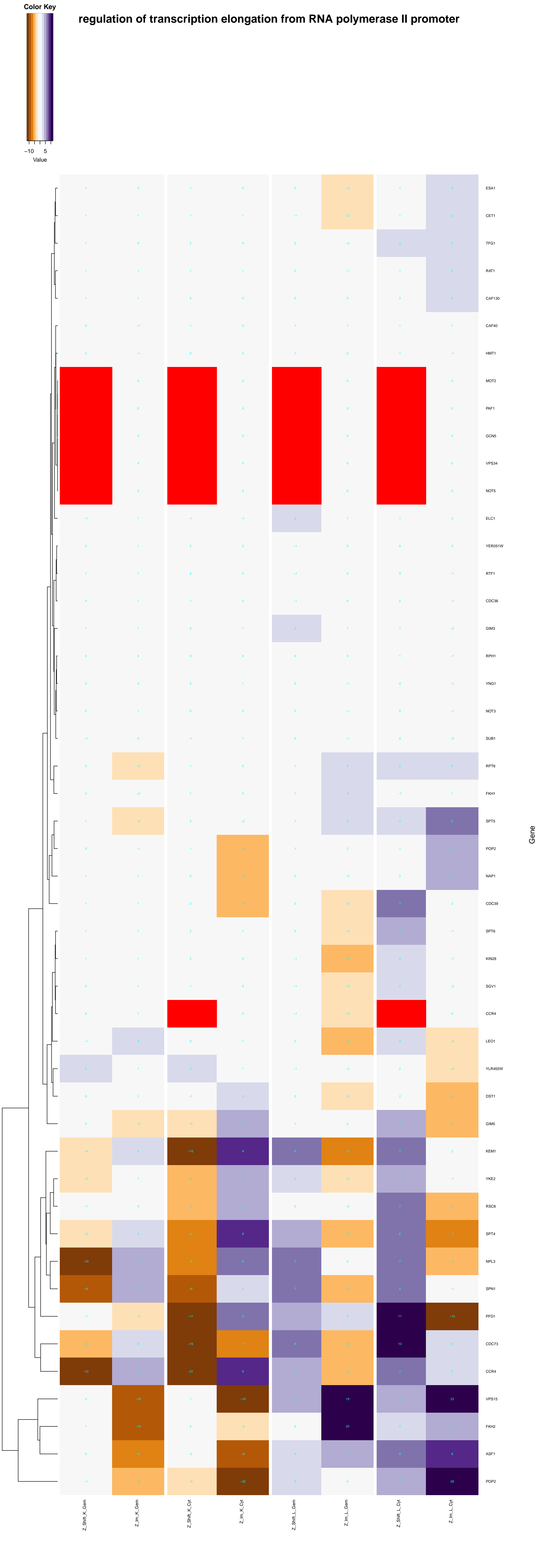
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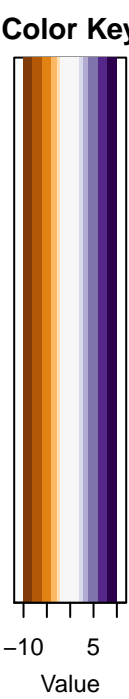
ation of transcription from RNA polymerase II promoter involved in meiotic cell cycle

-10 5

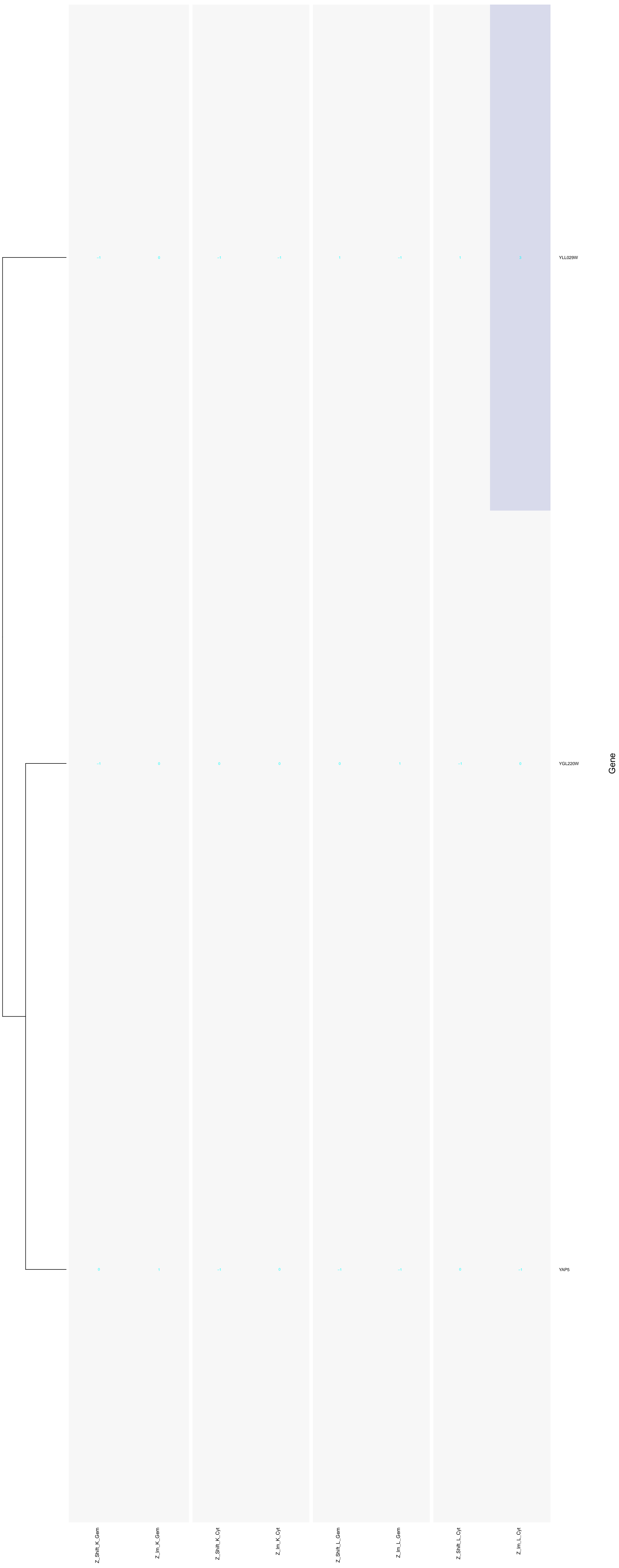
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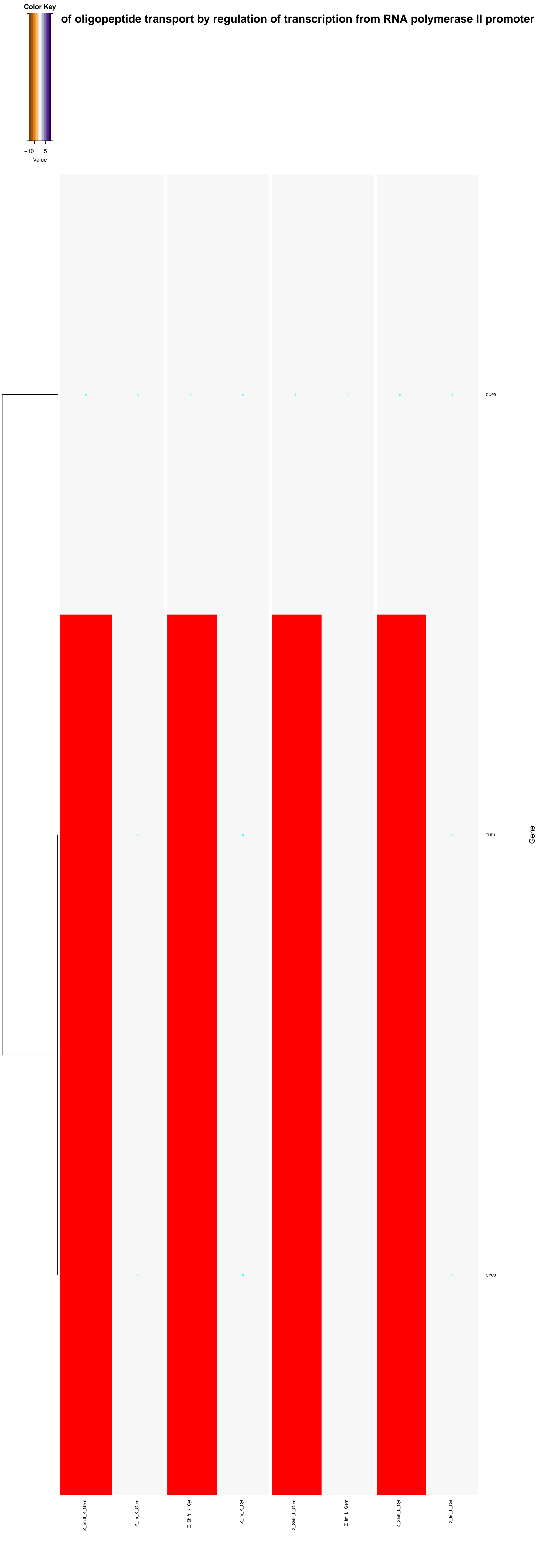


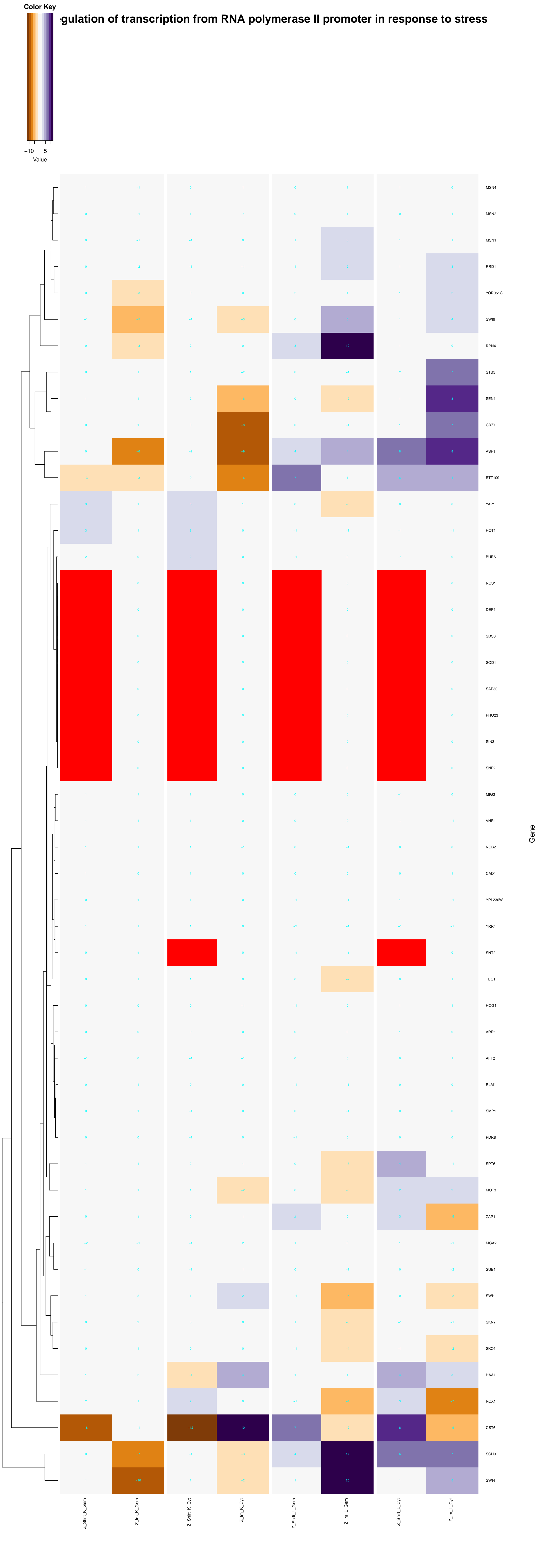




Regulation of transcription from RNA polymerase II promoter in response to iron



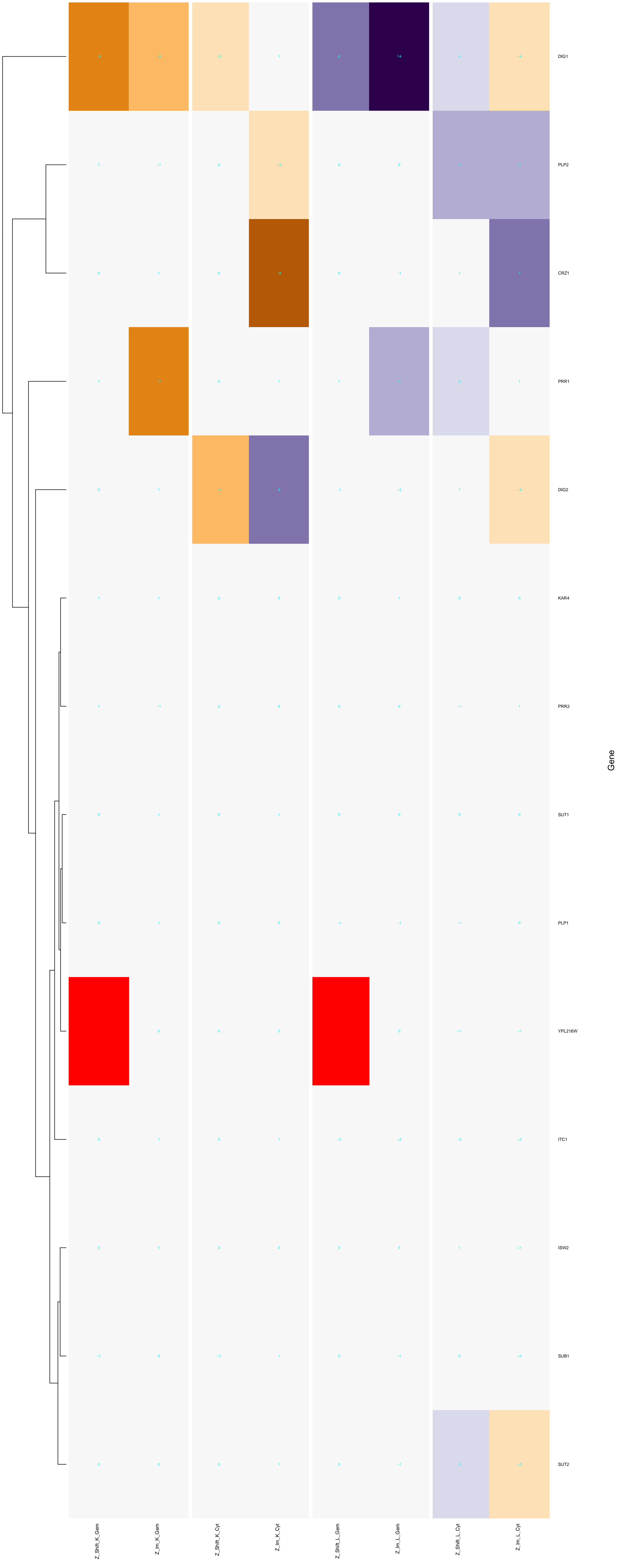








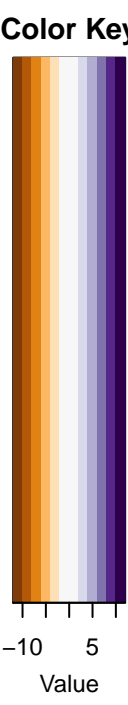
regulation of transcription from RNA polymerase II promoter by pheromones



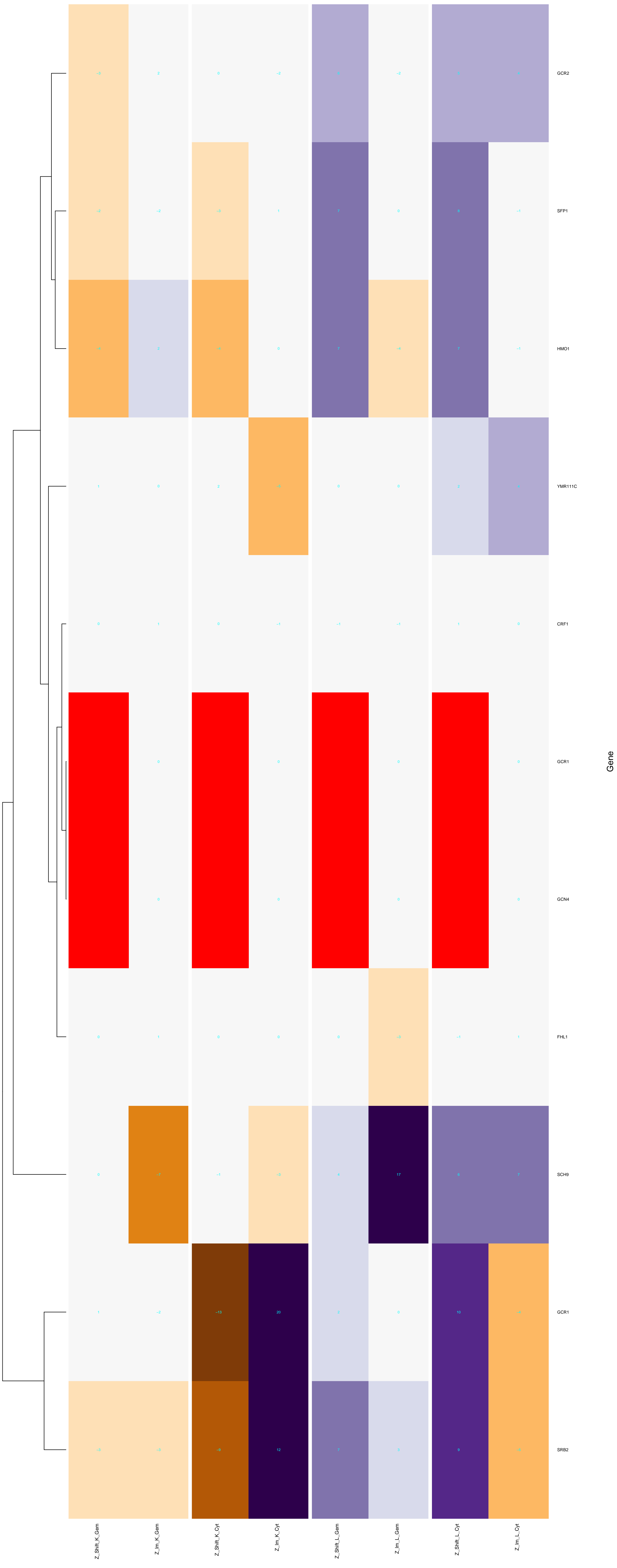


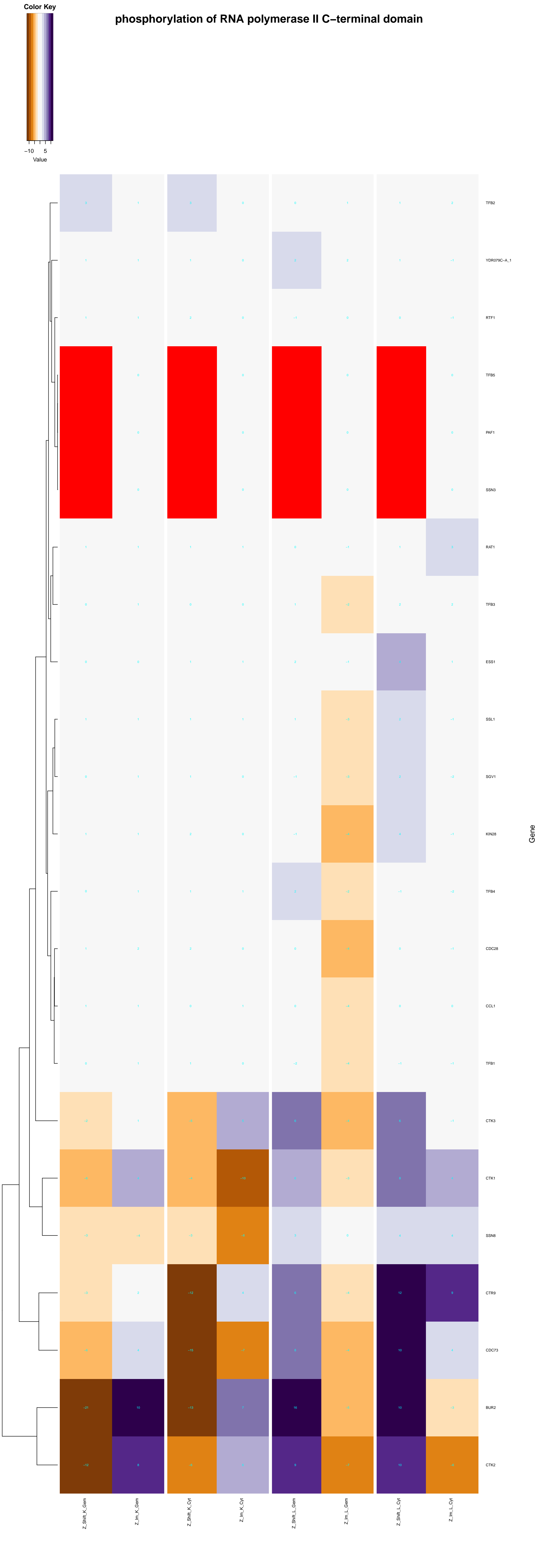
regulation of transcription initiation from RNA polymerase II promoter

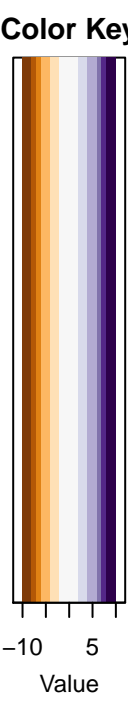




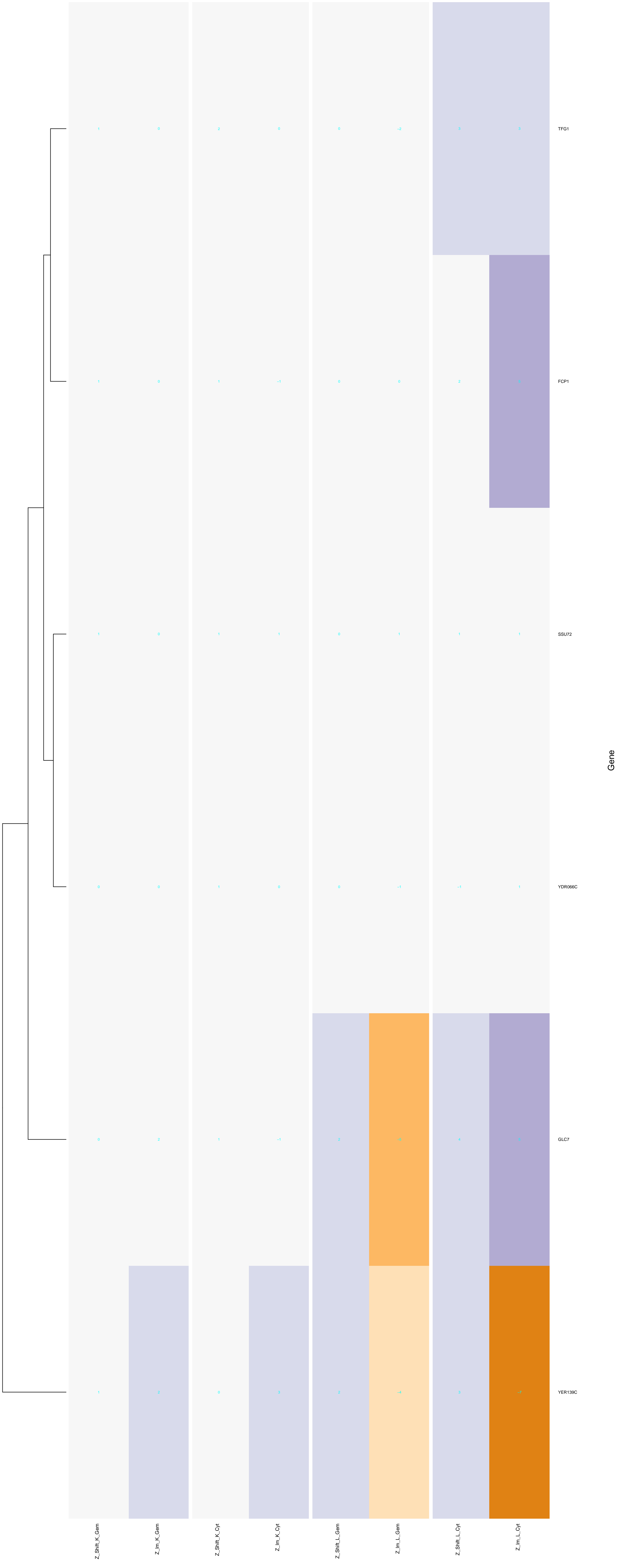
Regulation of ribosomal protein gene transcription from RNA polymerase II promoter

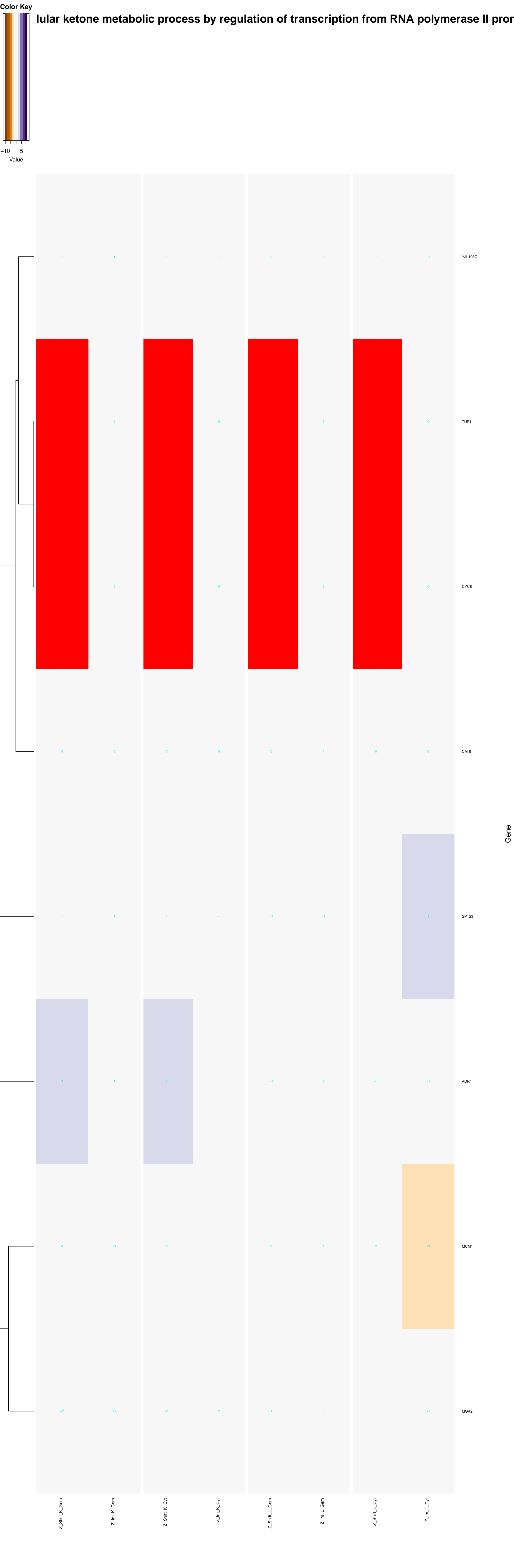






dephosphorylation of RNA polymerase II C-terminal domain



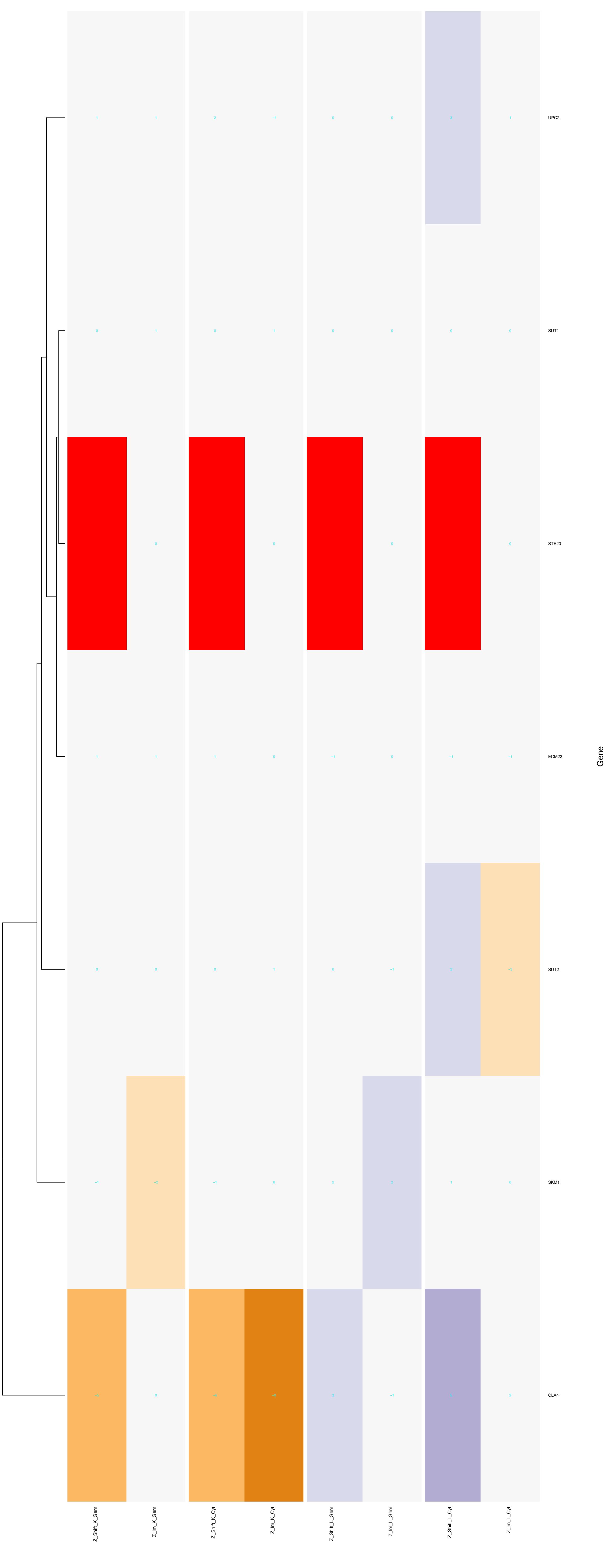


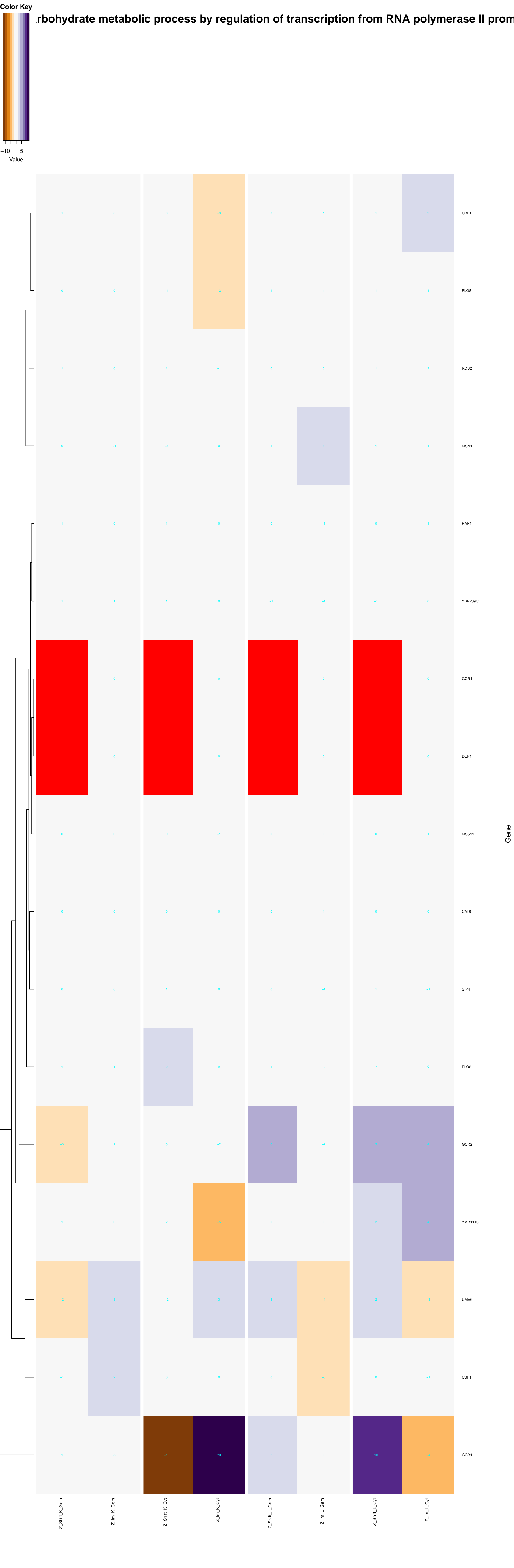
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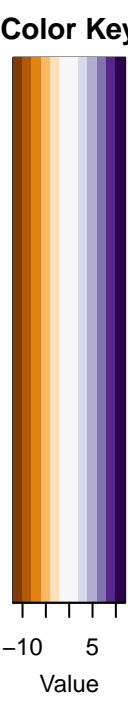
tion of lipid transport by regulation of transcription from RNA polymerase II promoter

-10 5

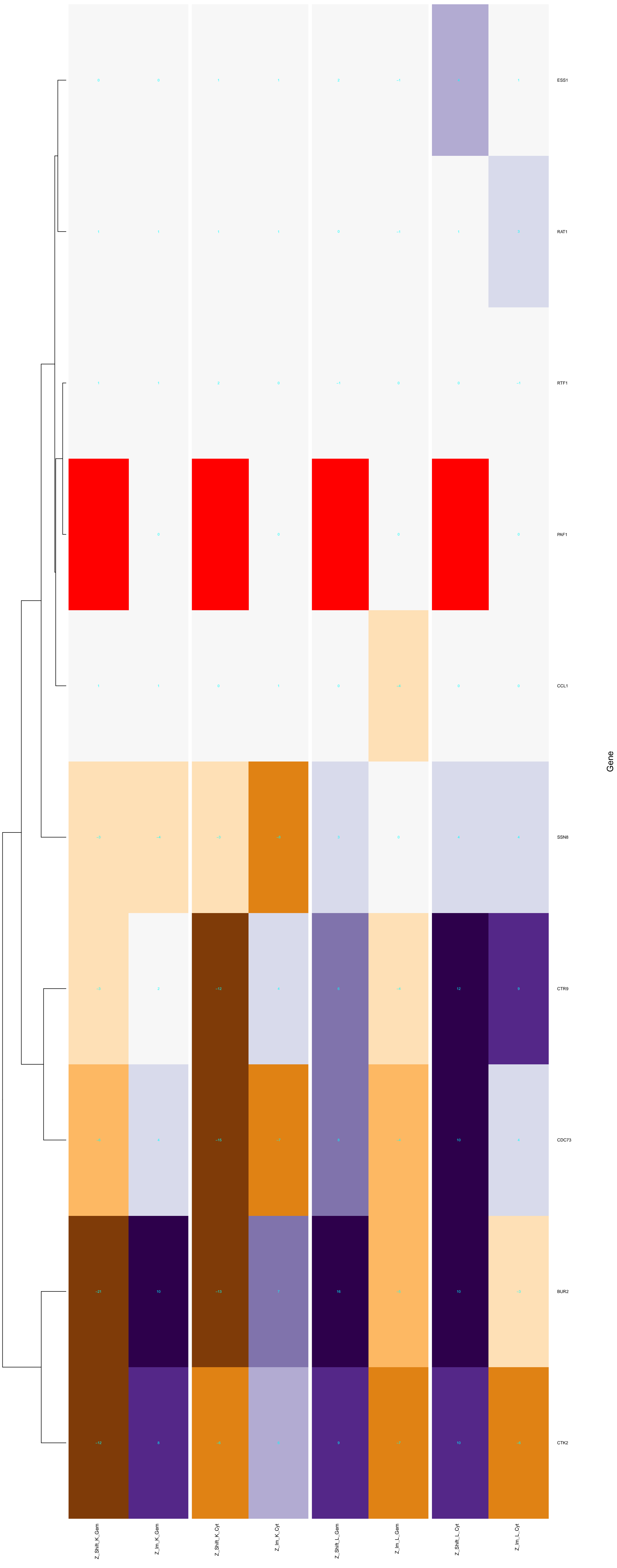
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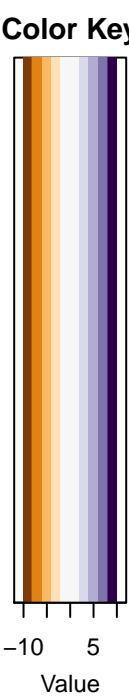




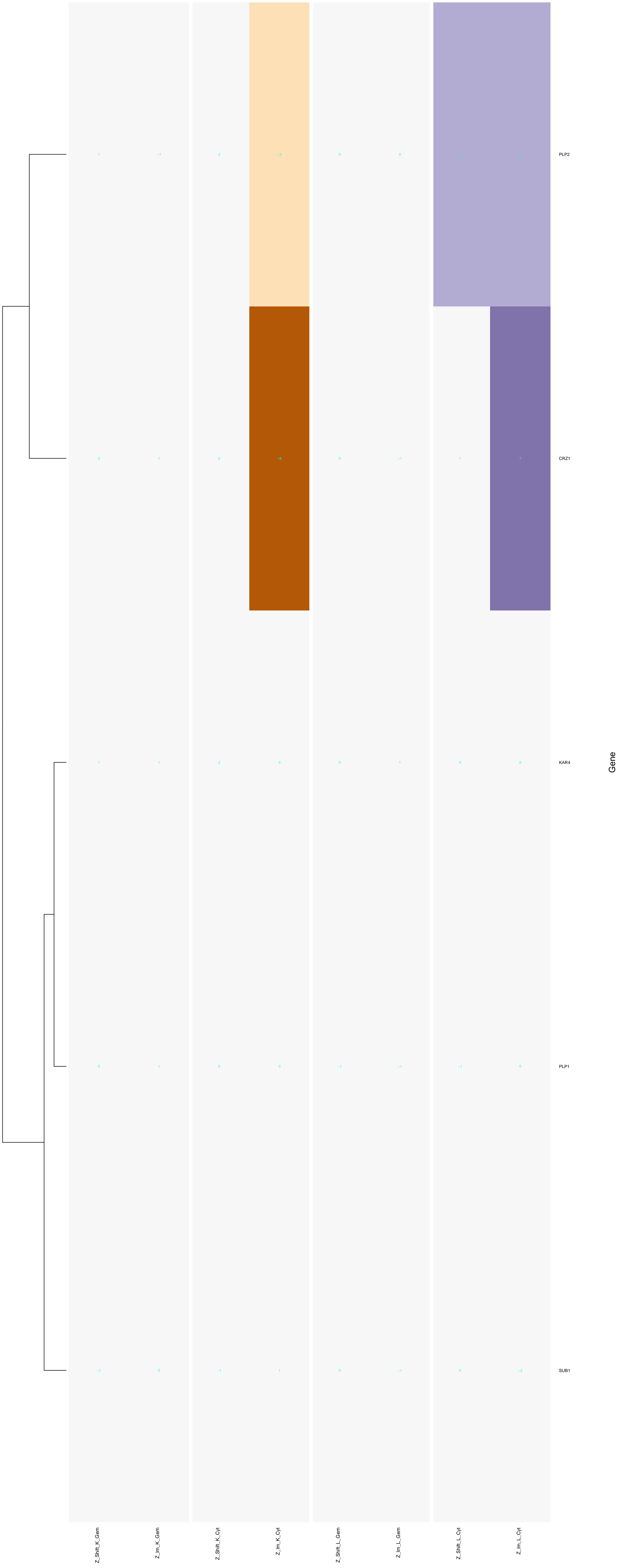


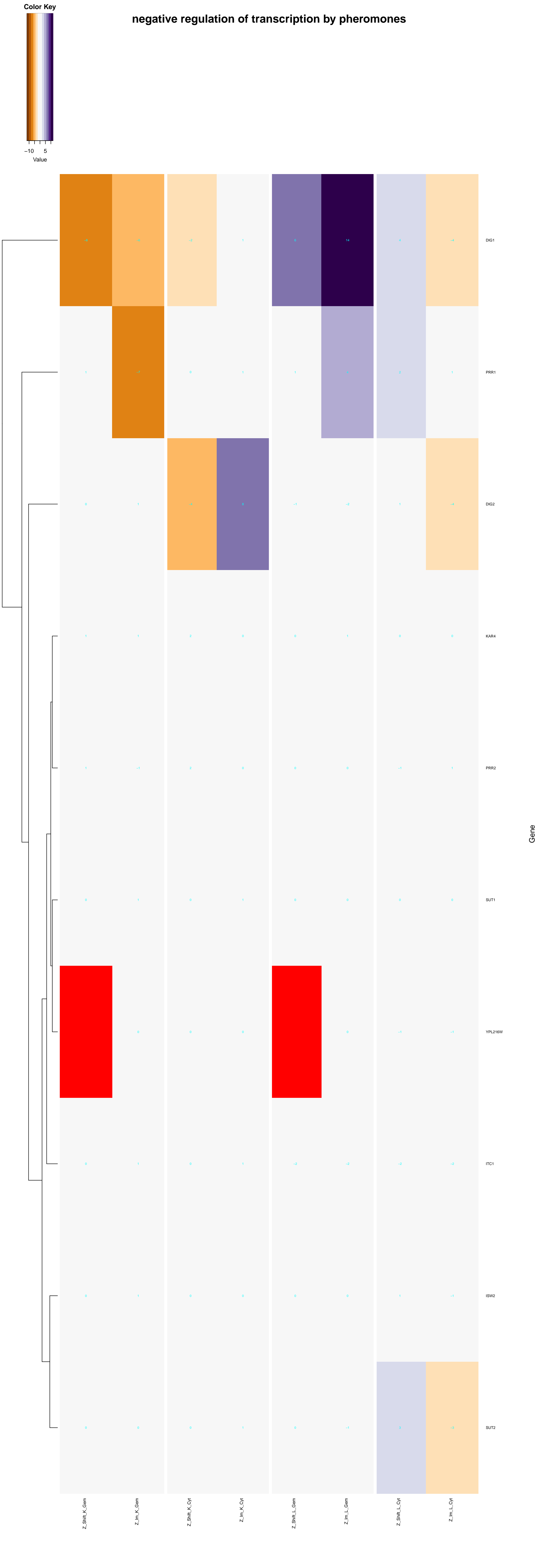
regulation of phosphorylation of RNA polymerase II C-terminal domain

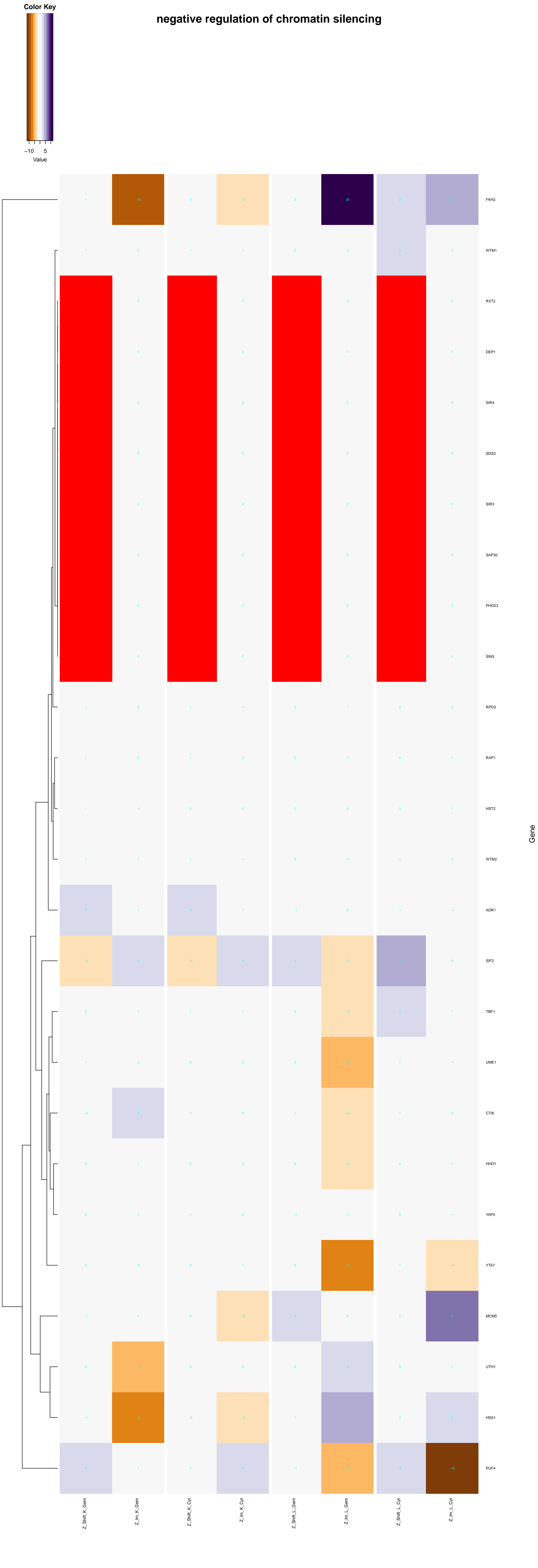


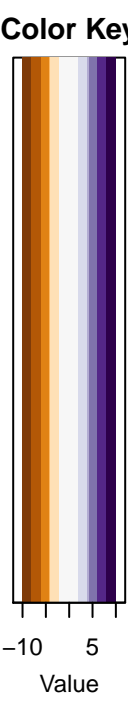


positive regulation of transcription by pheromones

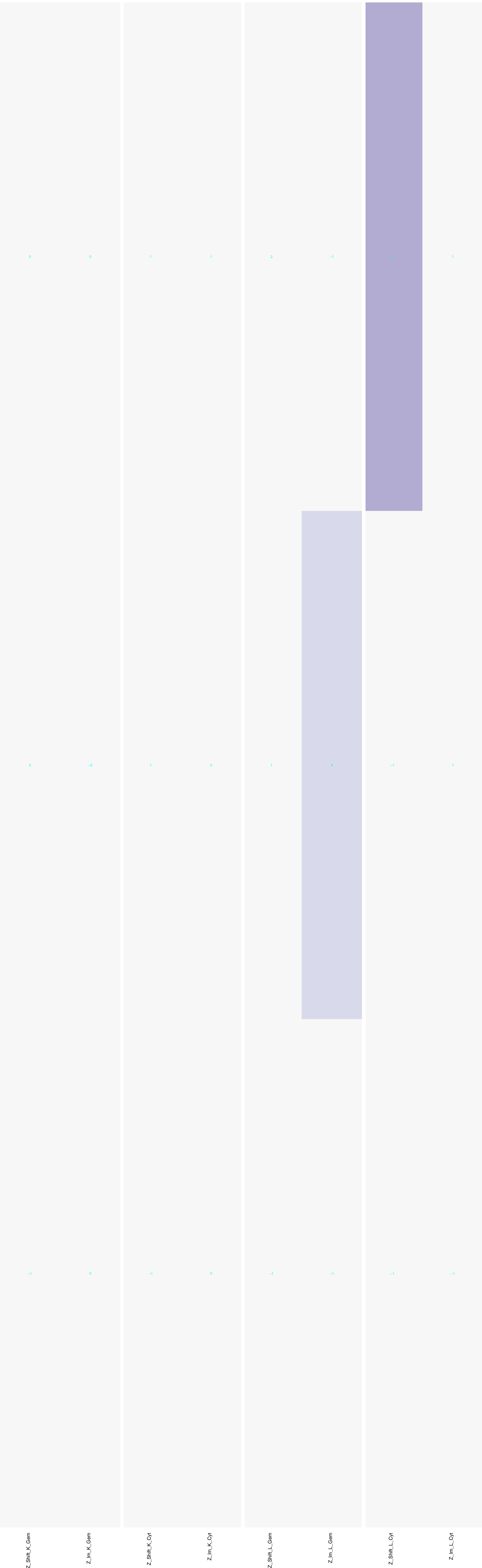




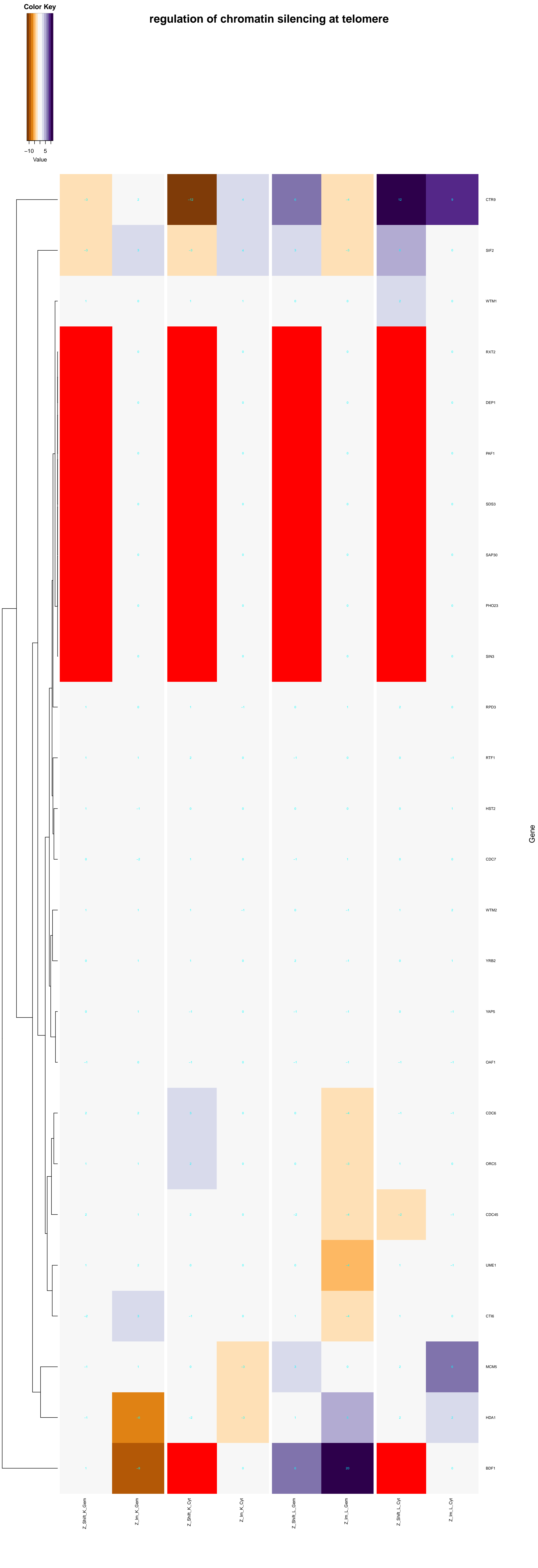


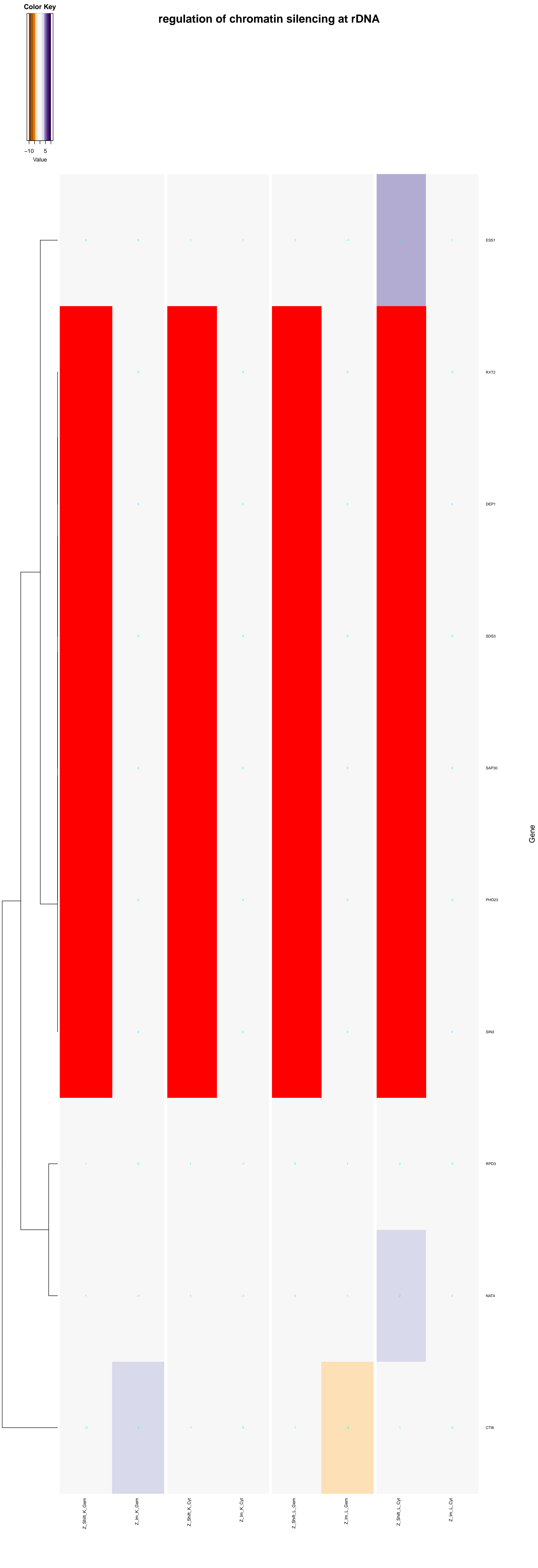


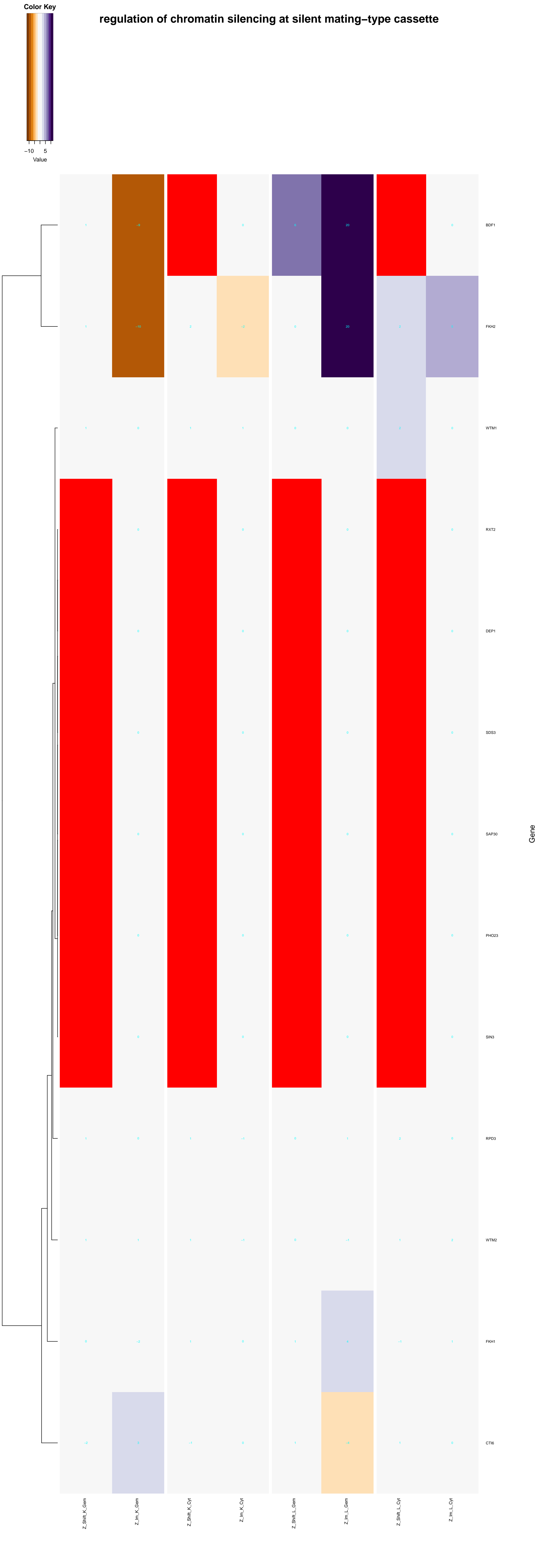
positive regulation of chromatin silencing

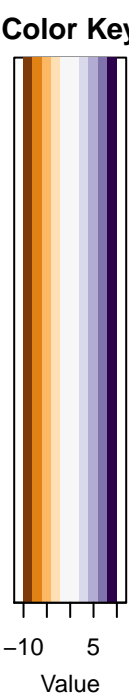


Gene

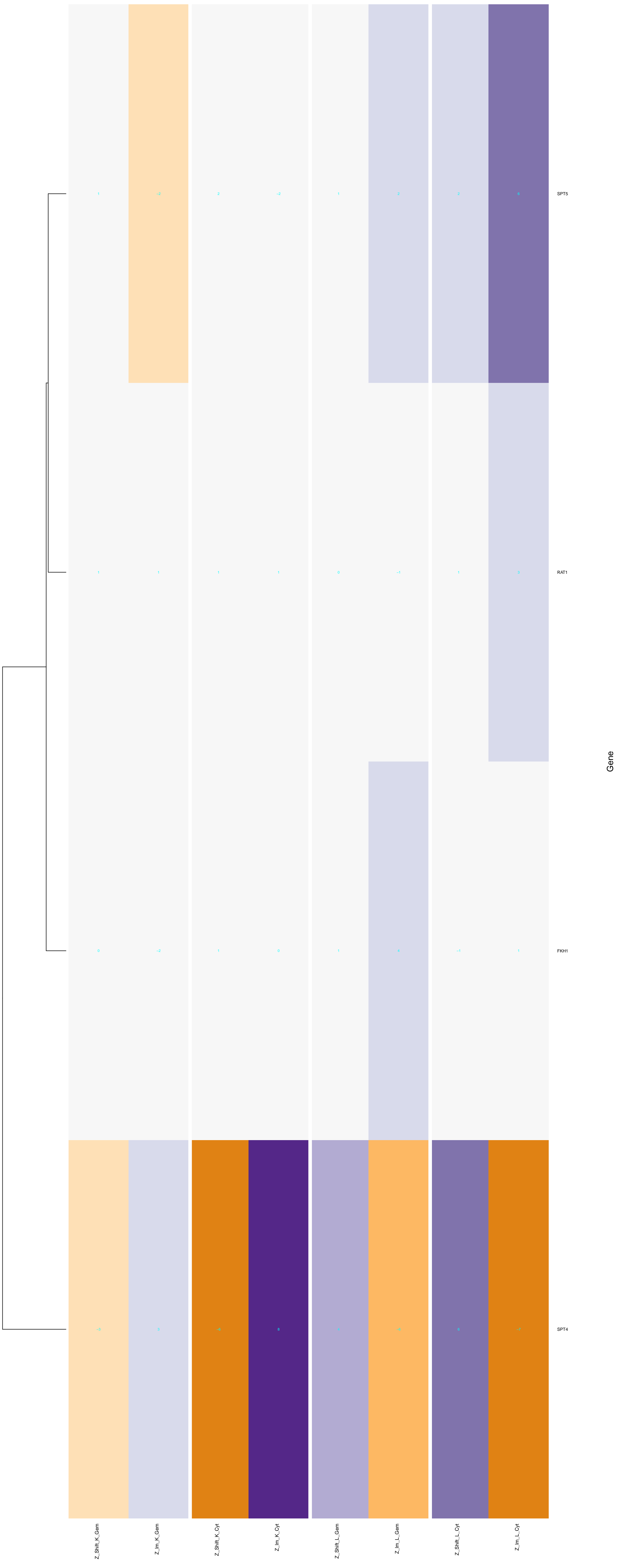


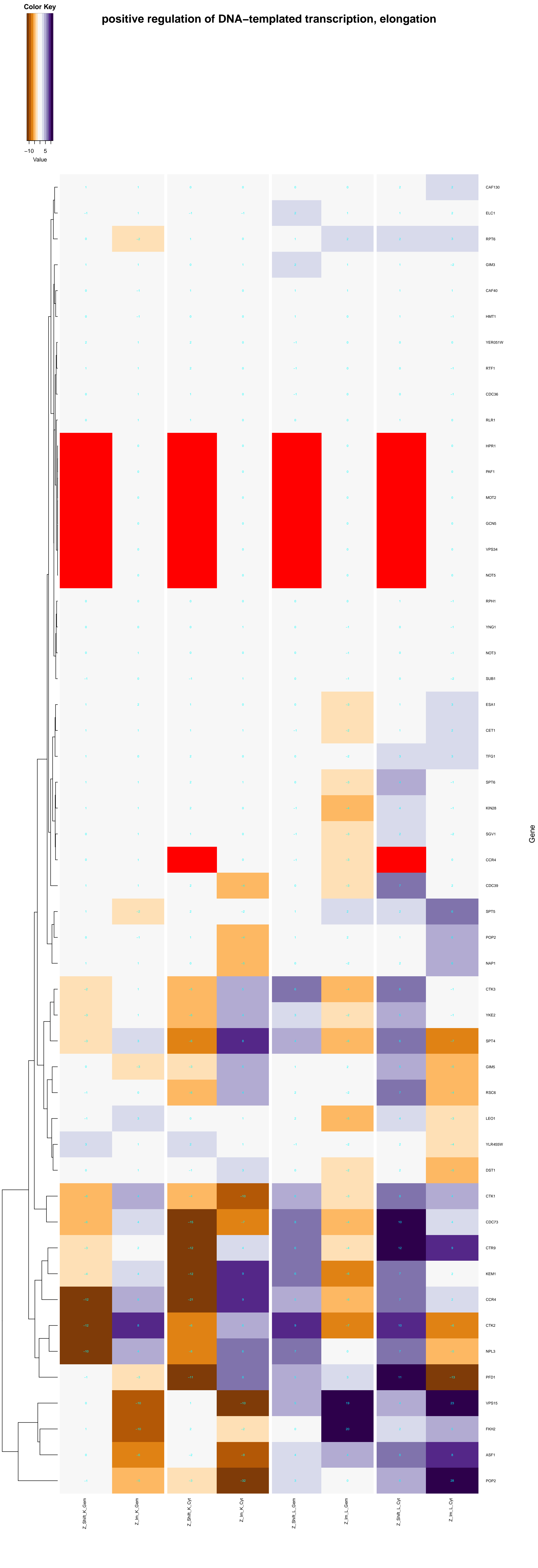


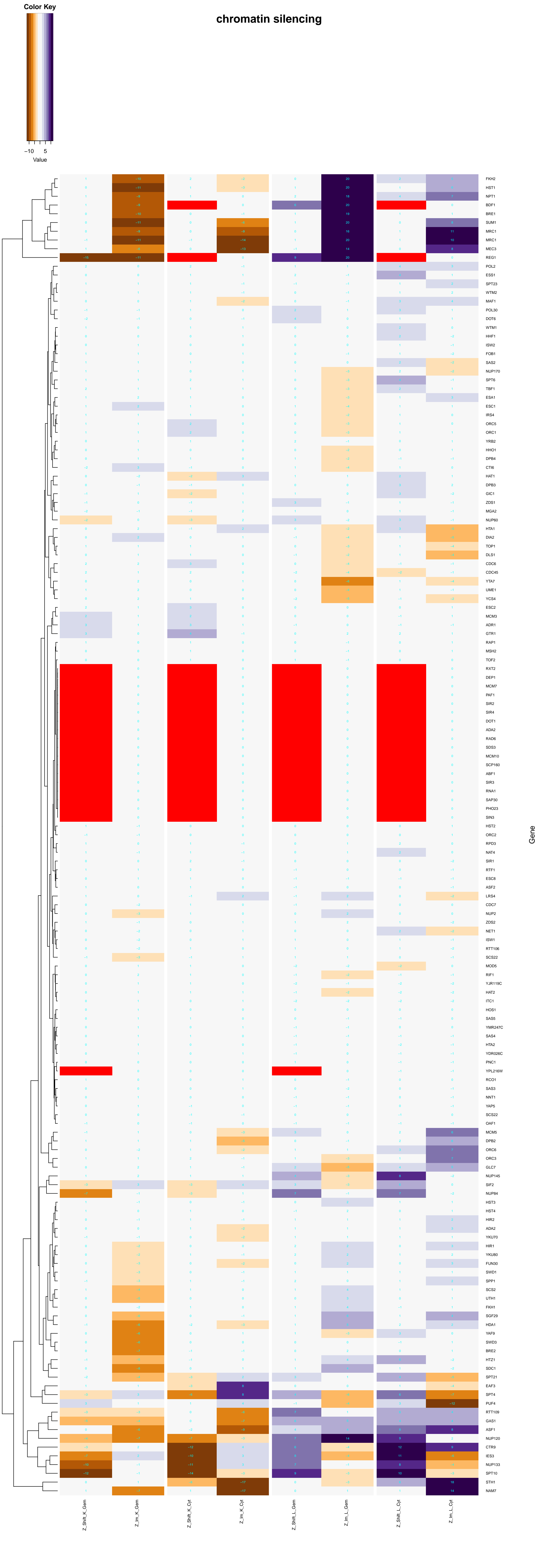


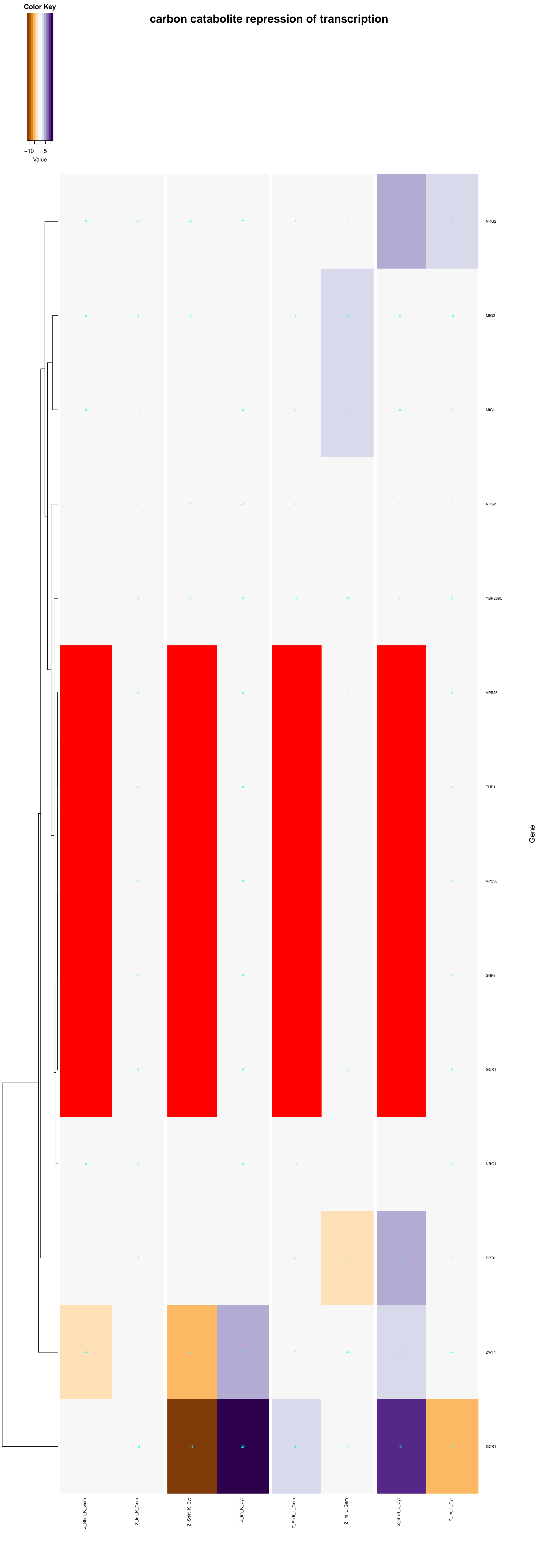


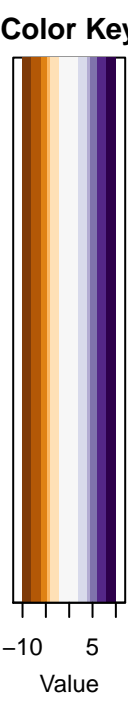
negative regulation of DNA-templated transcription, elongation



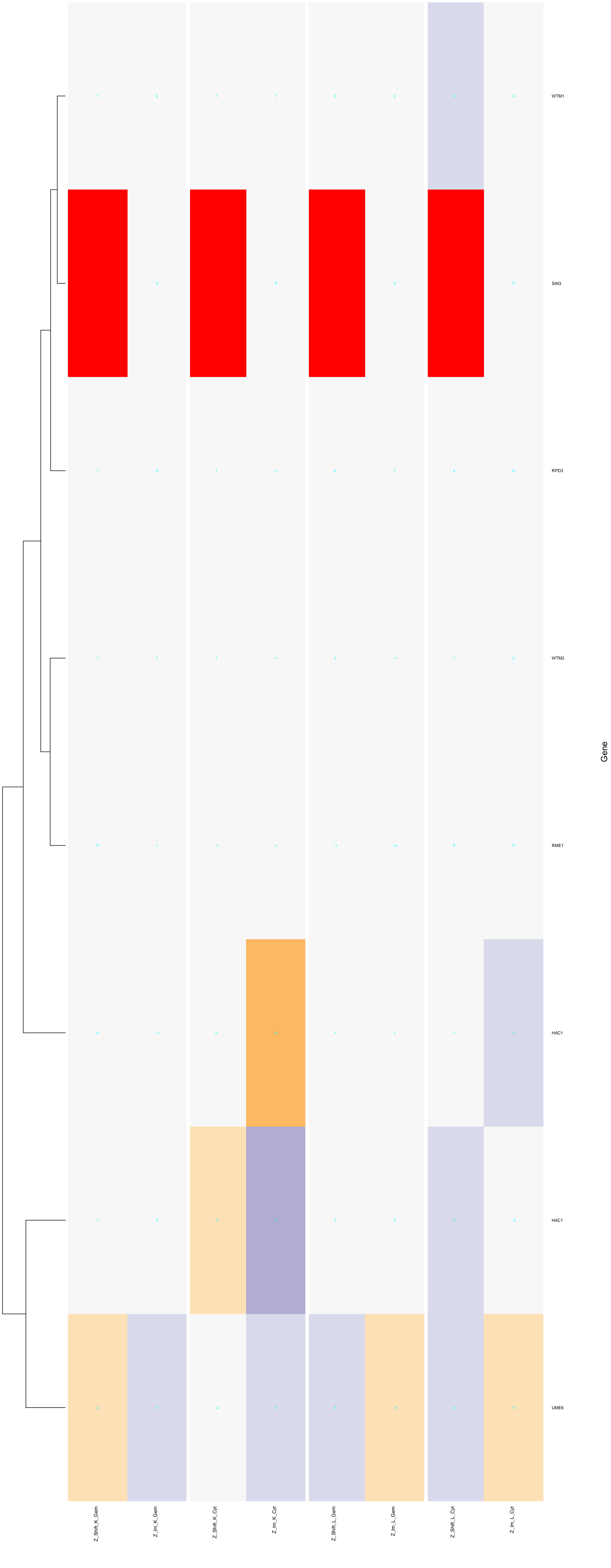


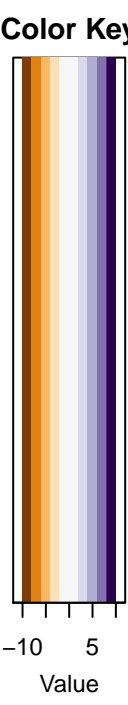




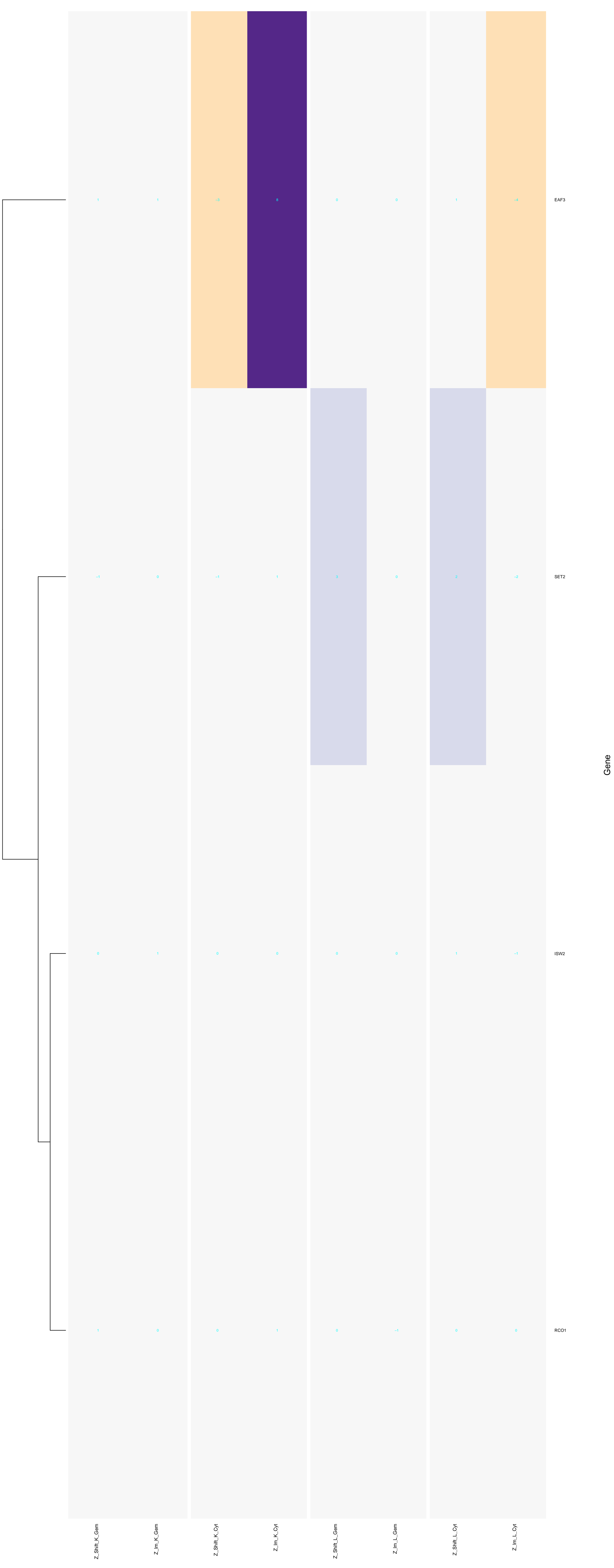


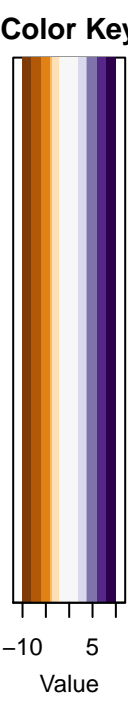
negative regulation of transcription involved in meiotic cell cycle



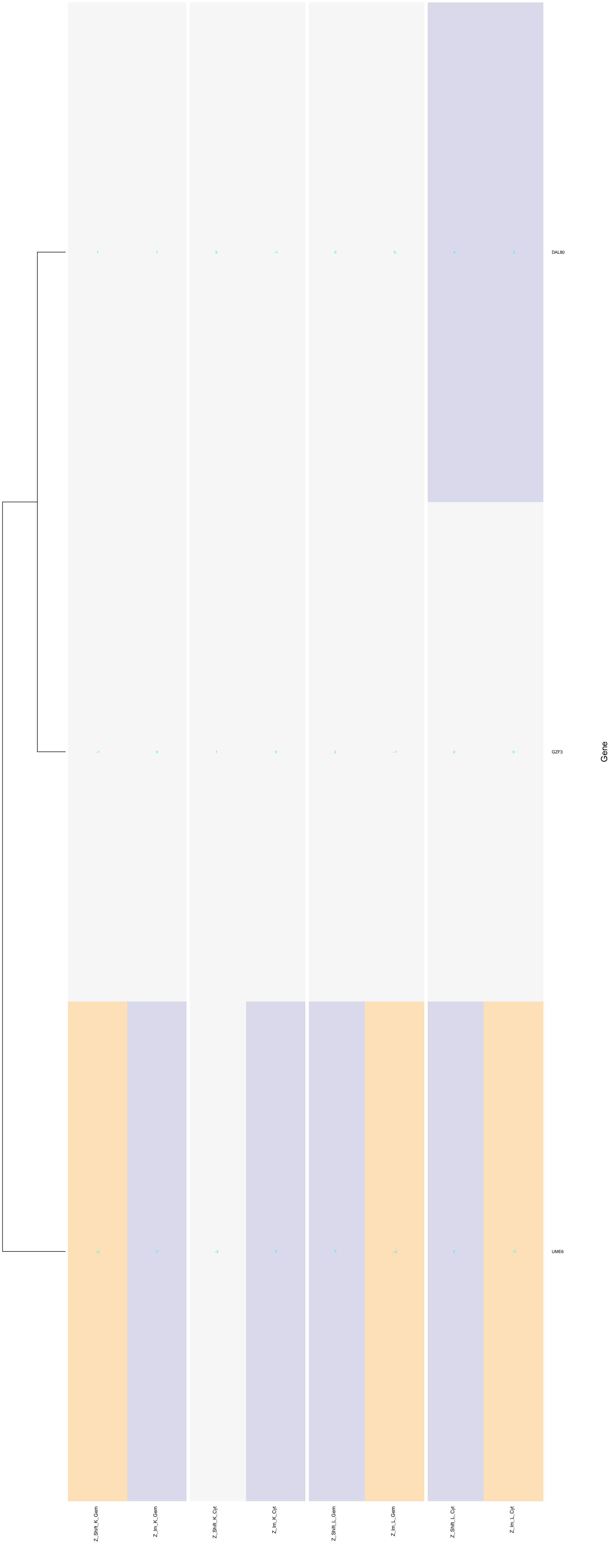


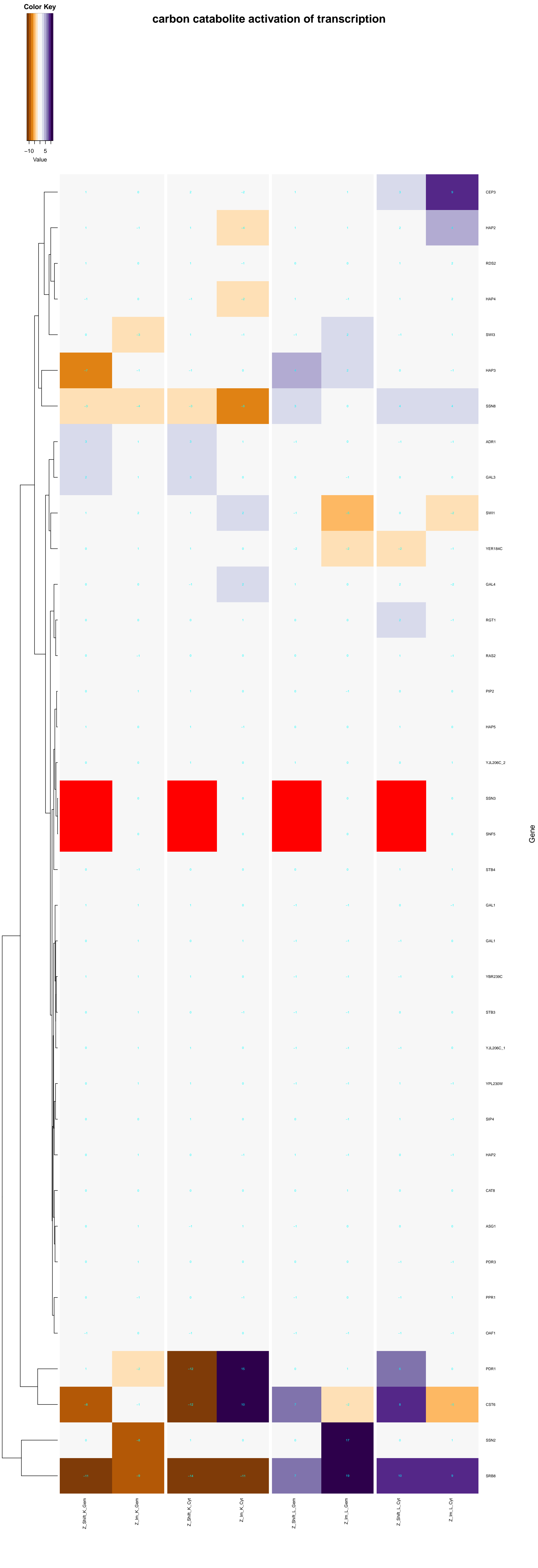
negative regulation of antisense RNA transcription

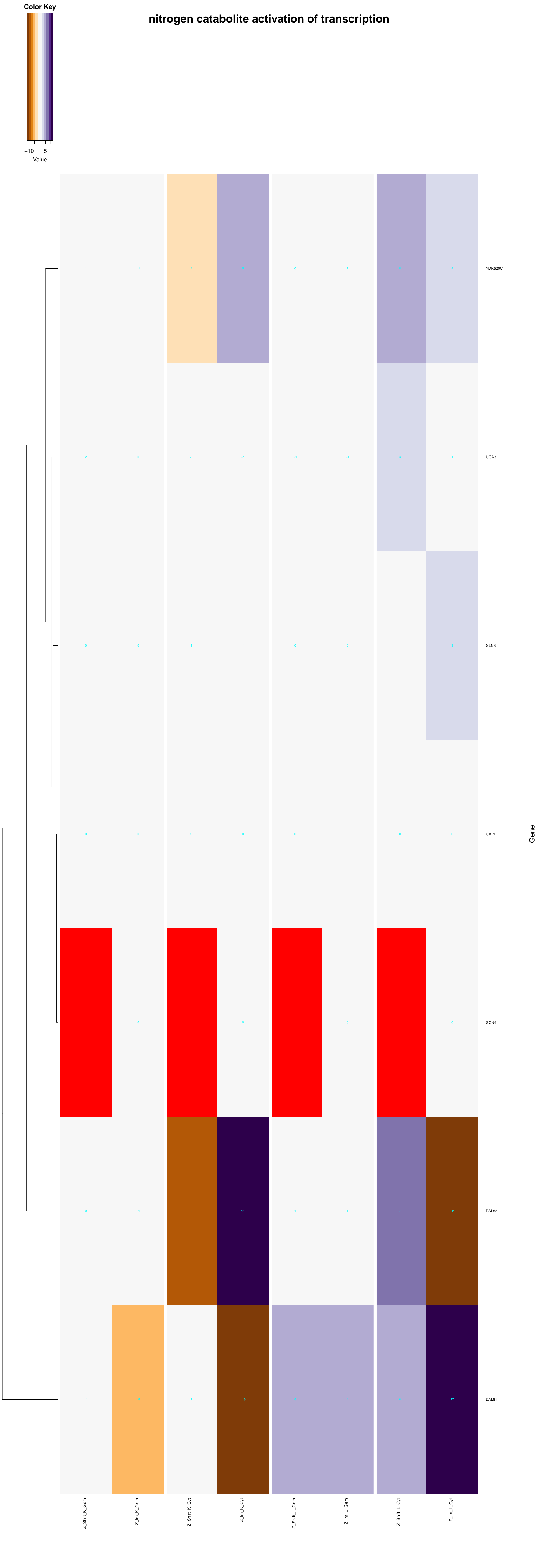


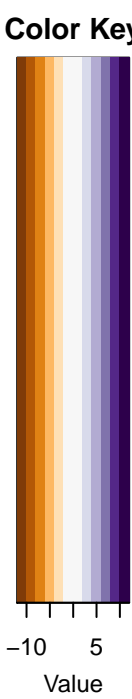


nitrogen catabolite repression of transcription

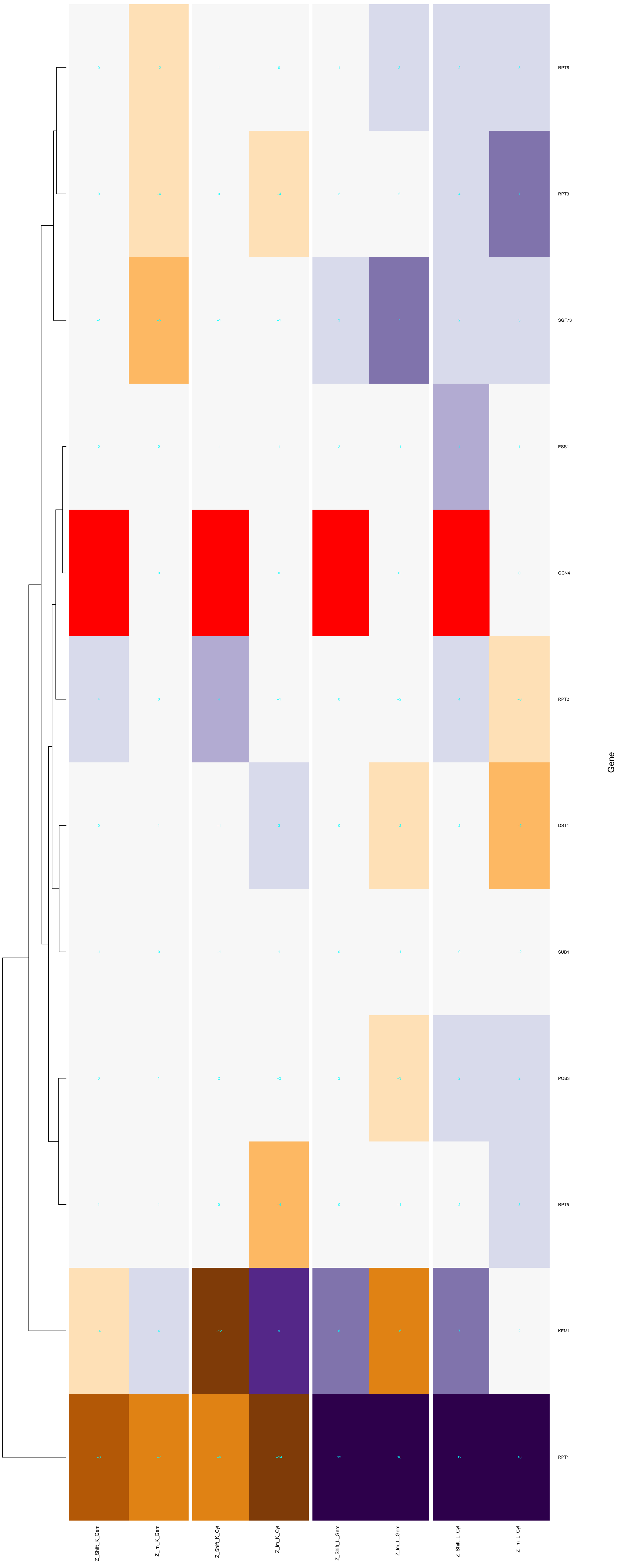


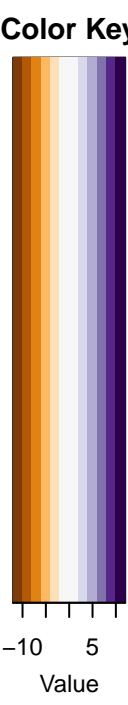




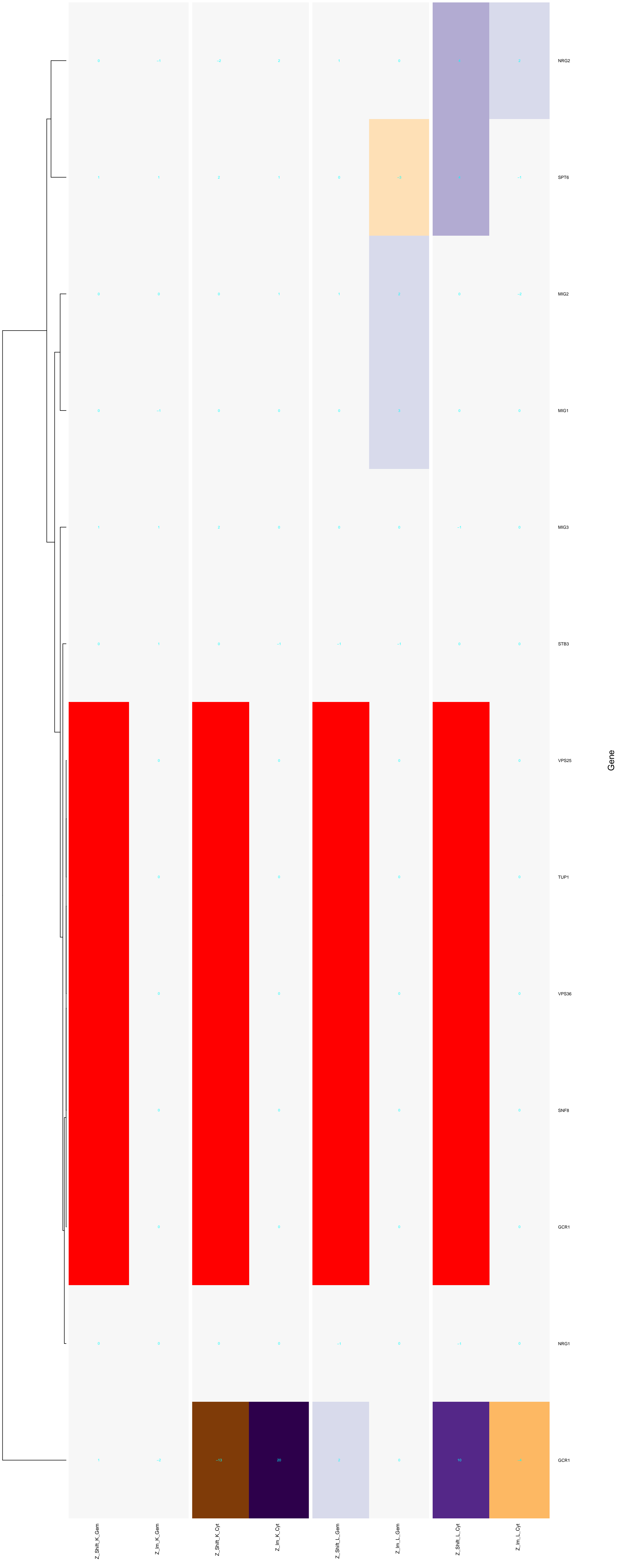


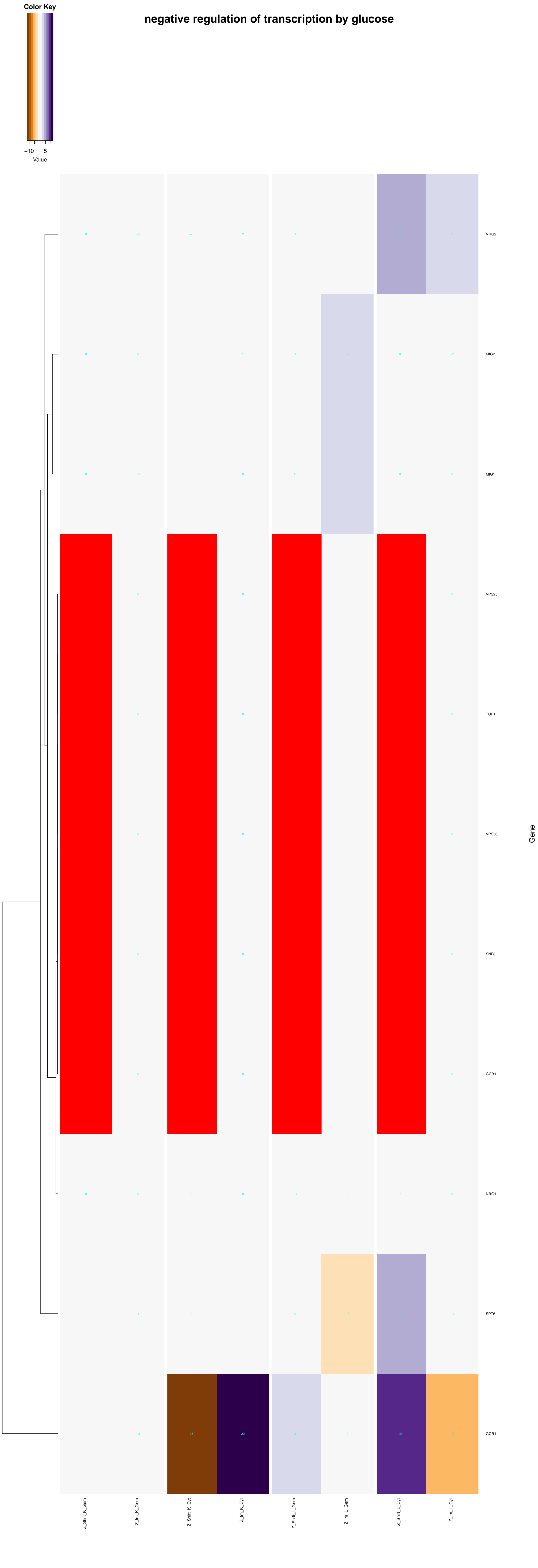
positive regulation of DNA-templated transcription, initiation

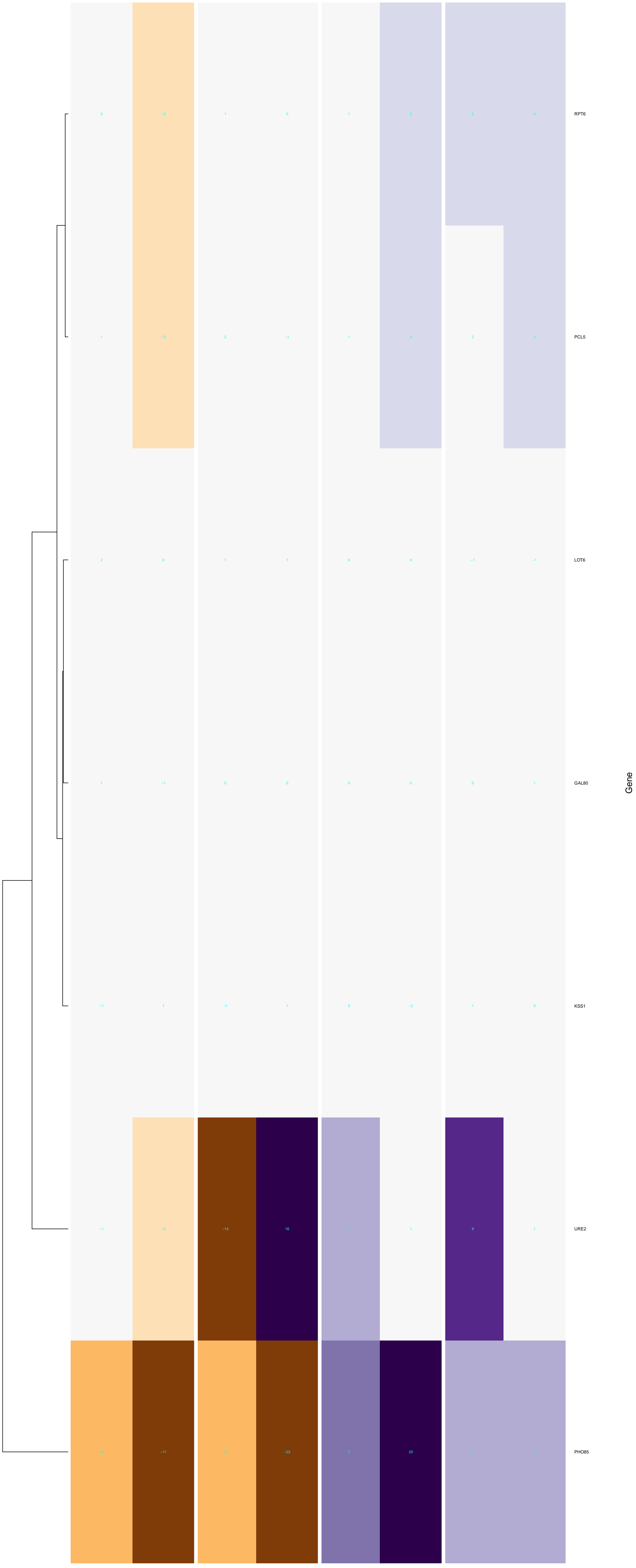
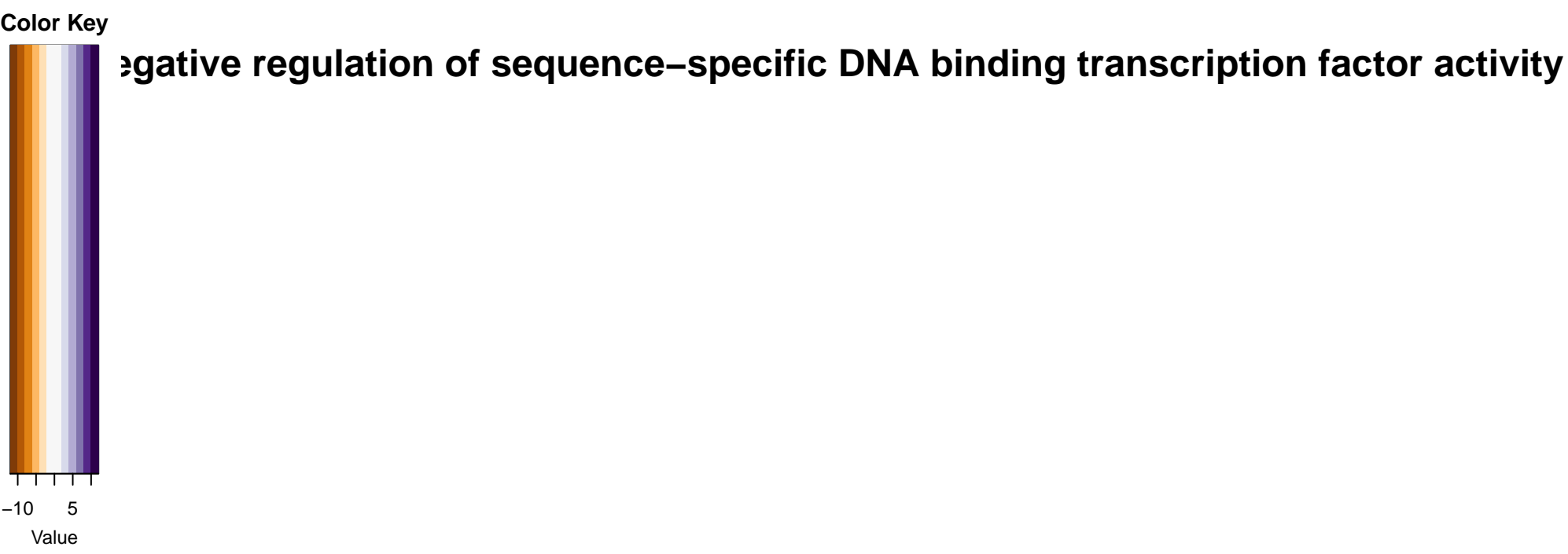


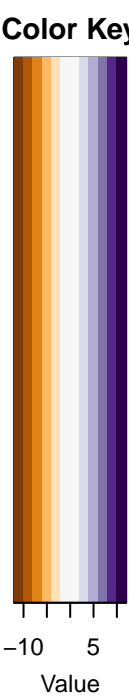


regulation of transcription from RNA polymerase II promoter by glucose

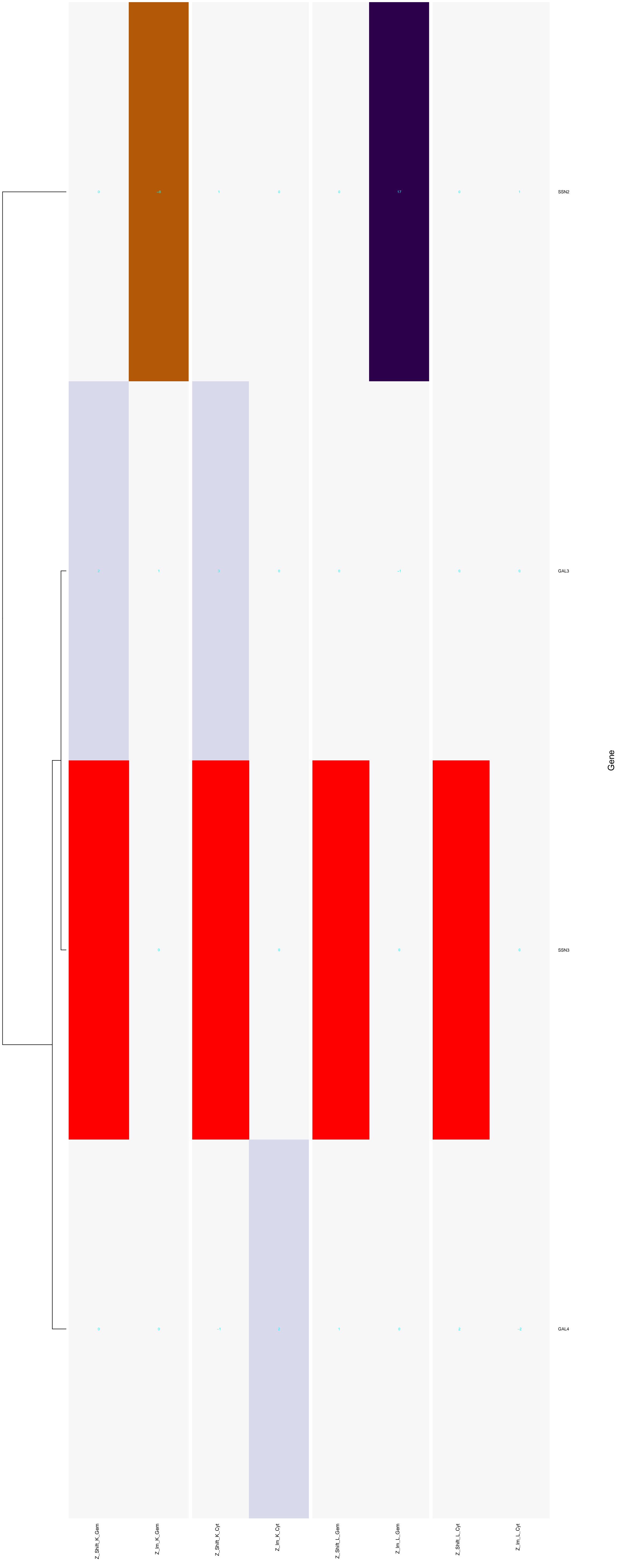


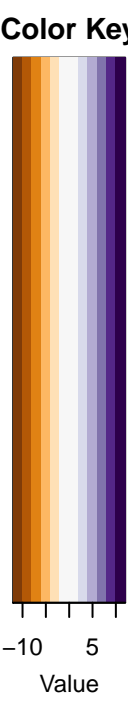




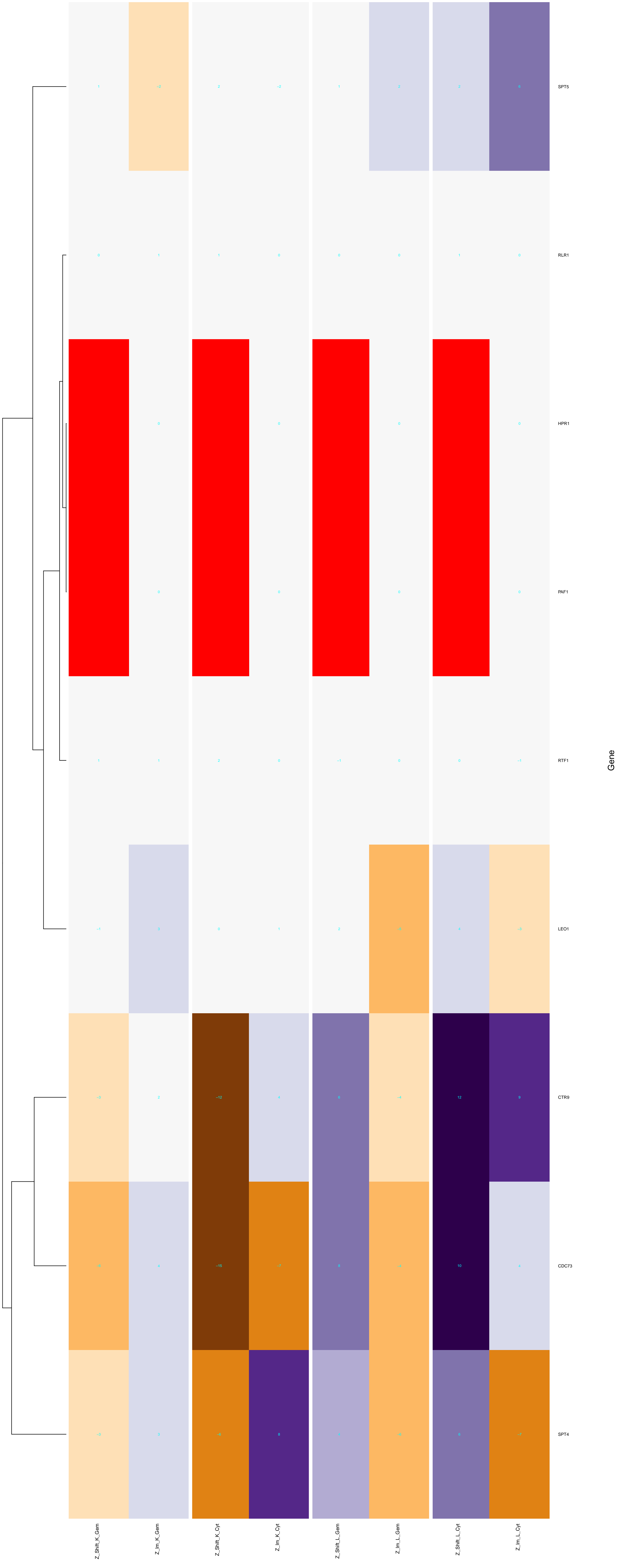


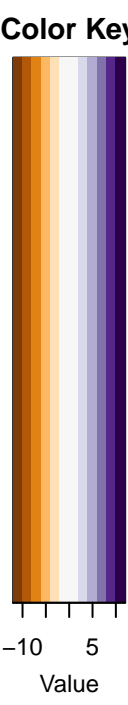
Positive regulation of transcription from RNA polymerase II promoter by galactose



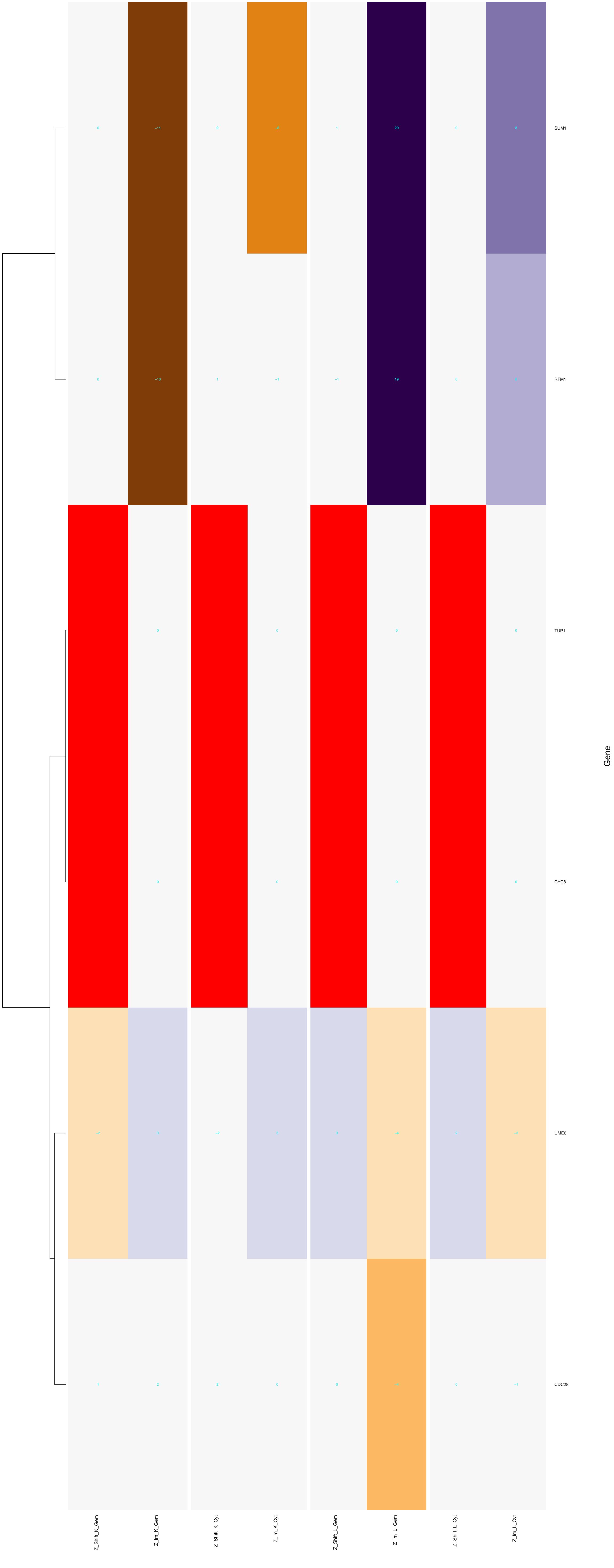


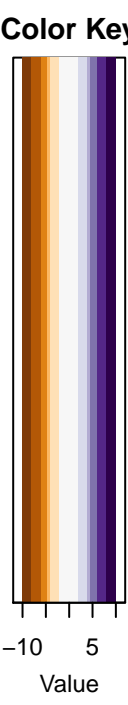
positive regulation of transcription elongation from RNA polymerase I promoter



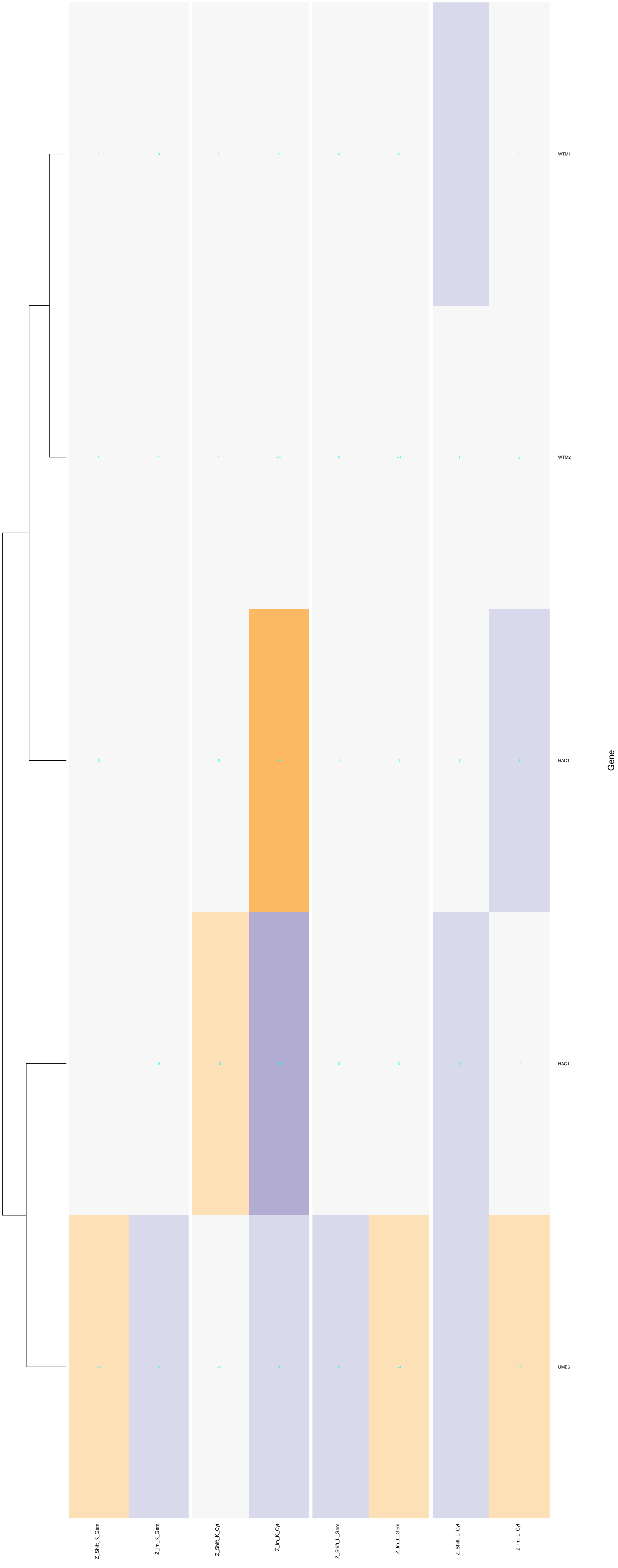


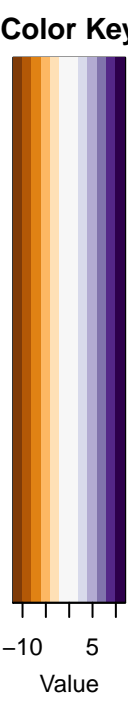
Regulation of transcription from RNA polymerase II promoter during mitotic cell cycle



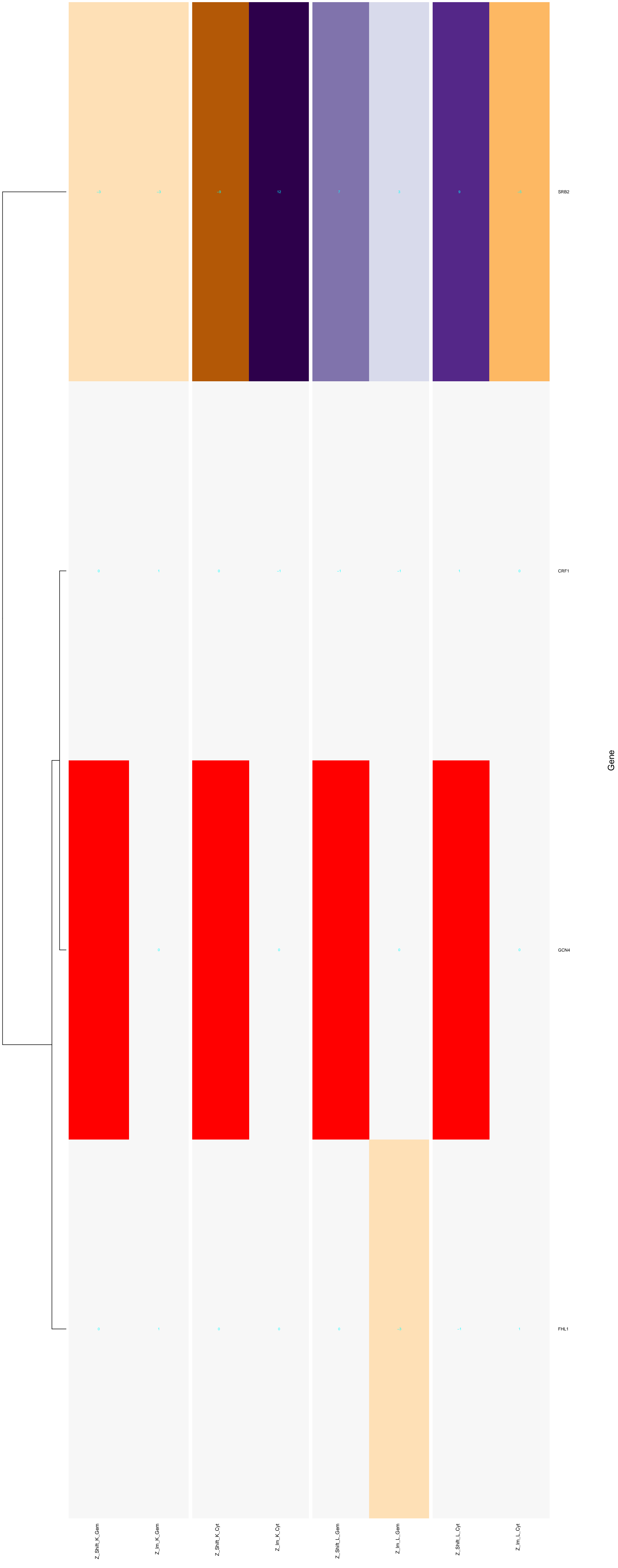


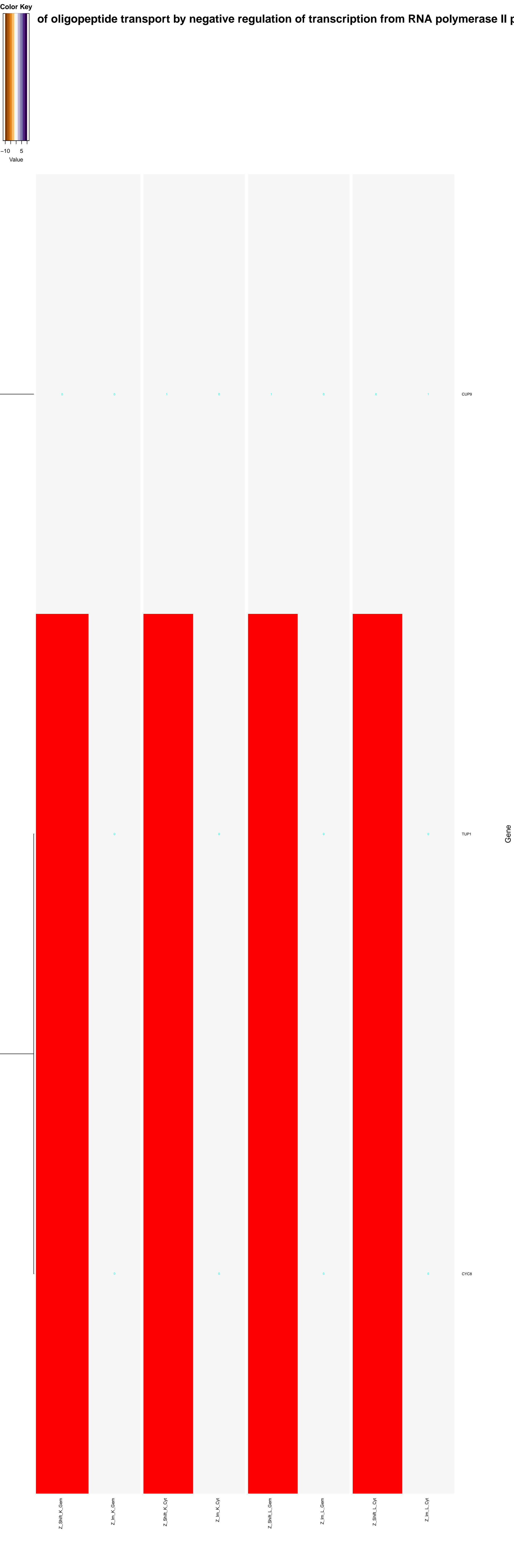
egulation of transcription from RNA polymerase II promoter involved in meiotic cell cycle

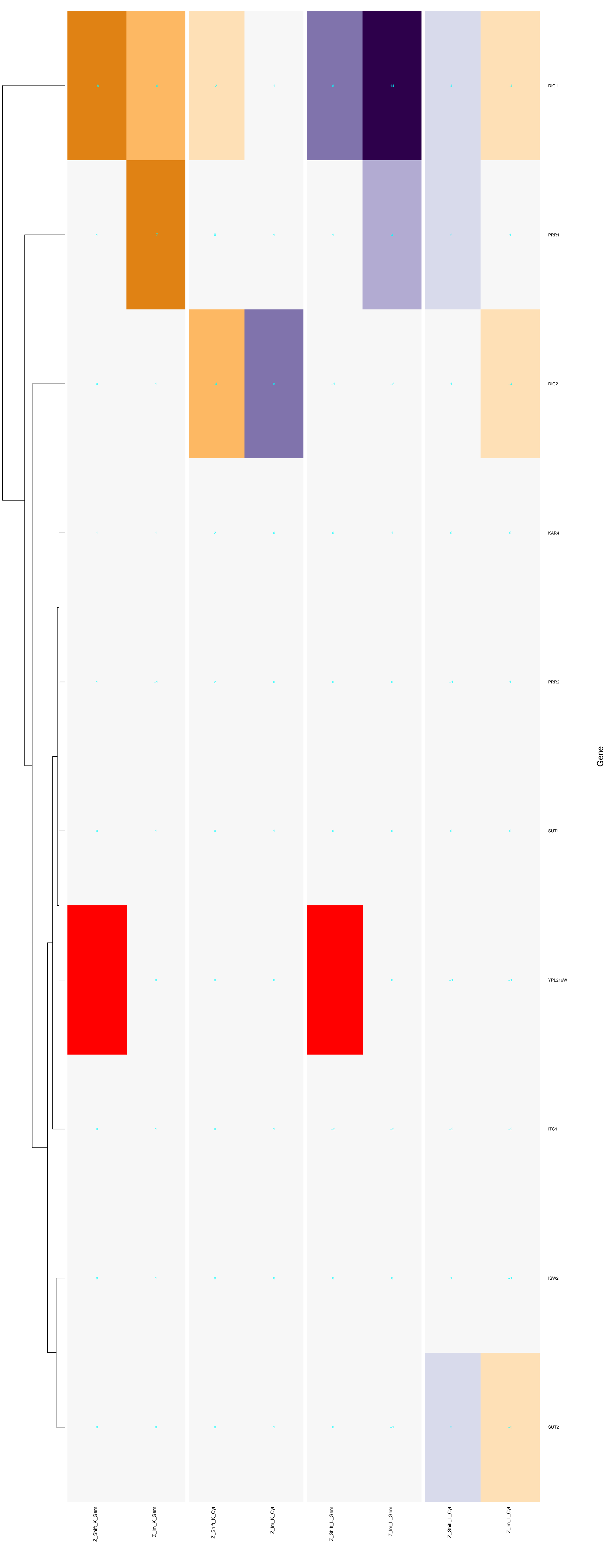
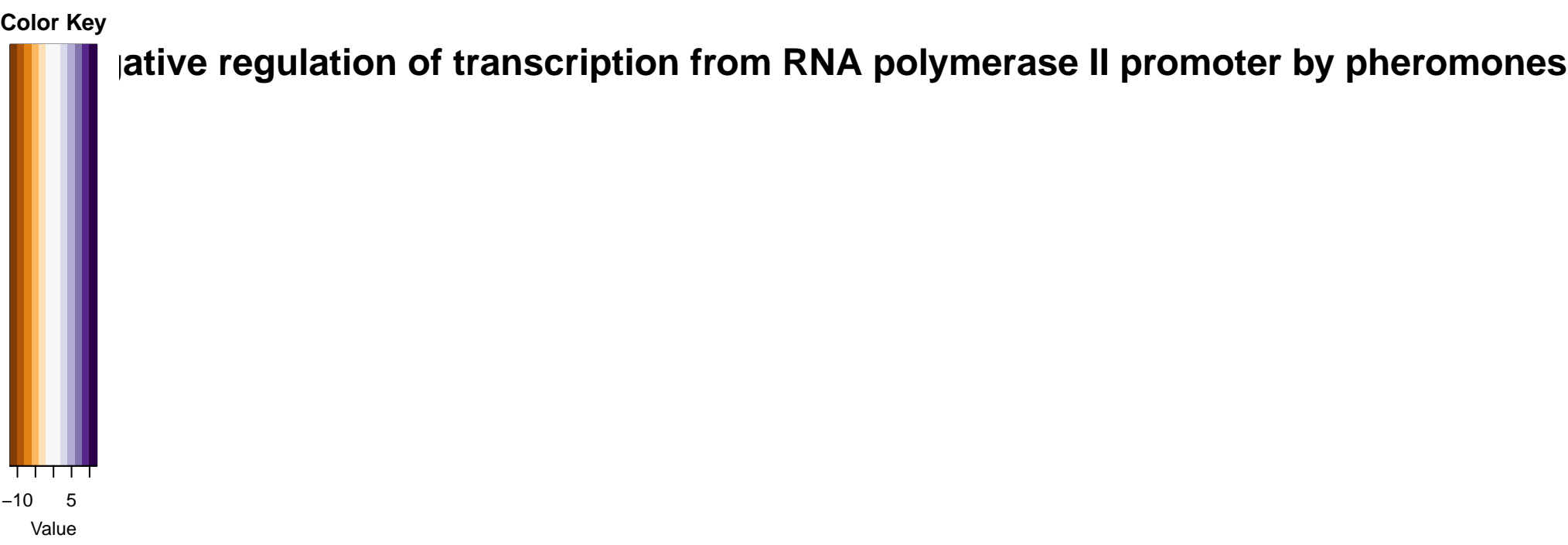


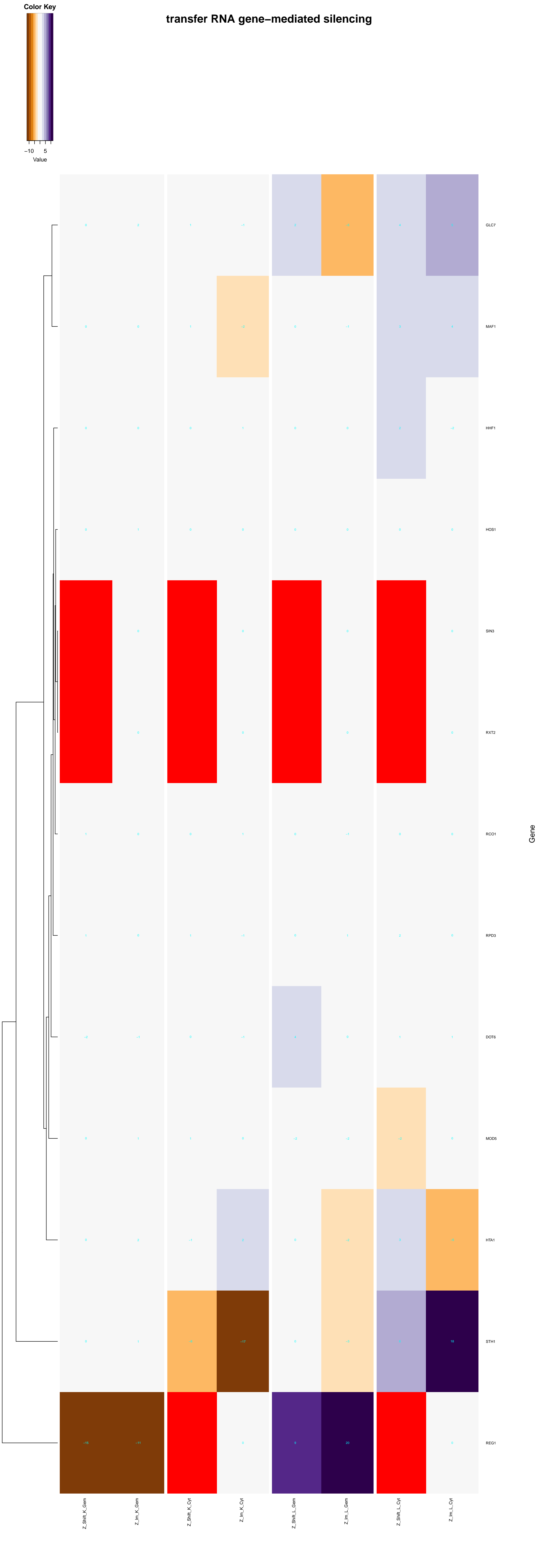


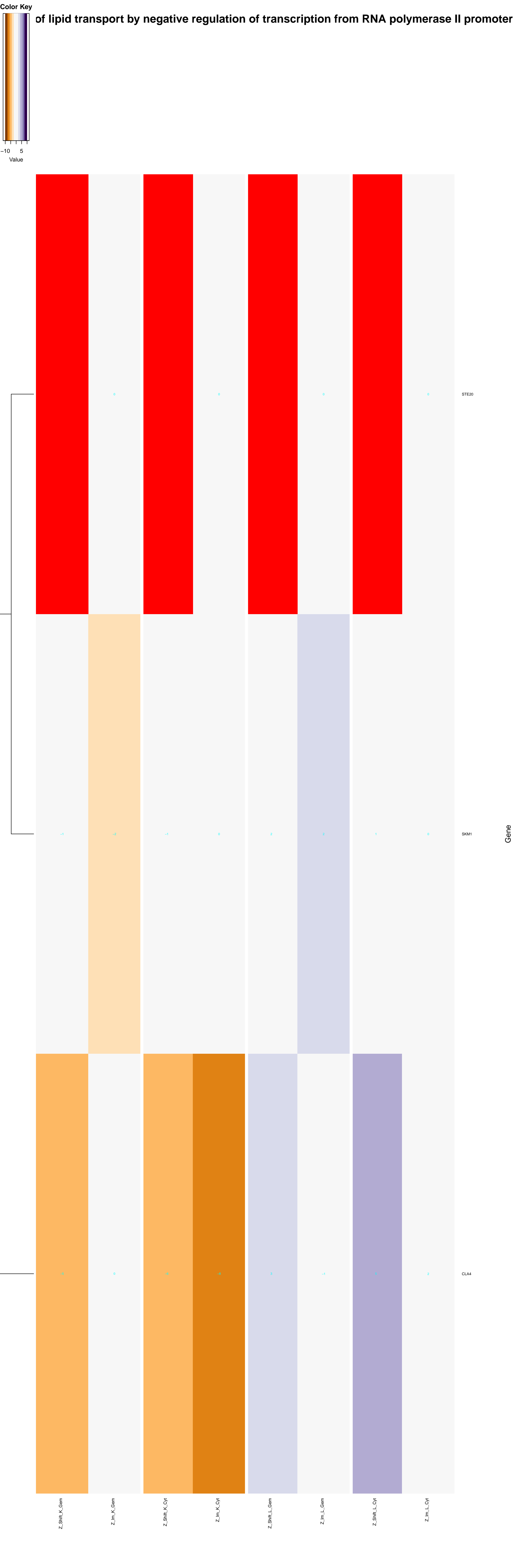
e regulation of ribosomal protein gene transcription from RNA polymerase II promoter

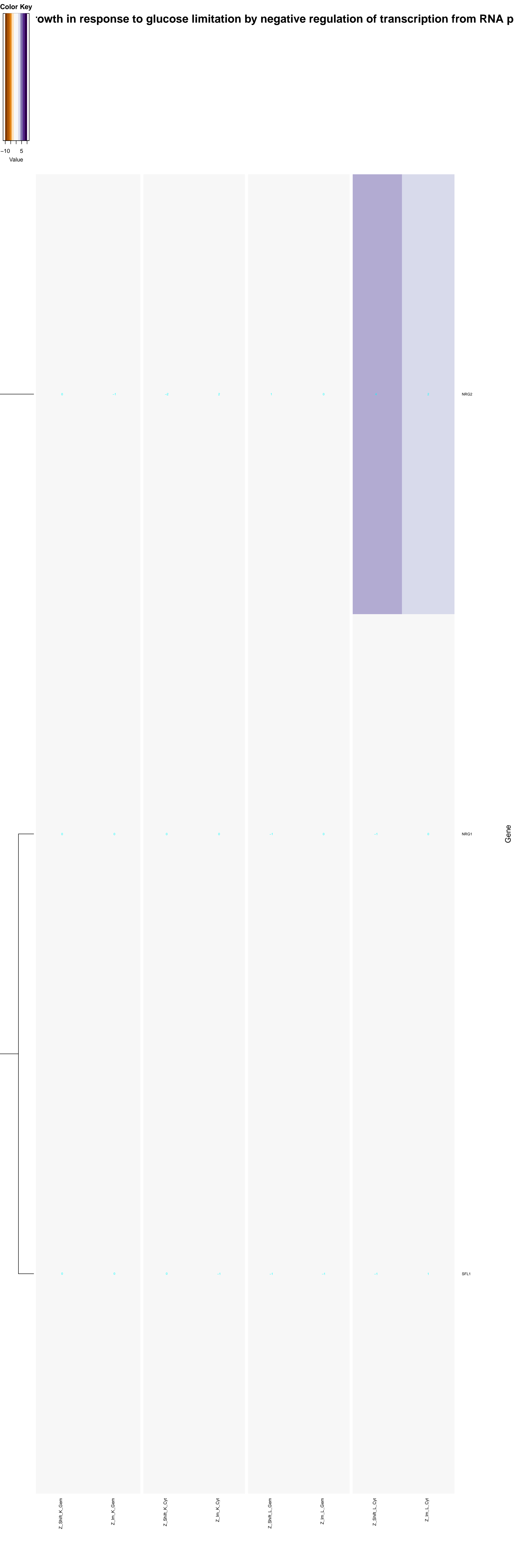


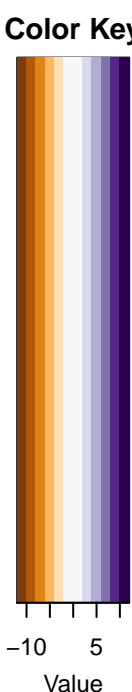




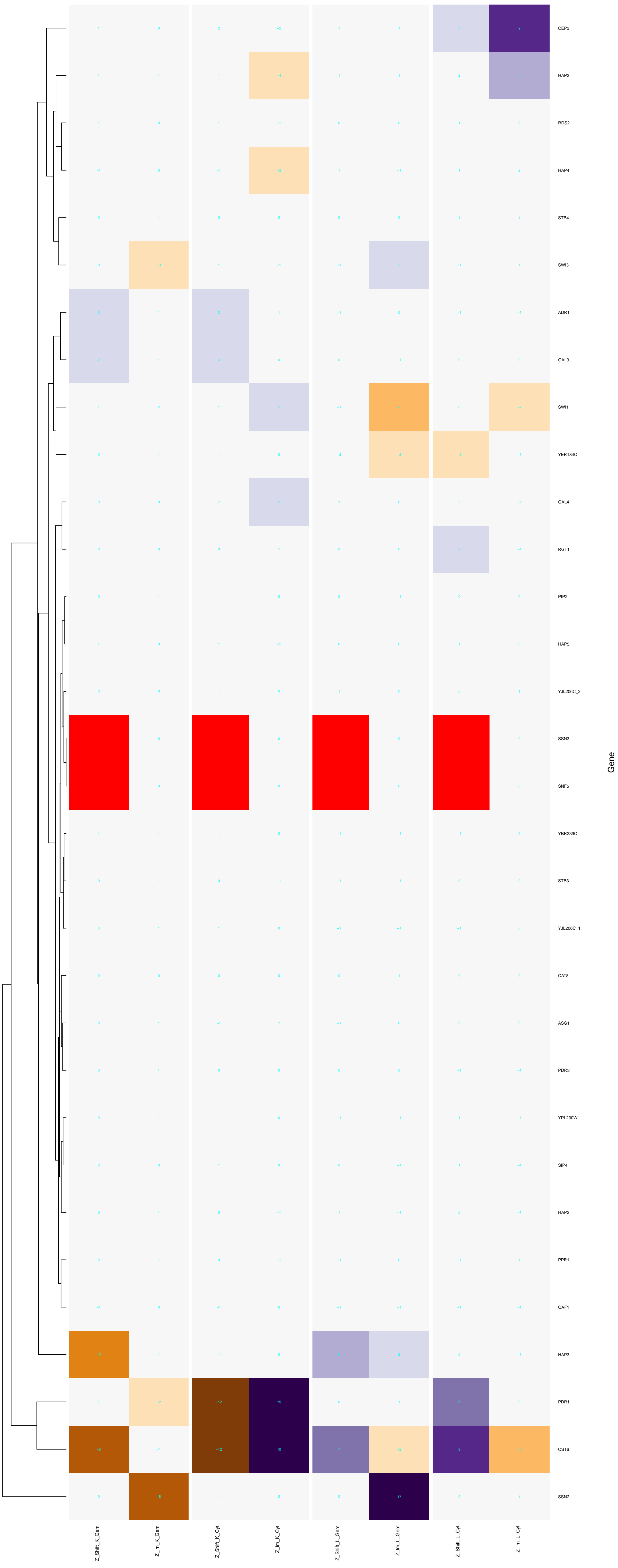


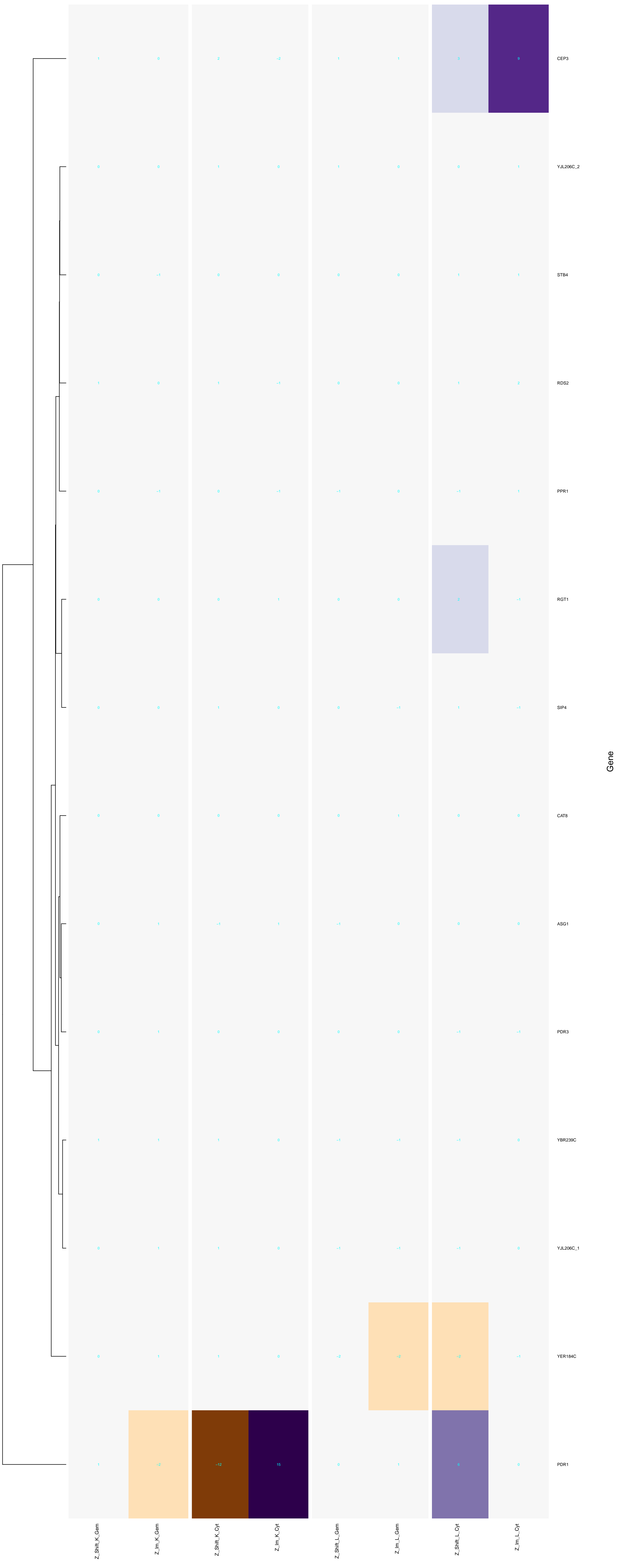
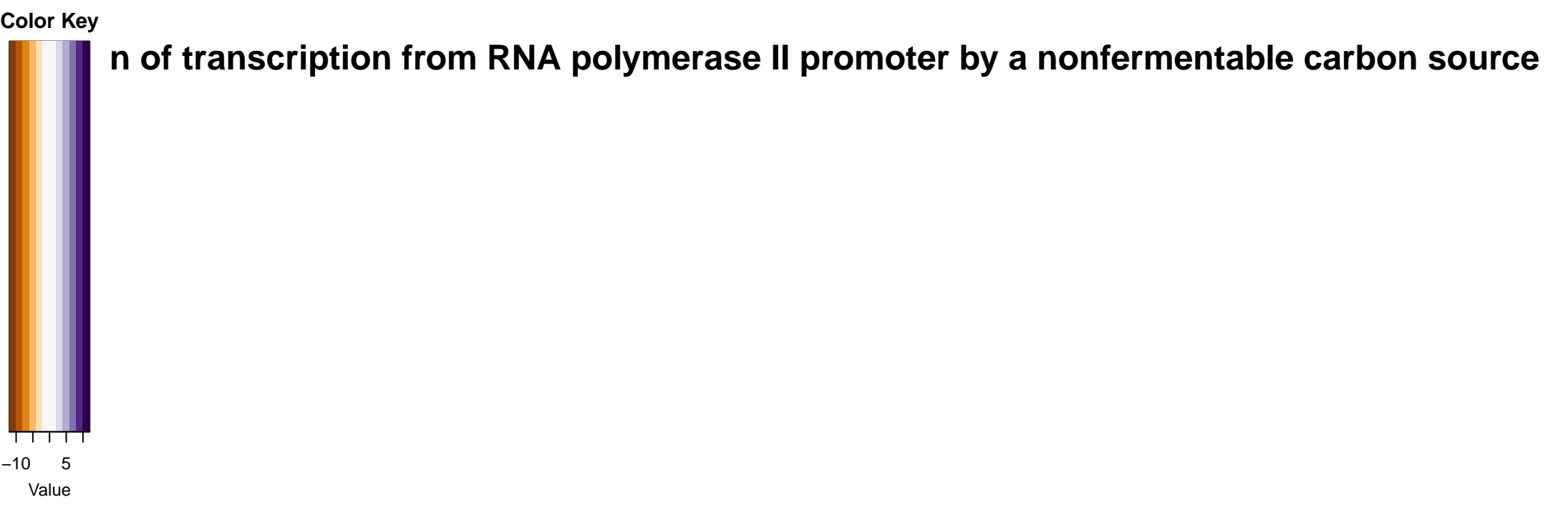






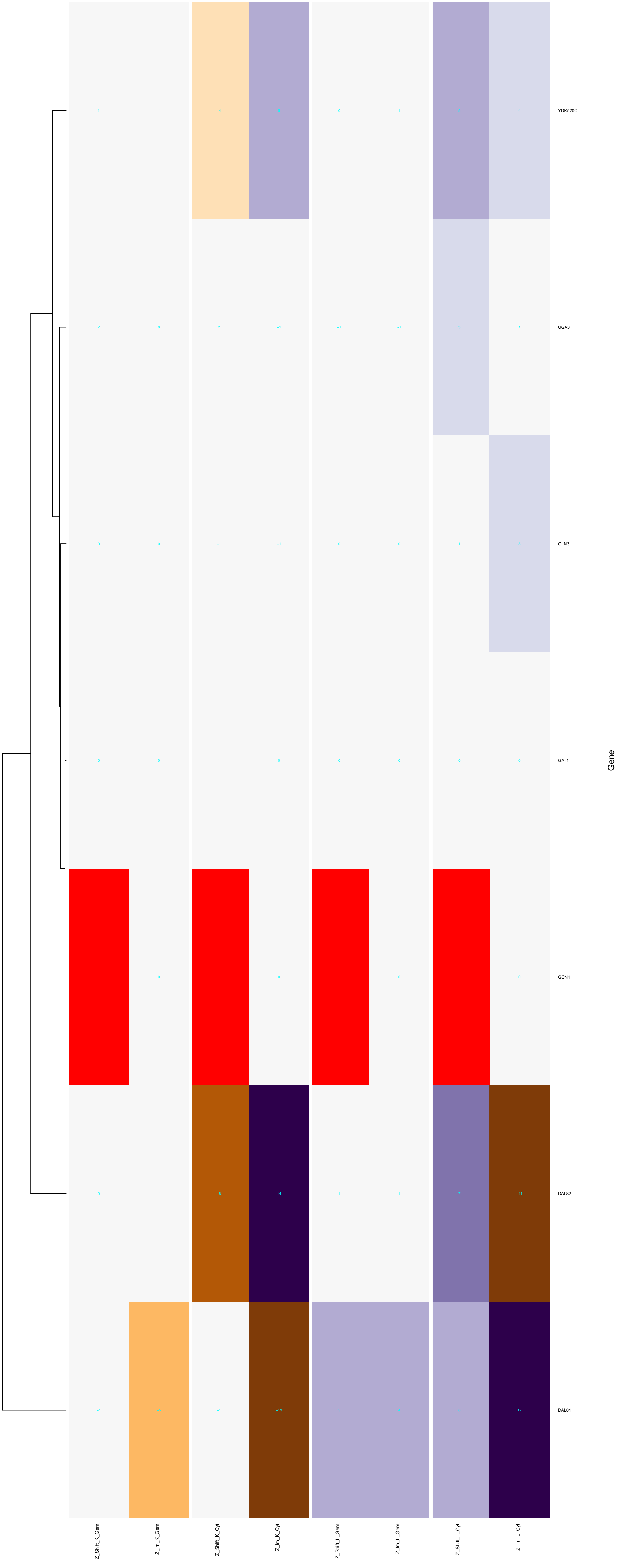
carbon catabolite activation of transcription from RNA polymerase II promoter

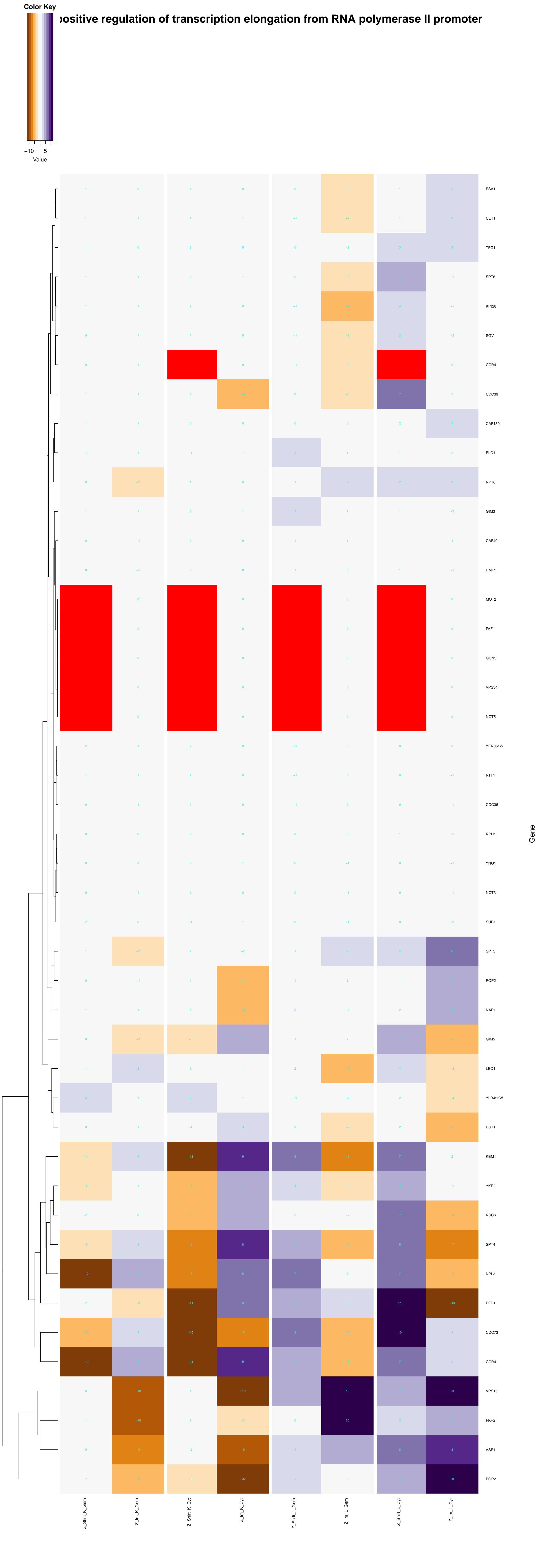


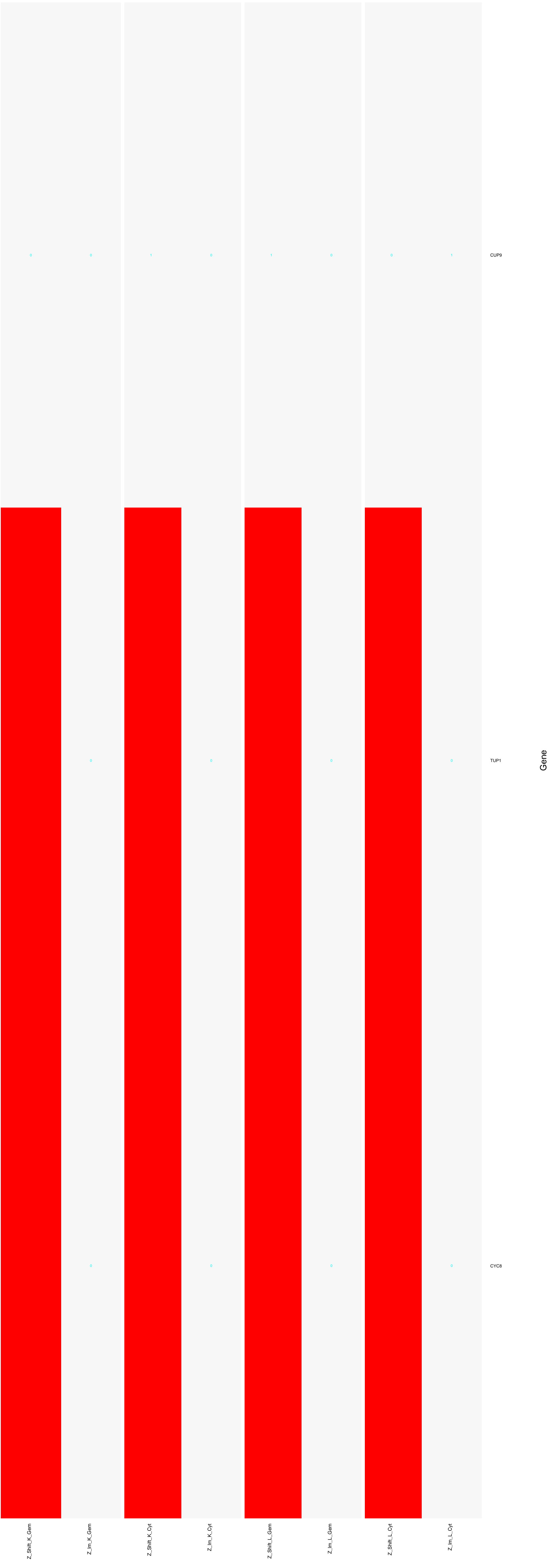
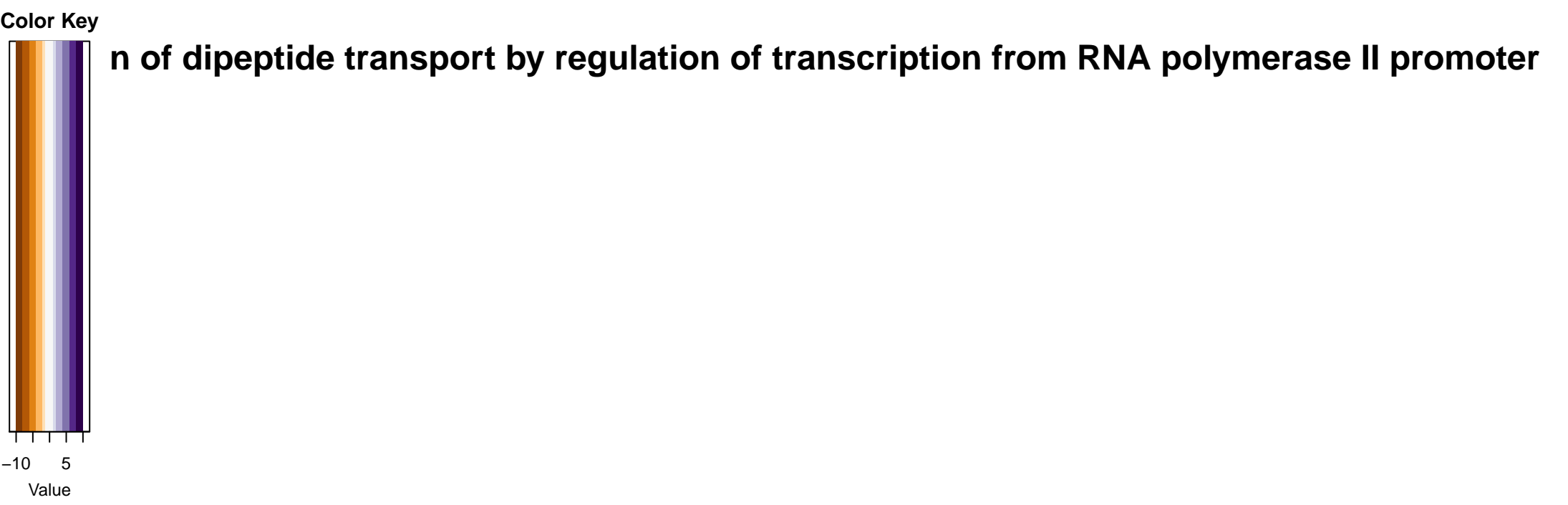


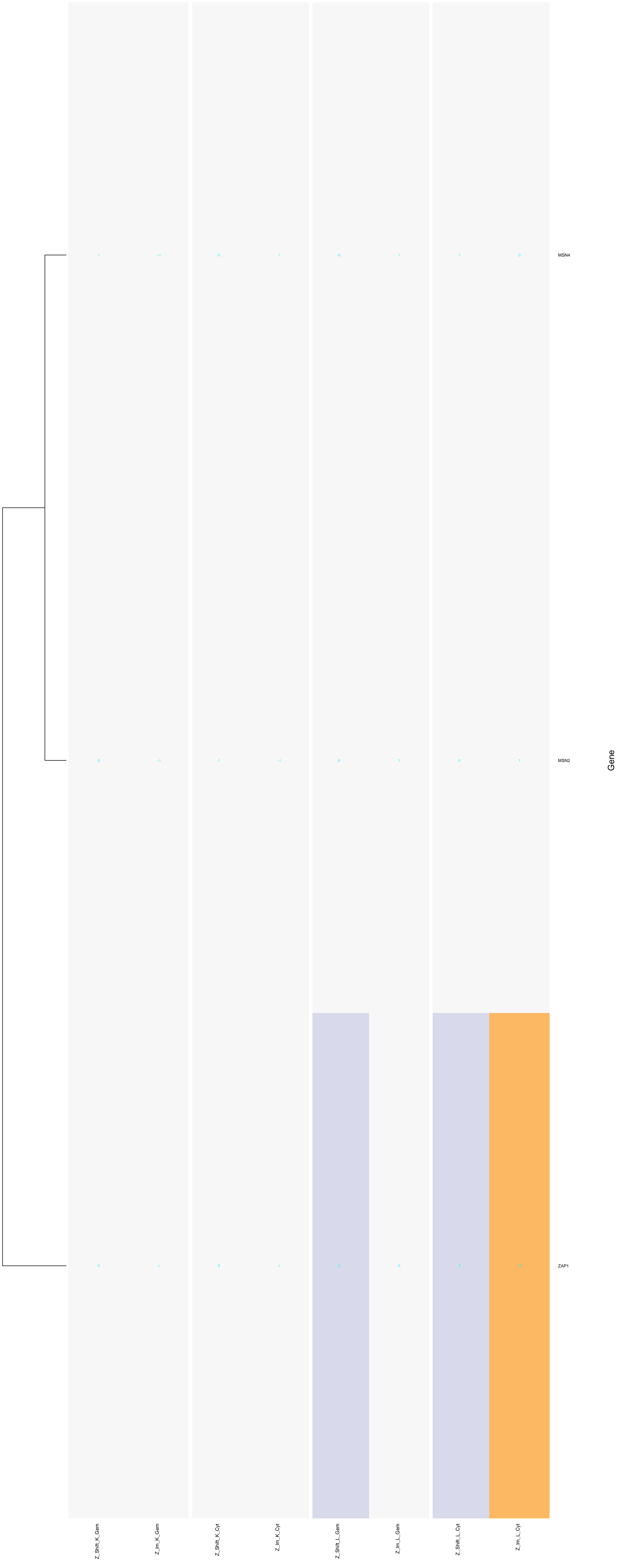
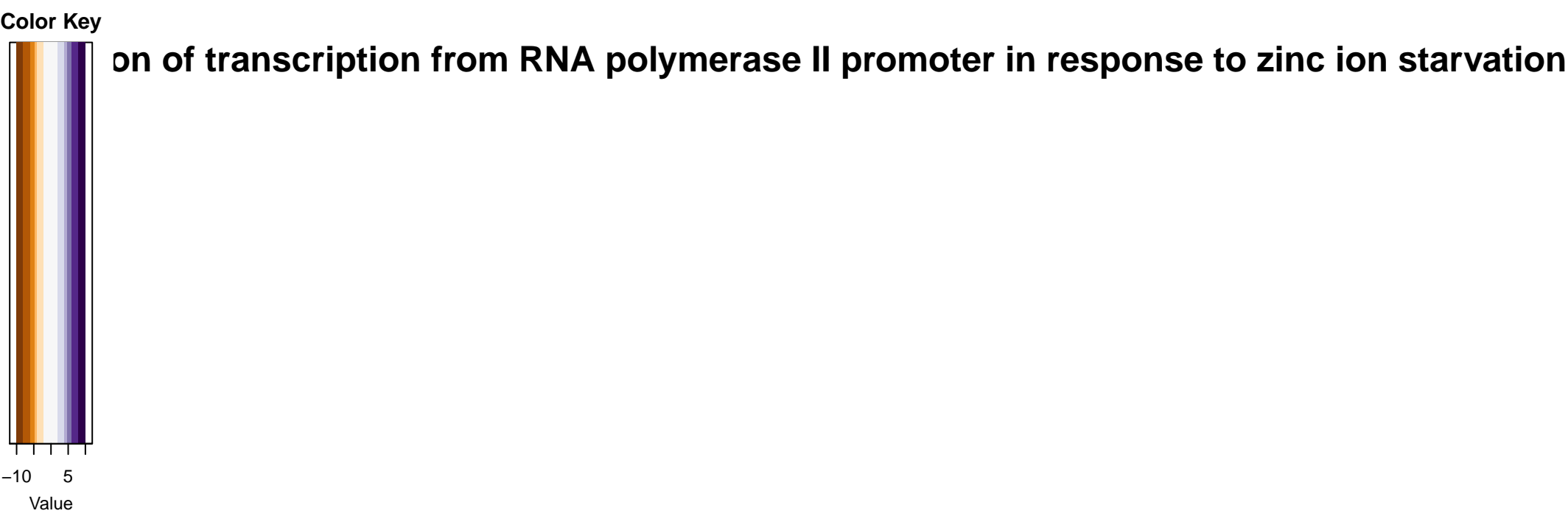


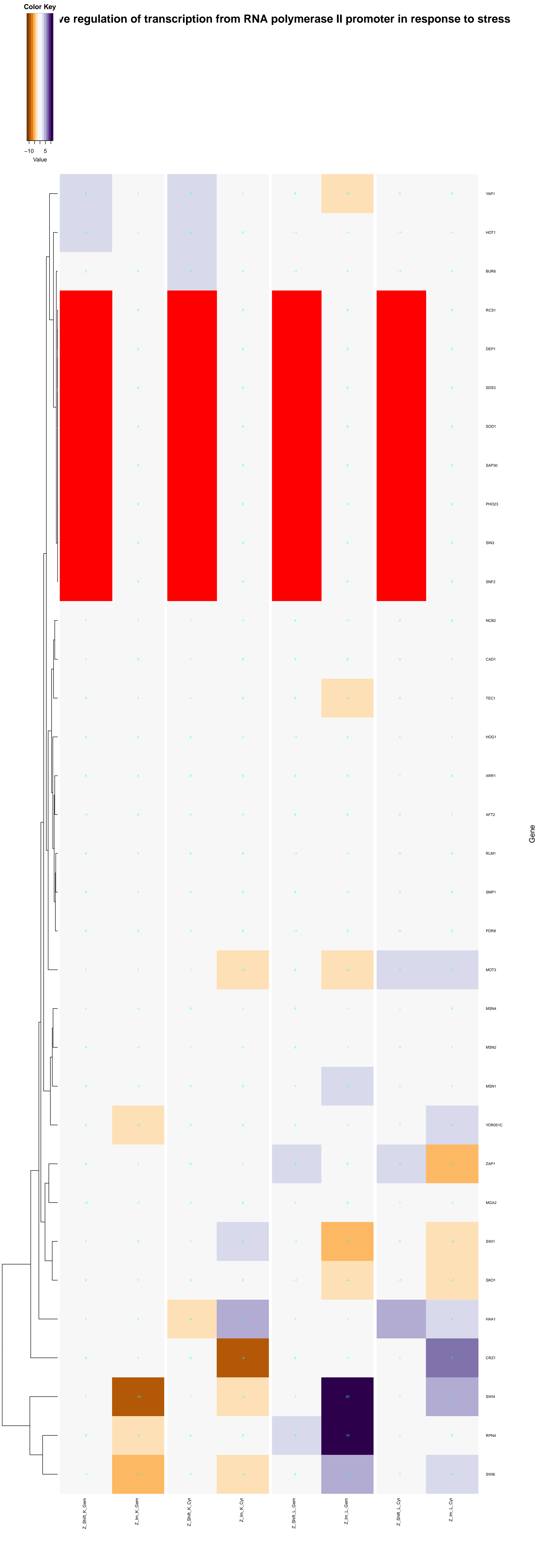
nitrogen catabolite activation of transcription from RNA polymerase II promoter









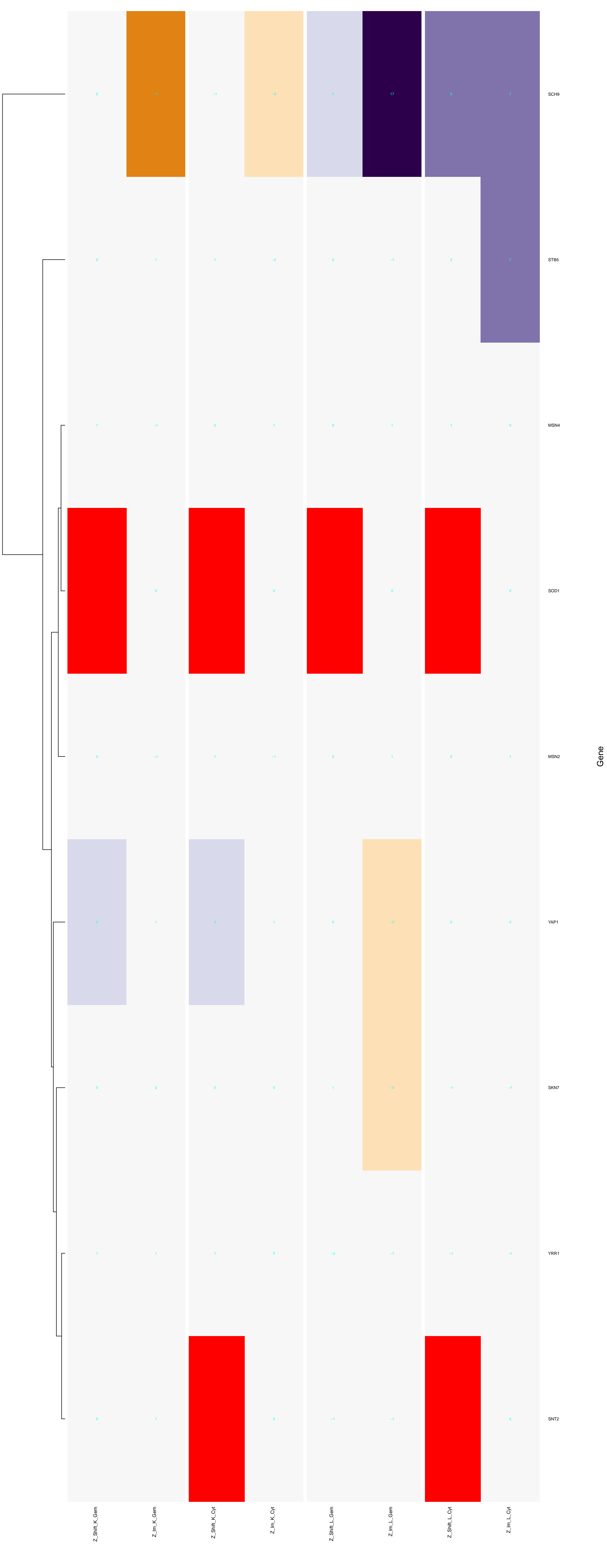


Color Key

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

-10 5

Value

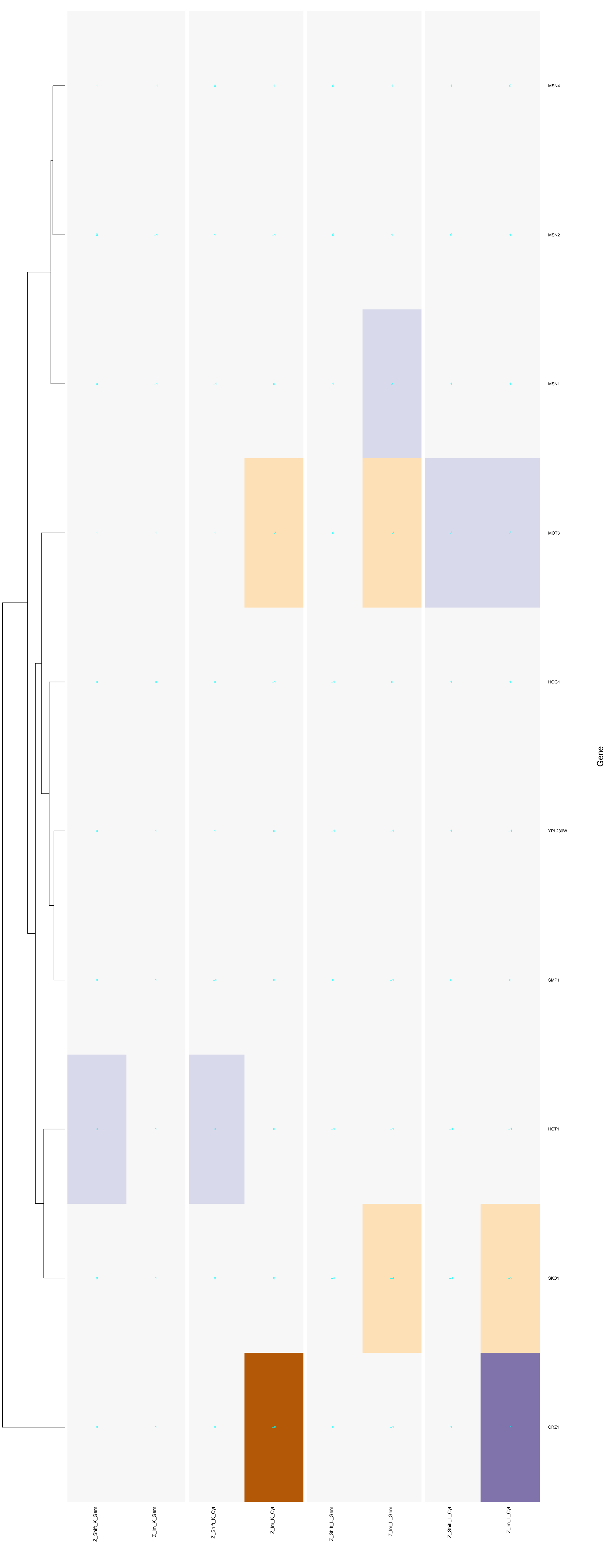


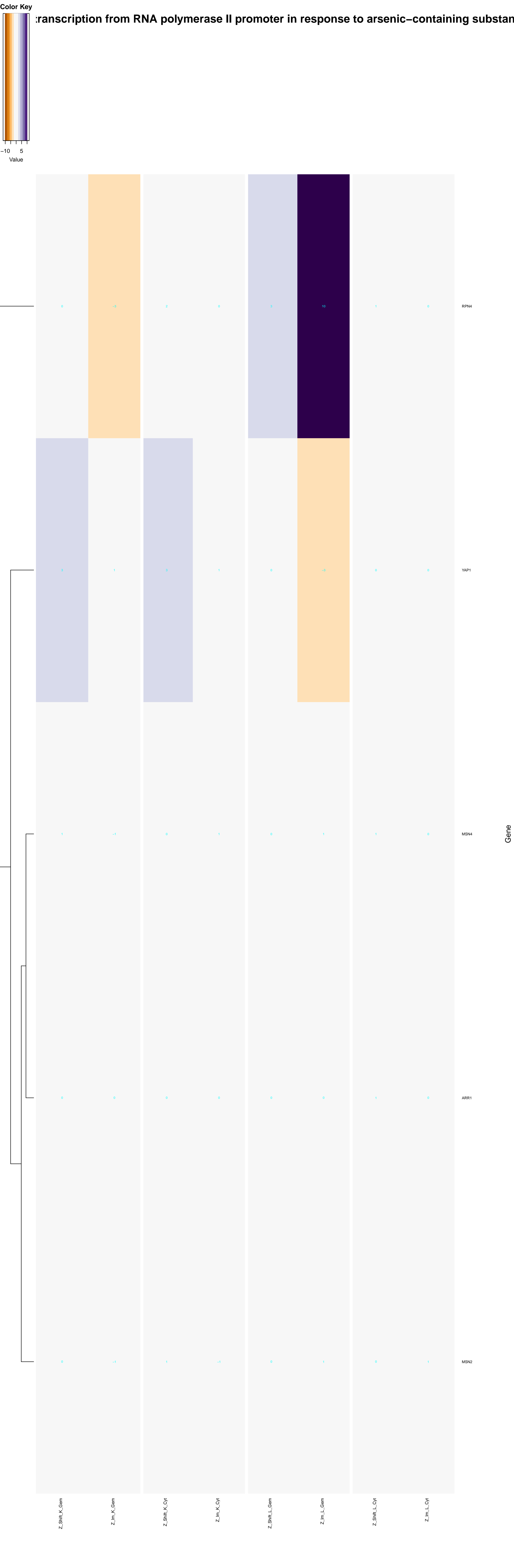
Color Key

Value

10 5

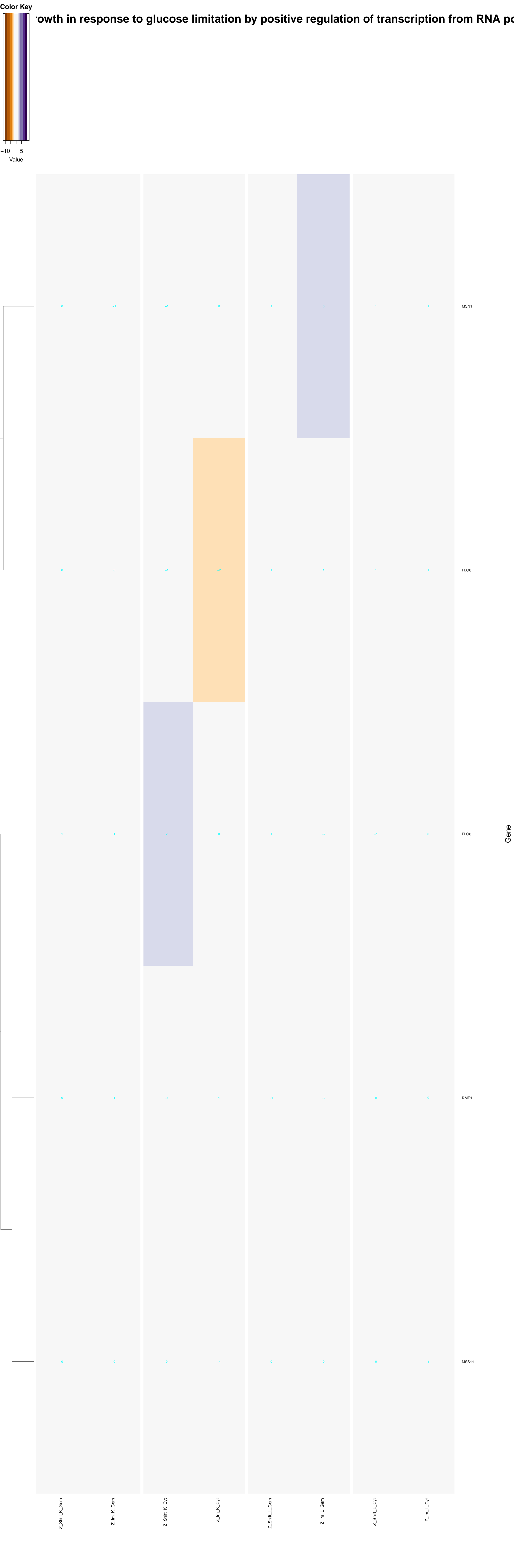
tion of transcription from RNA polymerase II promoter in response to osmotic stress

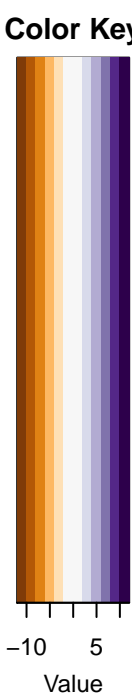




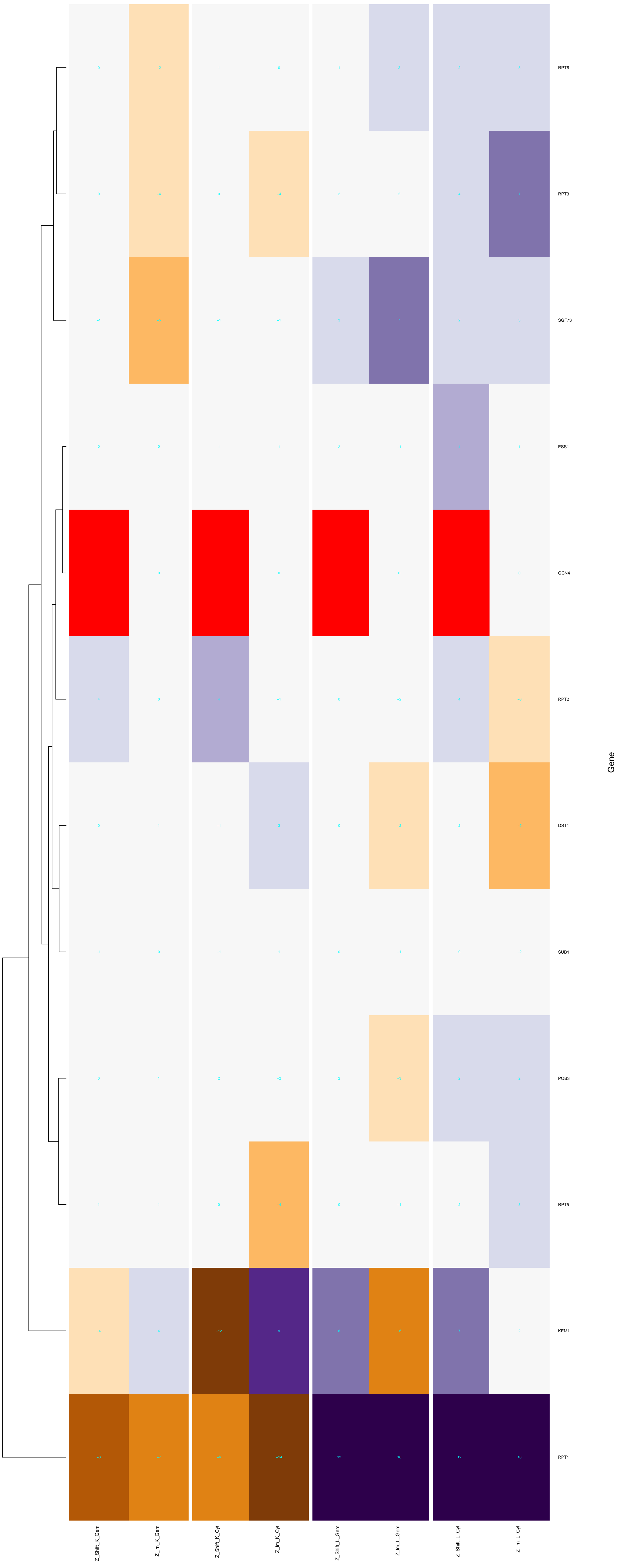


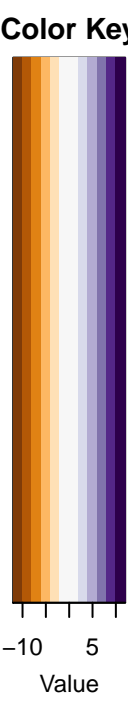




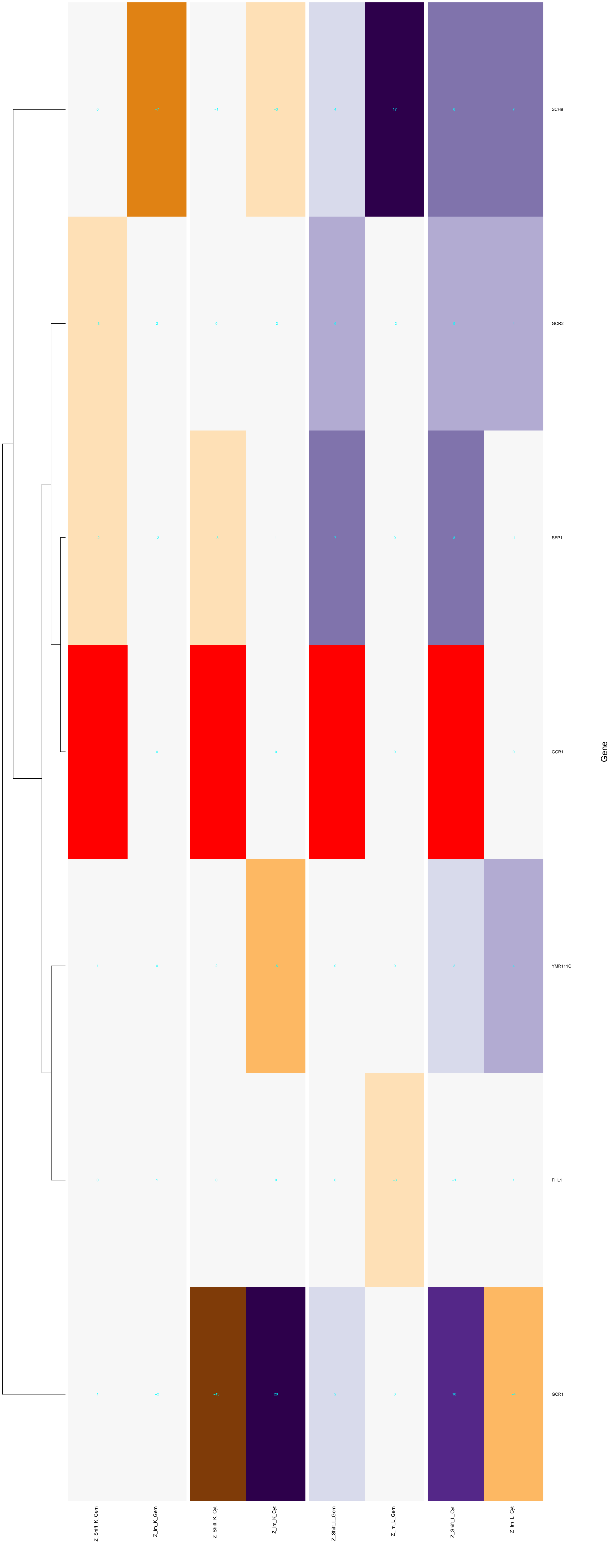


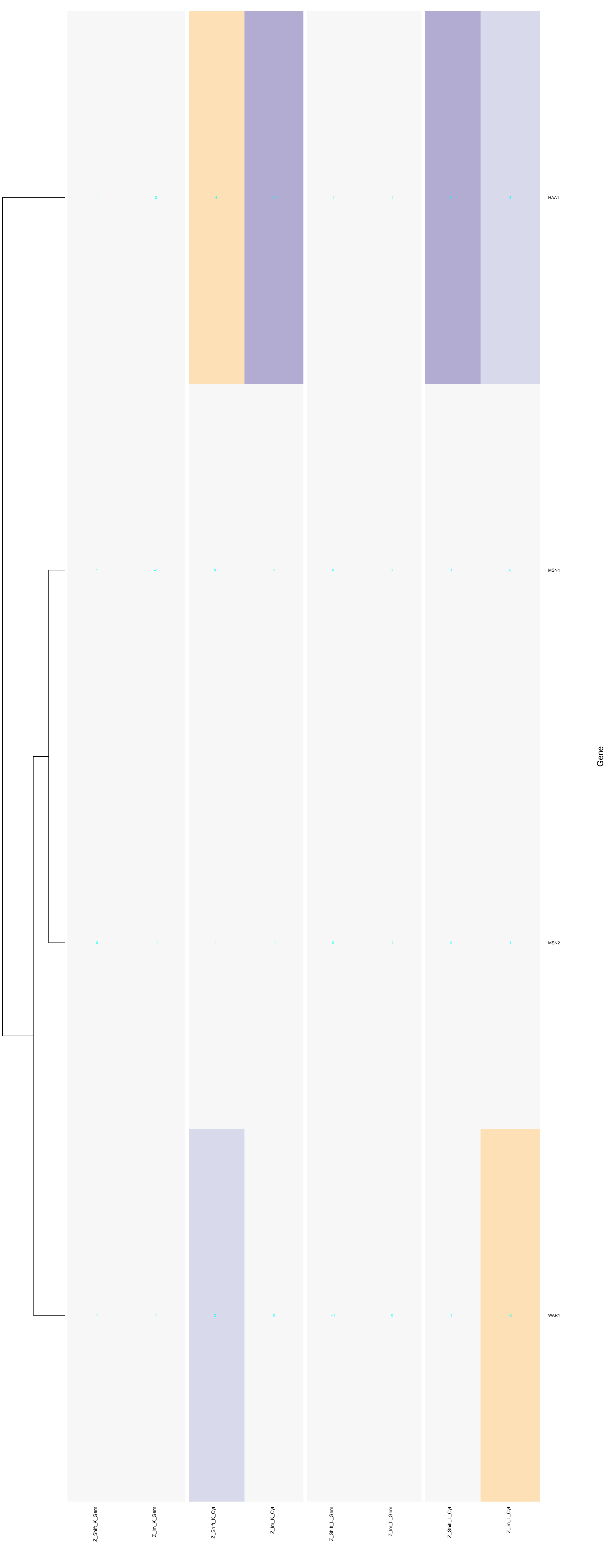
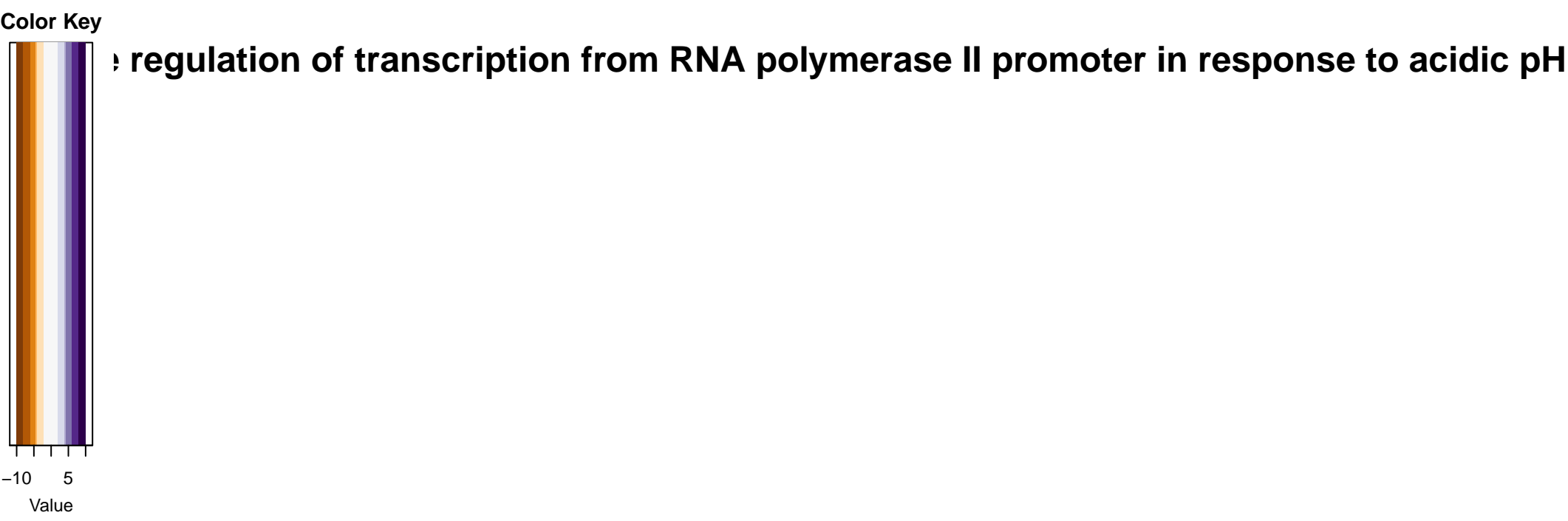
positive regulation of transcription initiation from RNA polymerase II promoter

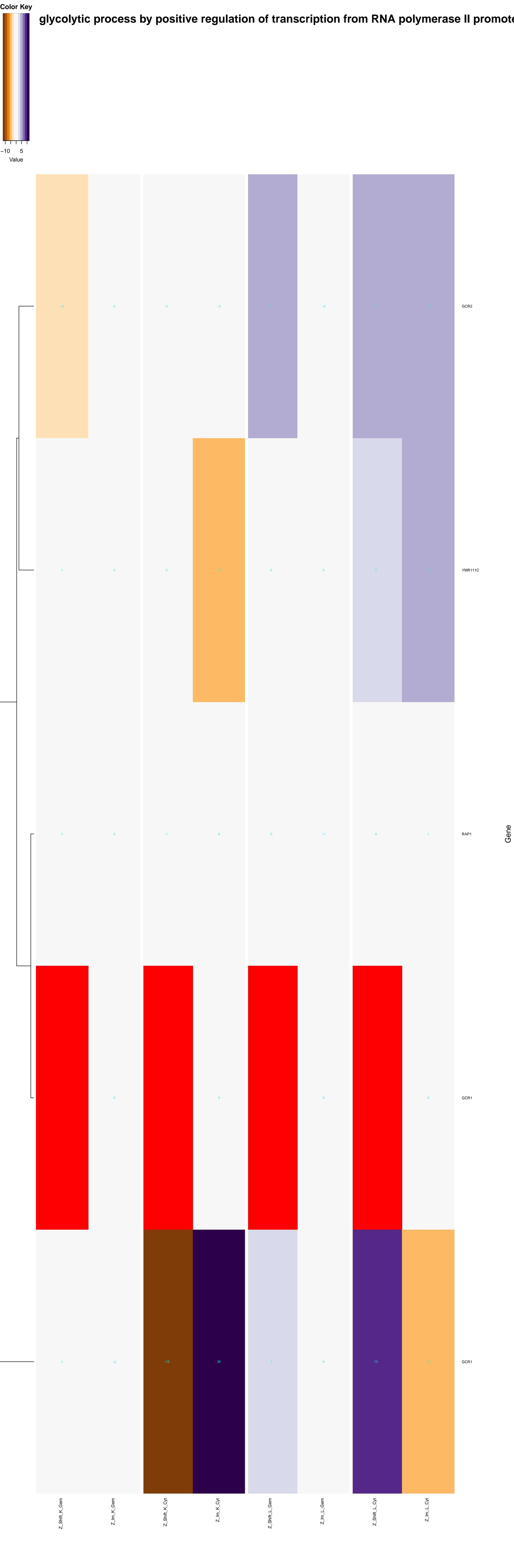


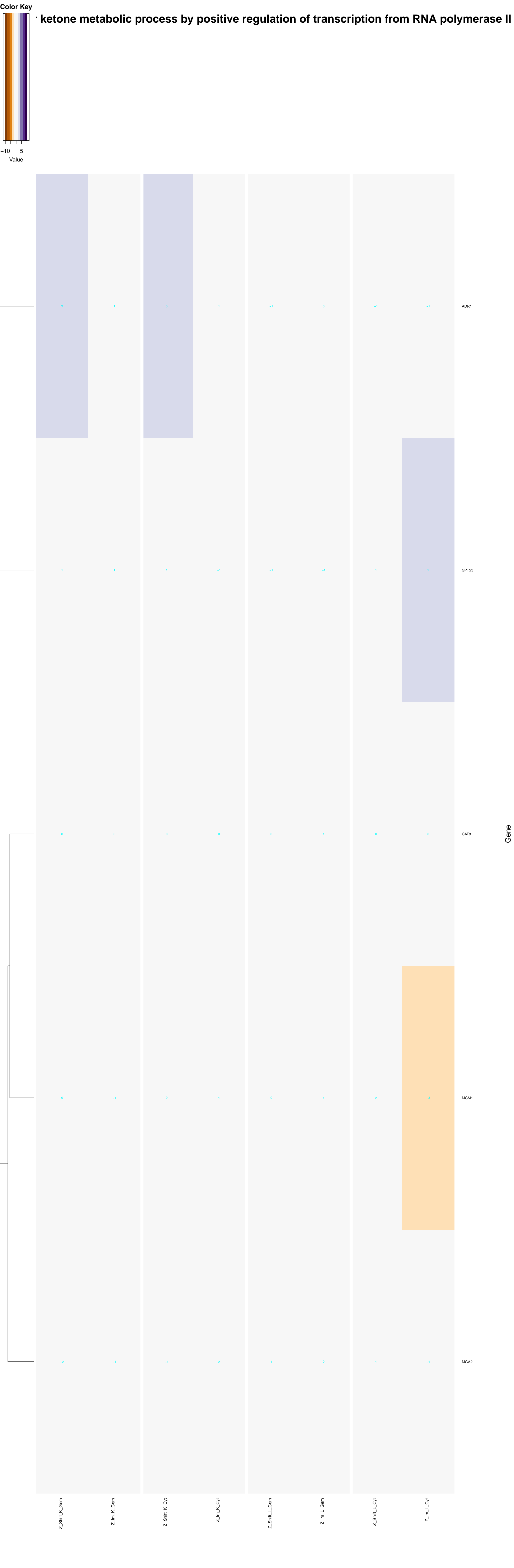


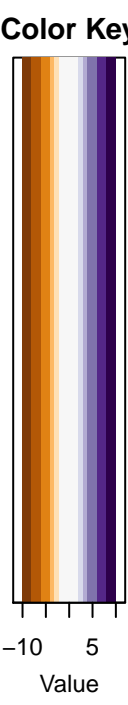
e regulation of ribosomal protein gene transcription from RNA polymerase II promoter



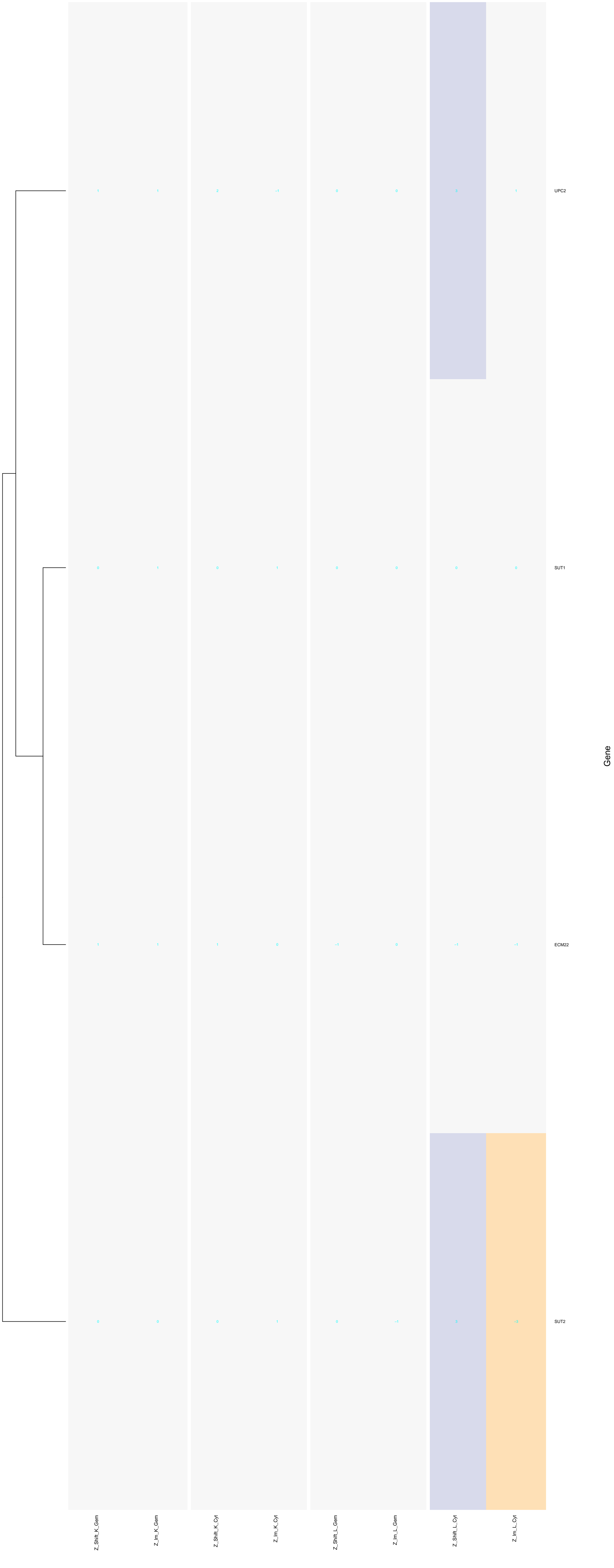


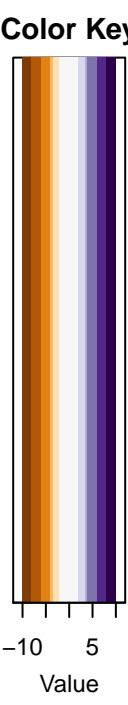




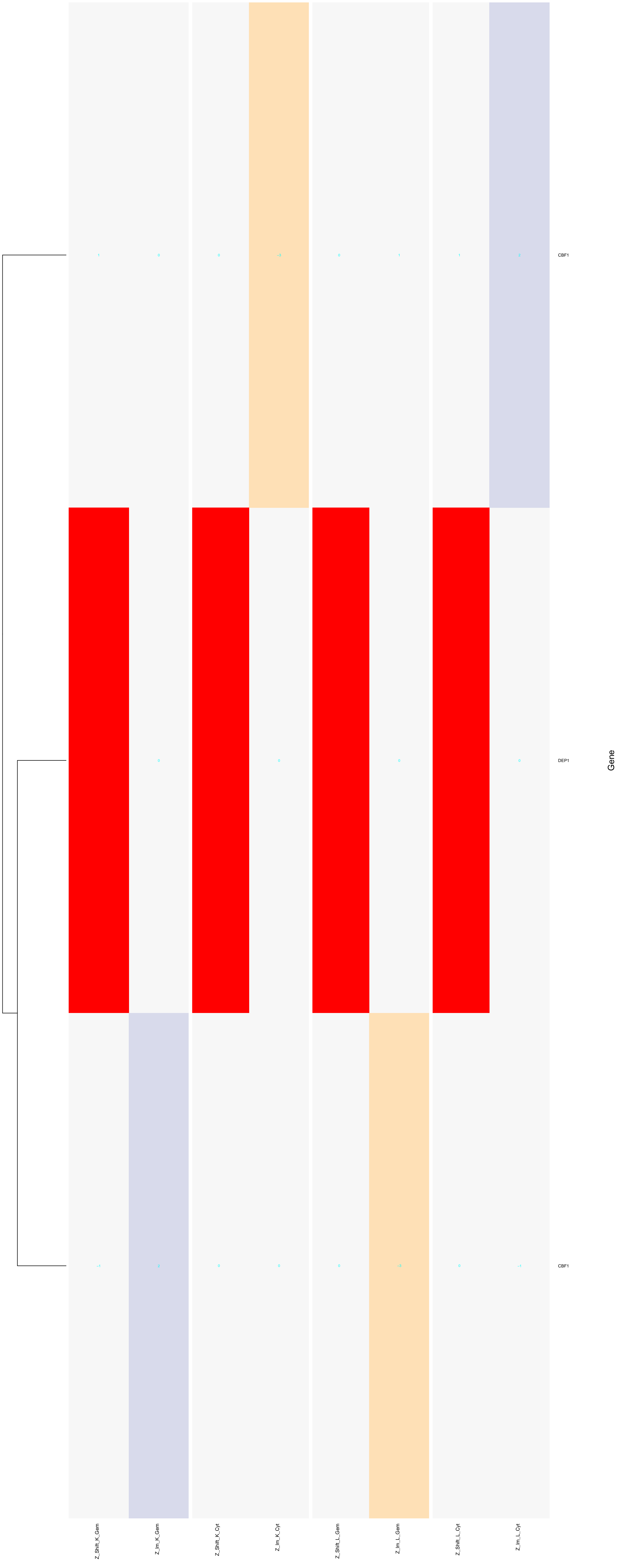


of lipid transport by positive regulation of transcription from RNA polymerase II promoter

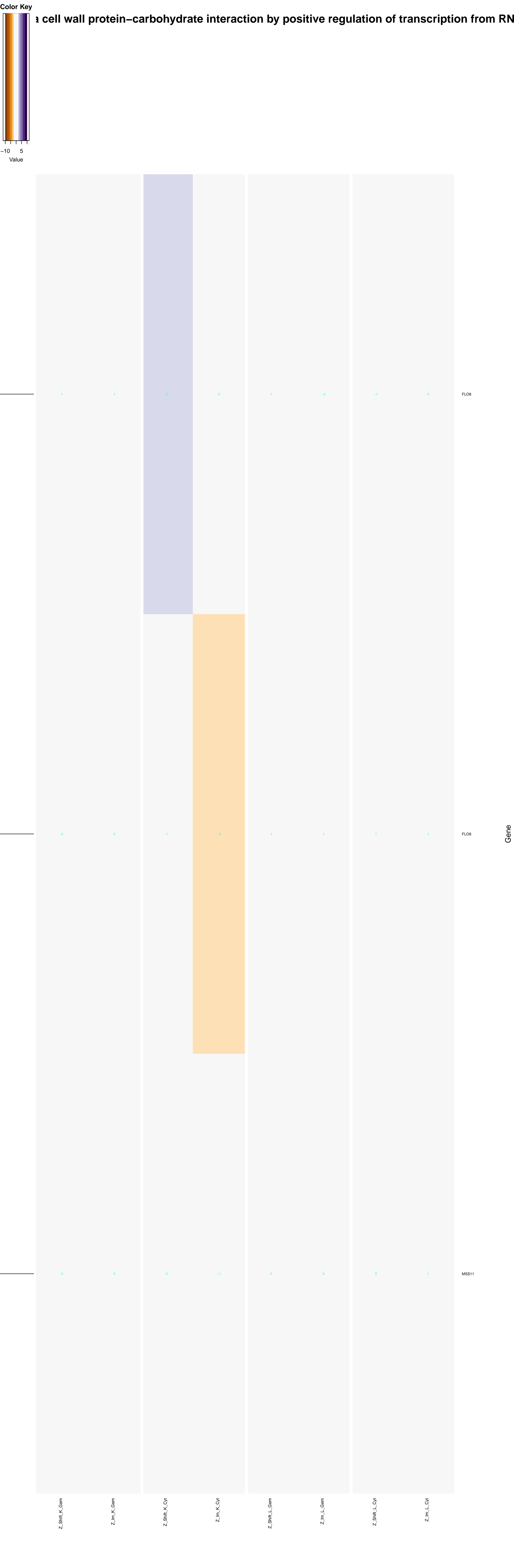


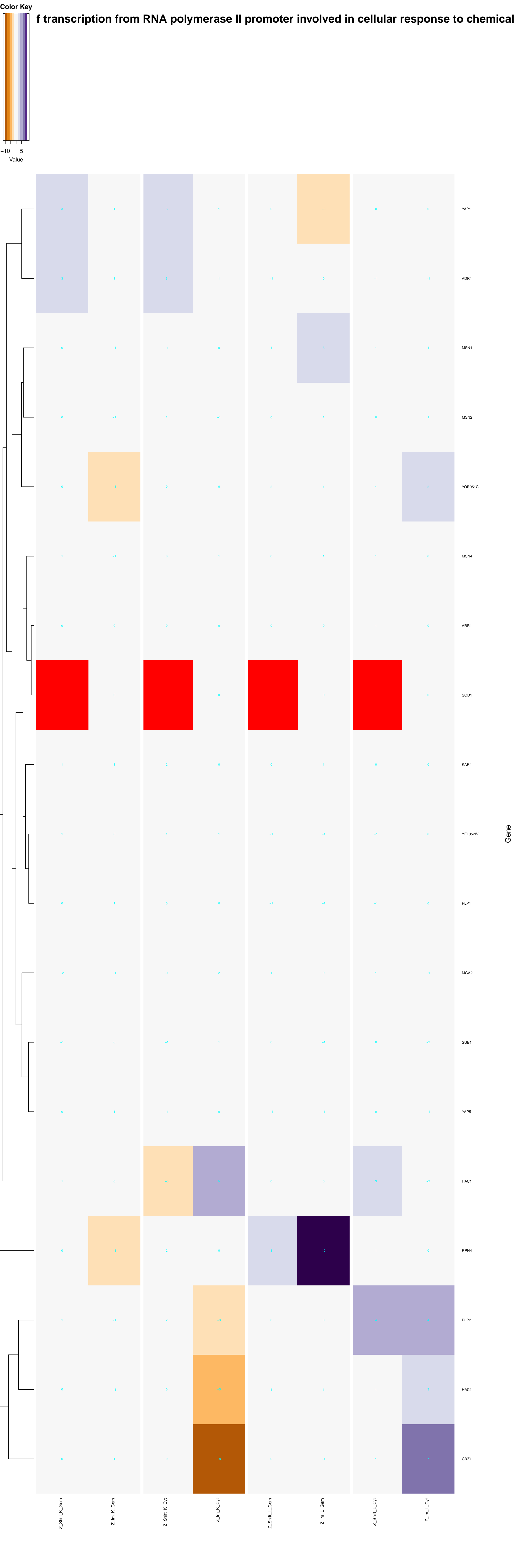


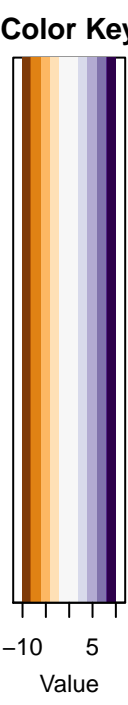
nositol biosynthetic process by positive regulation of transcription from RNA polymerase



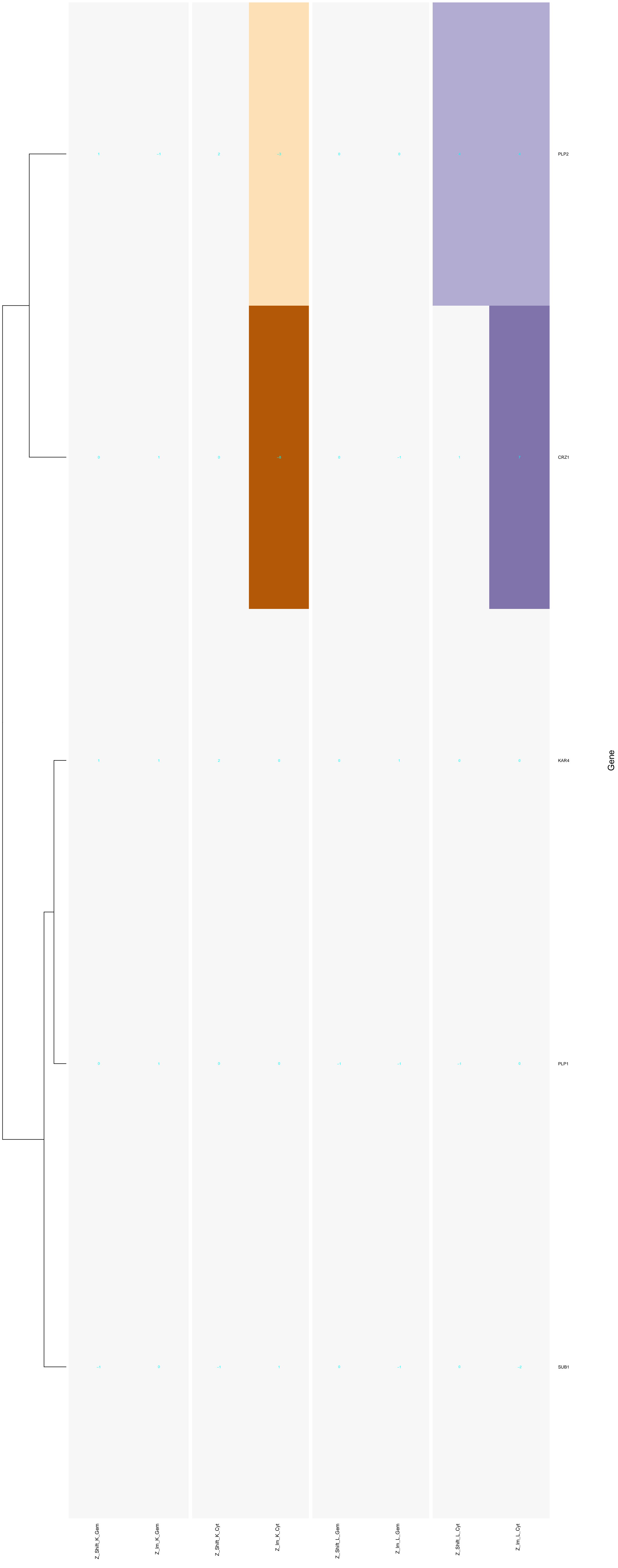






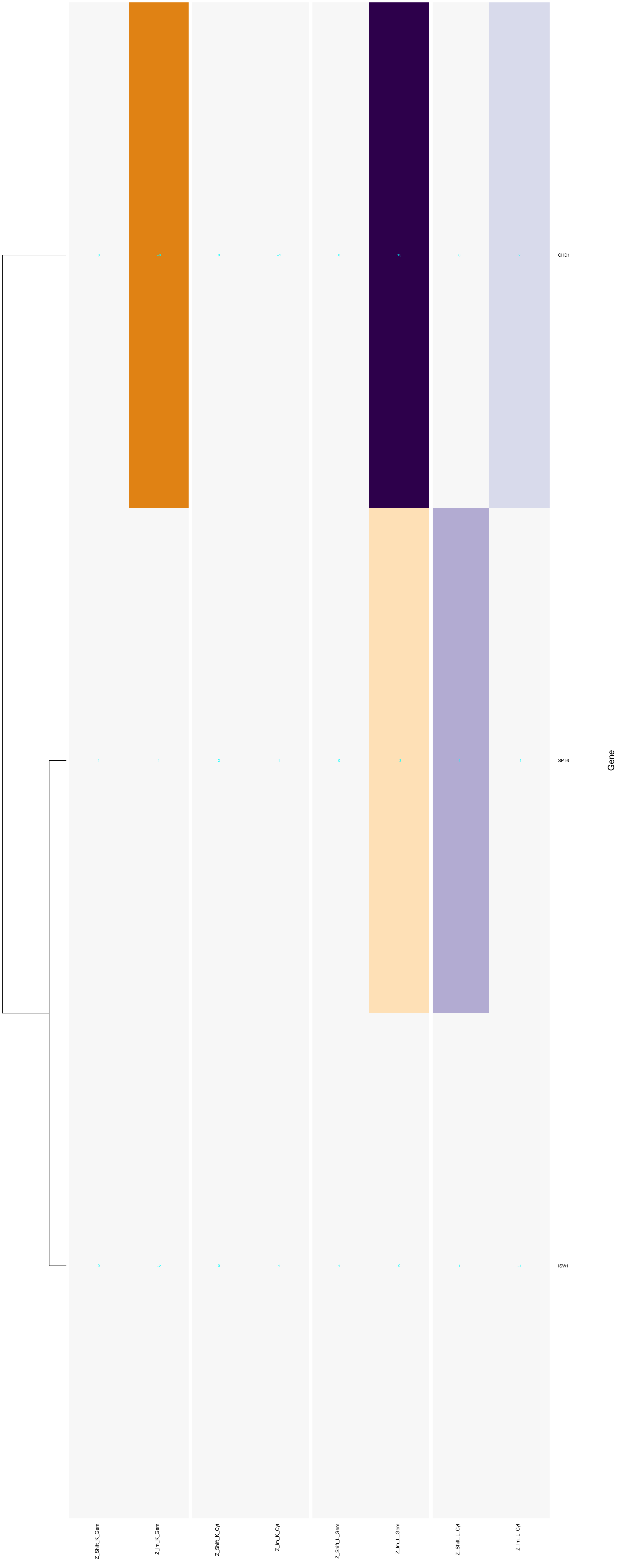


Positive regulation of transcription from RNA polymerase II promoter by pheromones





regulation of transcriptional start site selection at RNA polymerase II promoter



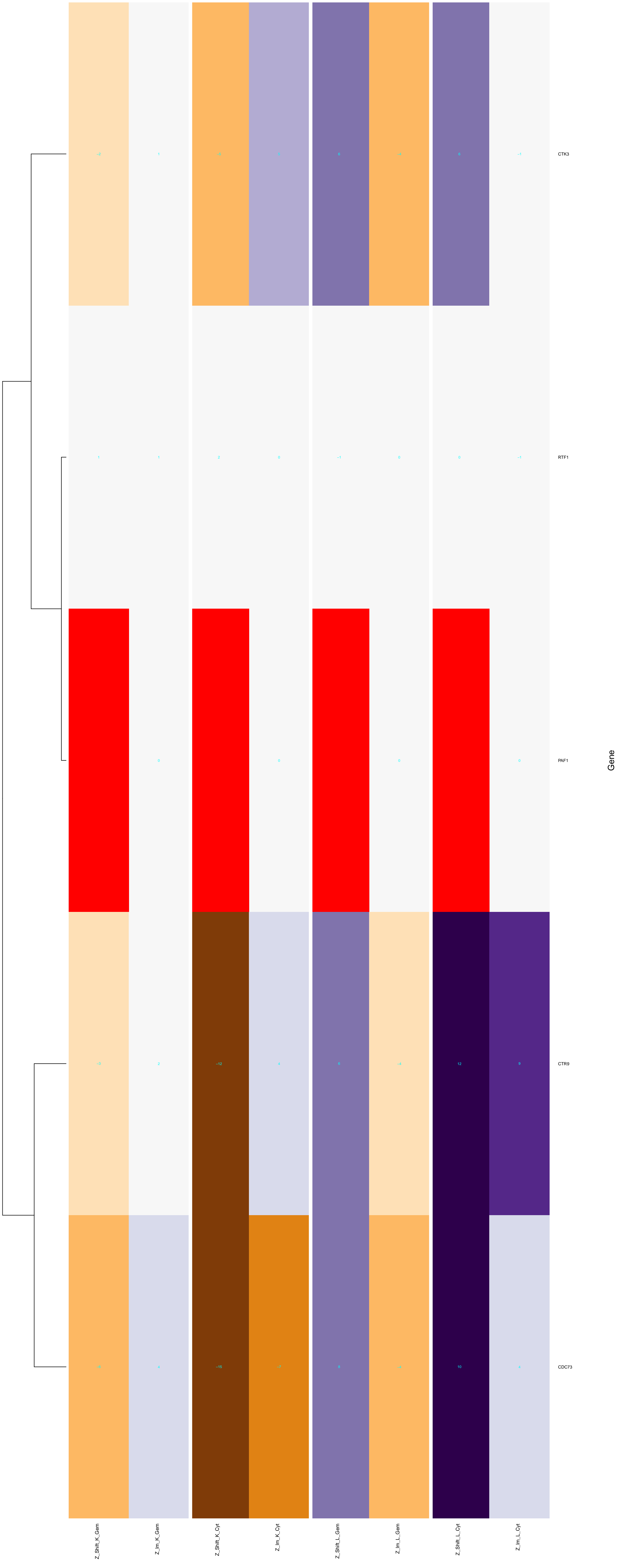


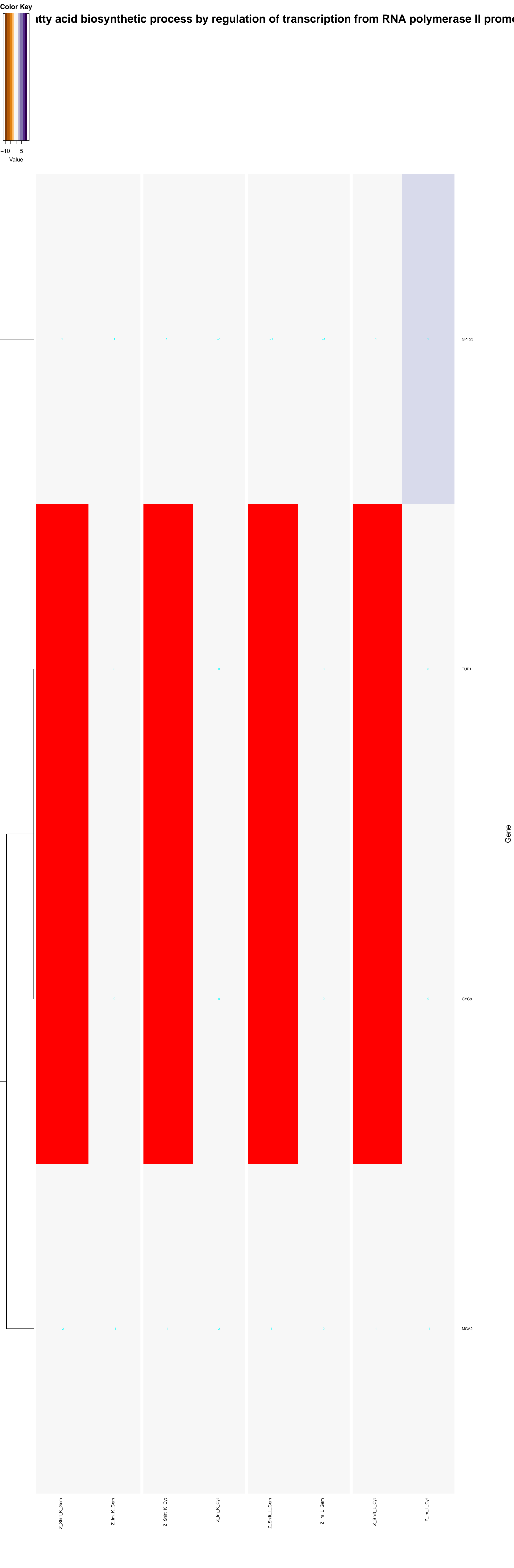
regulation of RNA polymerase II transcriptional preinitiation complex assembly





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



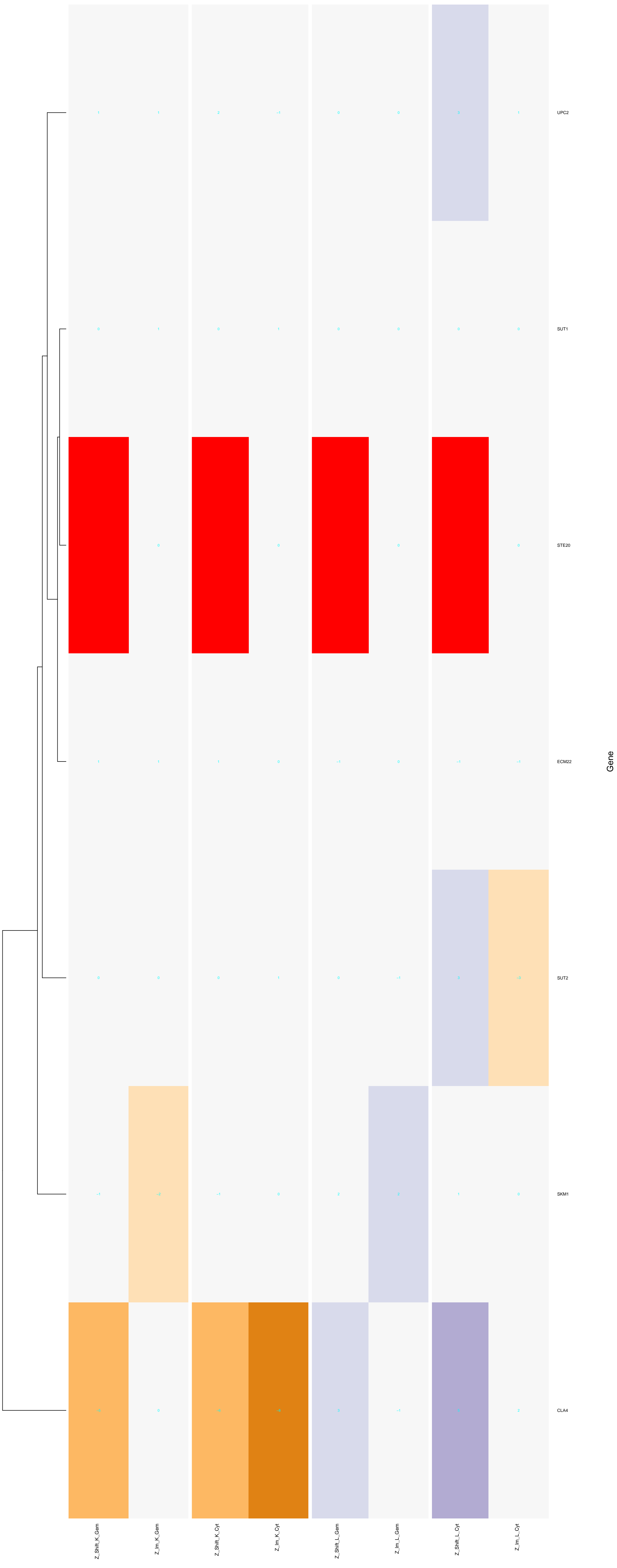


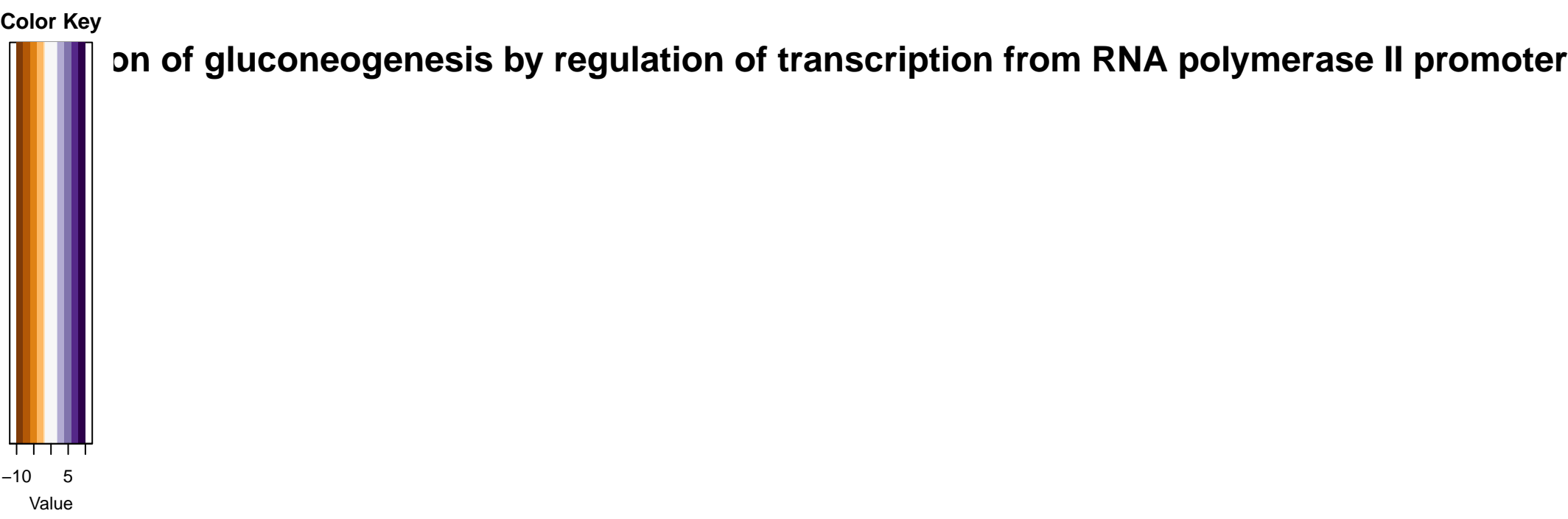
Color Key

ation of sterol import by regulation of transcription from RNA polymerase II promoter

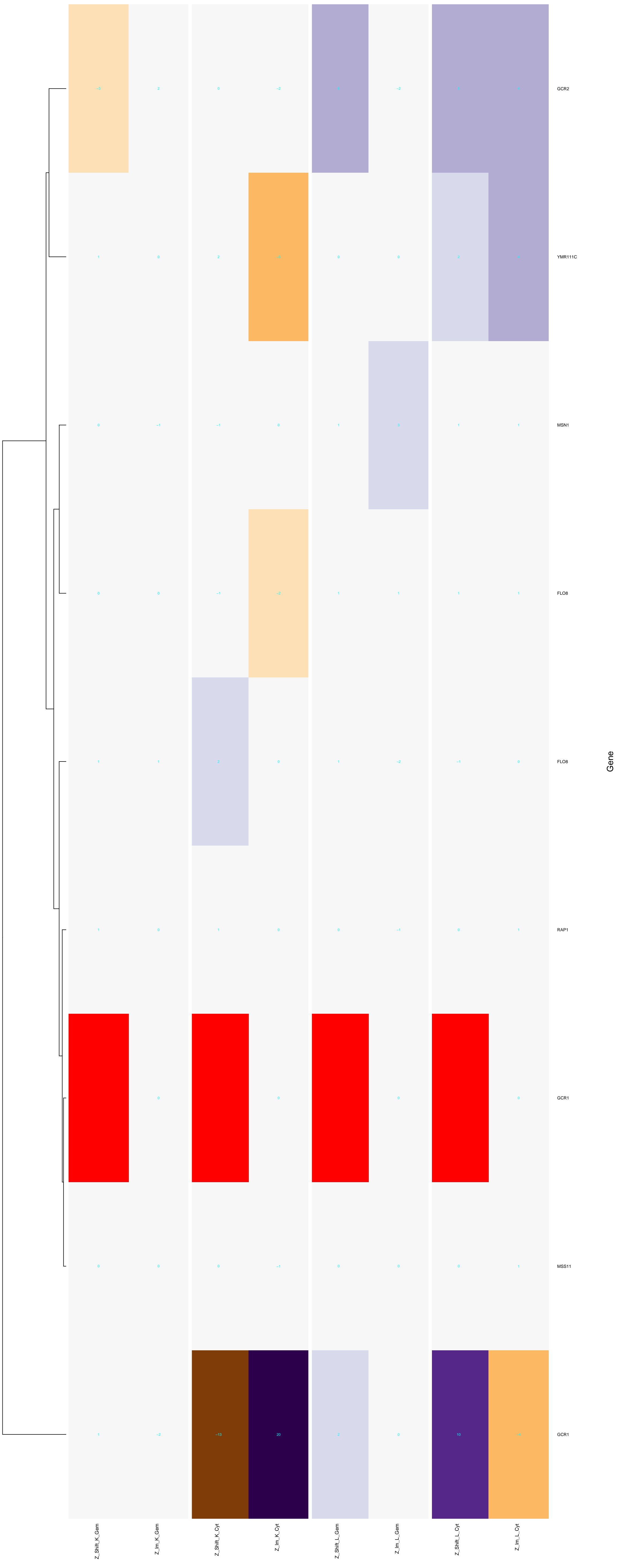
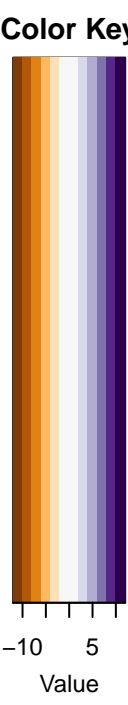
-10 5

Value



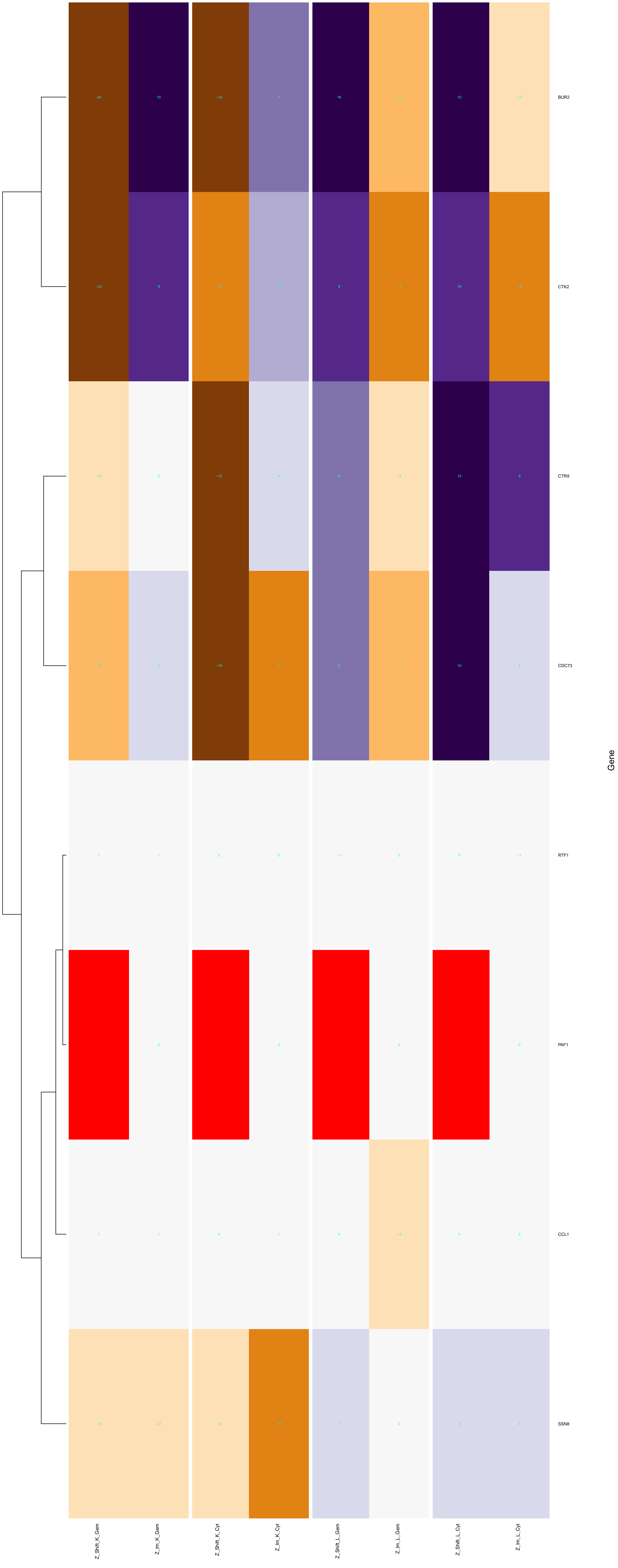


Gene



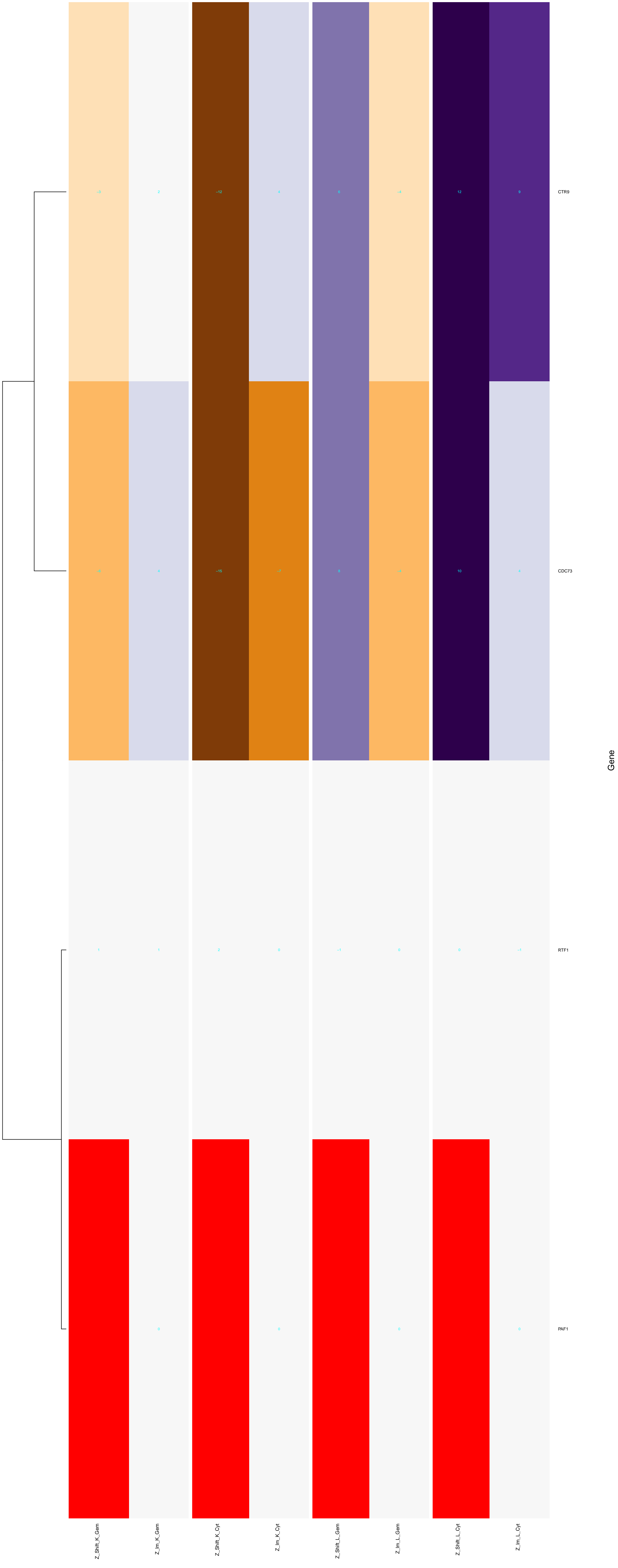


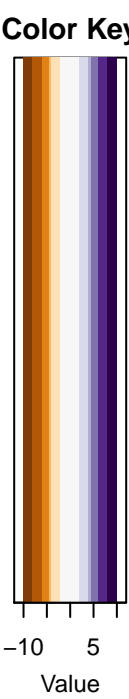
positive regulation of phosphorylation of RNA polymerase II C-terminal domain



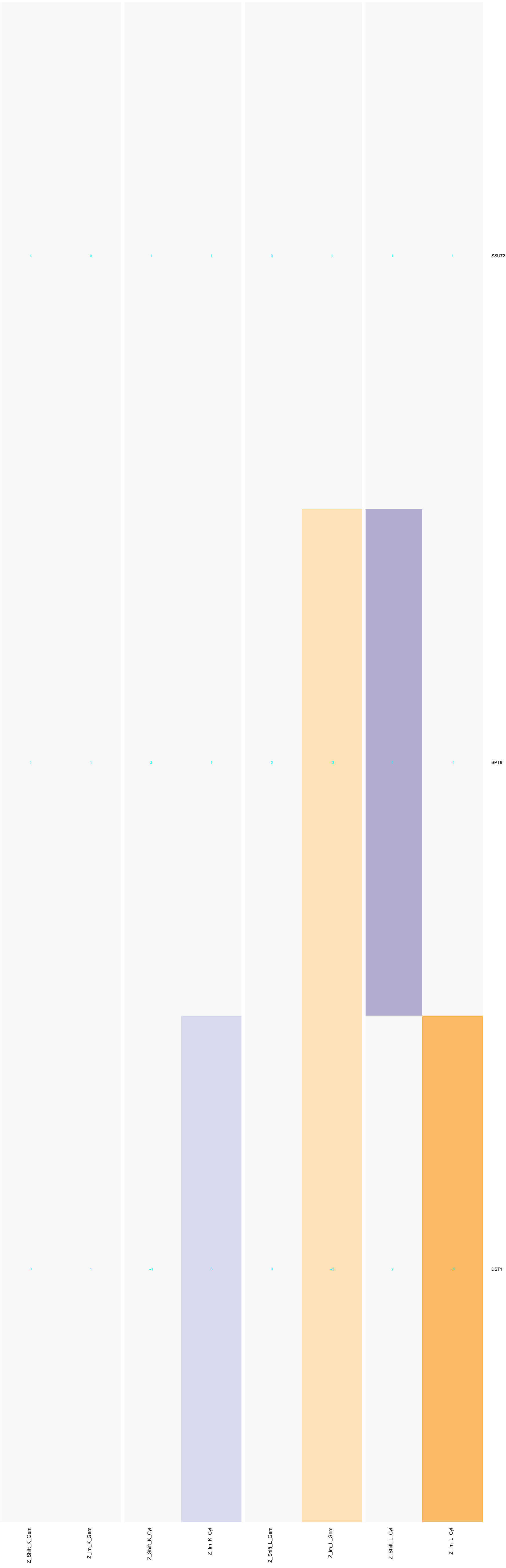


ation of phosphorylation of RNA polymerase II C-terminal domain serine 2 residues

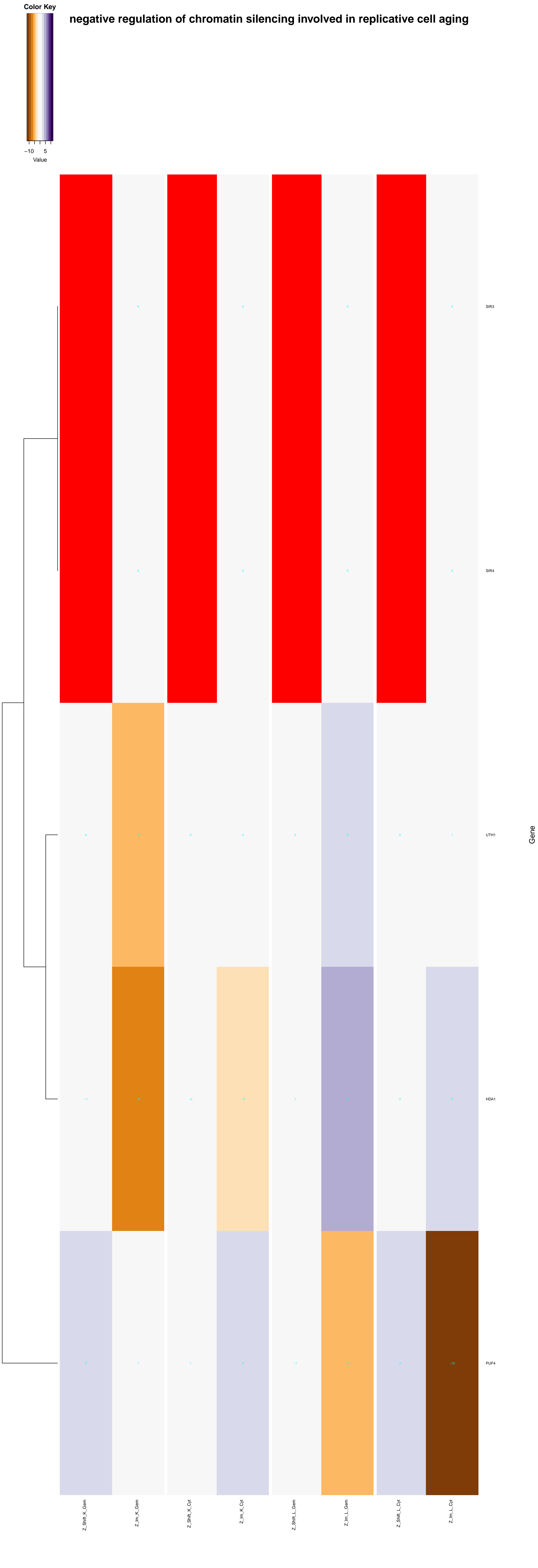


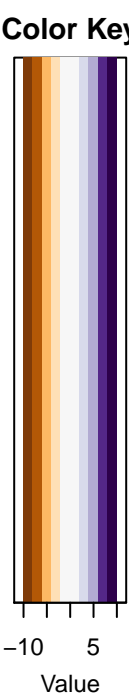


transcription antitermination

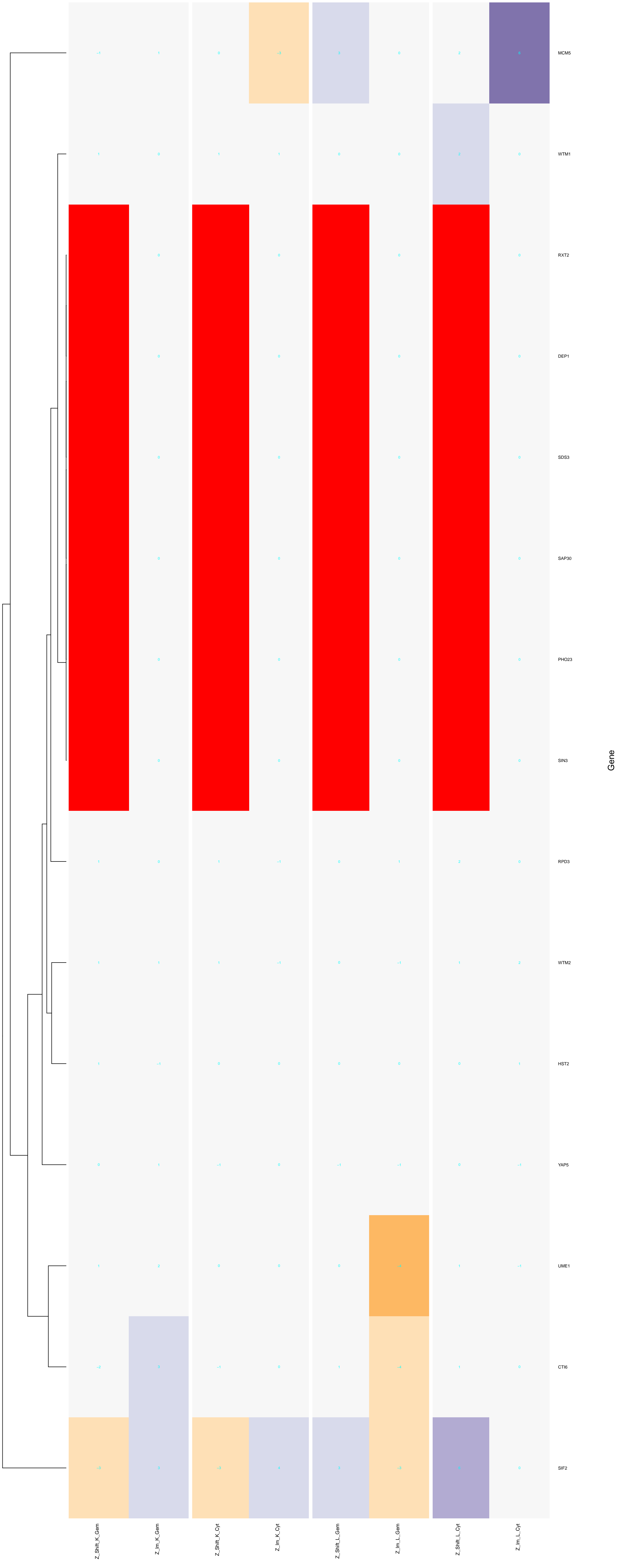


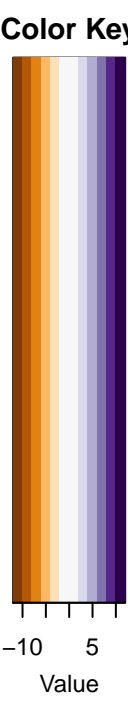
Gene



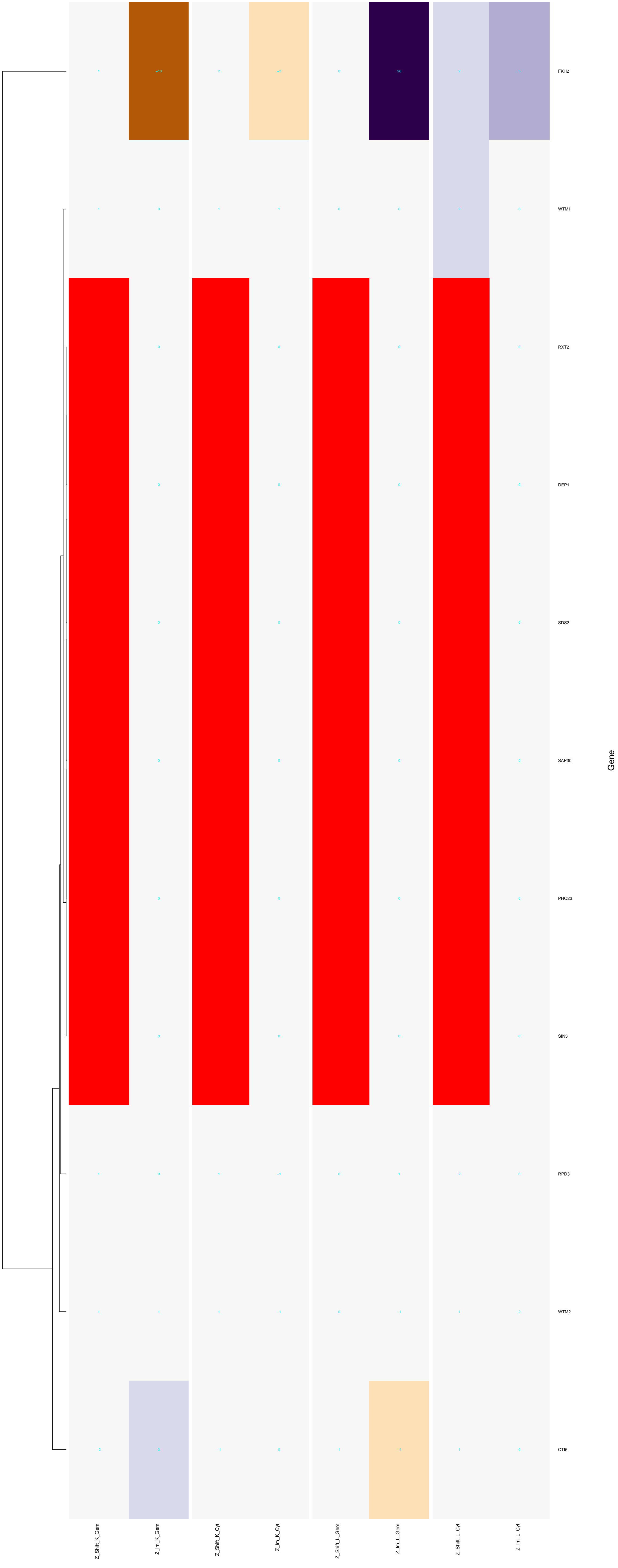


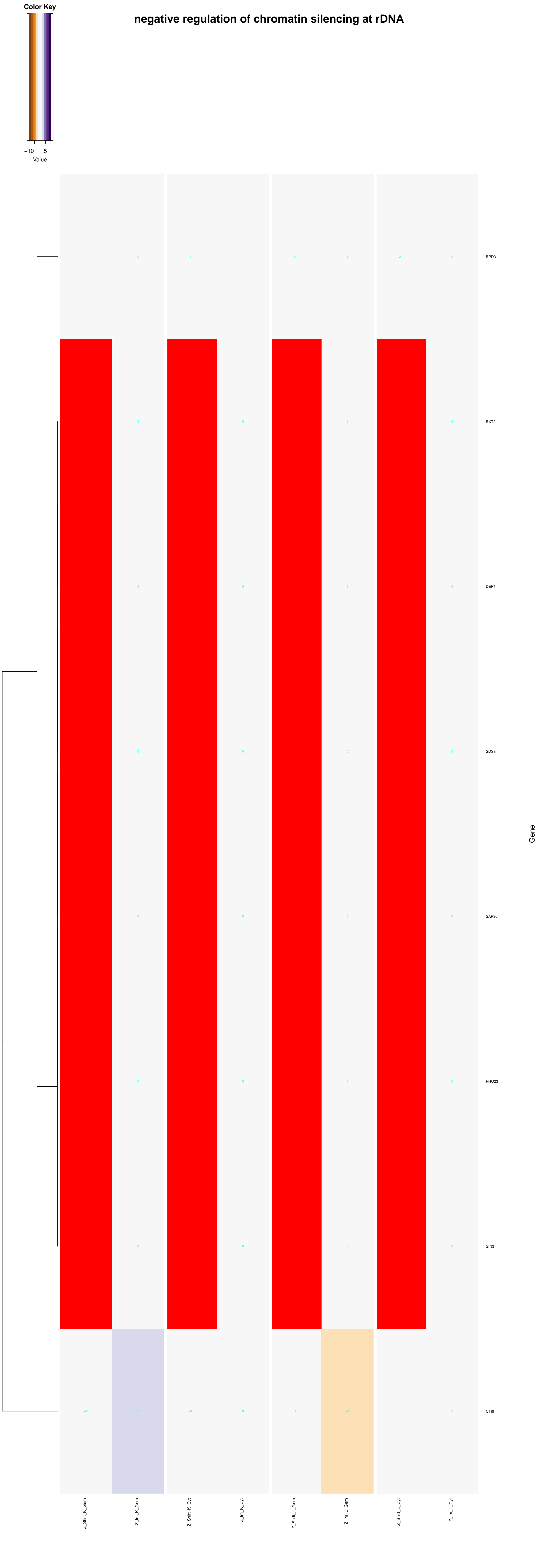
negative regulation of chromatin silencing at telomere

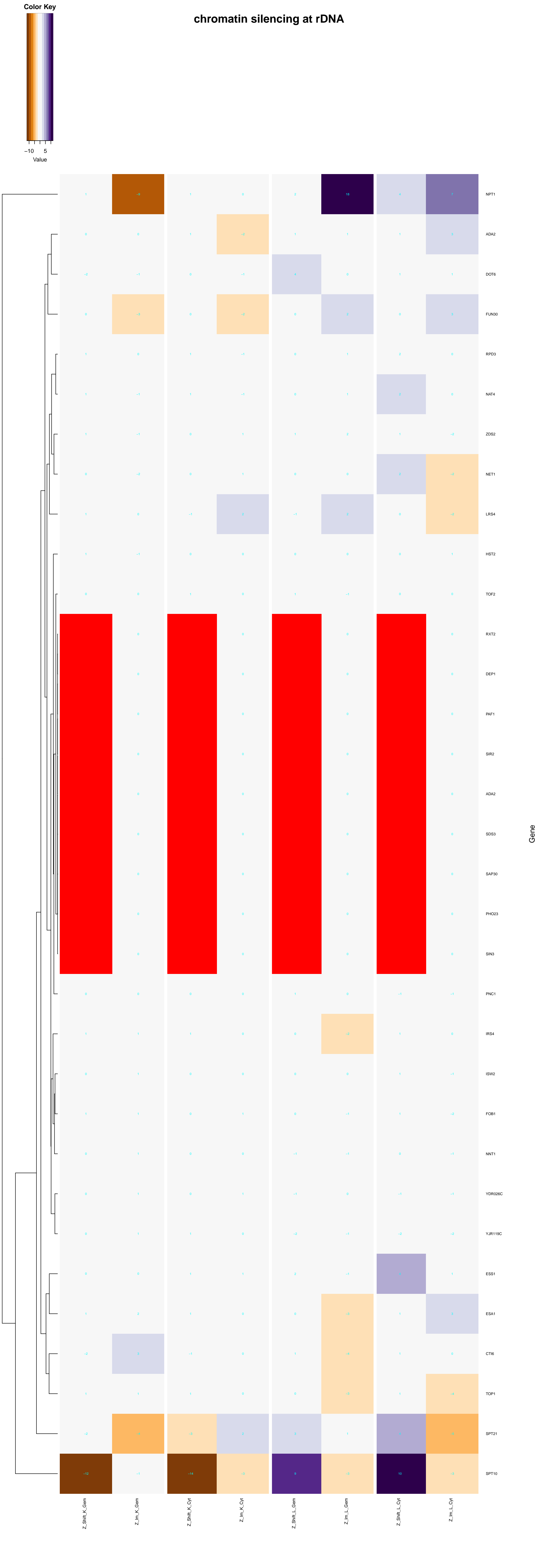


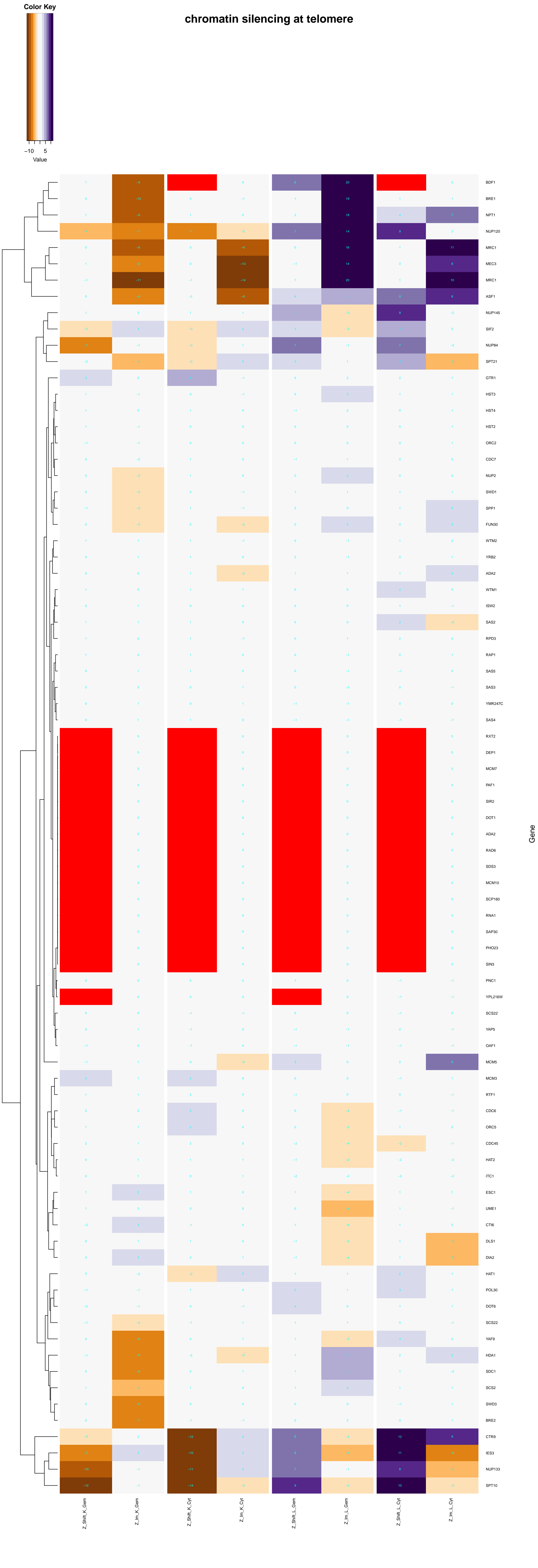


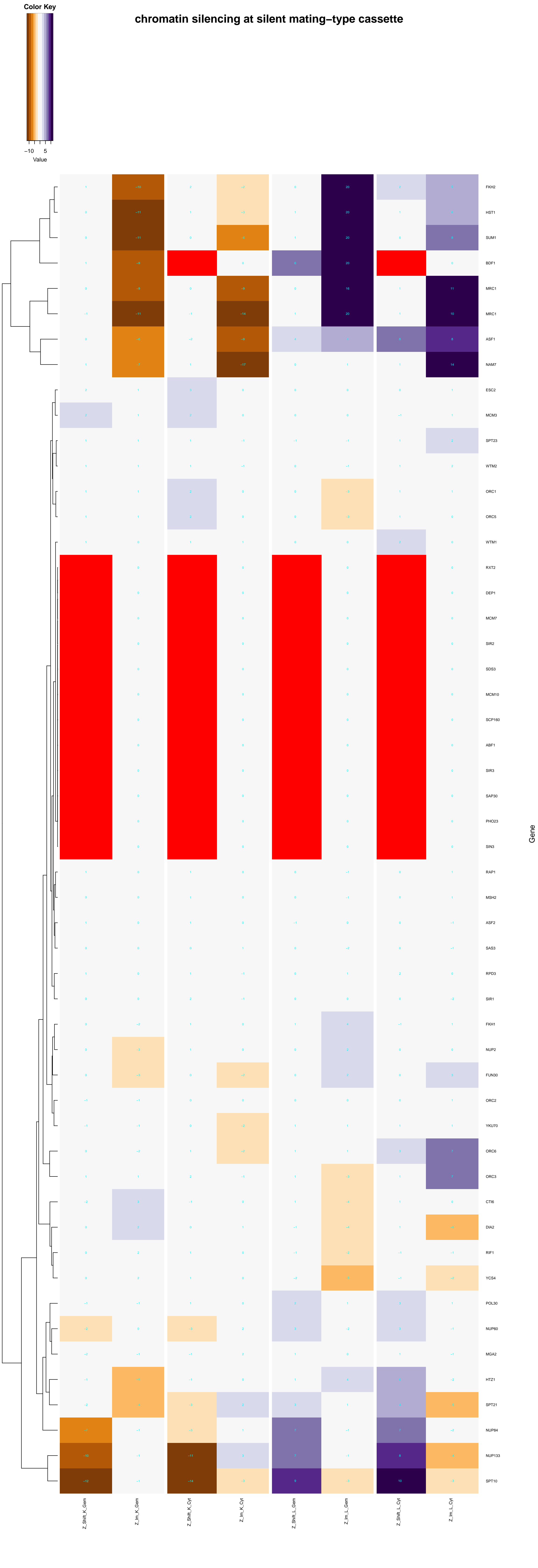
negative regulation of chromatin silencing at silent mating-type cassette

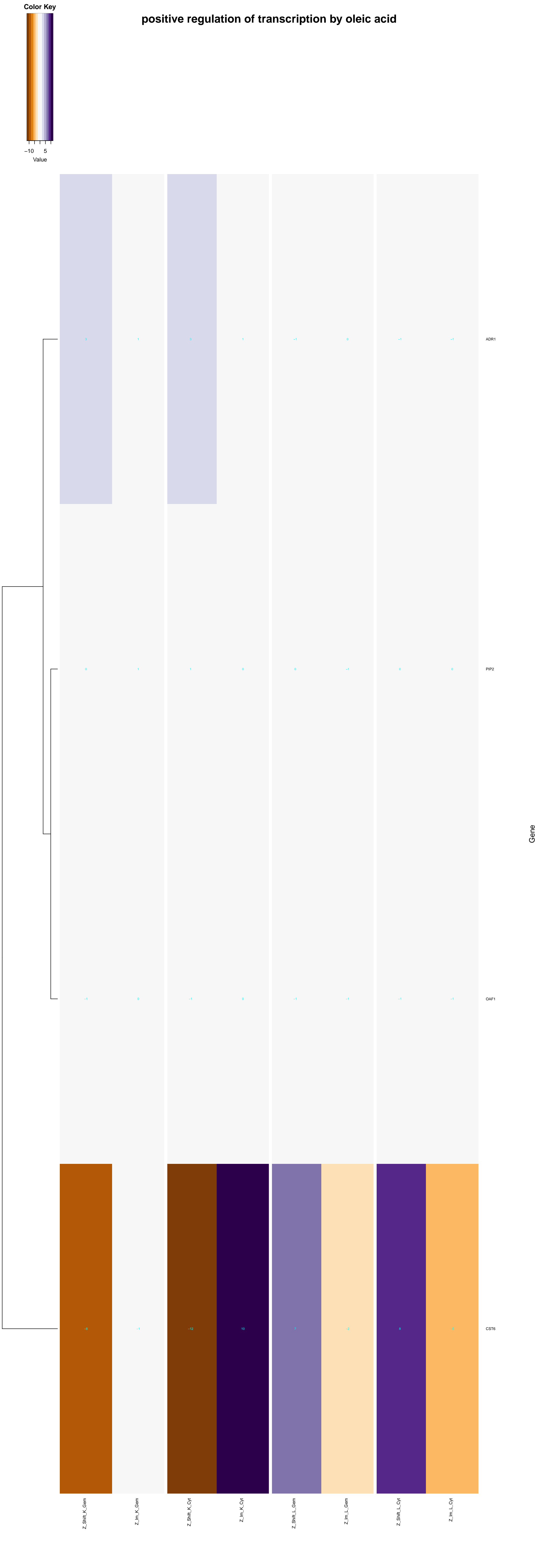


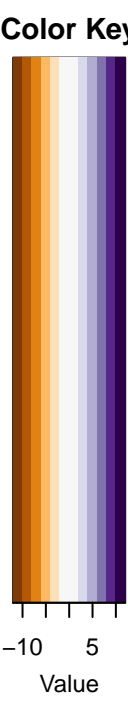




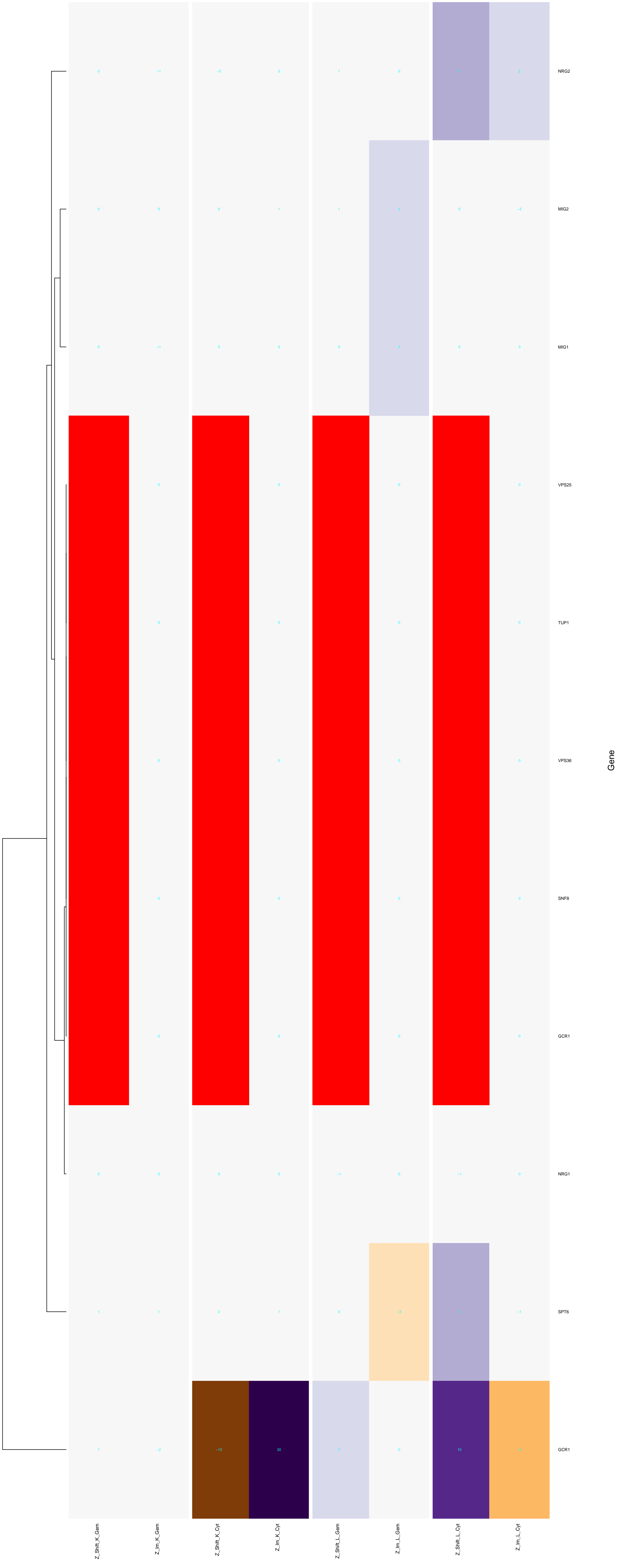


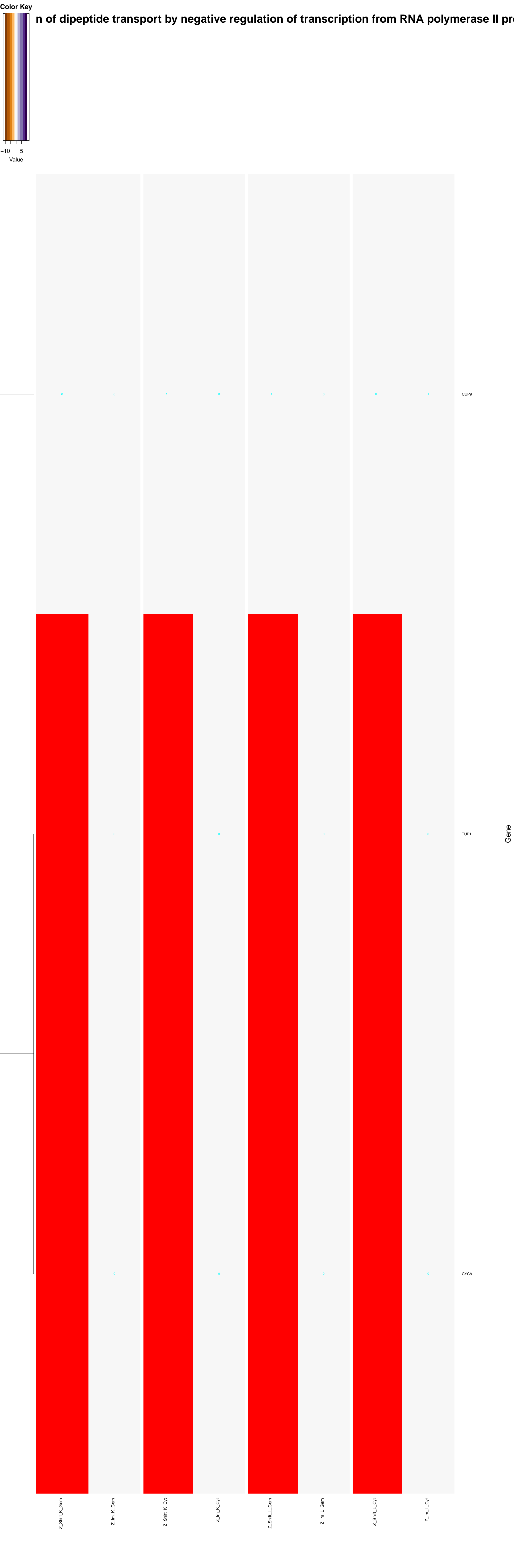


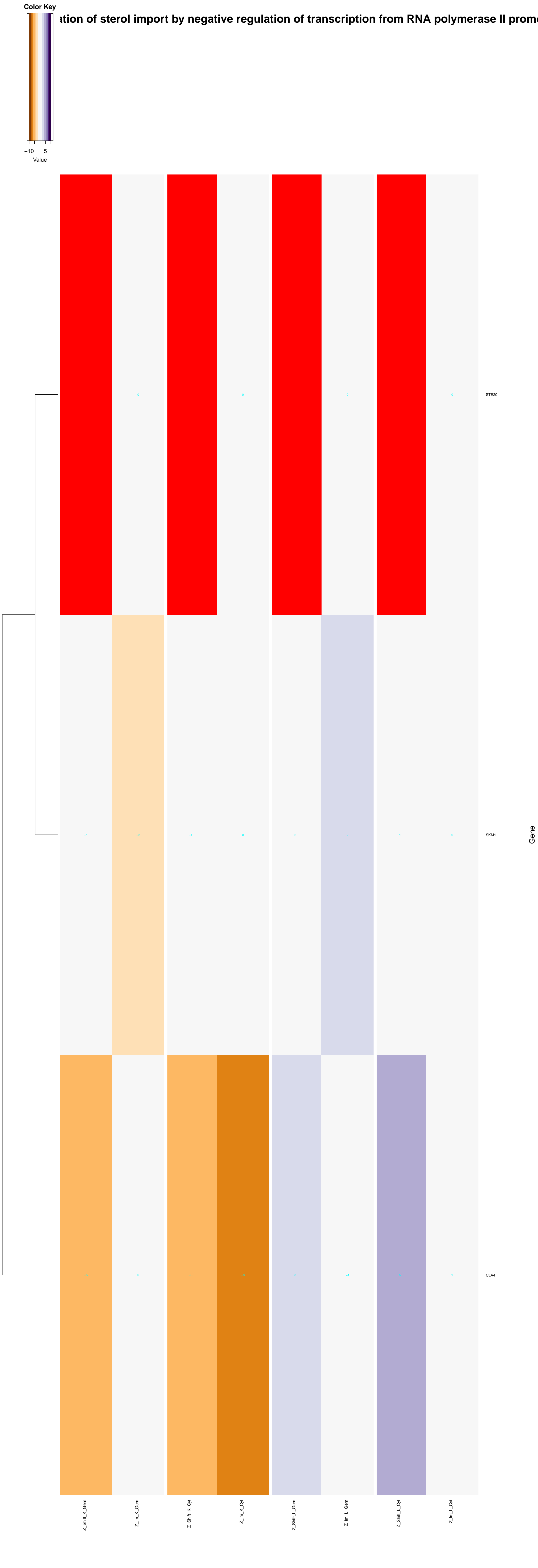


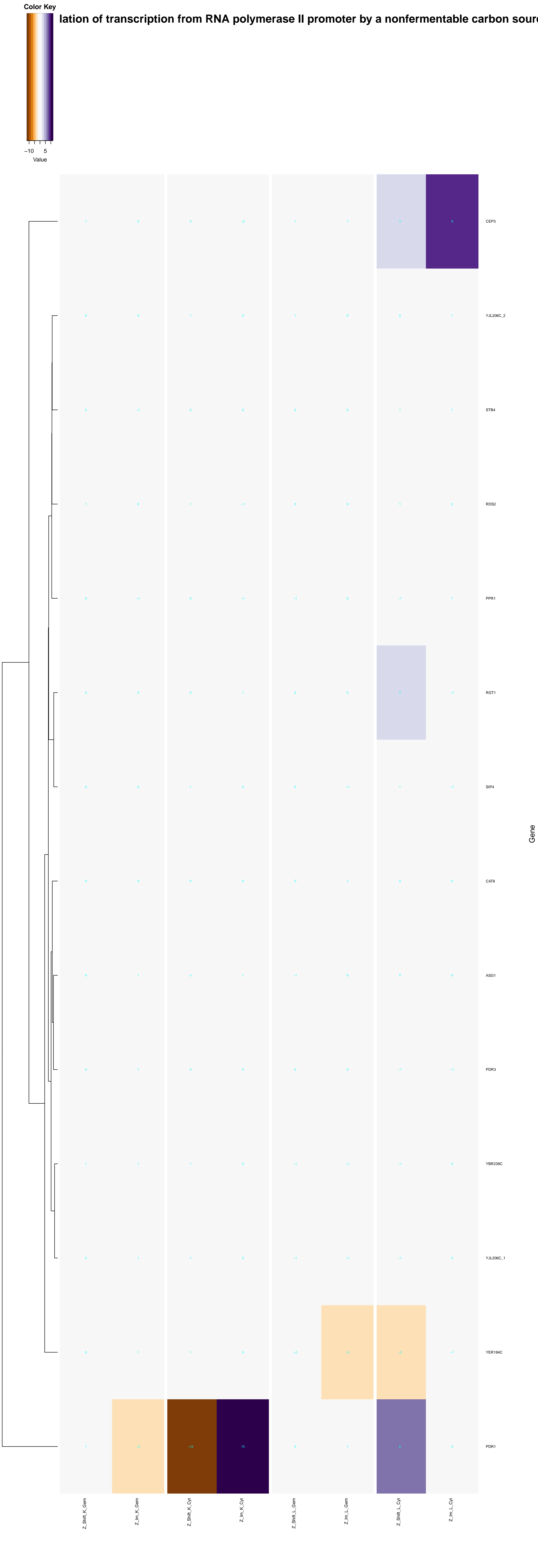


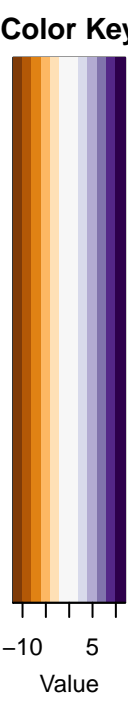
egative regulation of transcription from RNA polymerase II promoter by glucose



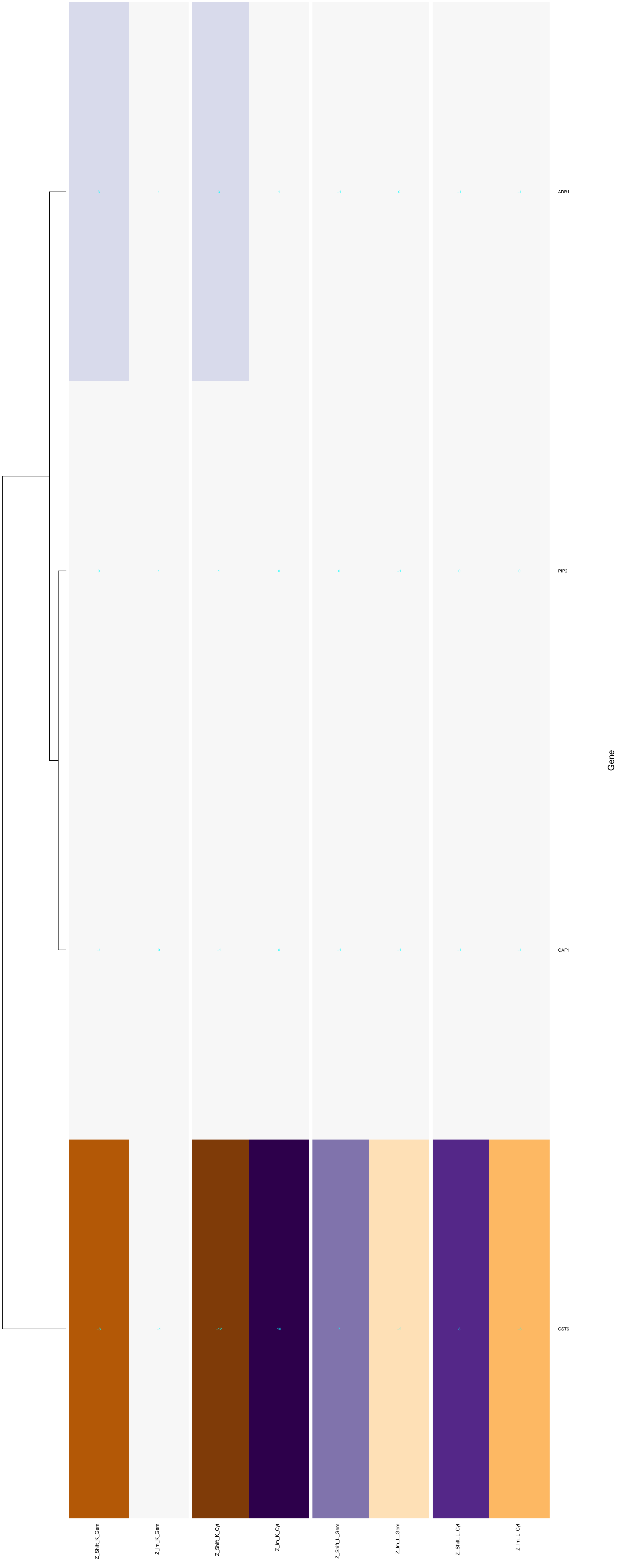


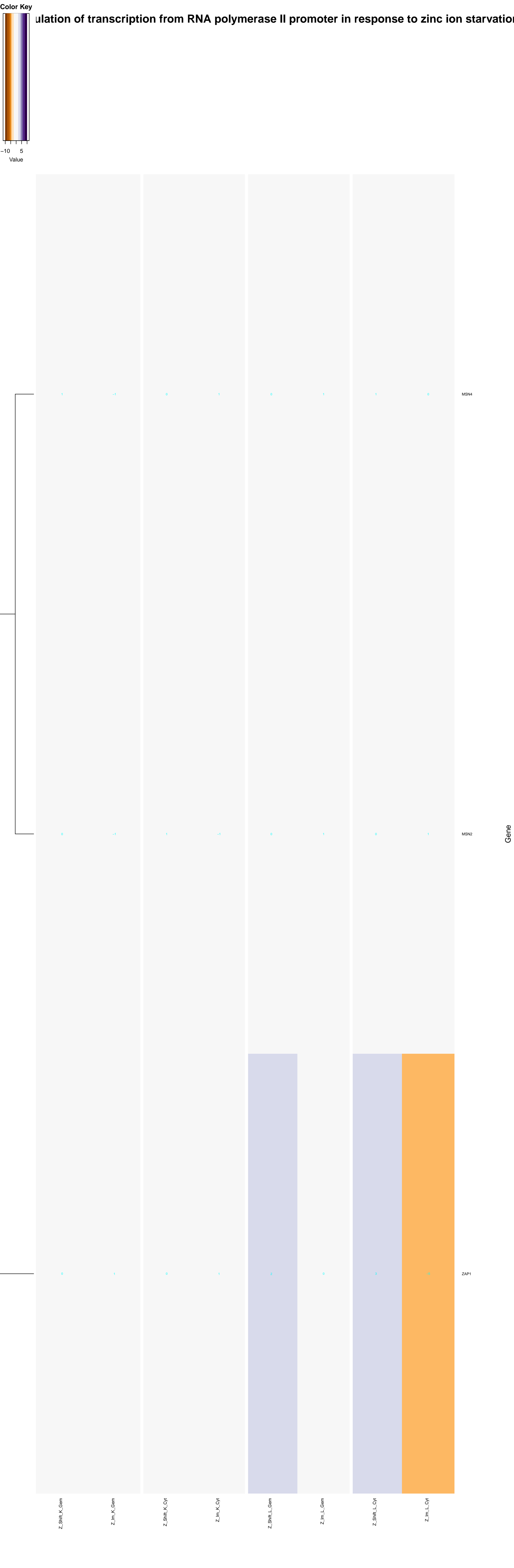


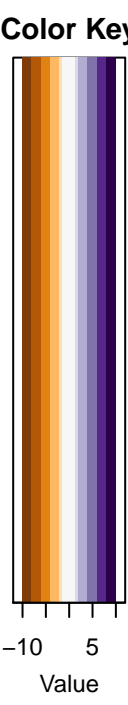




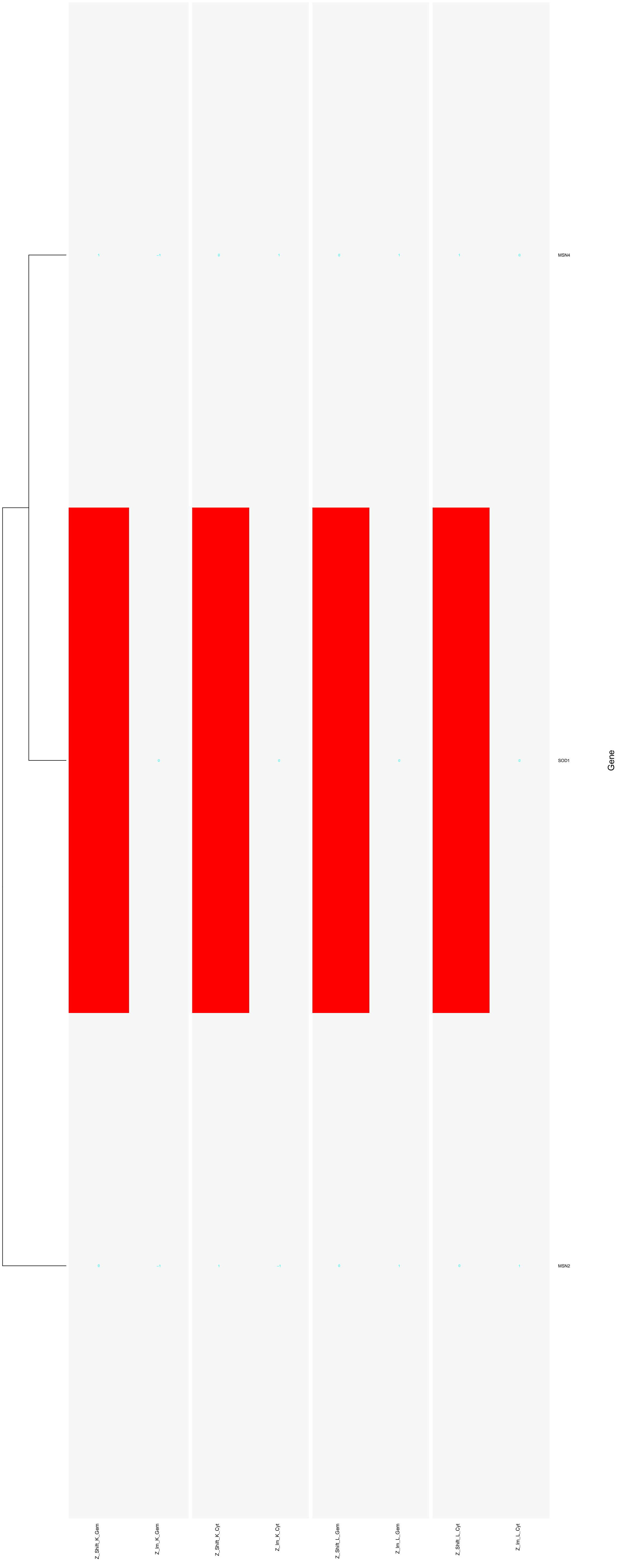
Positive regulation of transcription from RNA polymerase II promoter by oleic acid

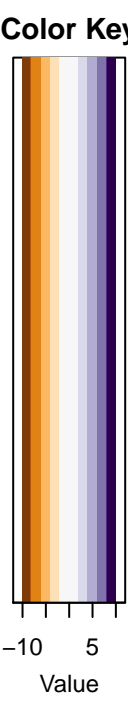




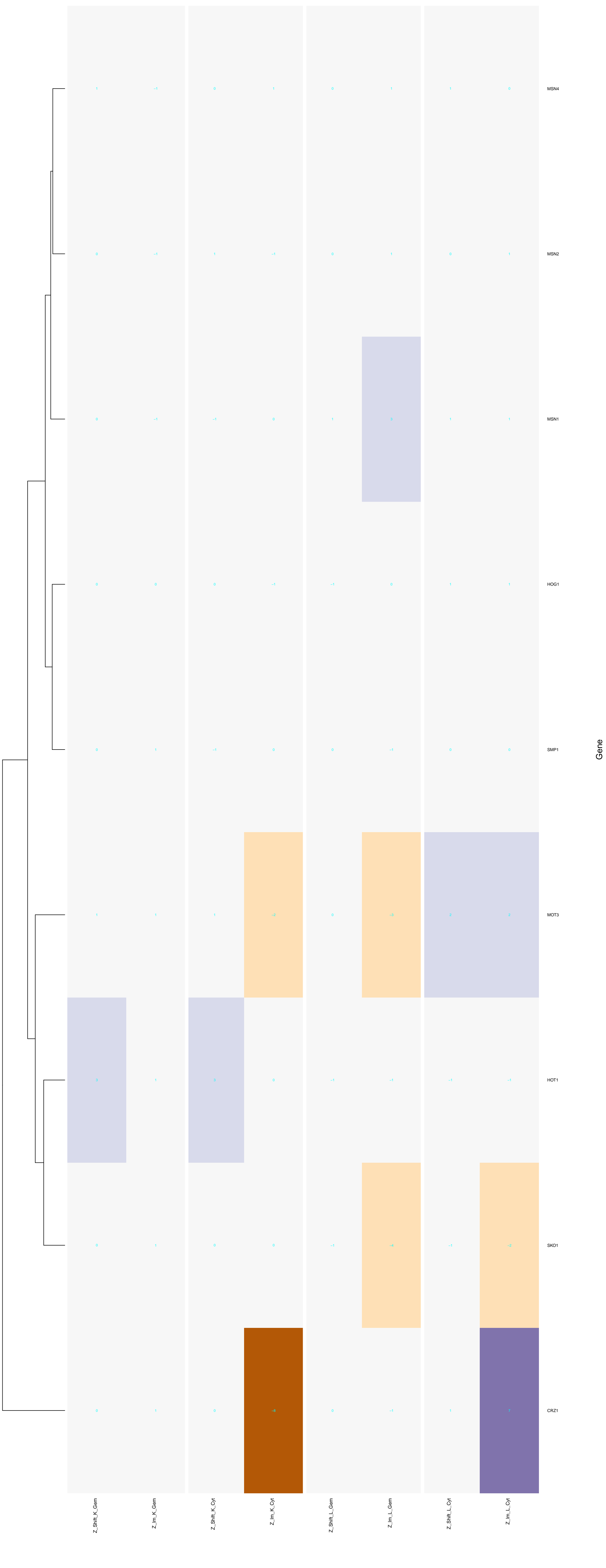


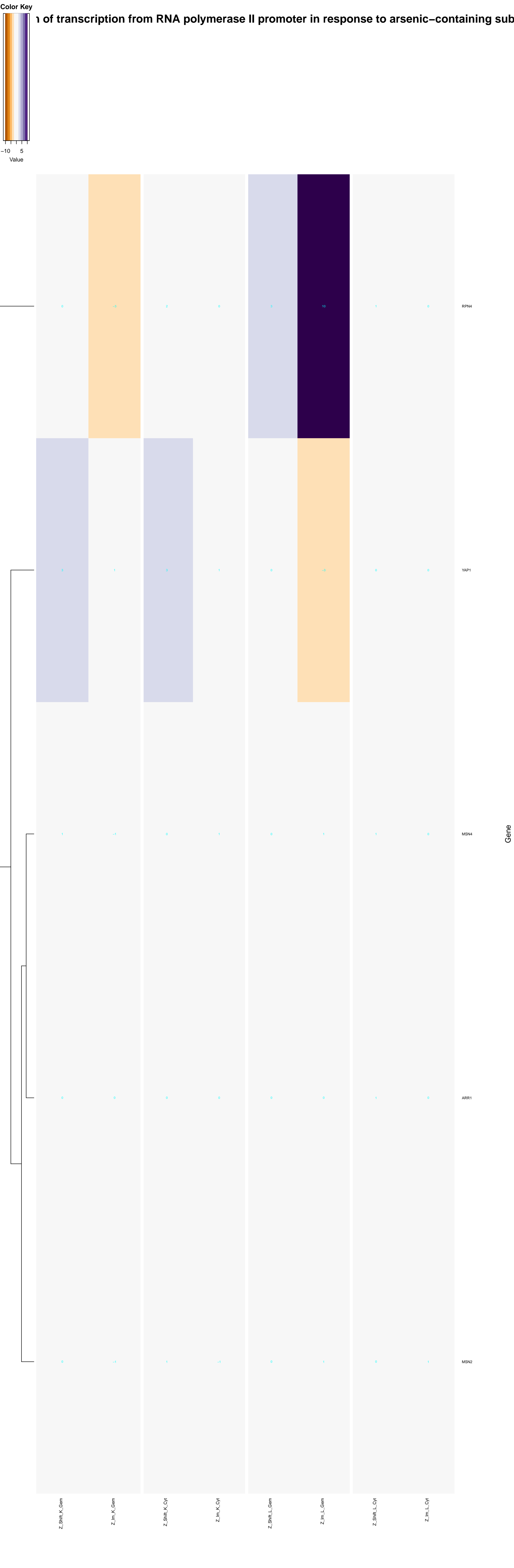
gulation of transcription from RNA polymerase II promoter in response to oxidative stress

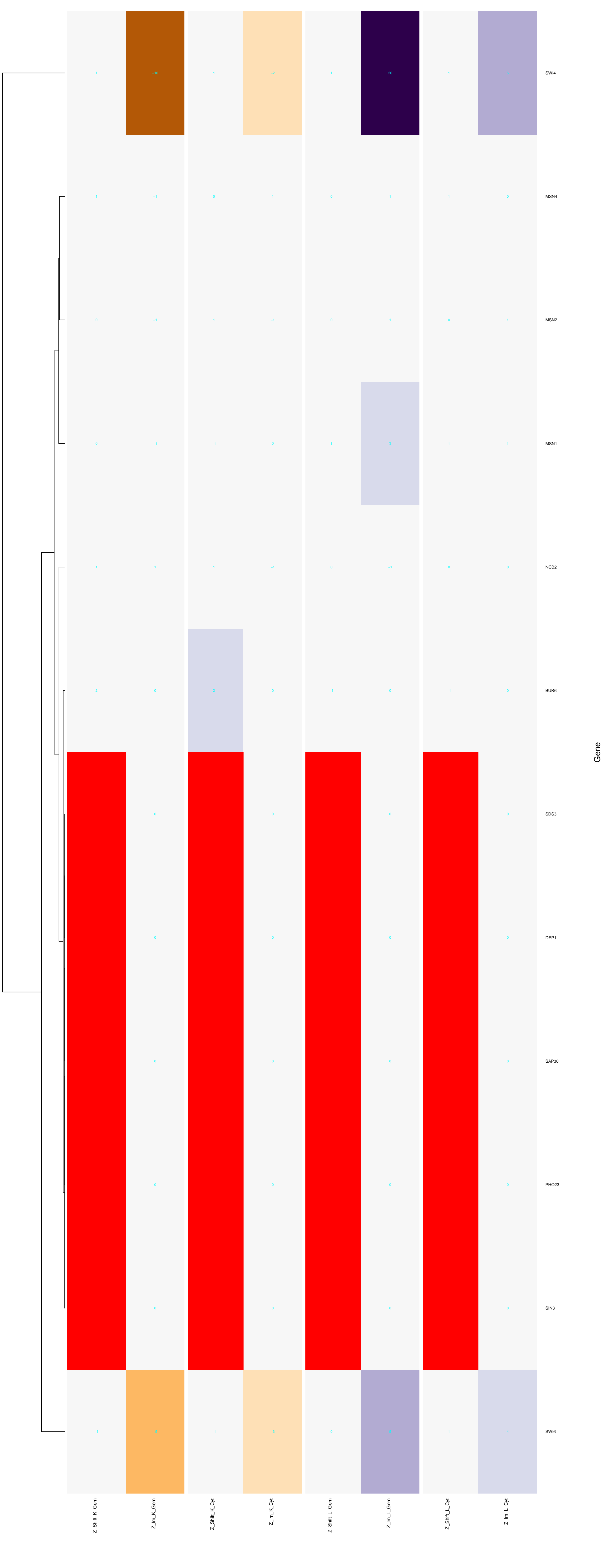
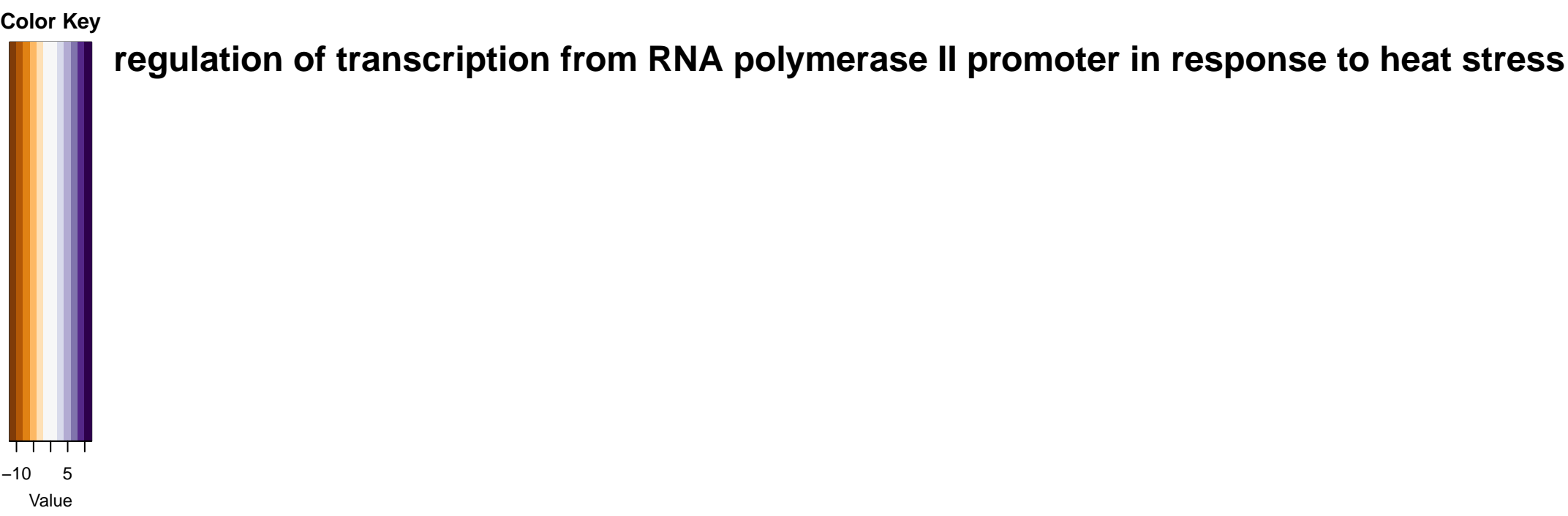




gulation of transcription from RNA polymerase II promoter in response to osmotic stress





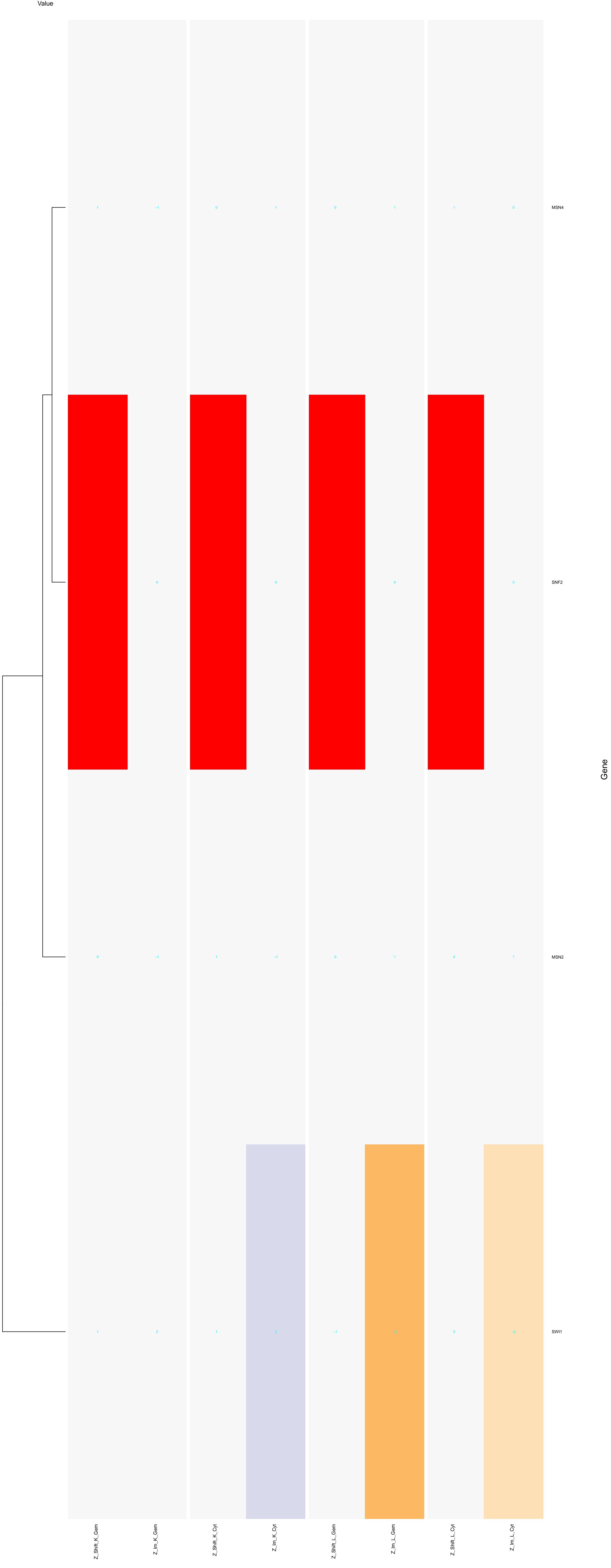


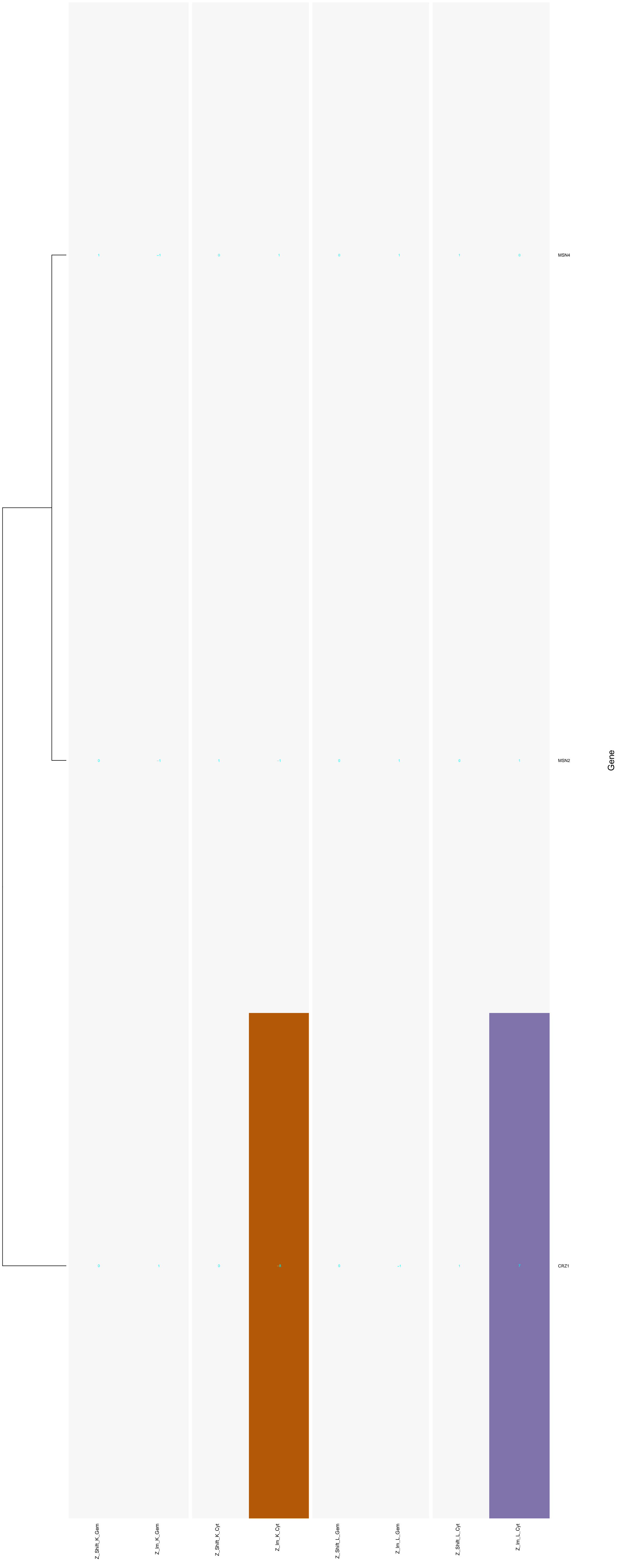
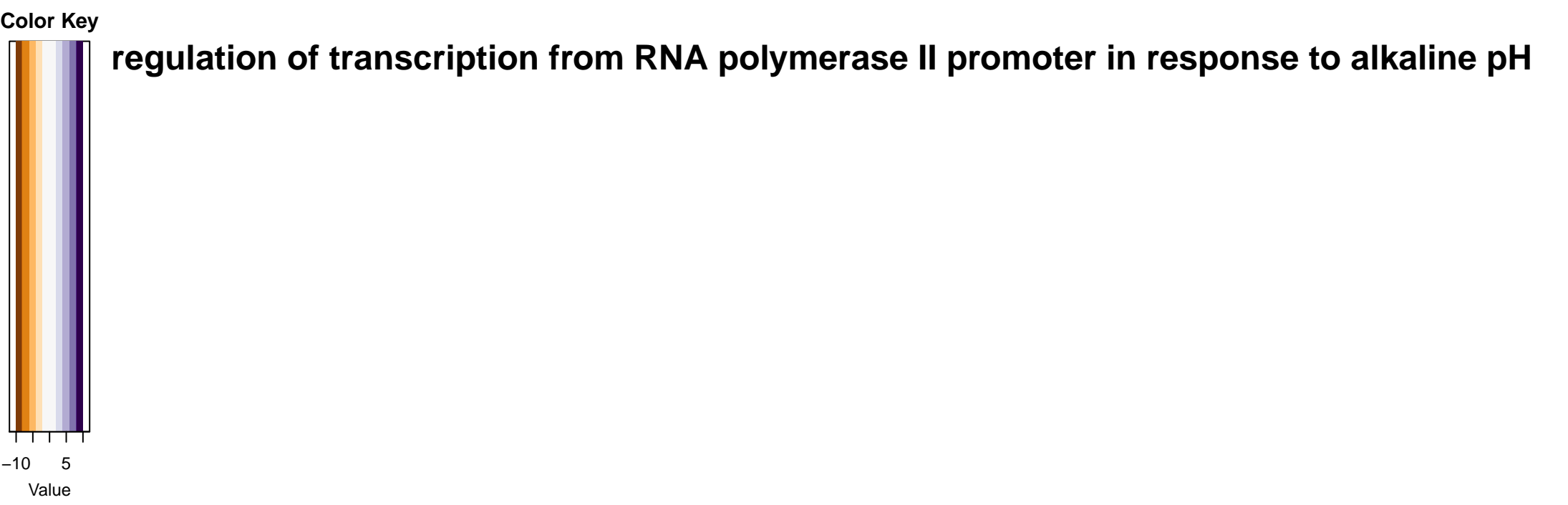
Color Key

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

-10 5

Value



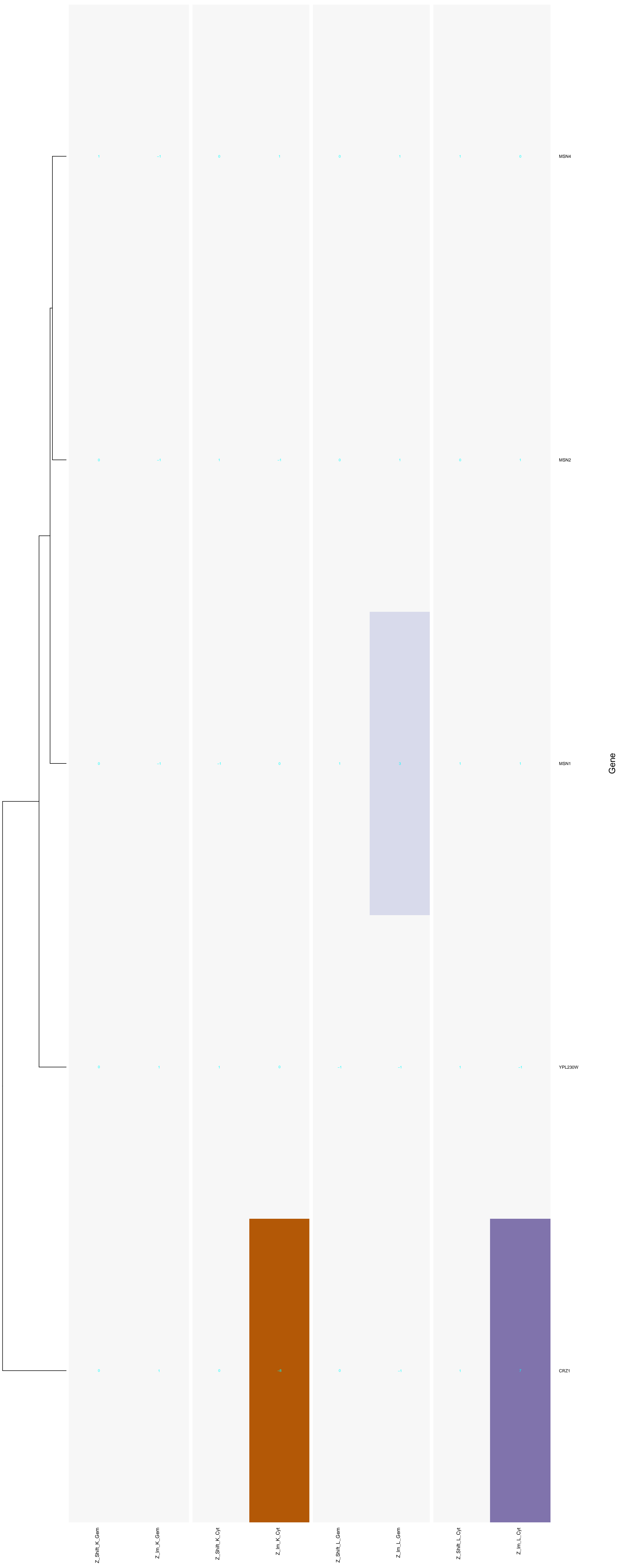


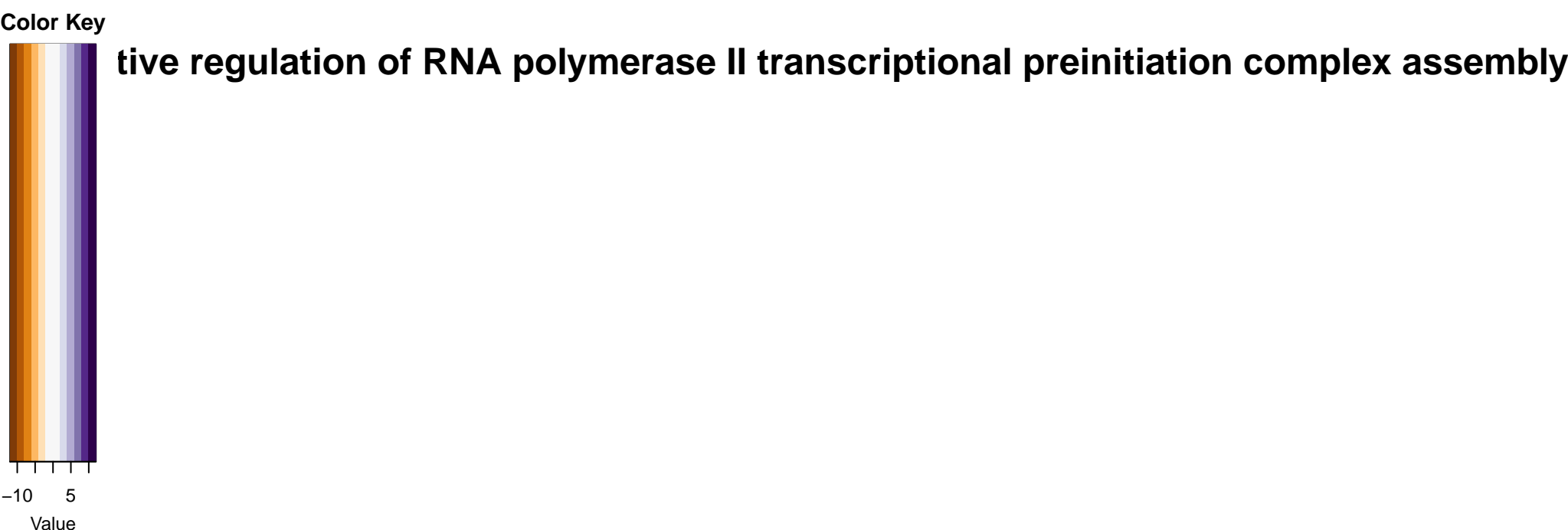
Color Key

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

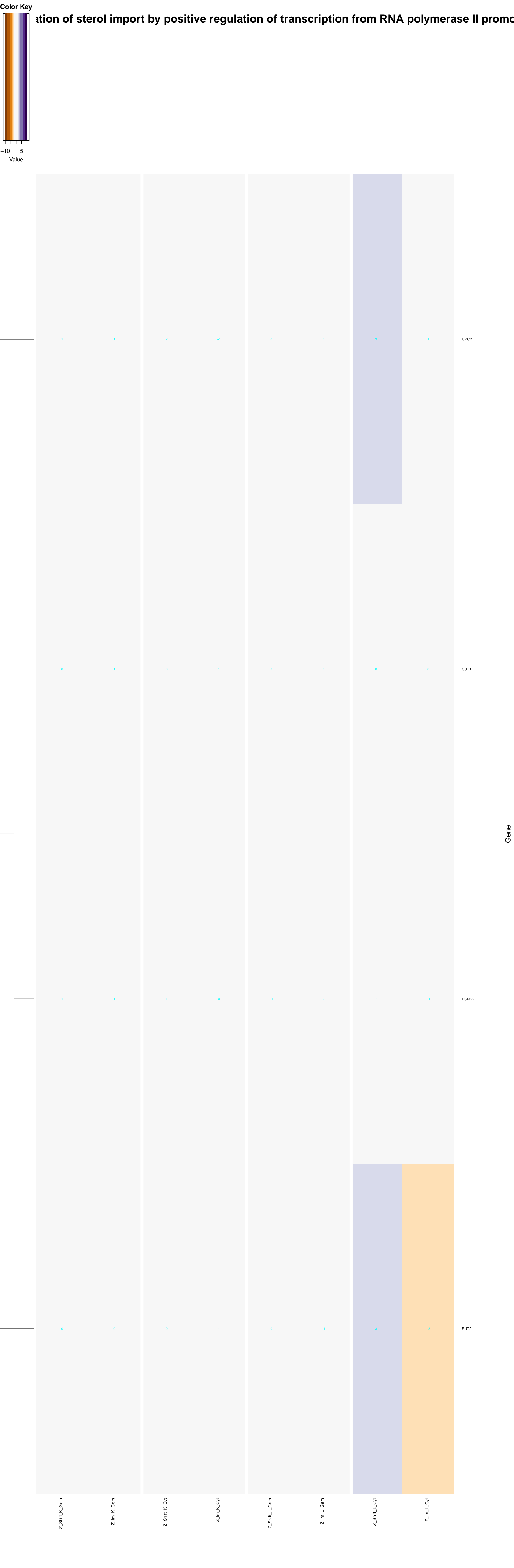
-10 5

Value





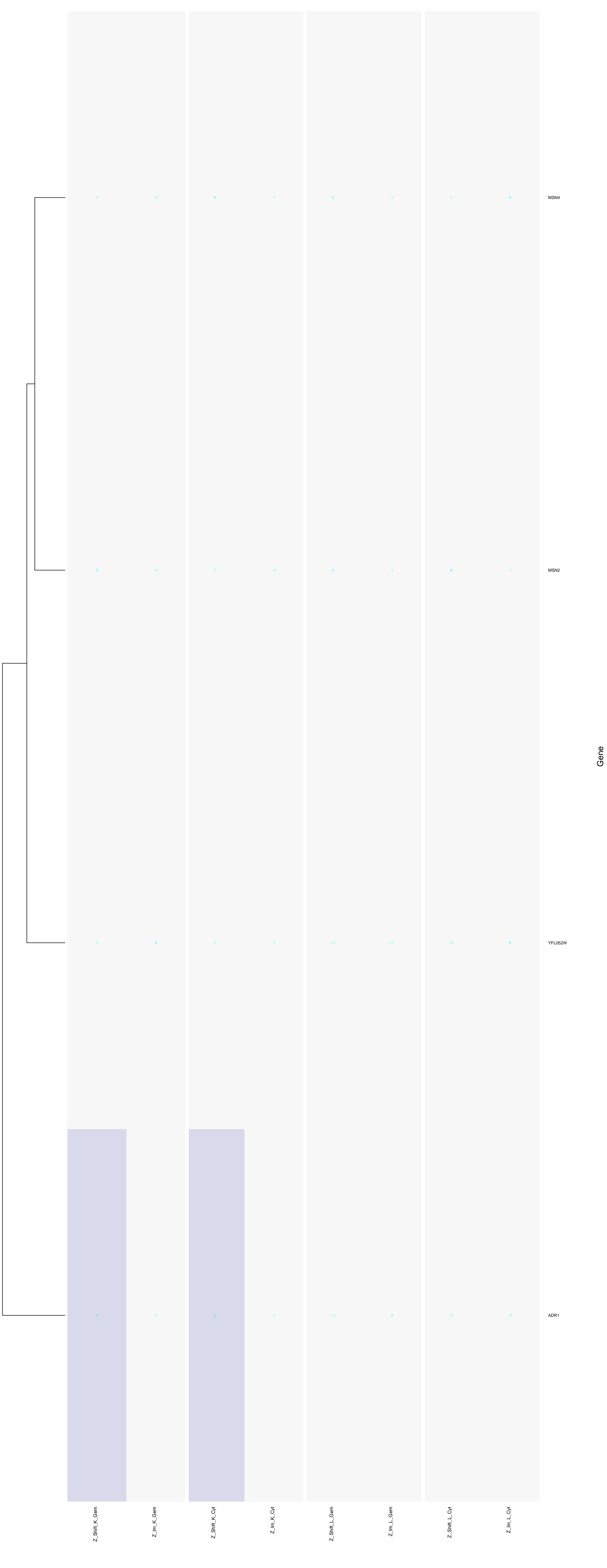
Gene

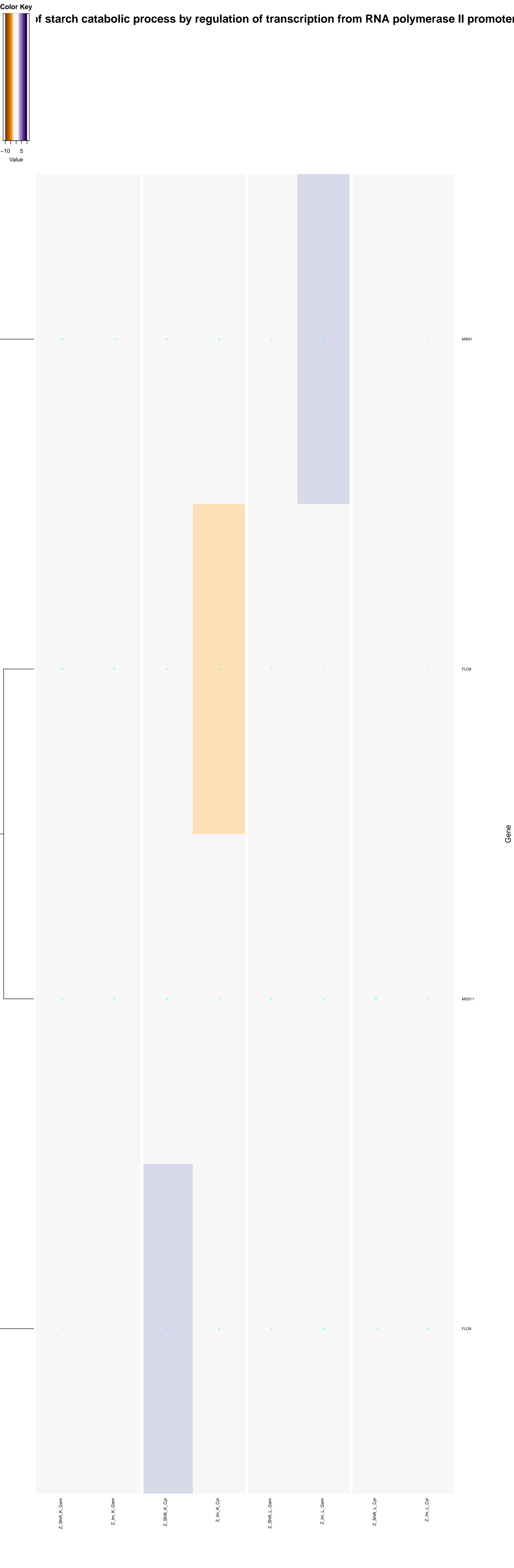


Color Key

Value

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



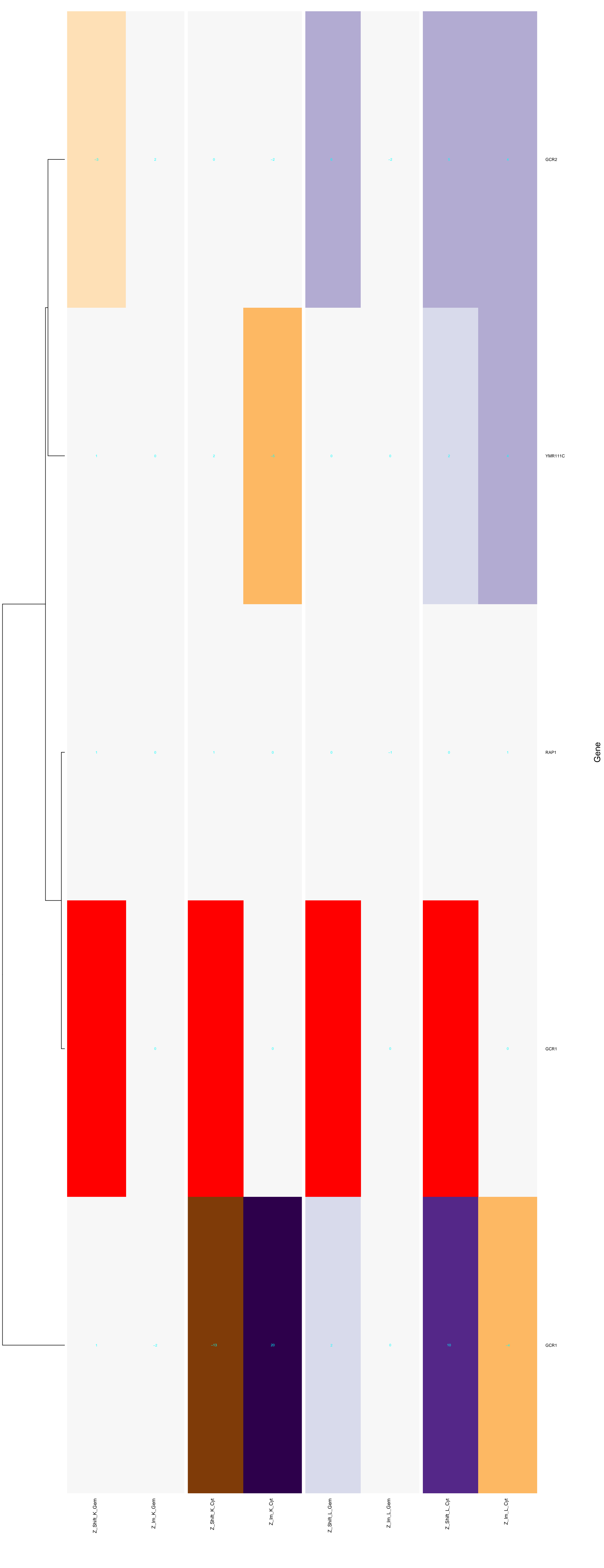


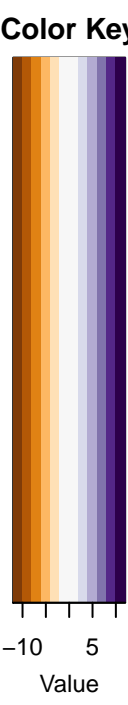
Color Key

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

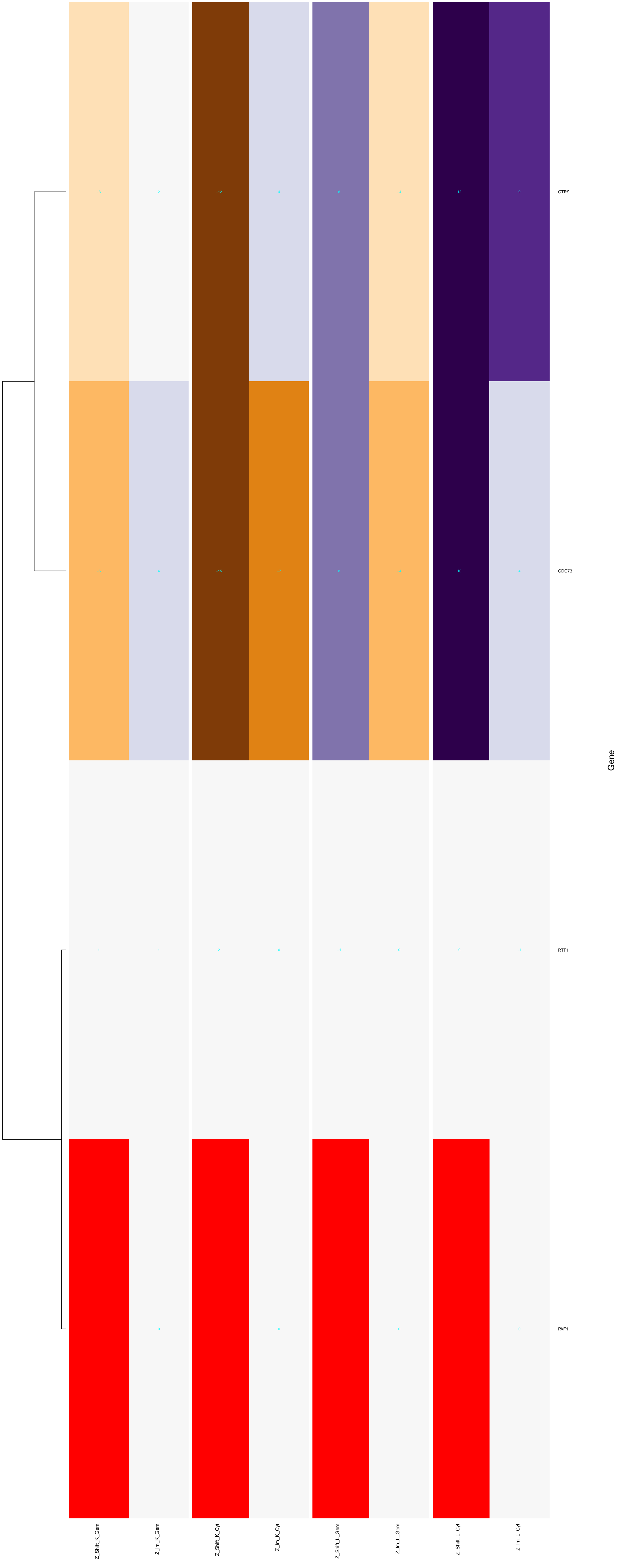
-10 5

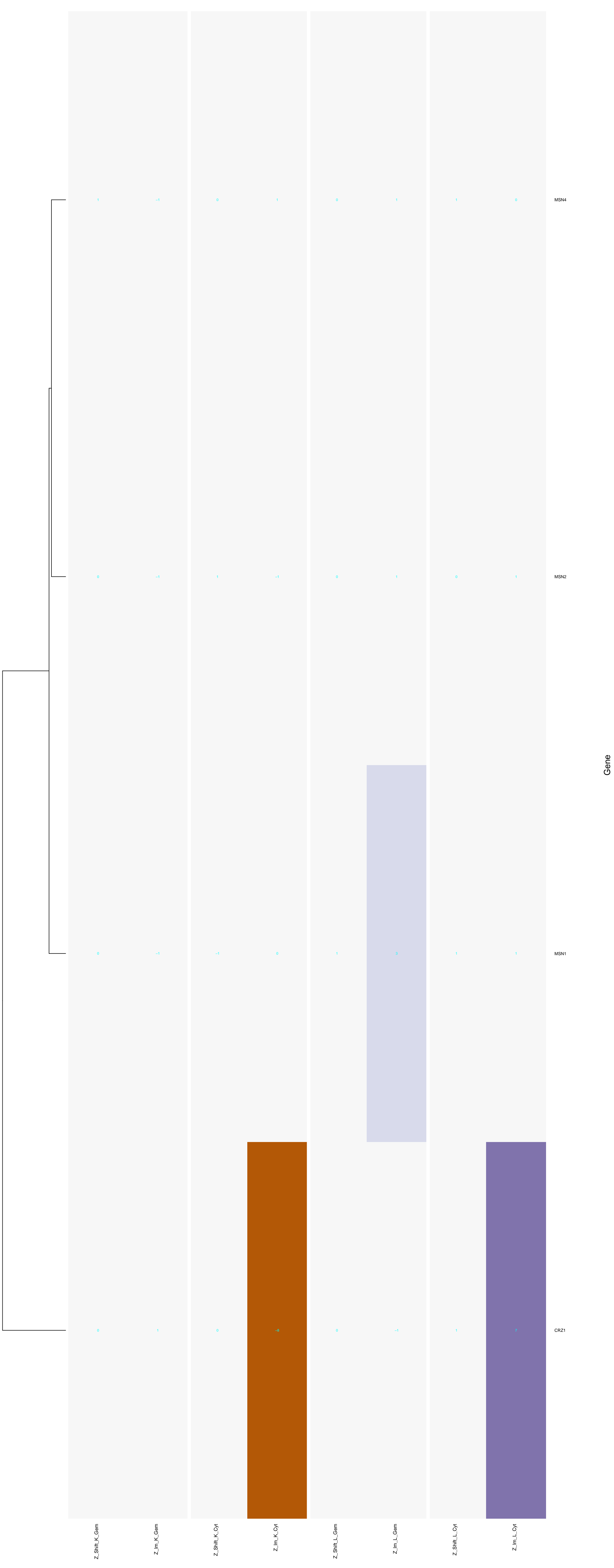
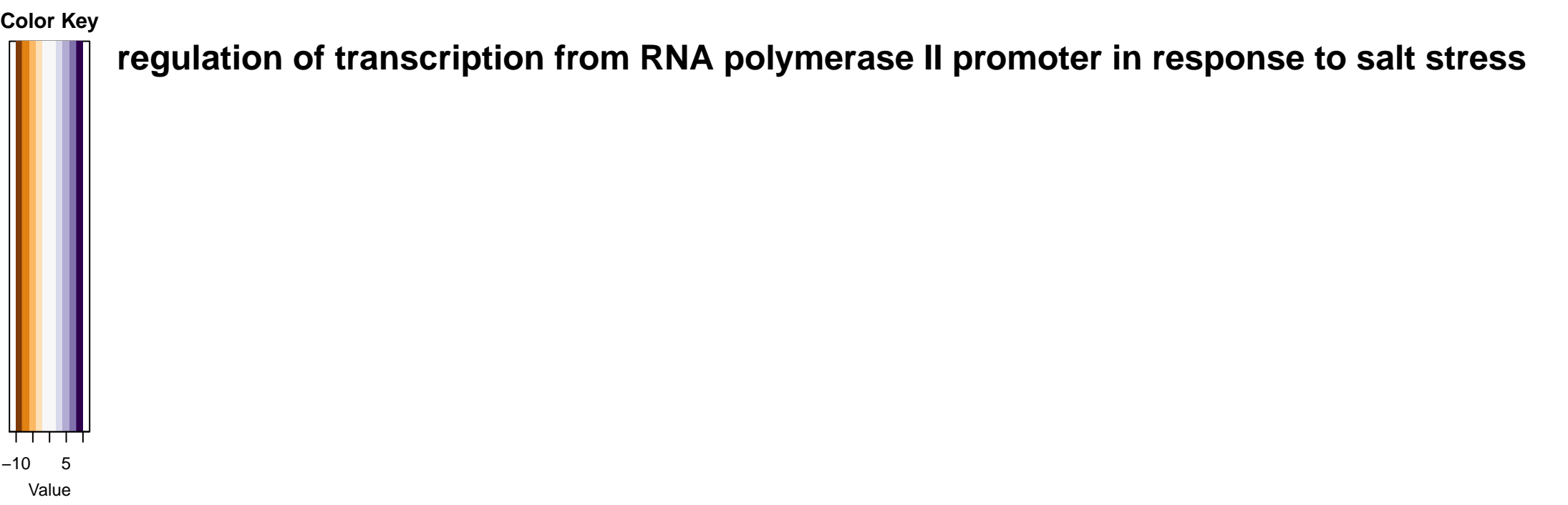
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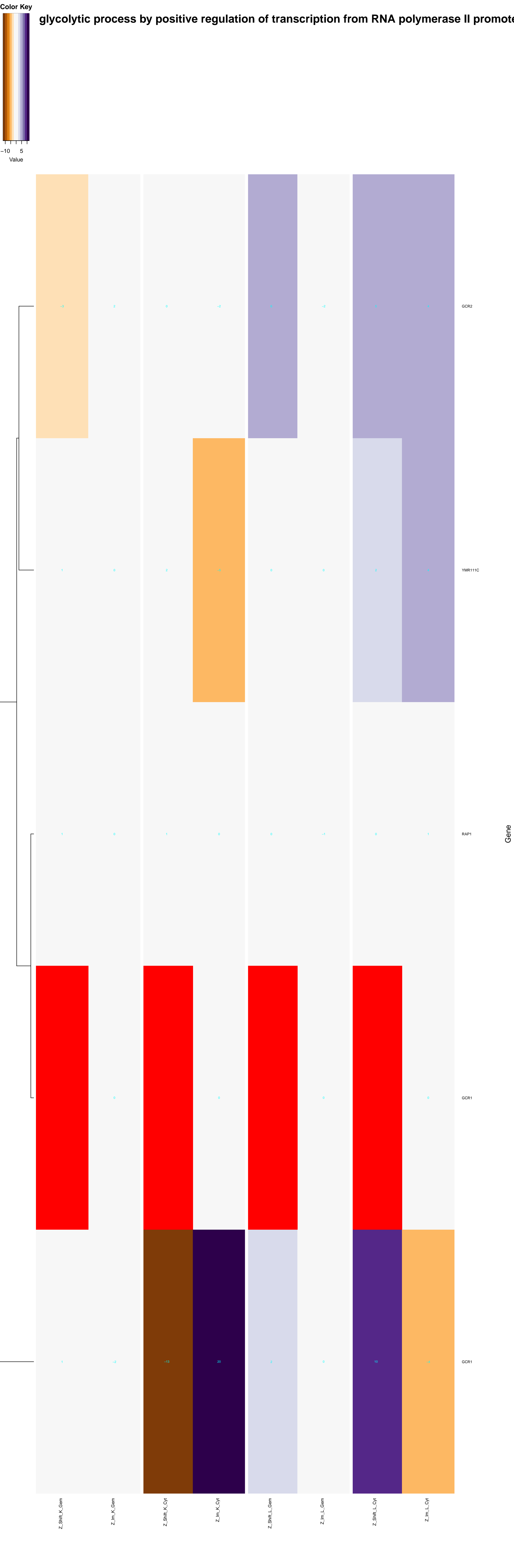


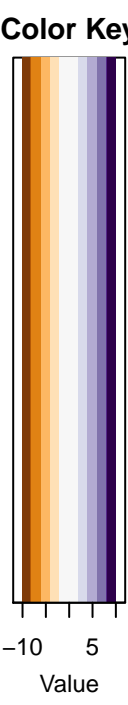
egulation of phosphorylation of RNA polymerase II C-terminal domain serine 2 residues











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

