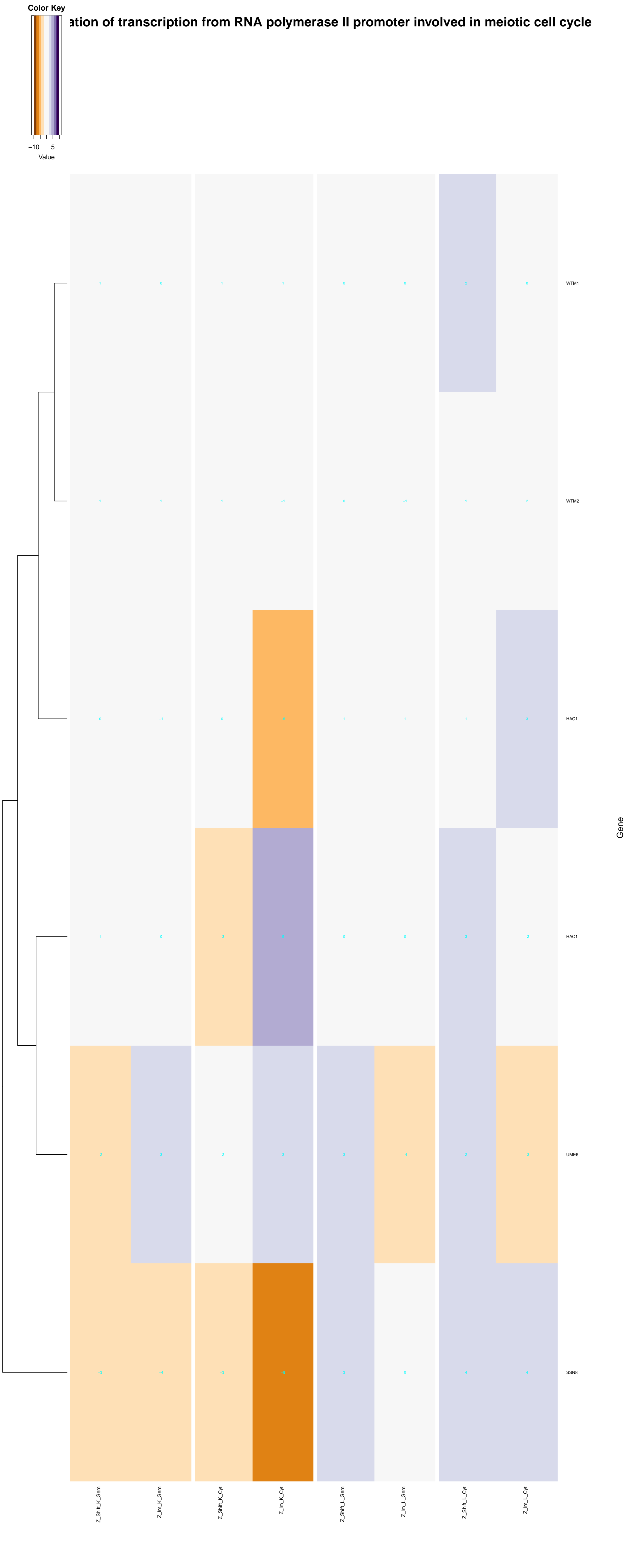
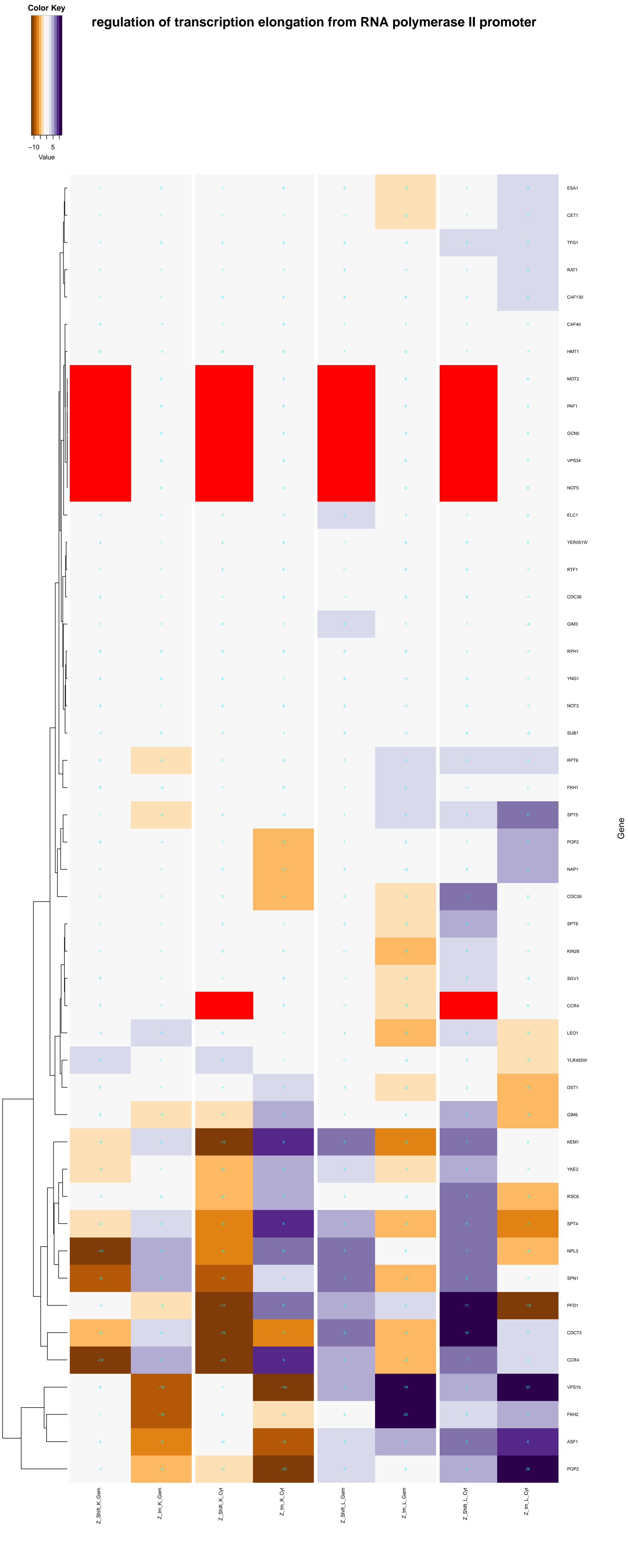
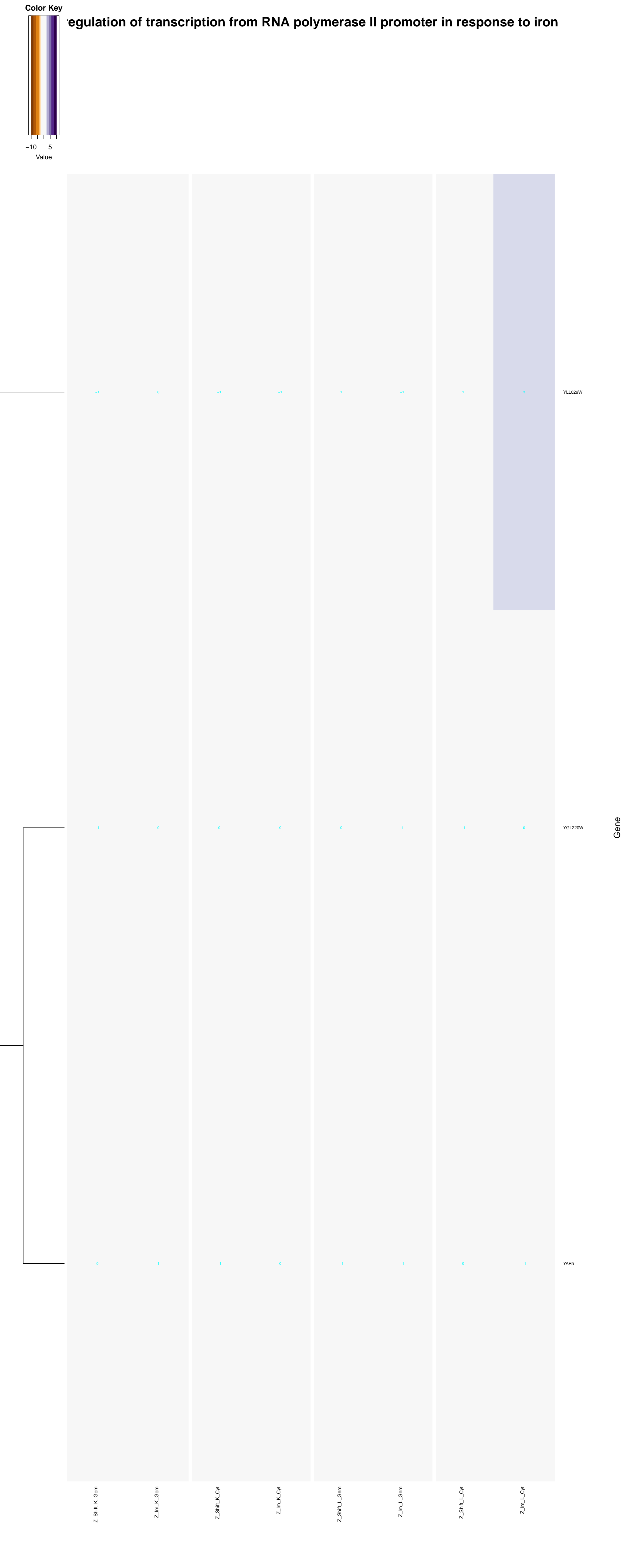
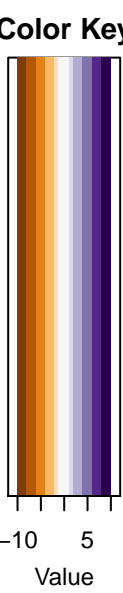


ation of transcription from RNA polymerase II promoter involved in meiotic cell cycle

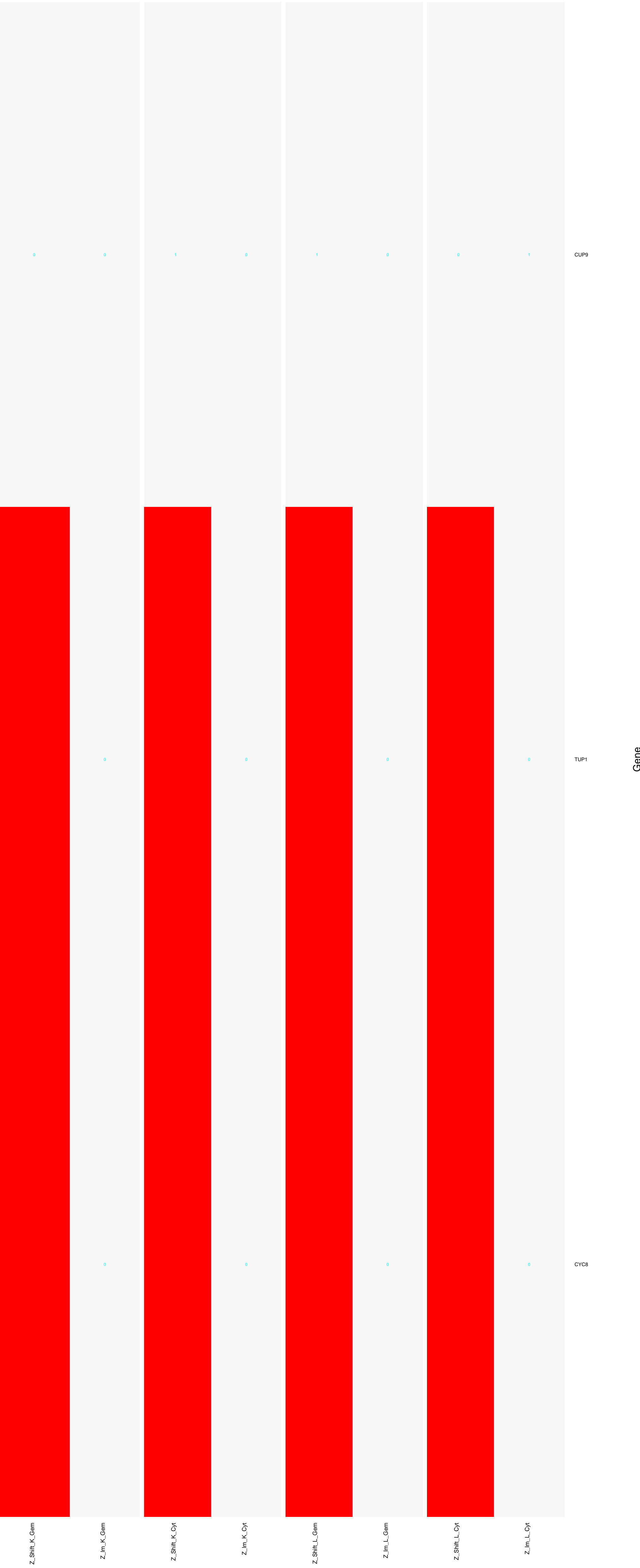




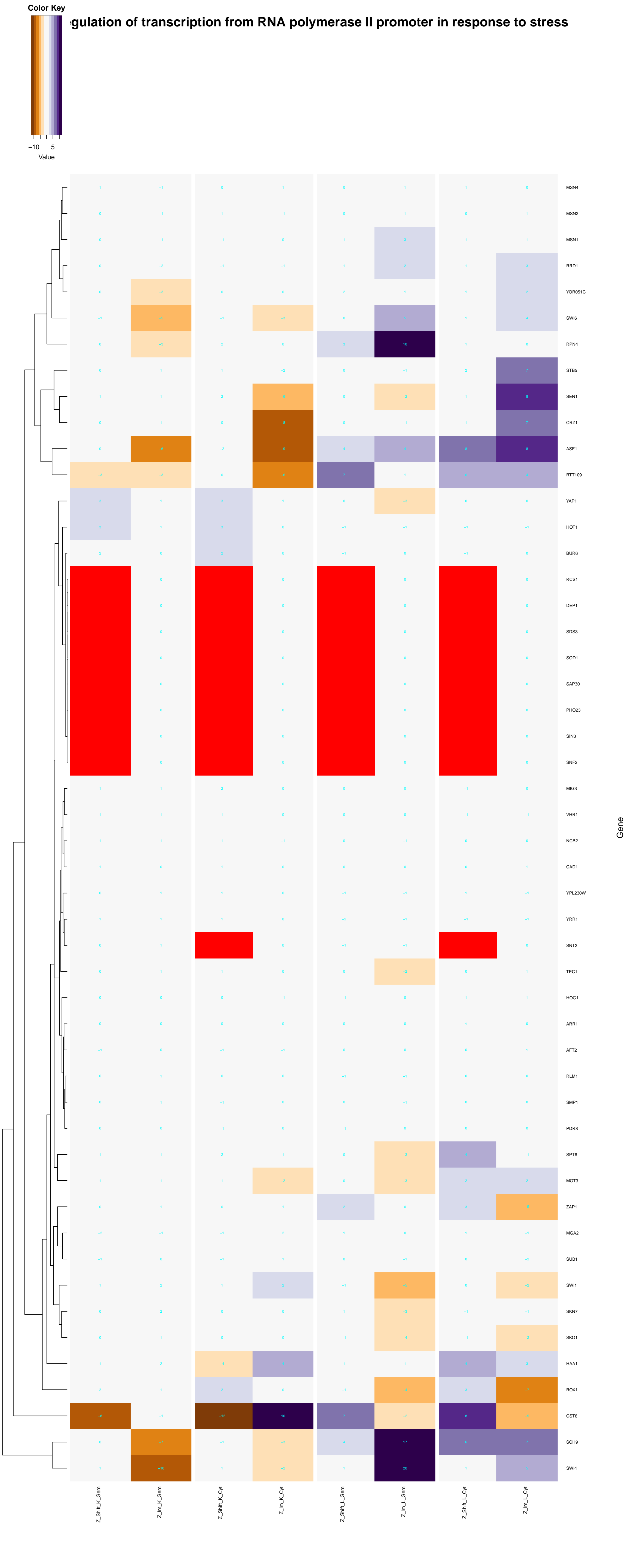




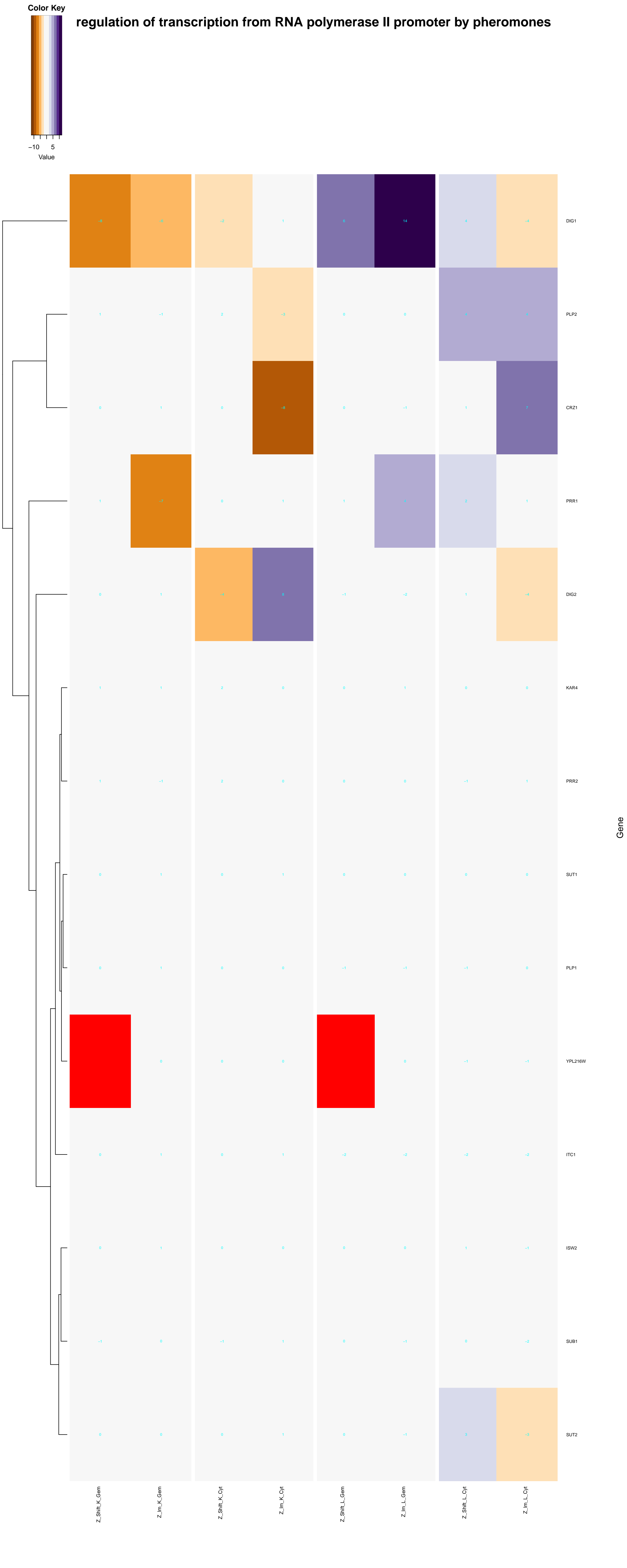
of oligopeptide transport by regulation of transcription from RNA polymerase II promoter

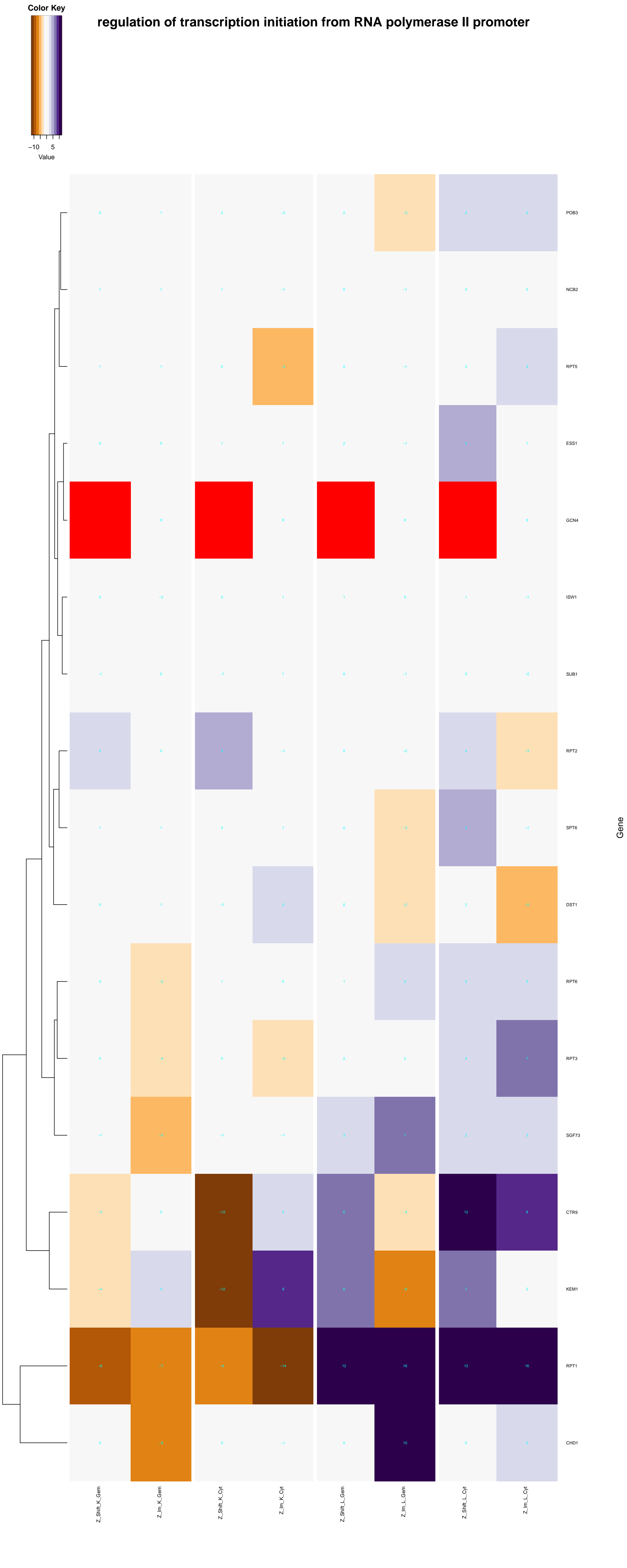


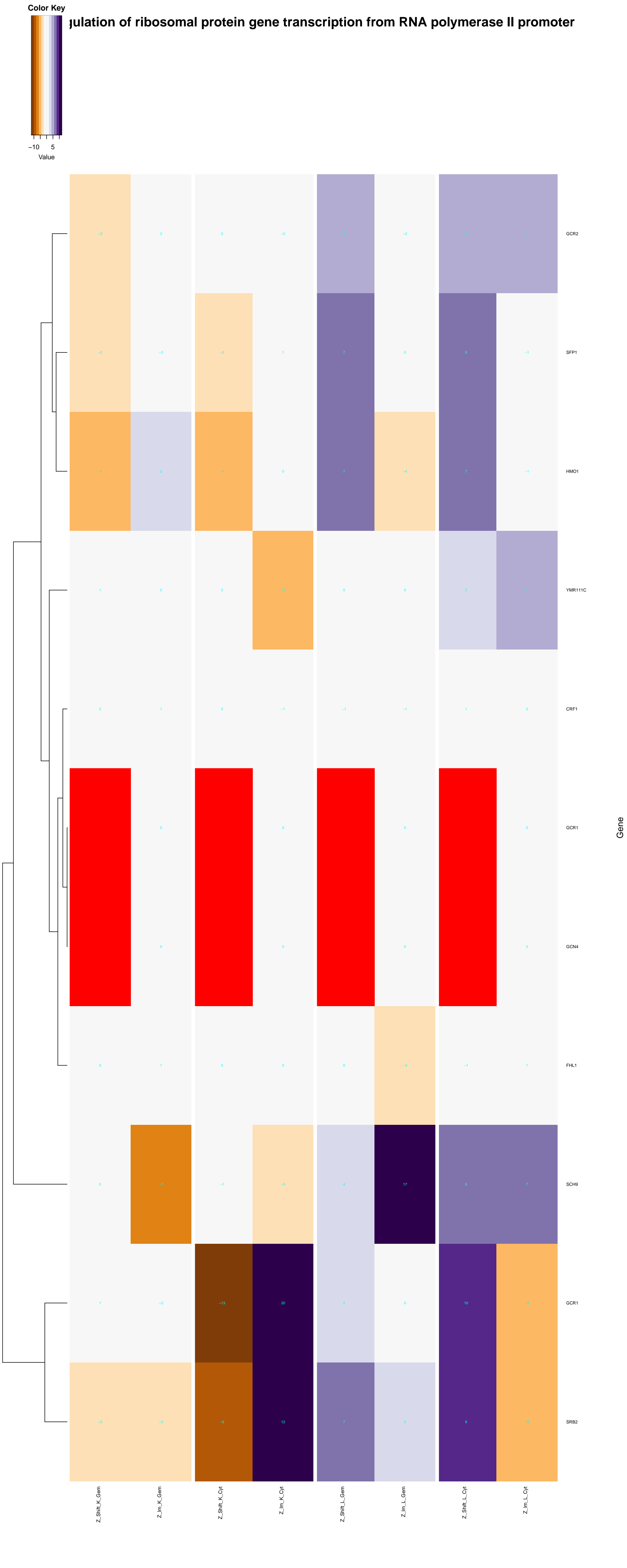


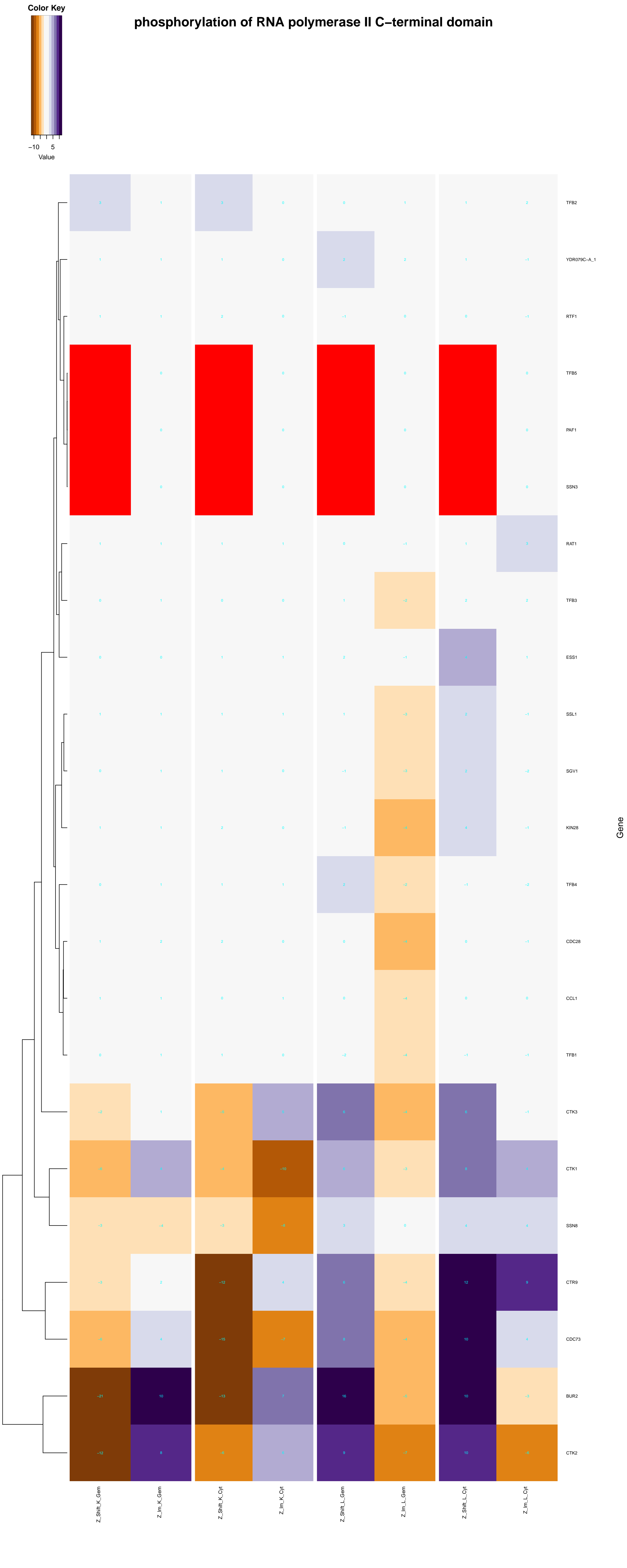


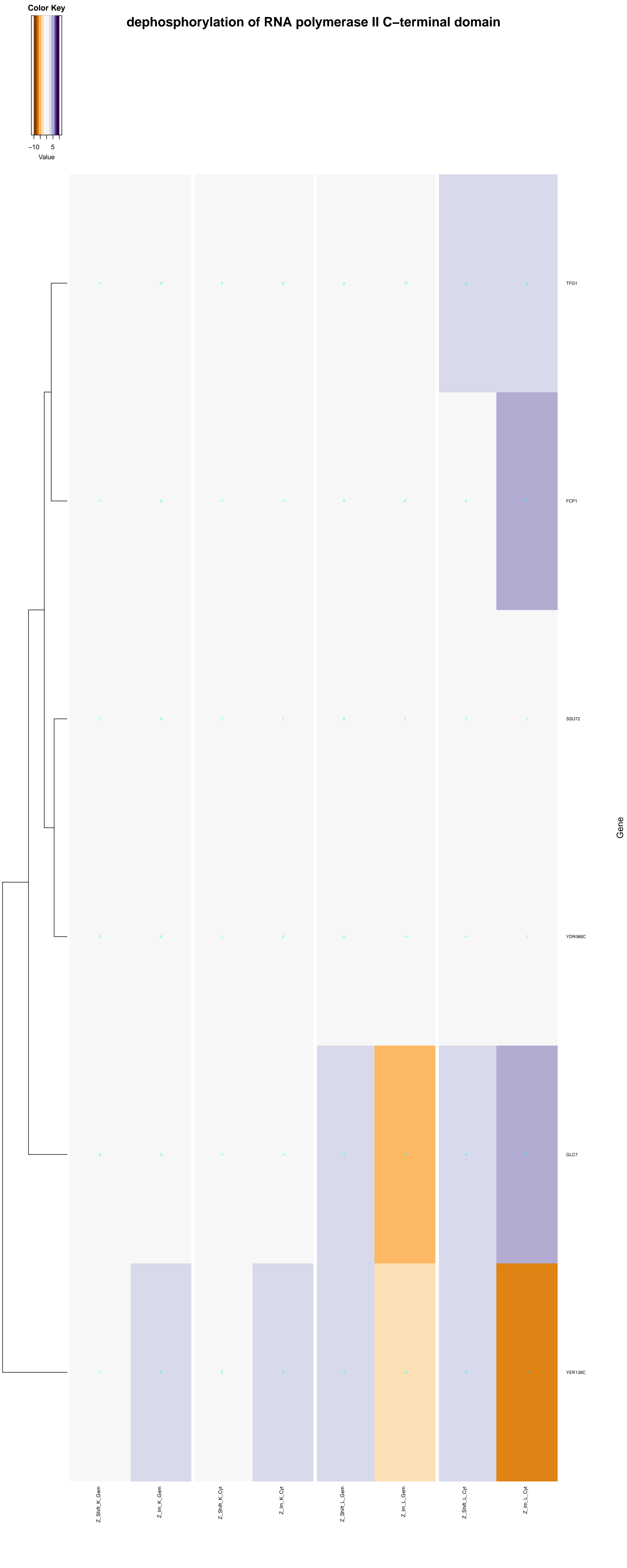


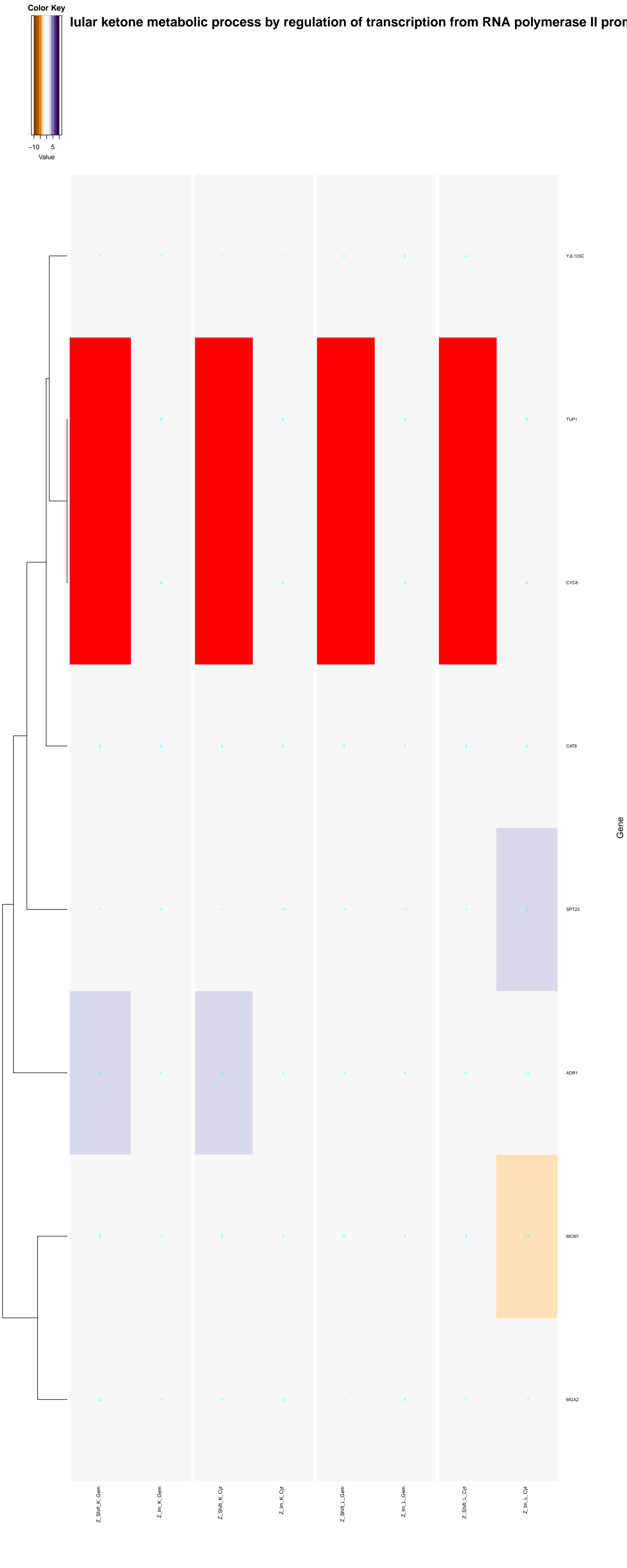




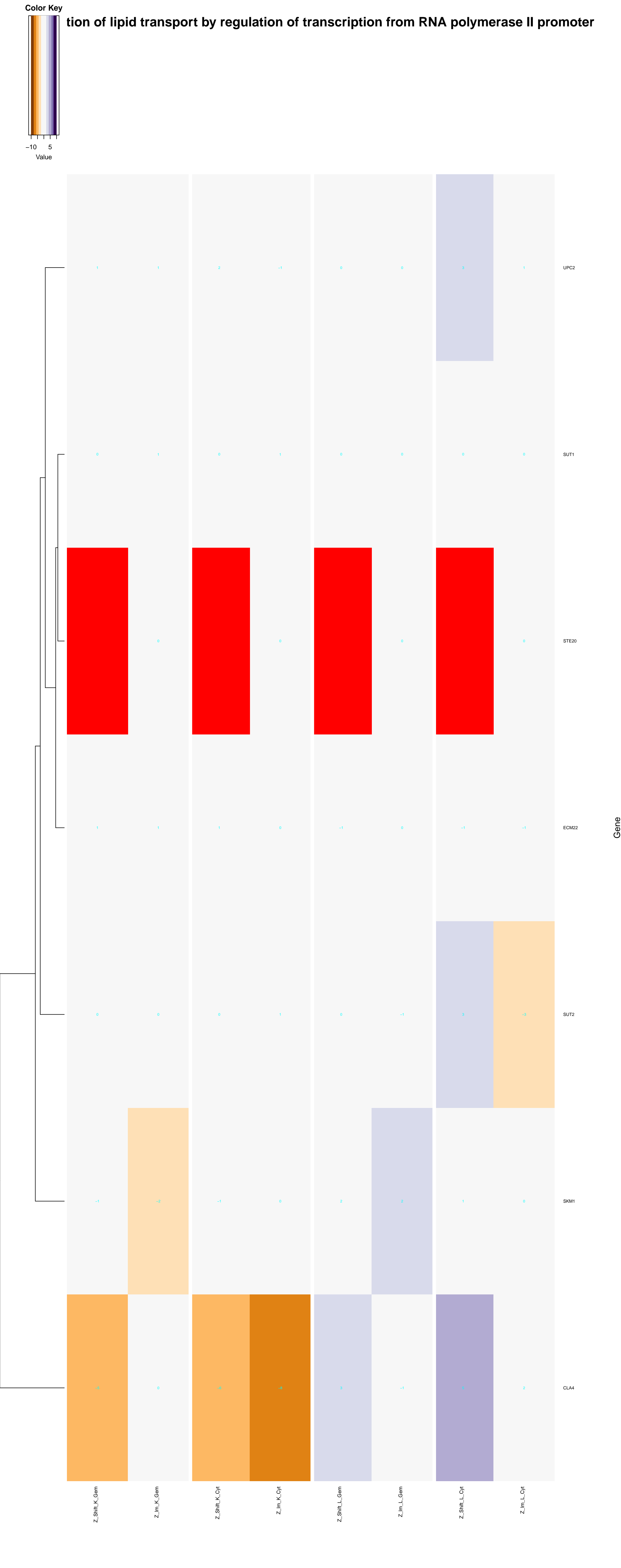


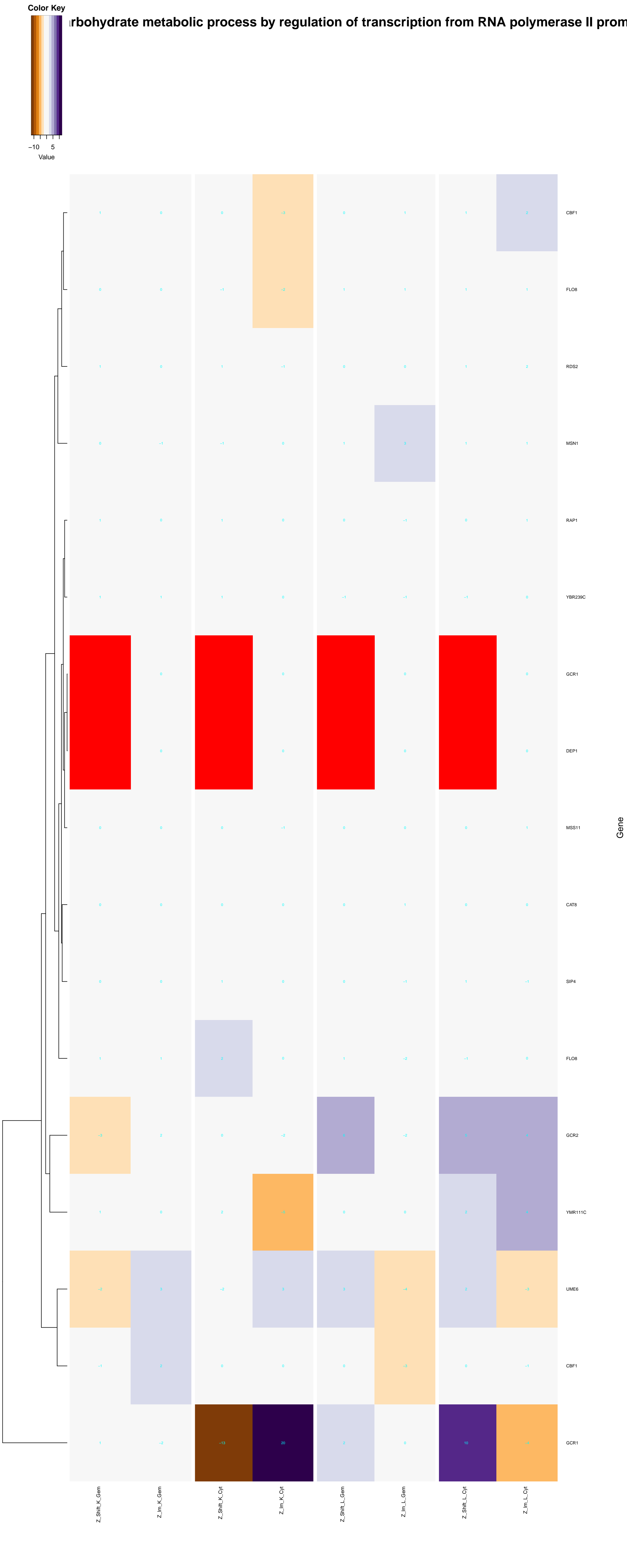


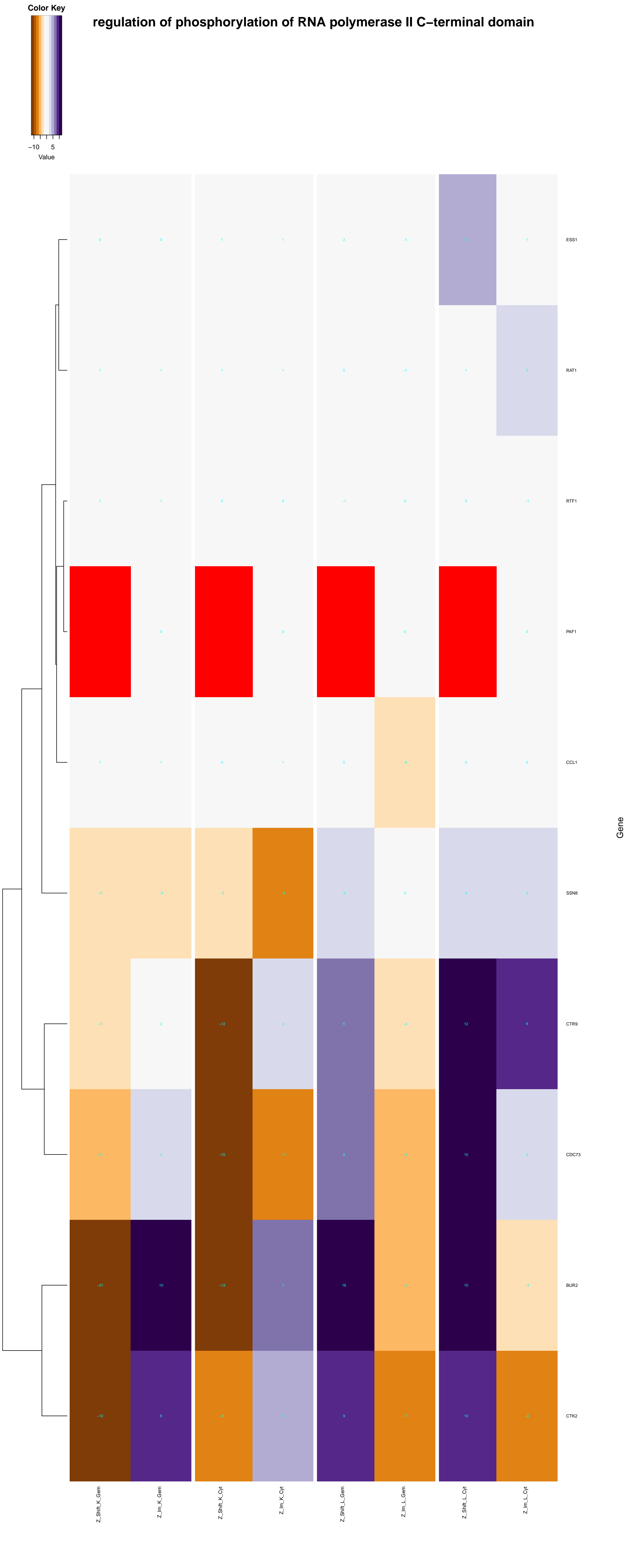




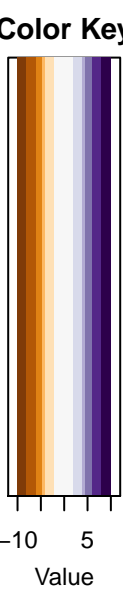




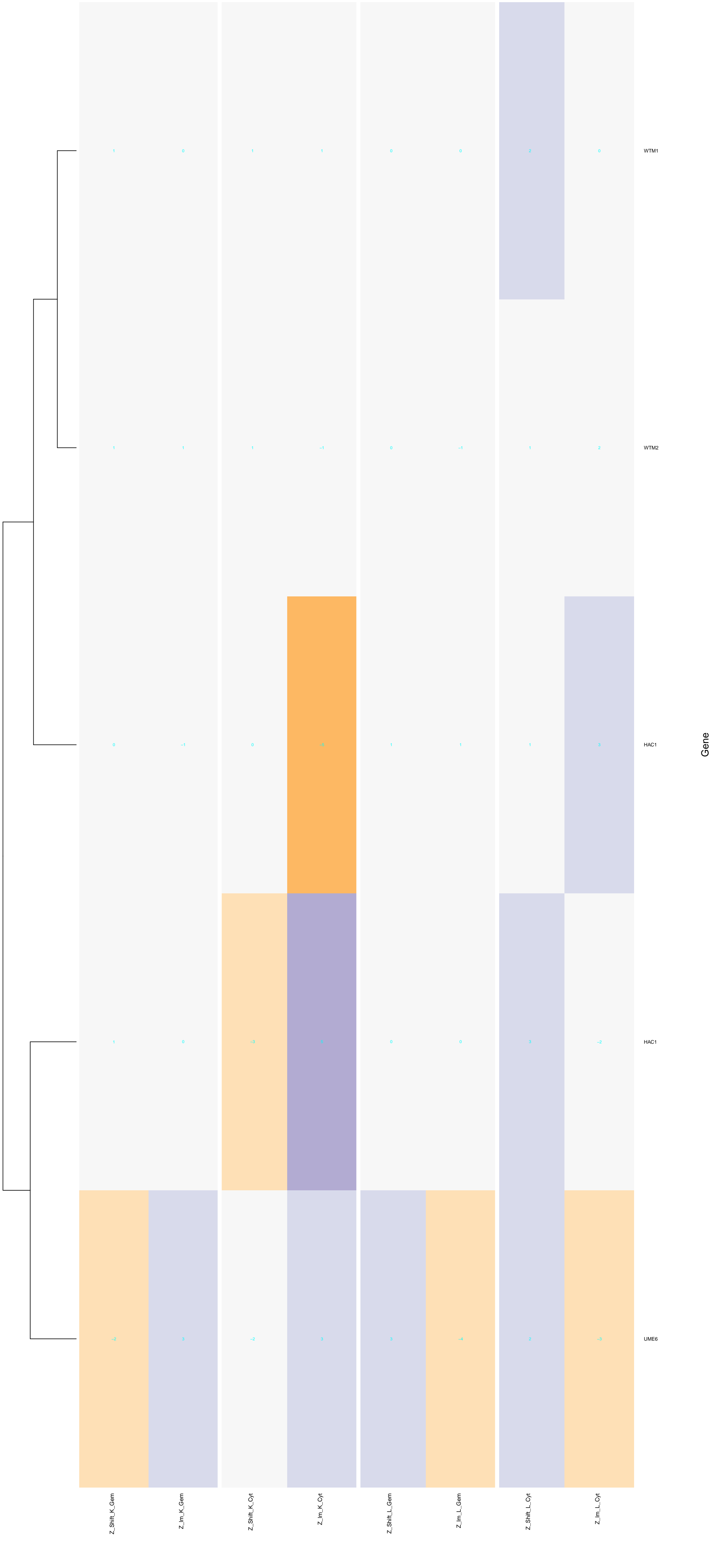


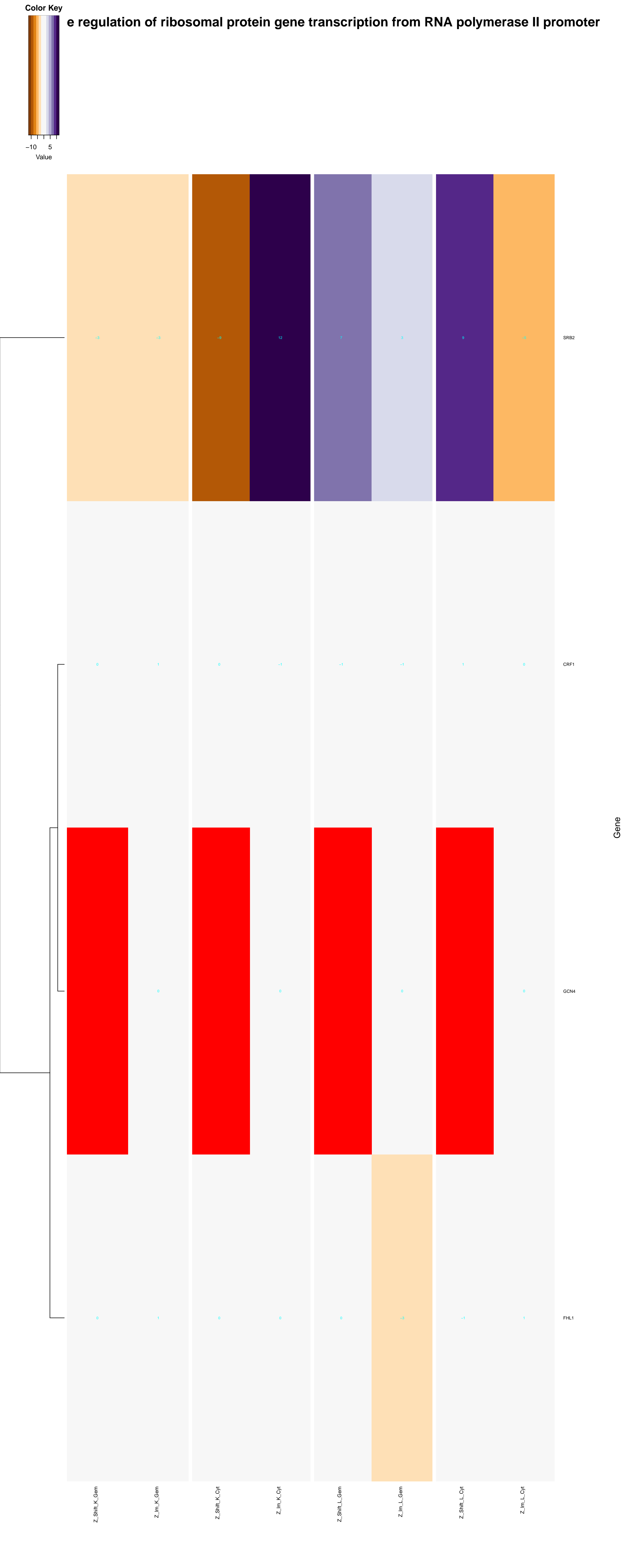


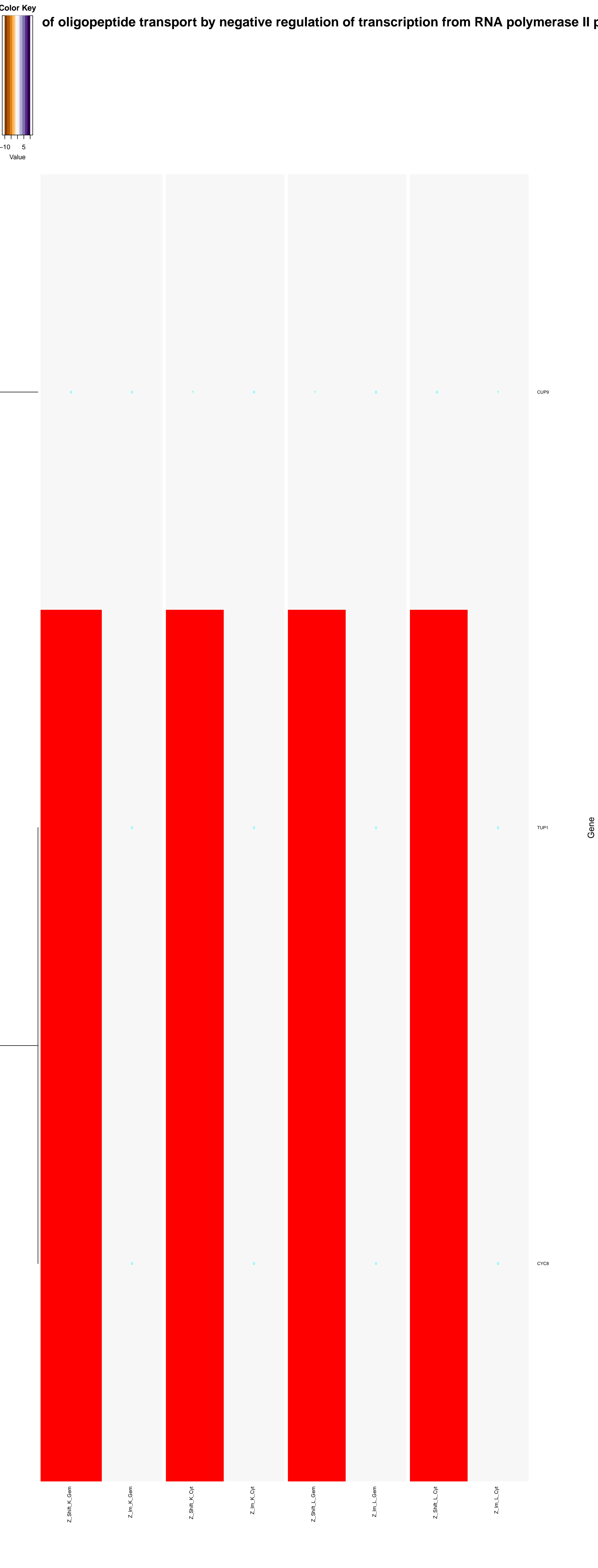


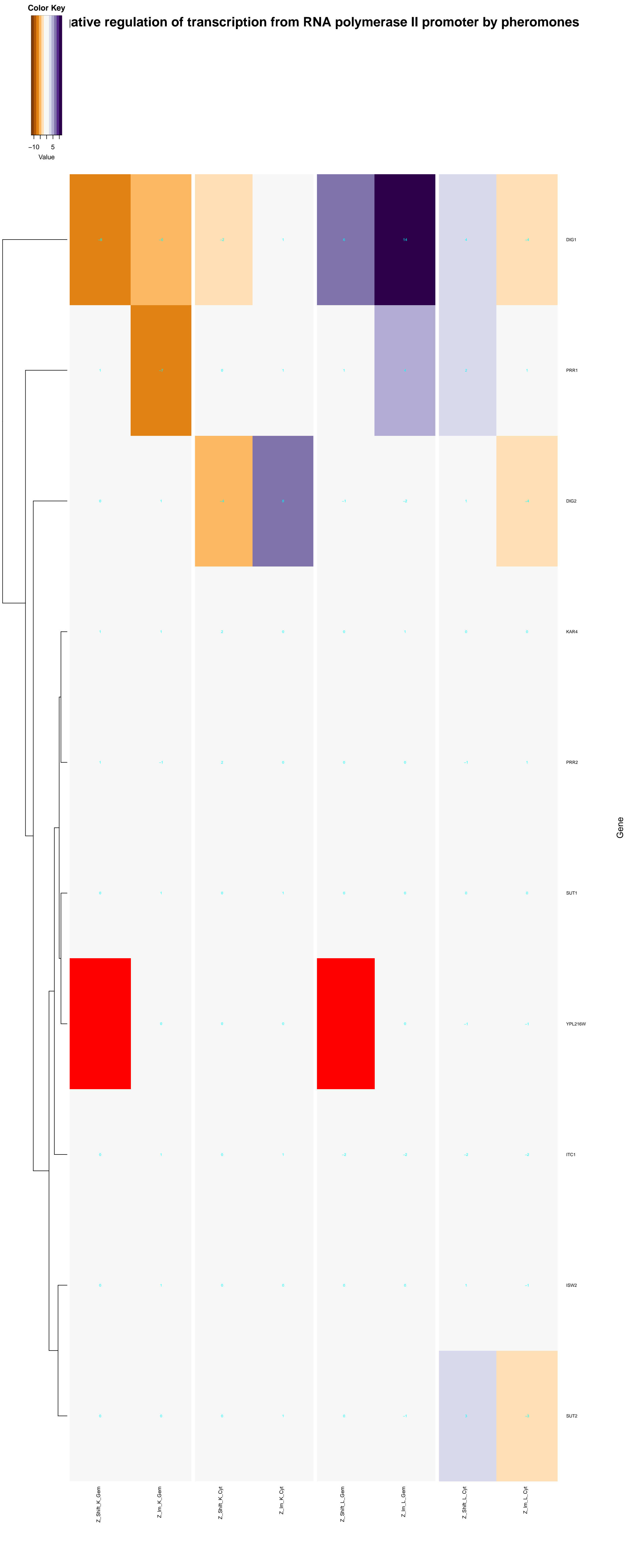


egulation of transcription from RNA polymerase II promoter involved in meiotic cell cycle

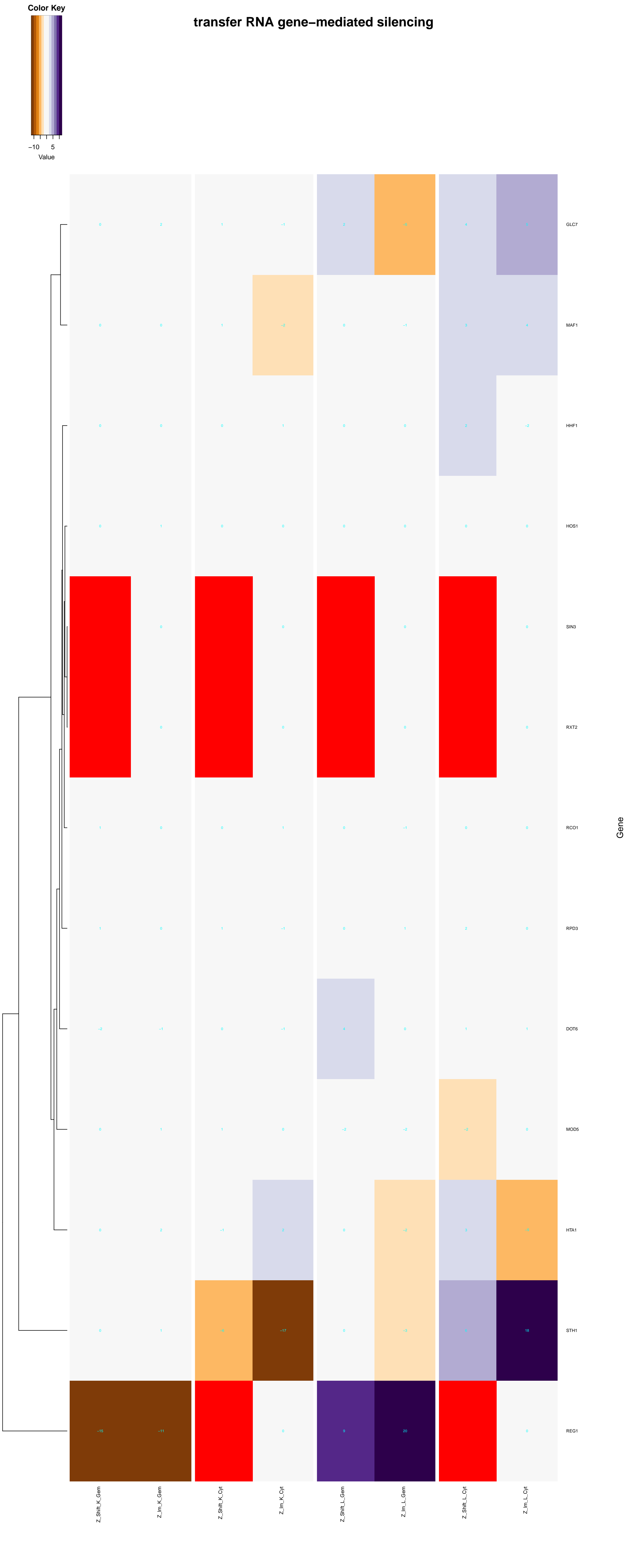


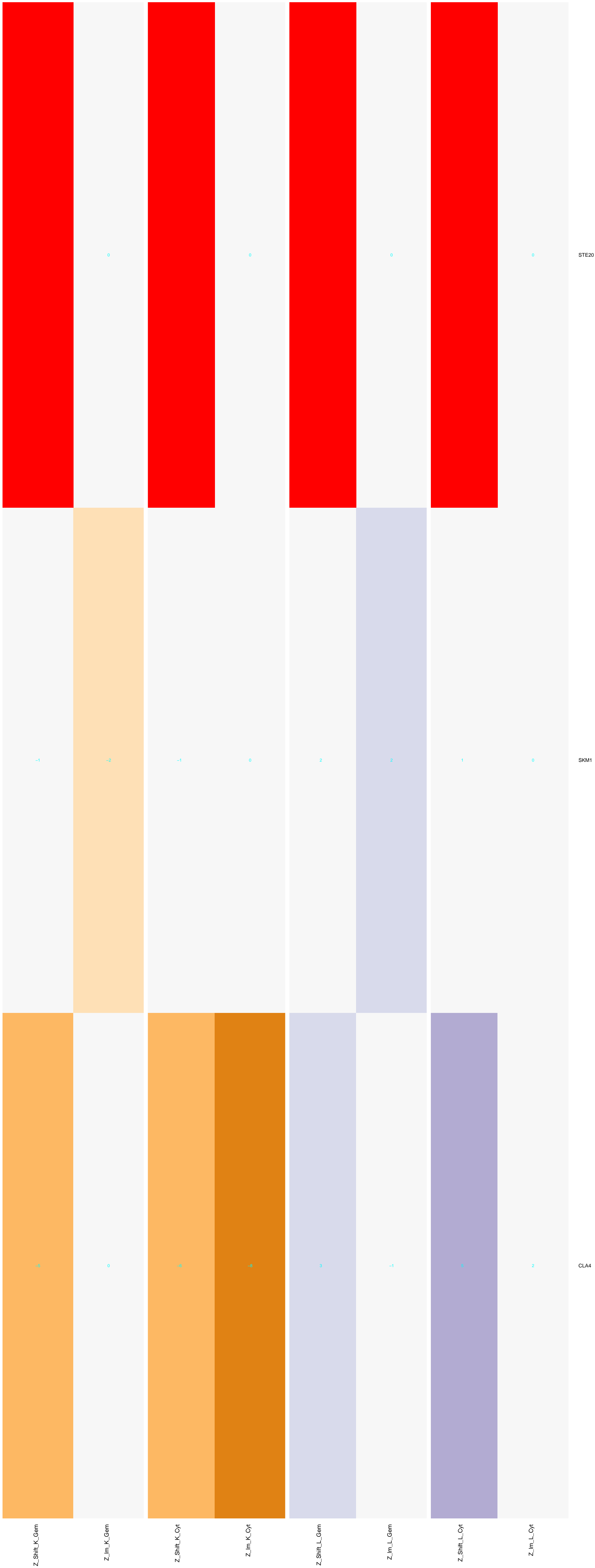
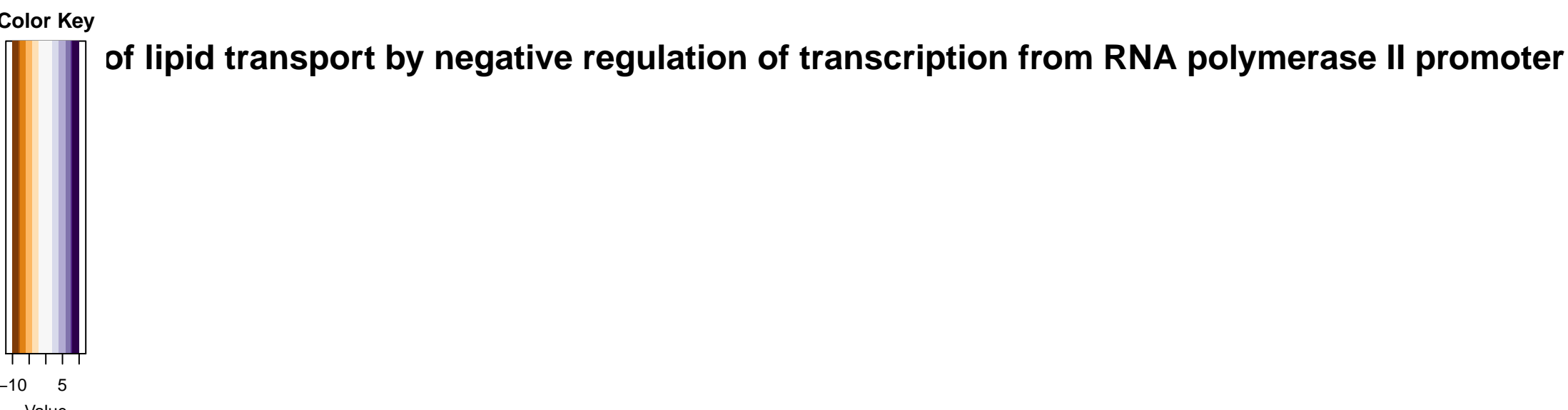




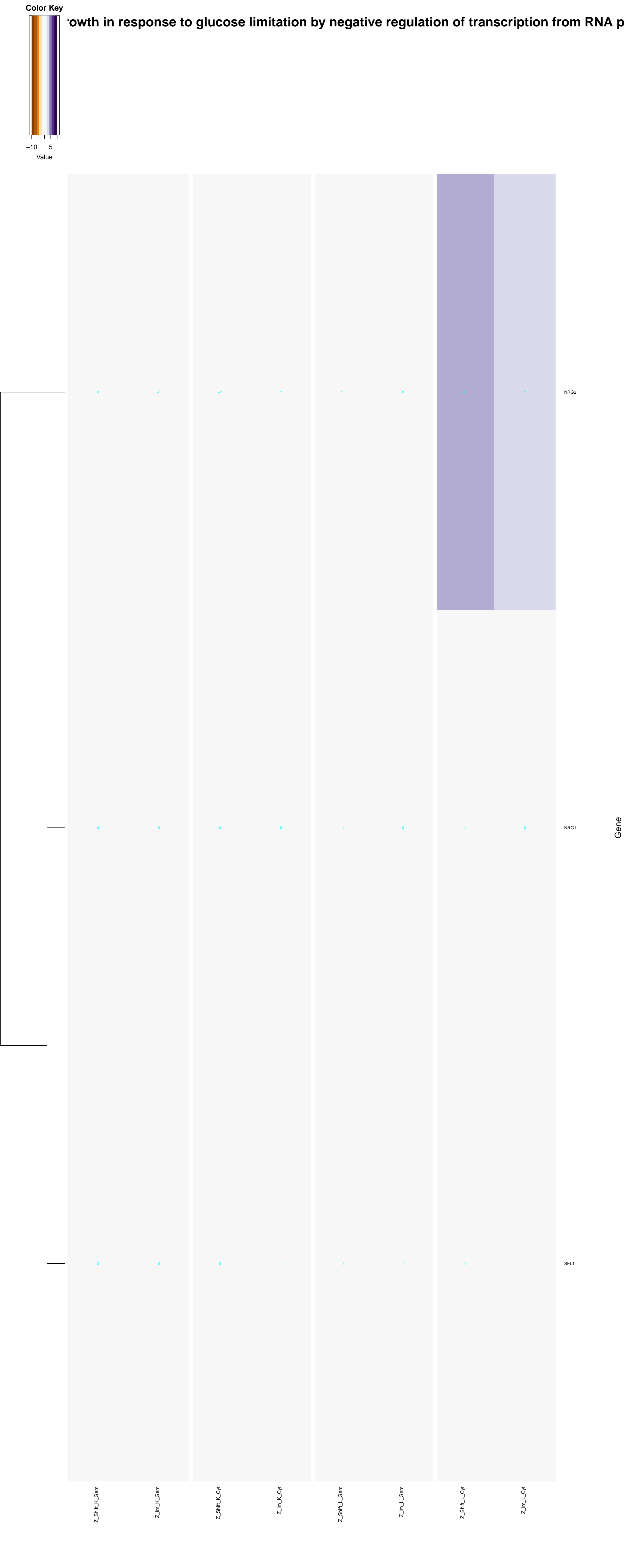


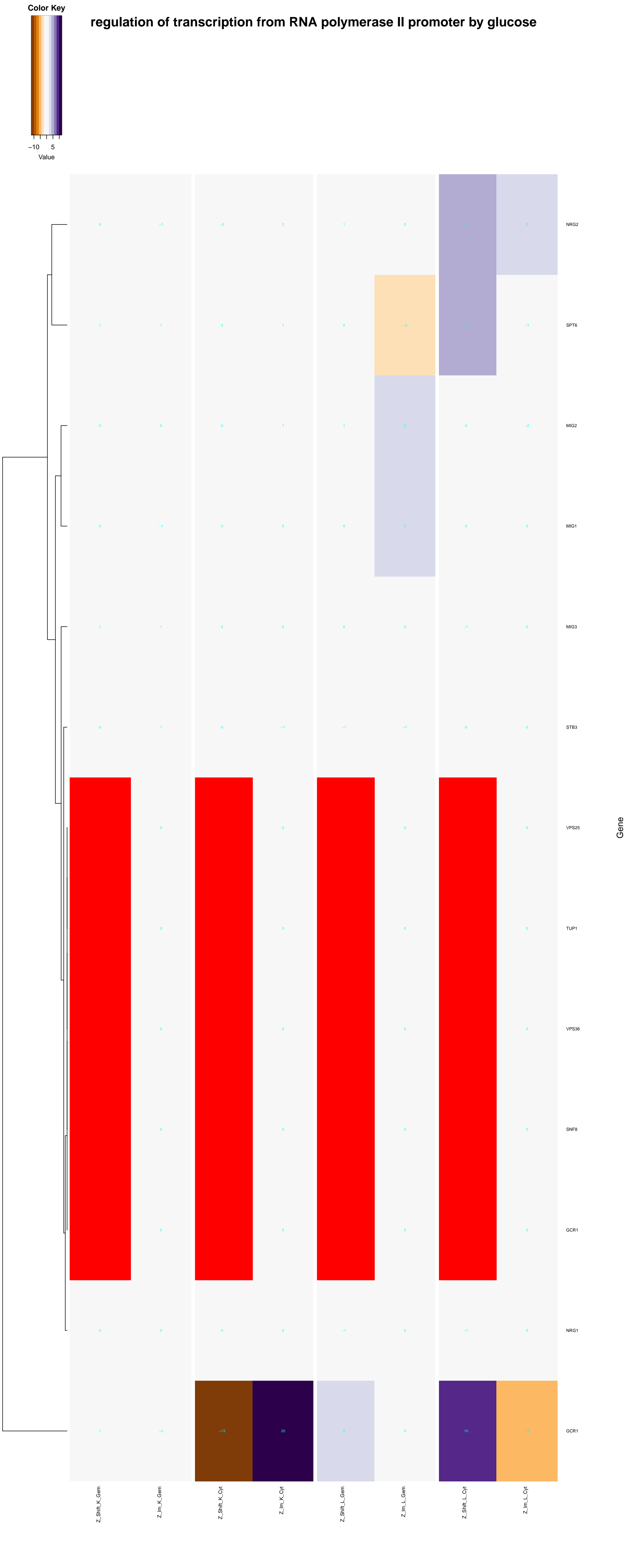






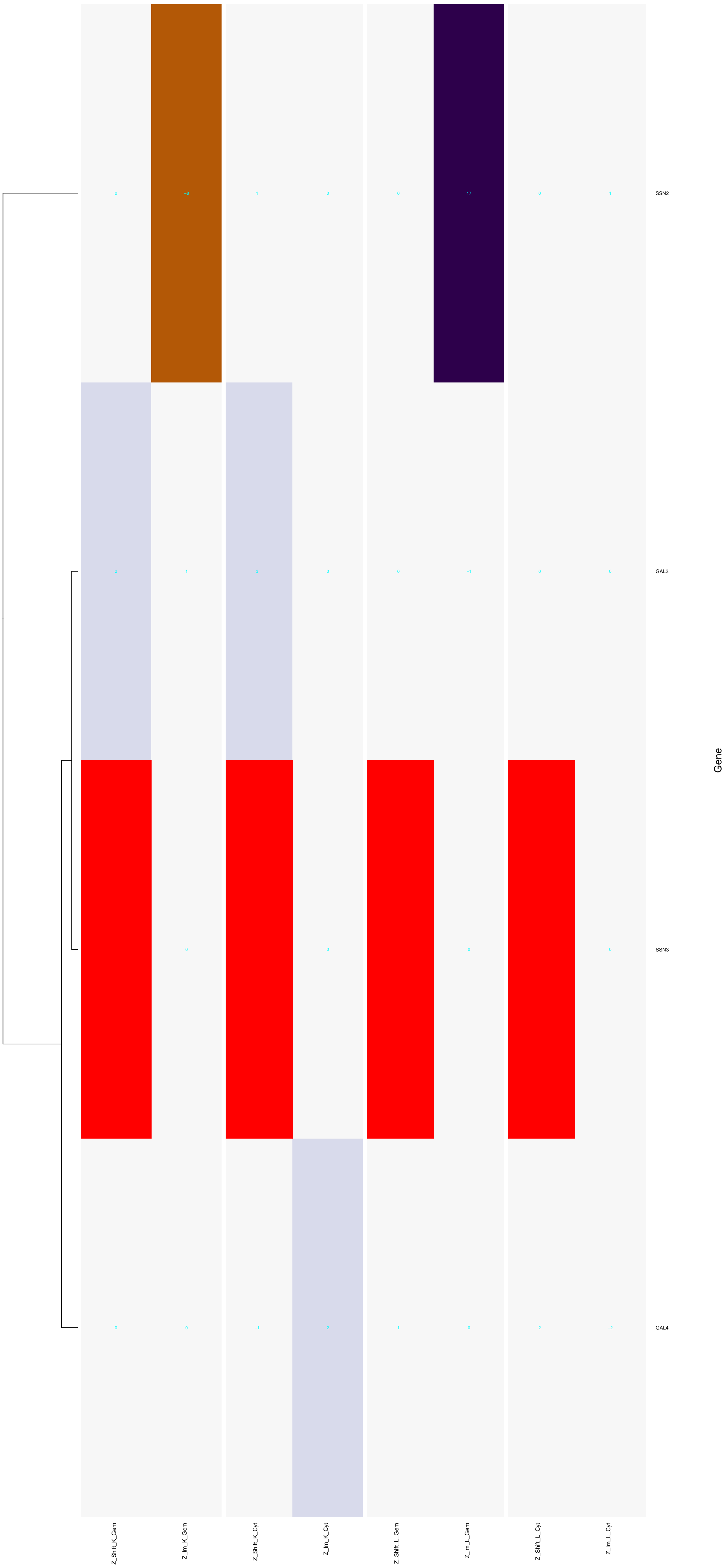
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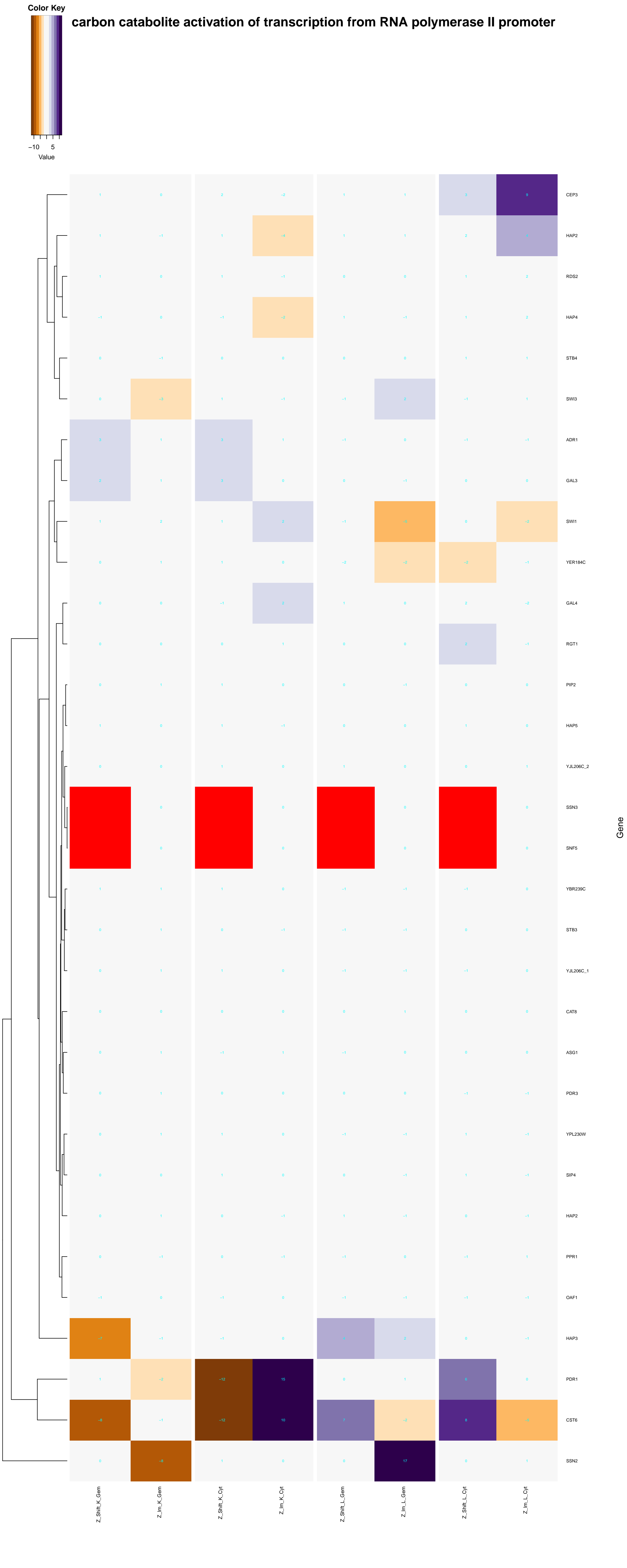


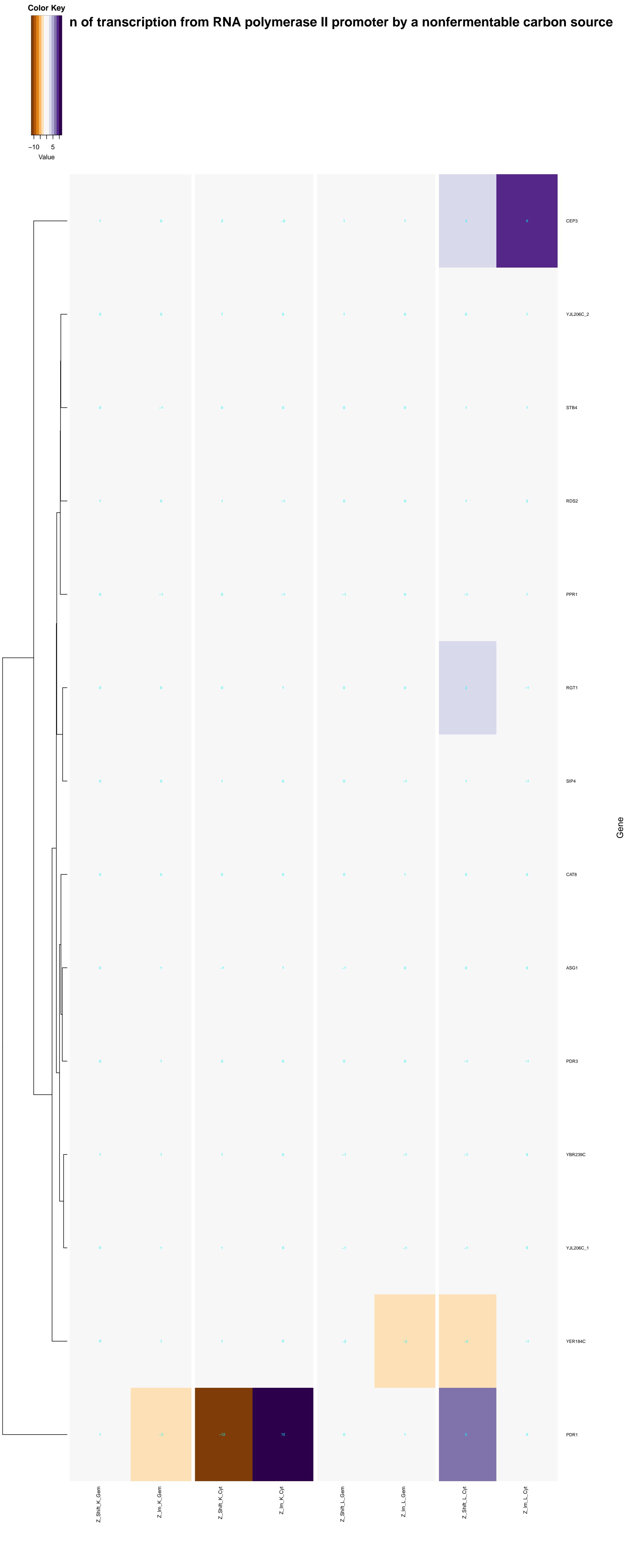


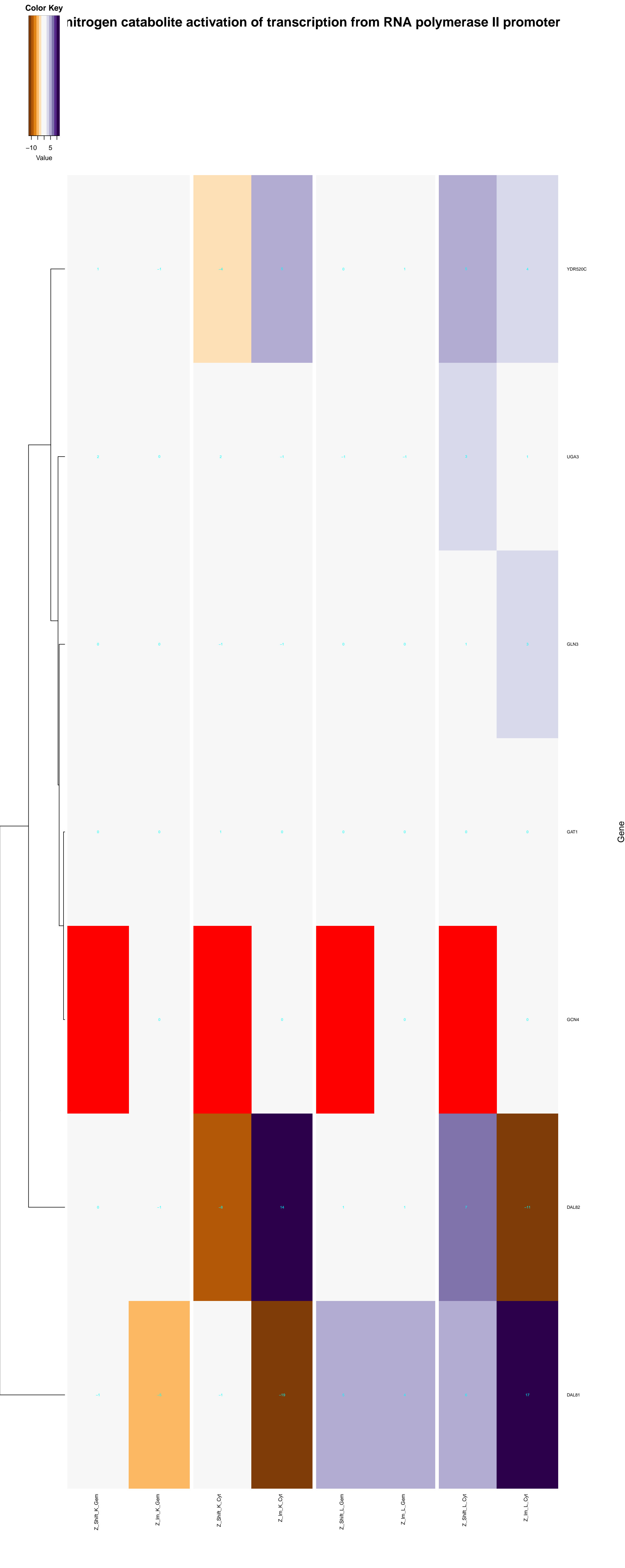


regulation of transcription from RNA polymerase II promoter by galactose

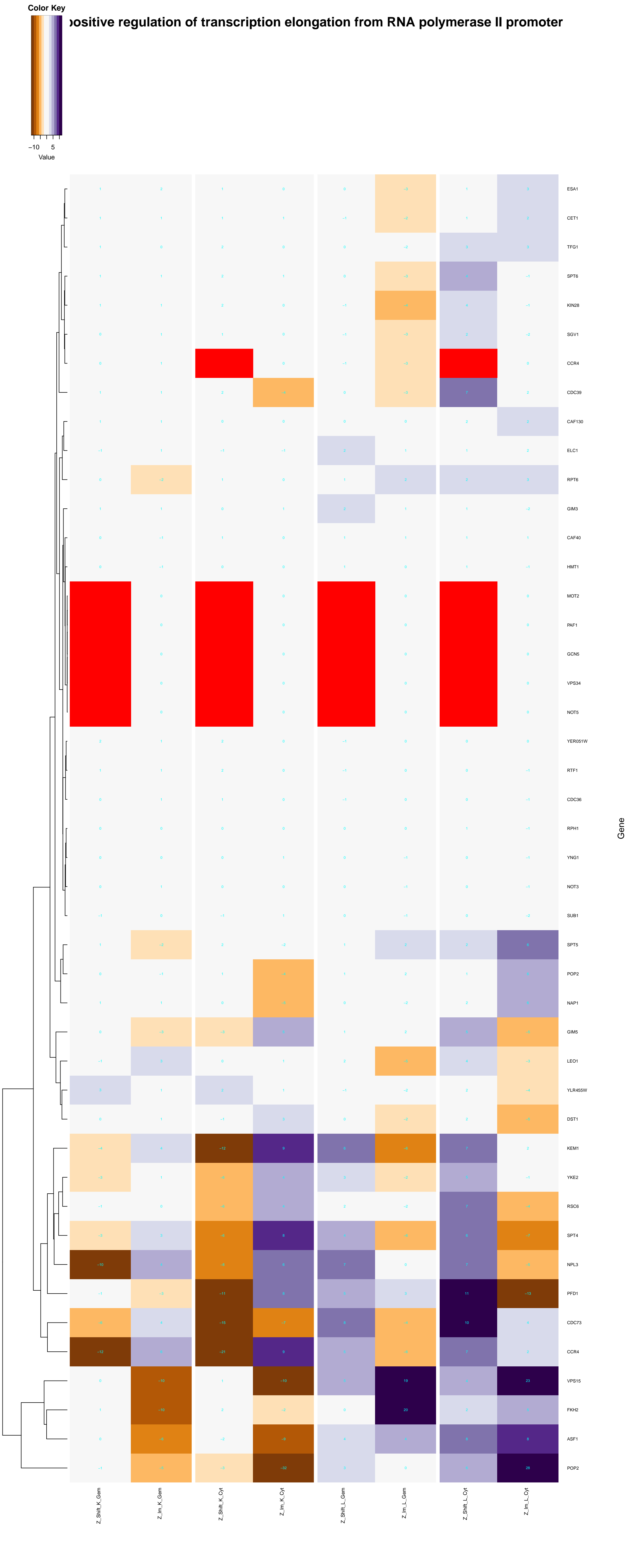










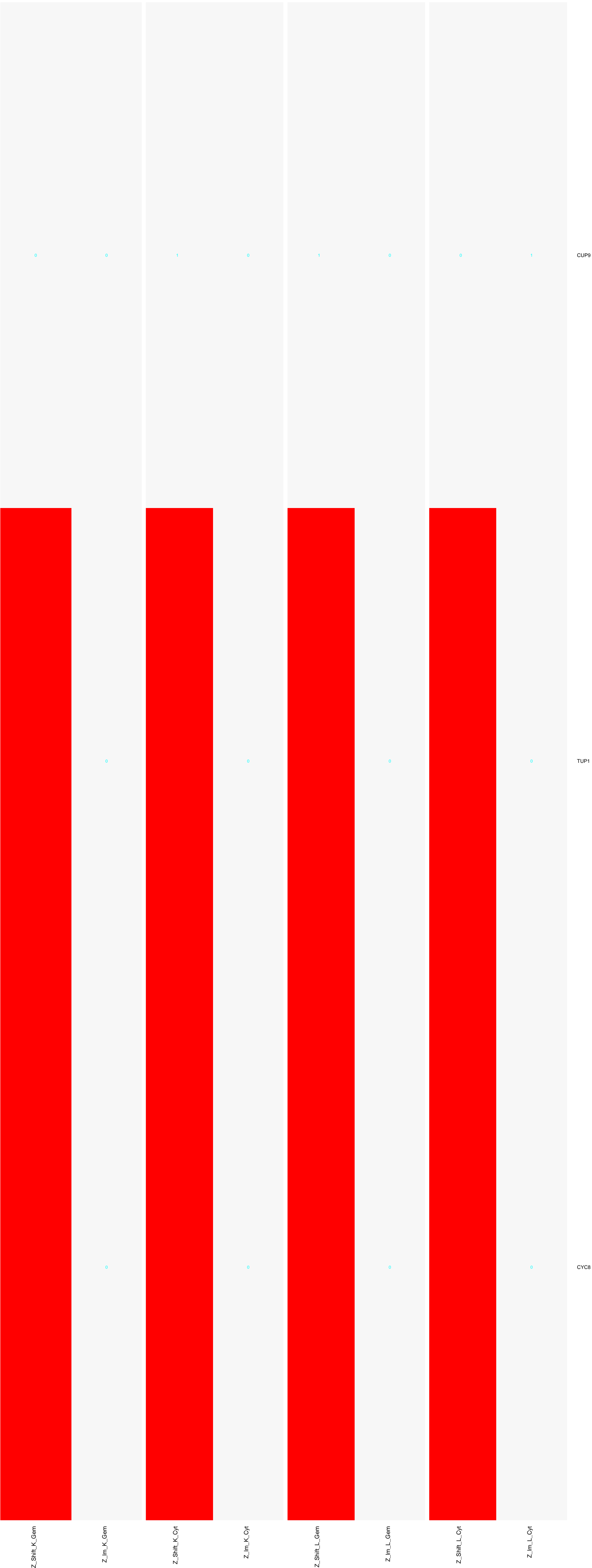


Color Key

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

Value

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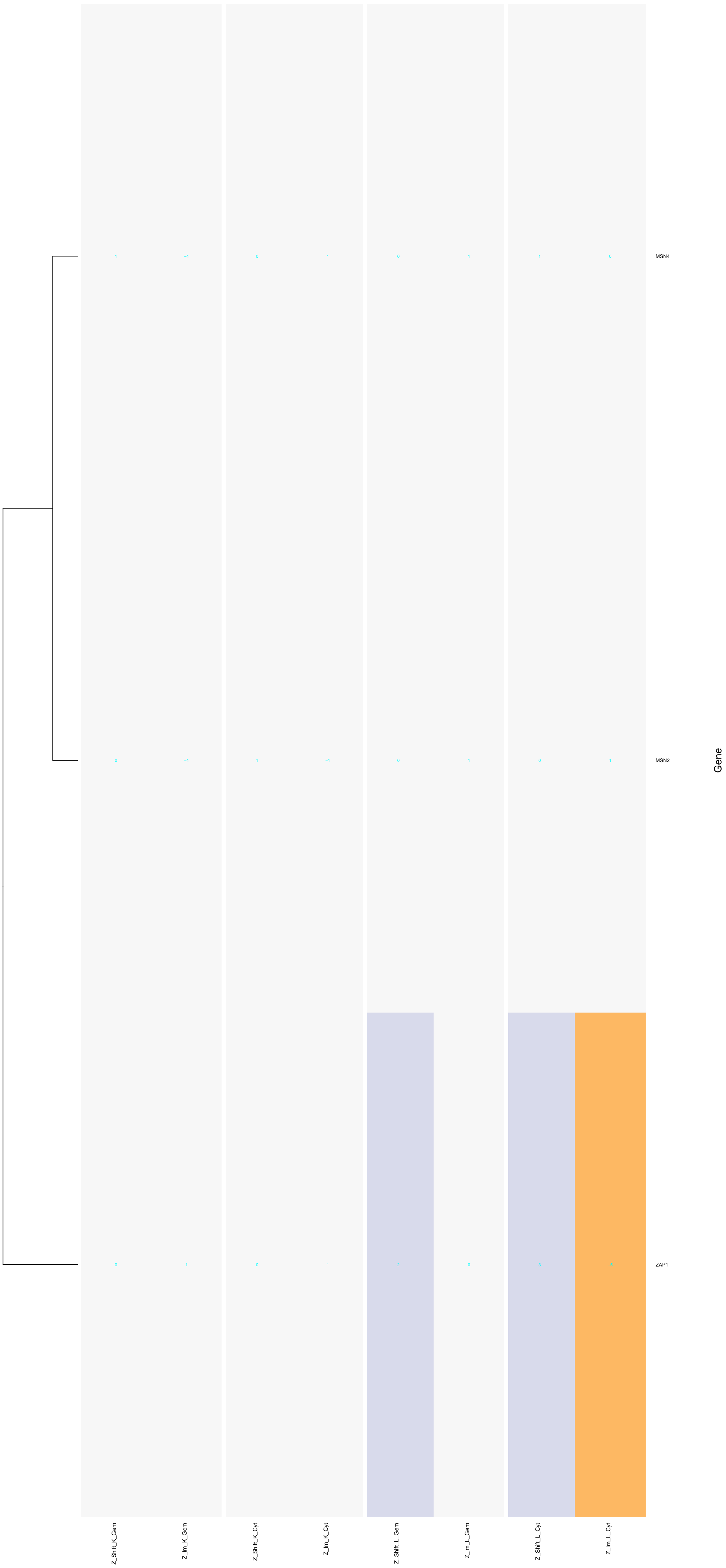
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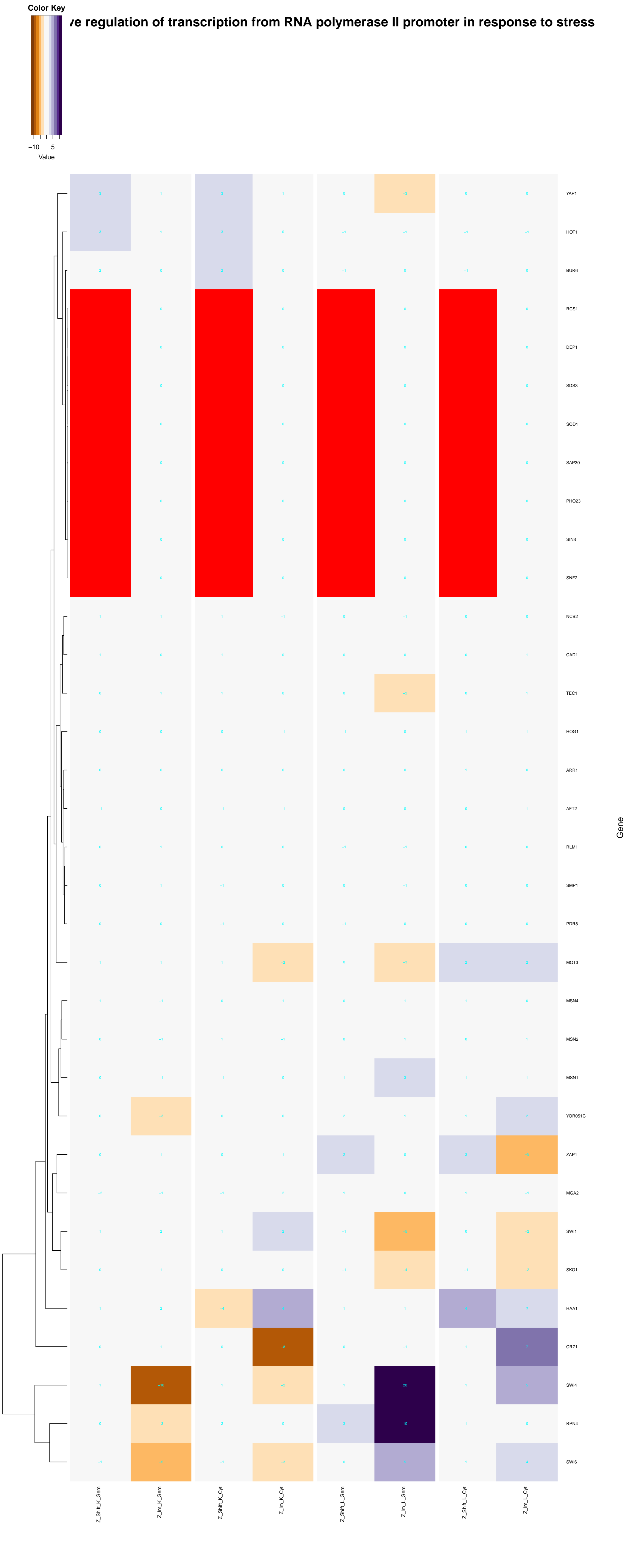
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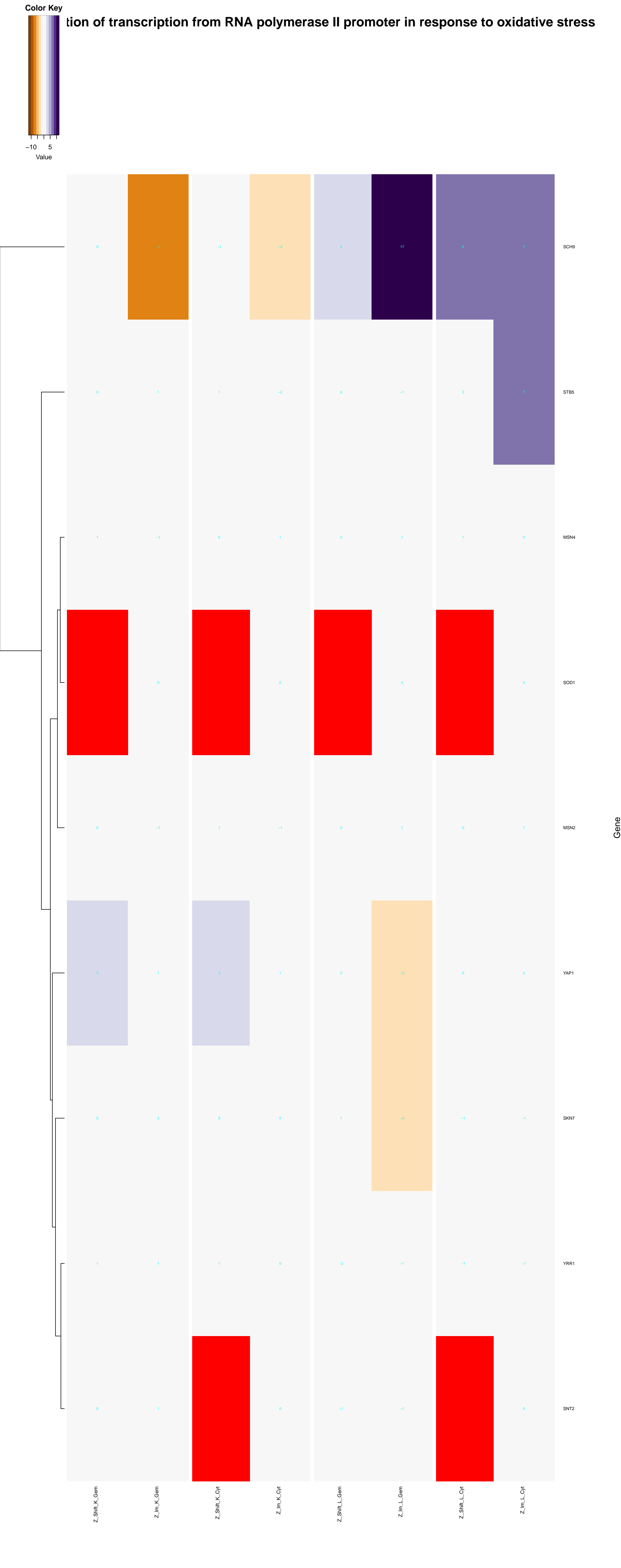
on of transcription from RNA polymerase II promoter in response to zinc ion starvation

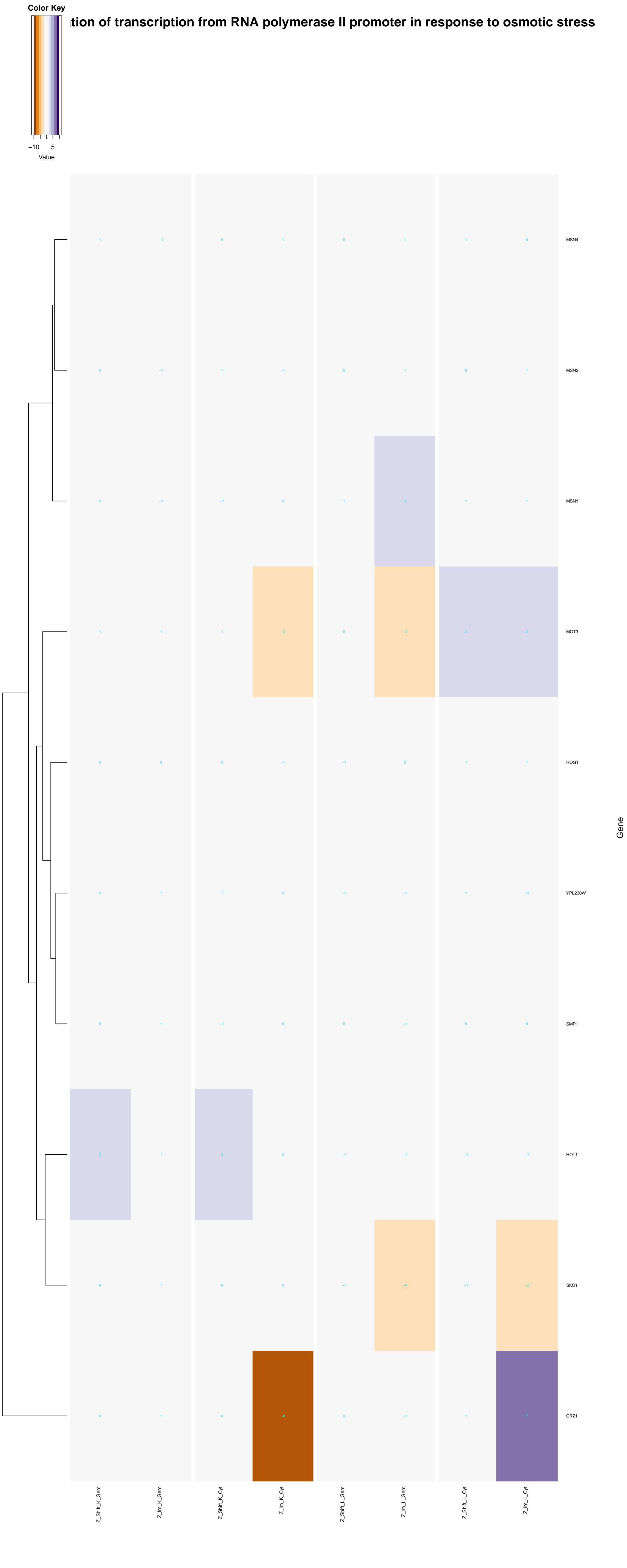
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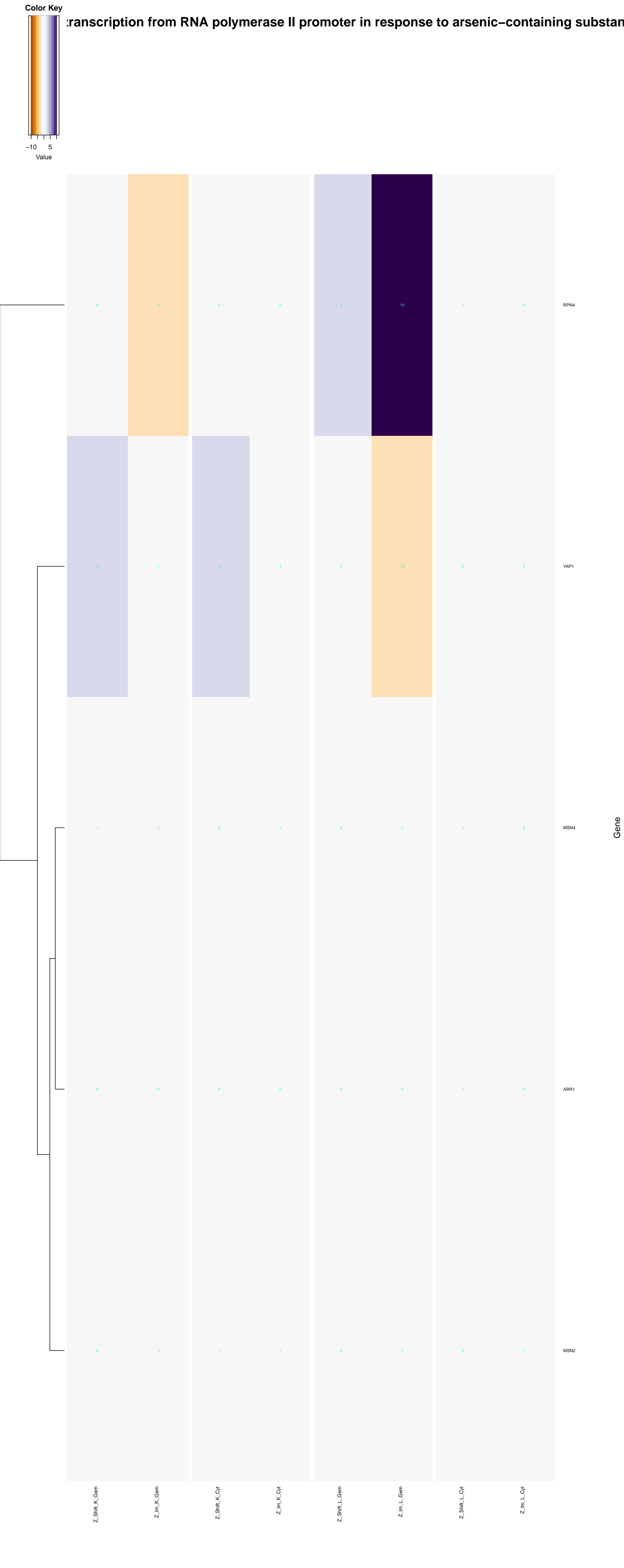
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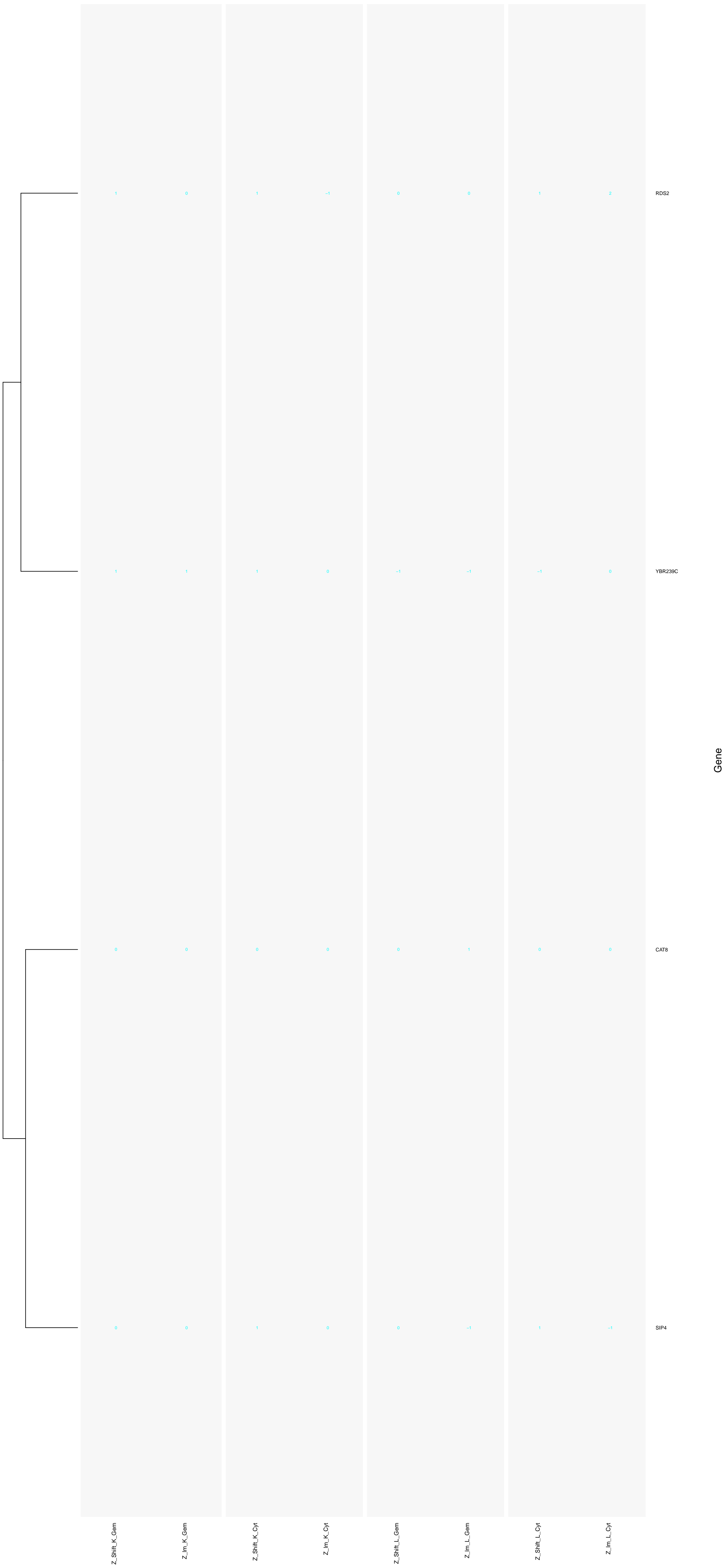
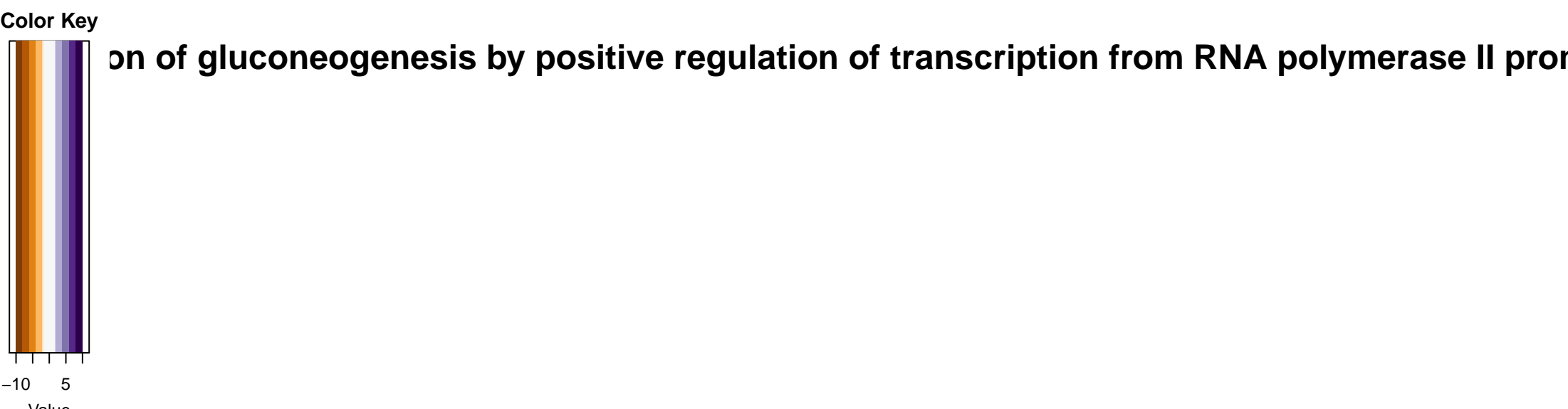




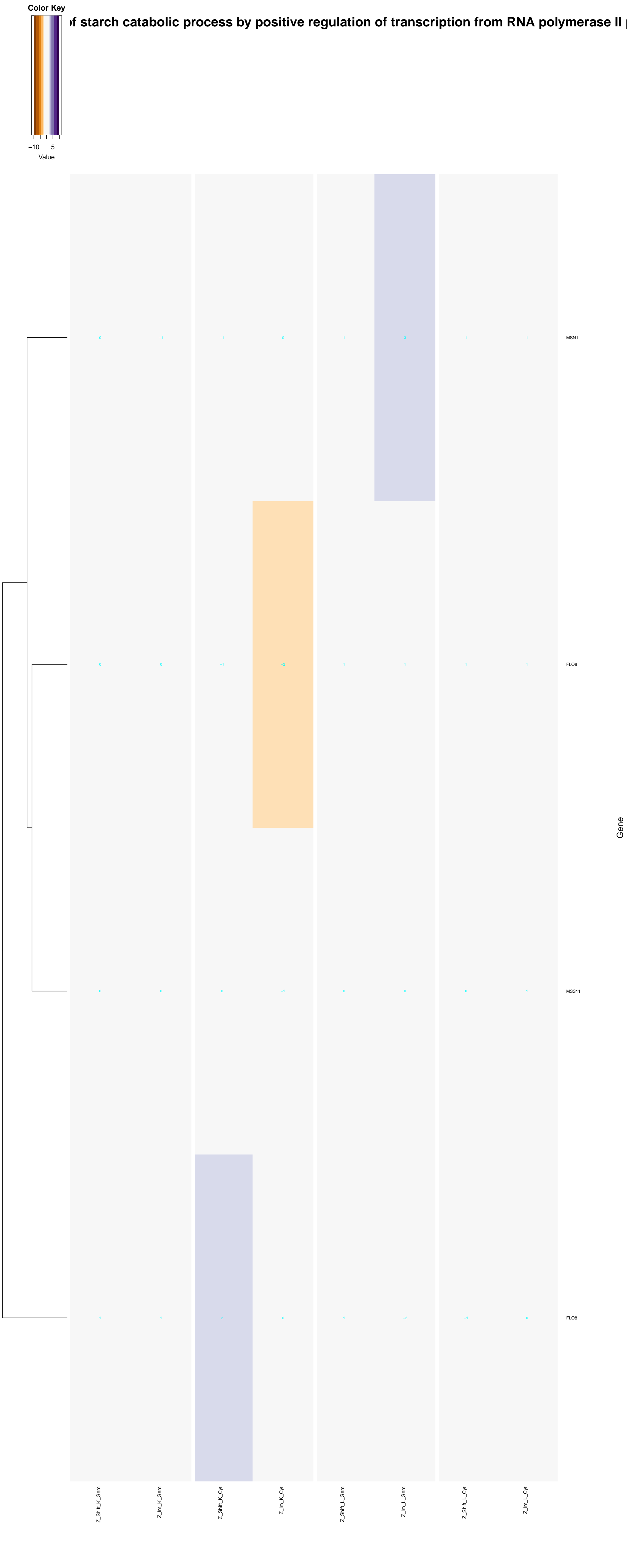


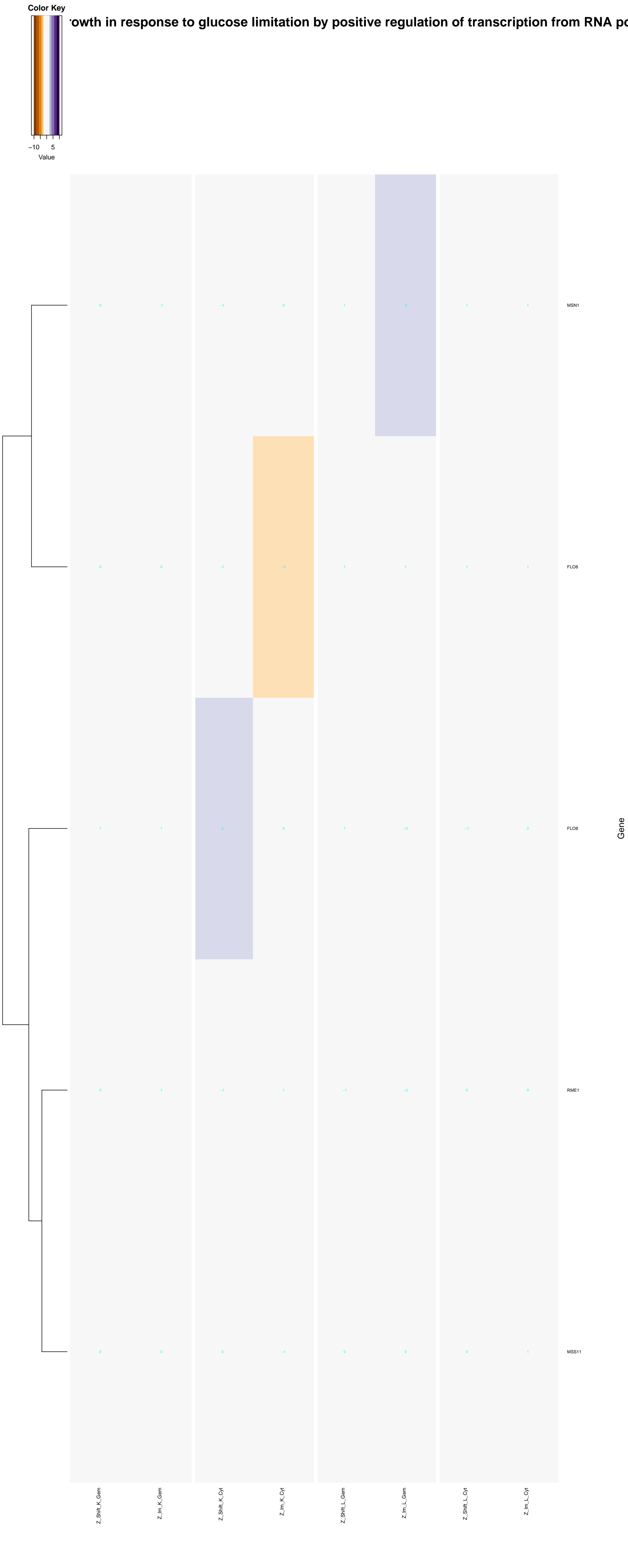






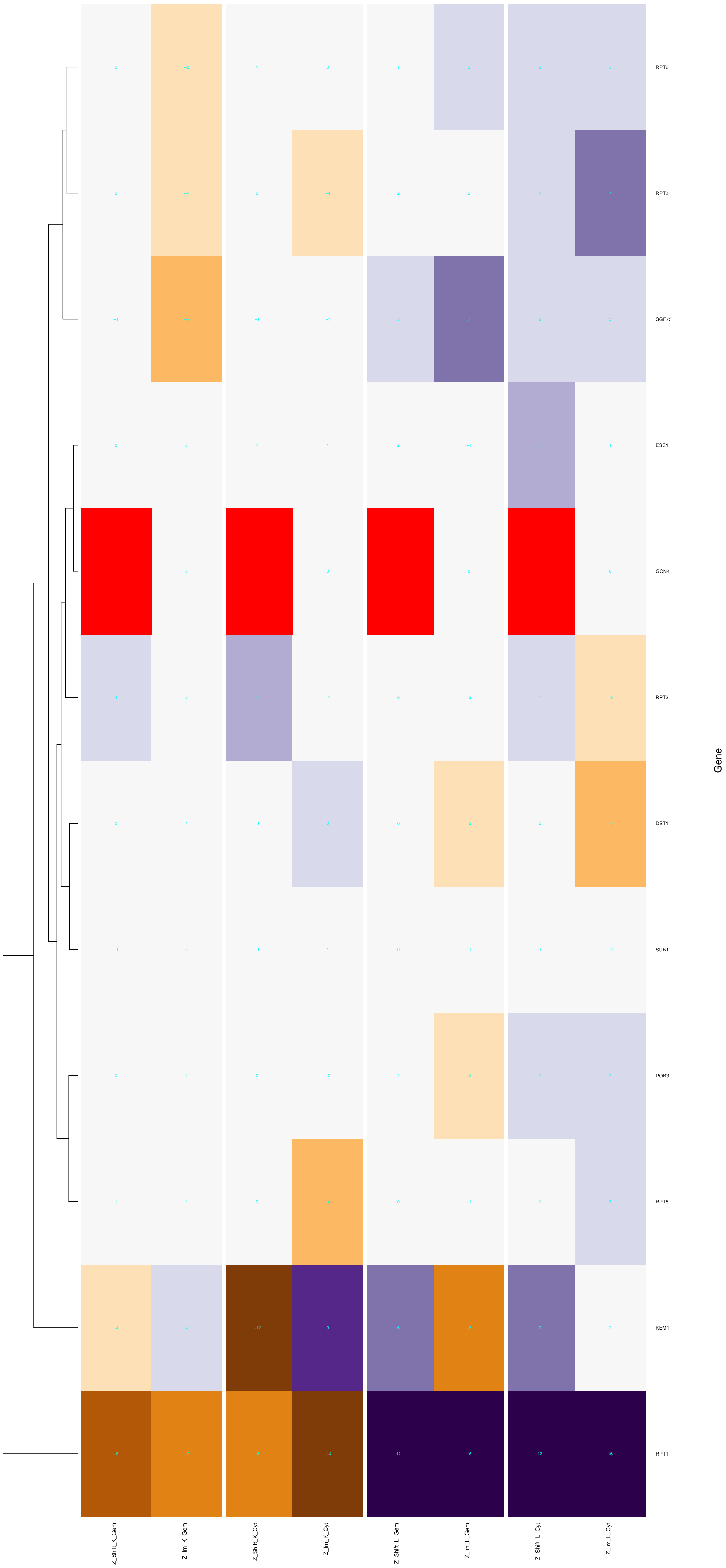


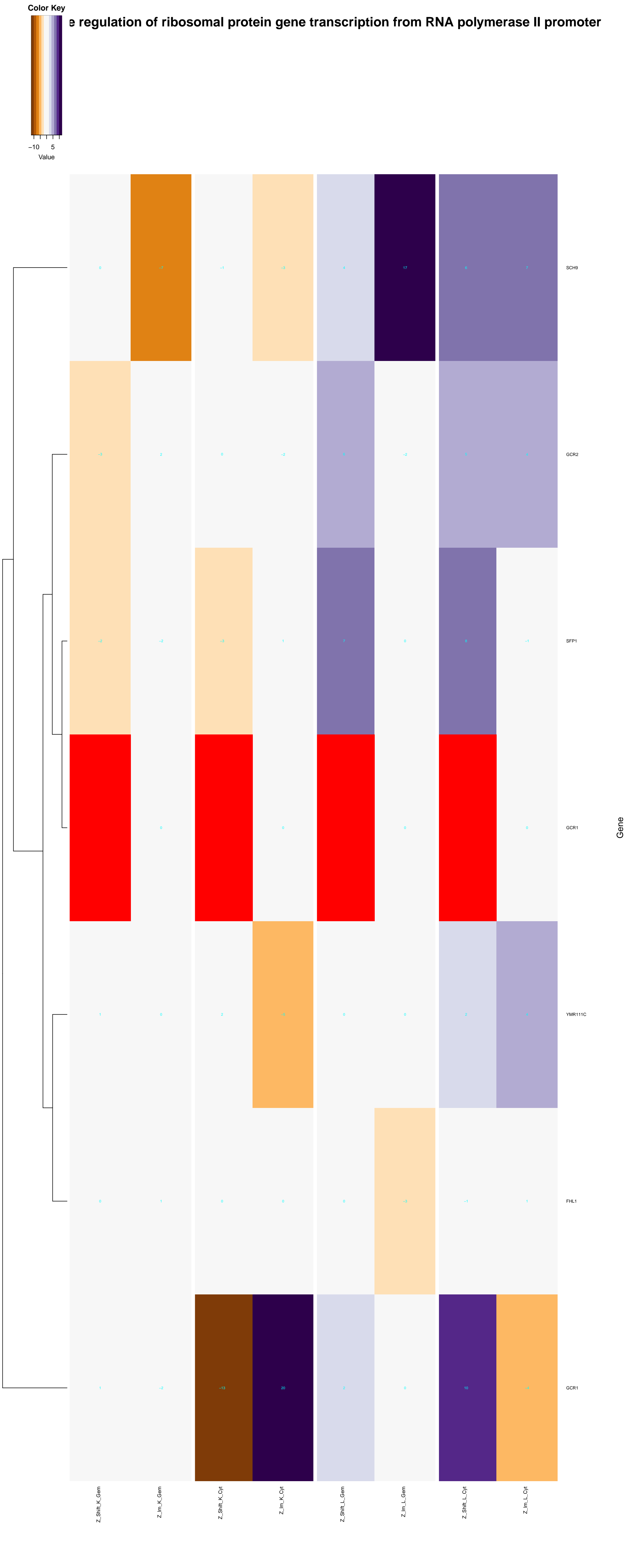


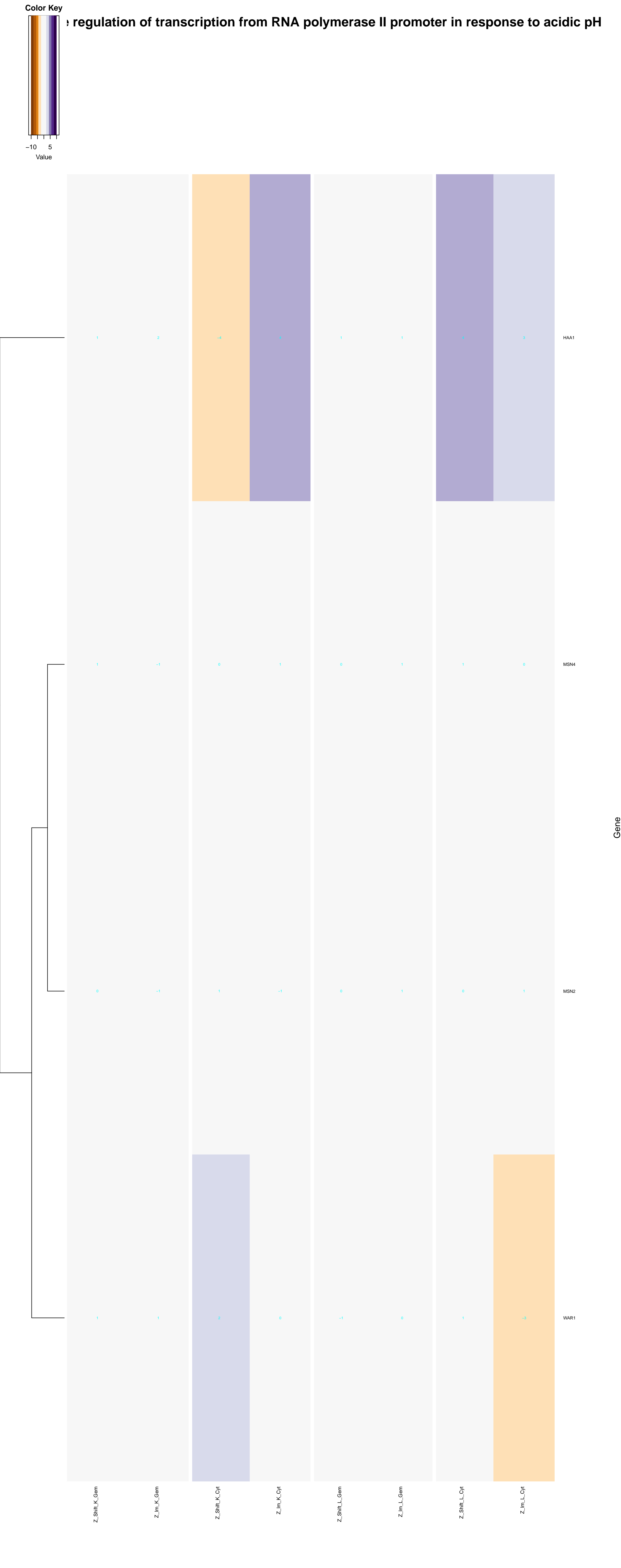


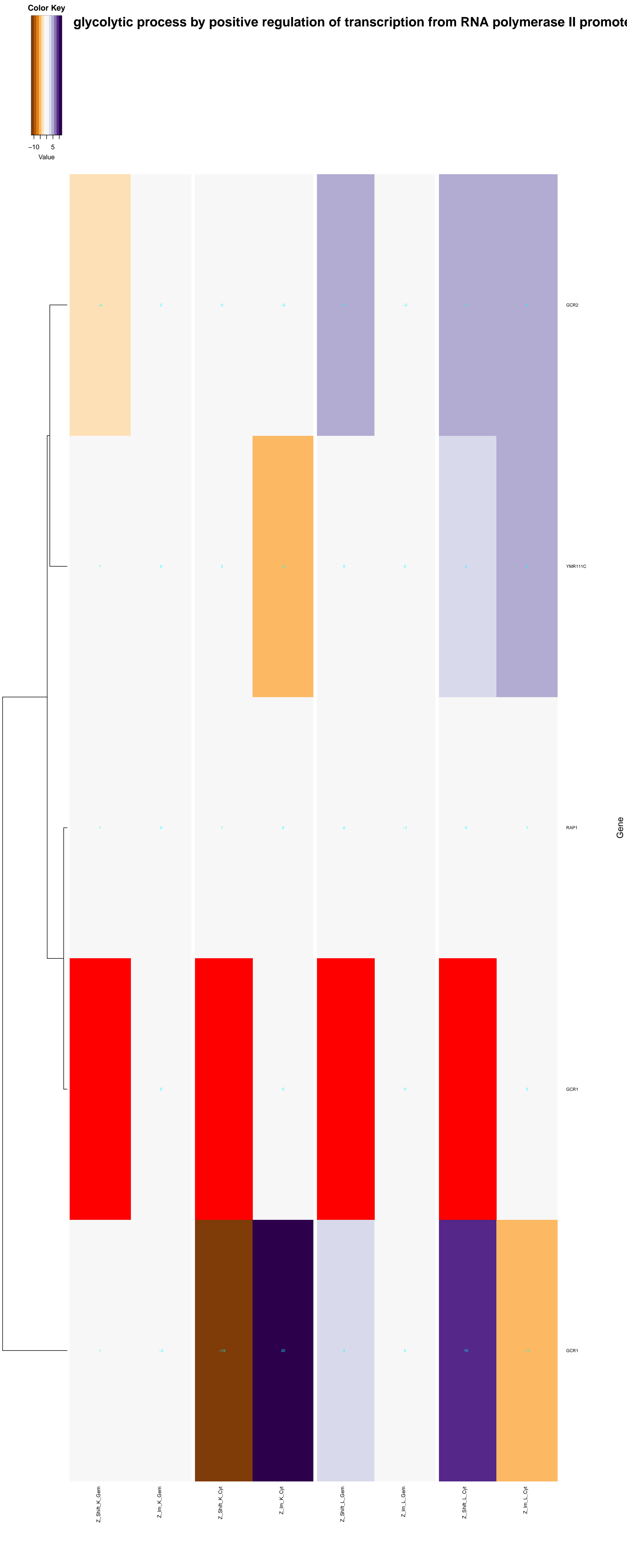


positive regulation of transcription initiation from RNA polymerase II promoter

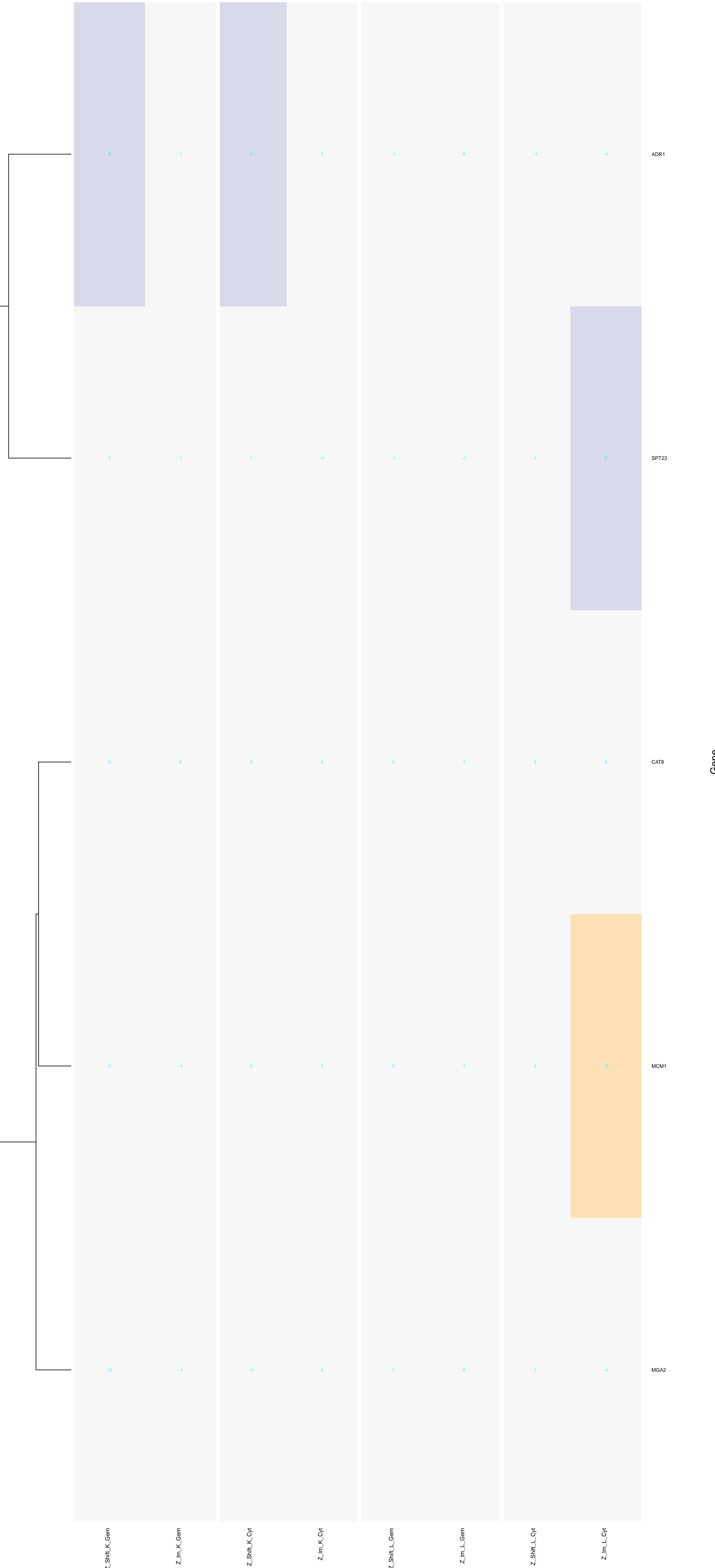
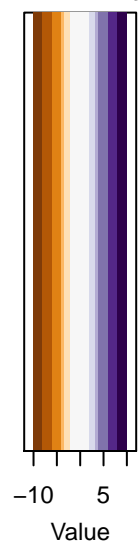


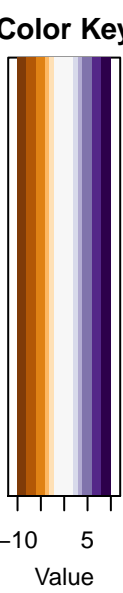




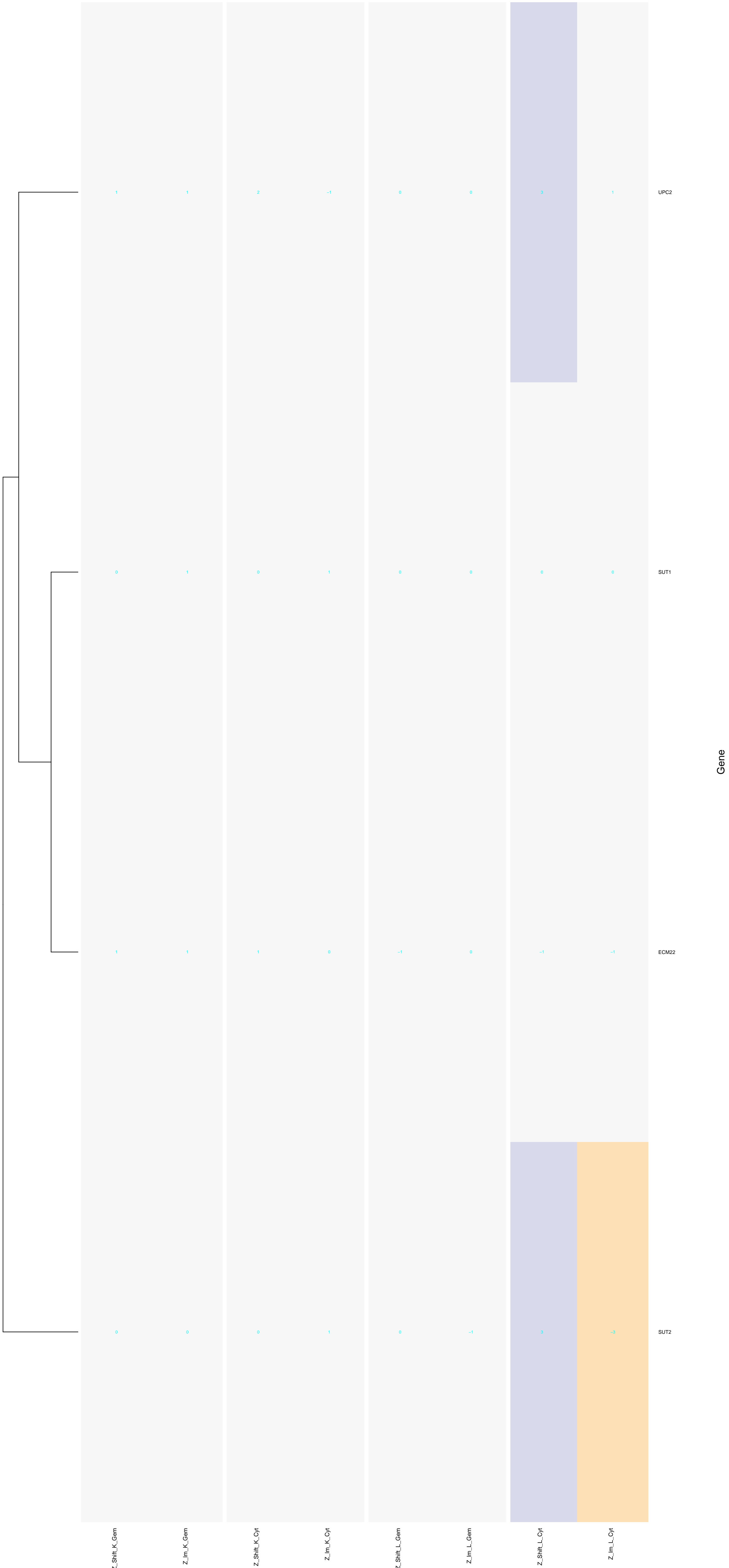


ketone metabolic process by positive regulation of transcription from RNA polymerase II





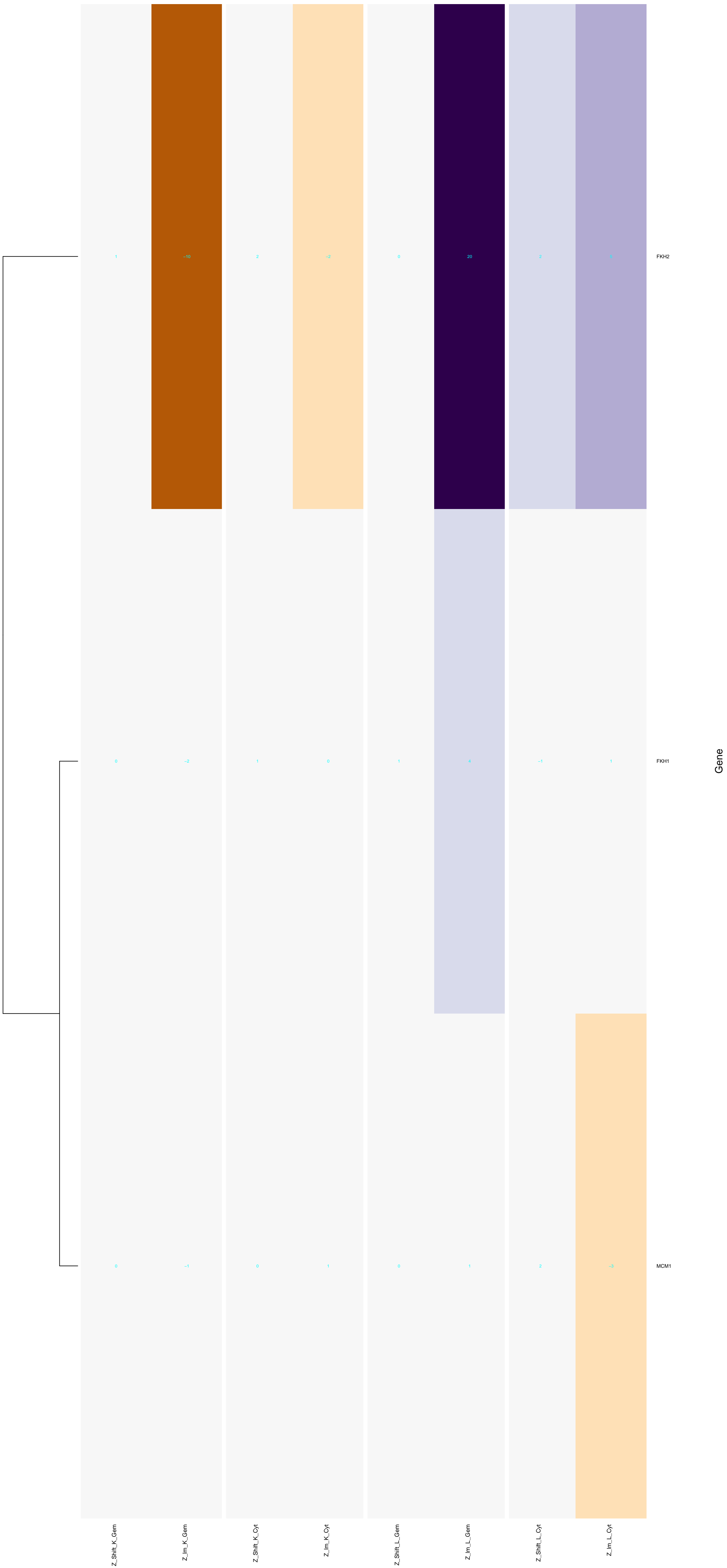
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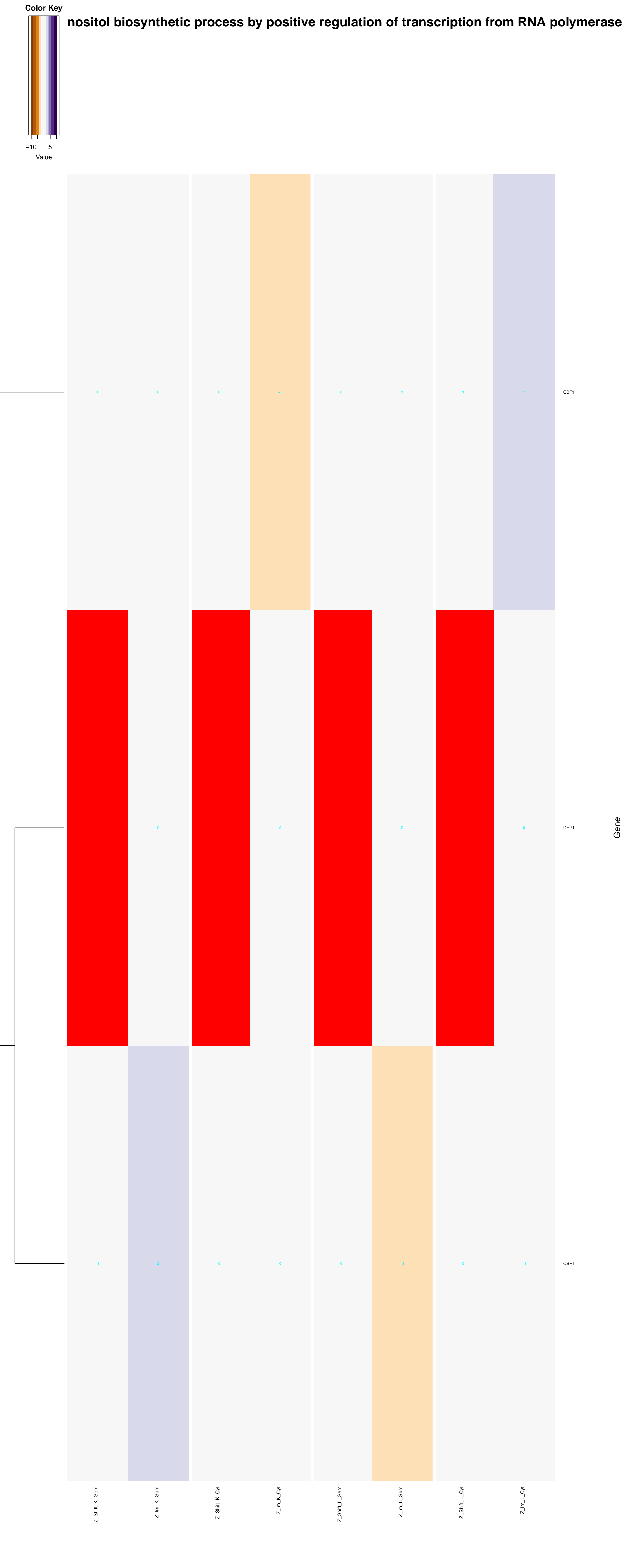






Positive regulation of transcription involved in G2/M transition of mitotic cell cycle



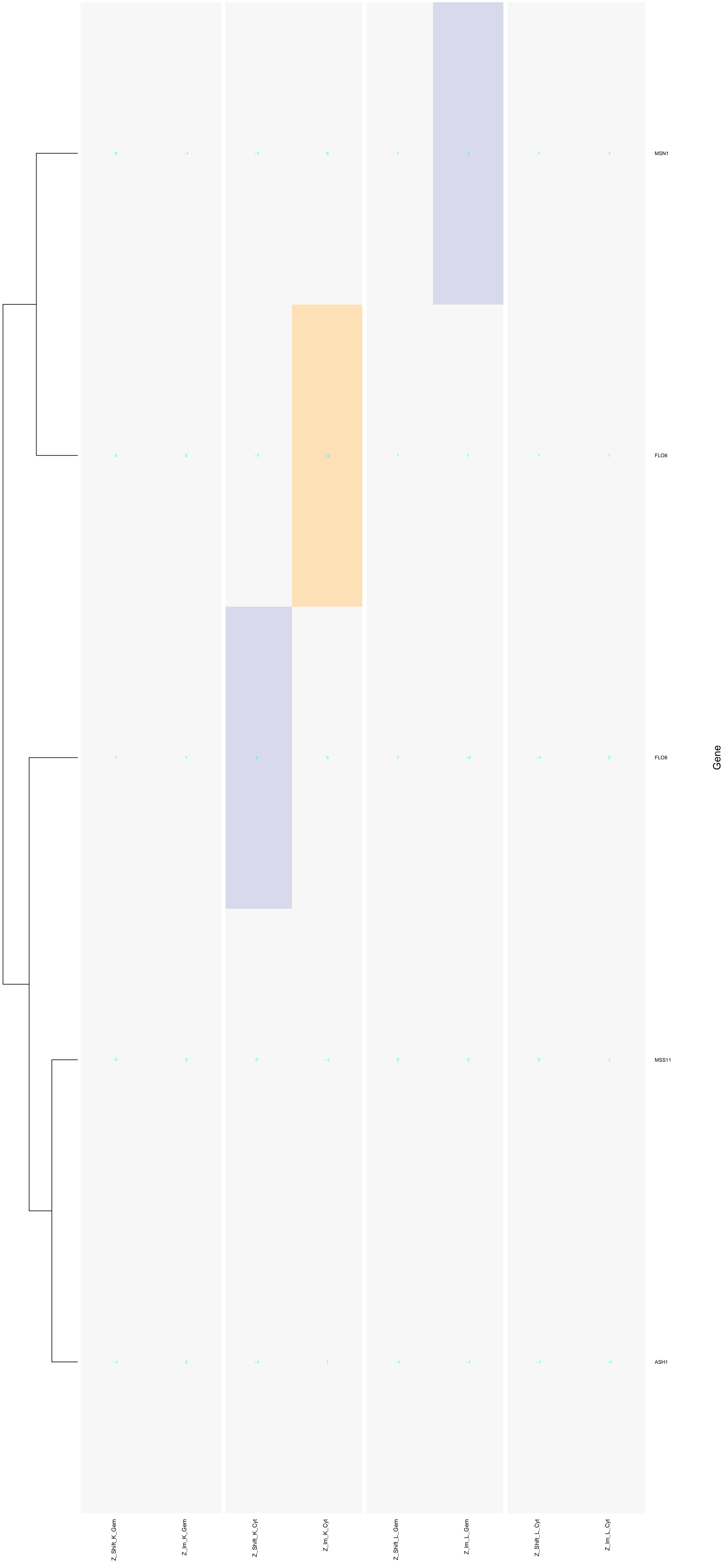


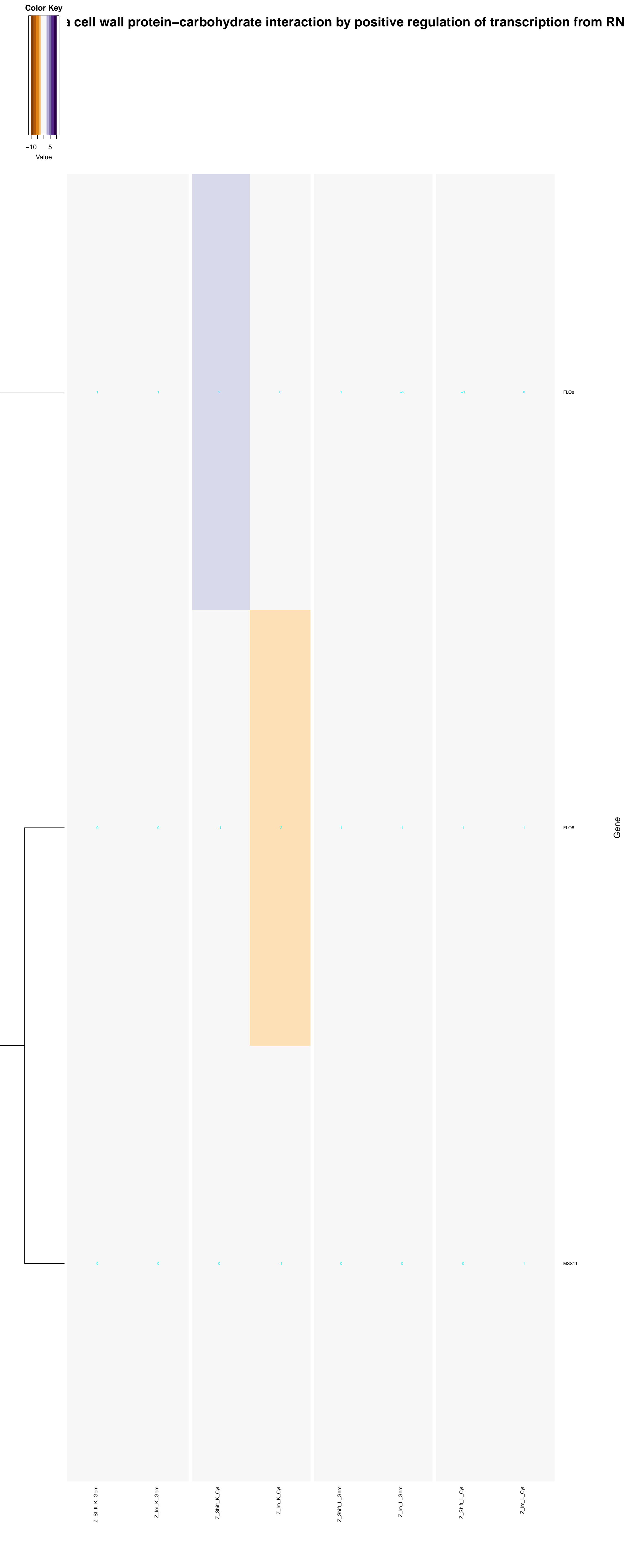
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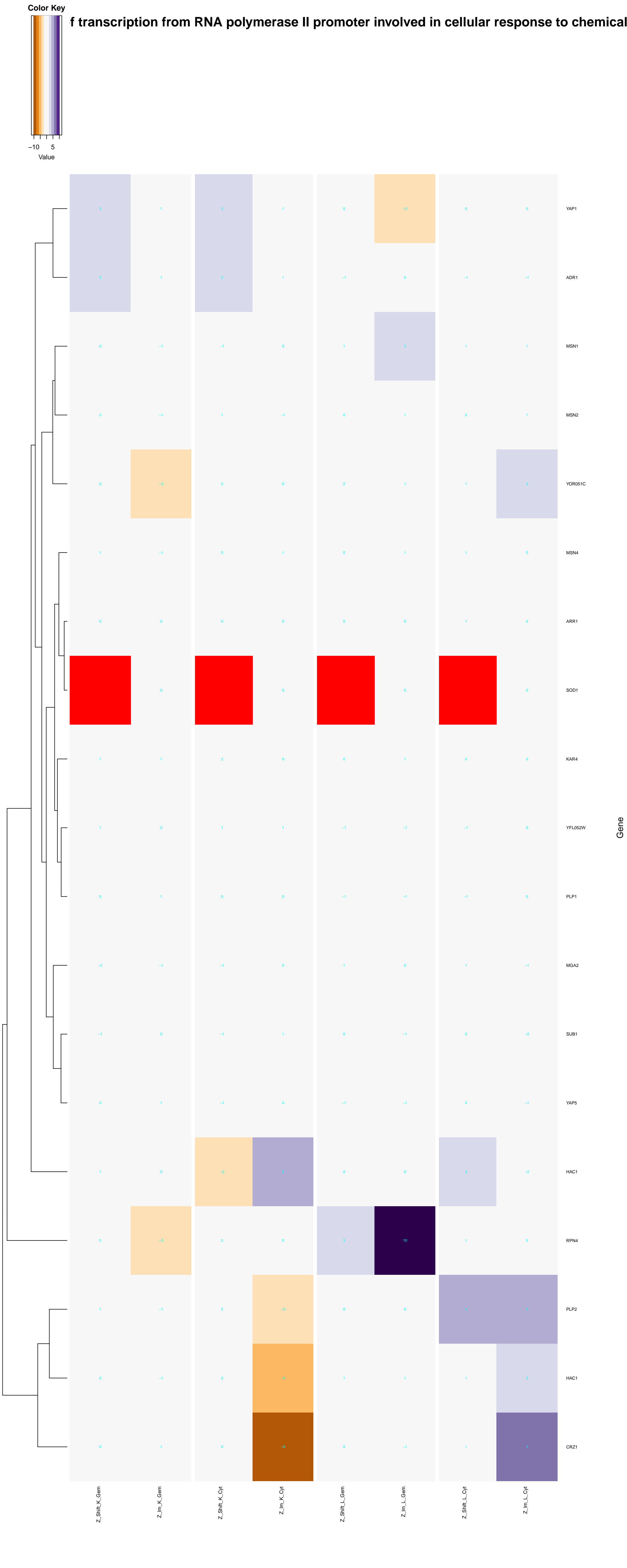
of pseudohyphal growth by positive regulation of transcription from RNA polymerase II pr

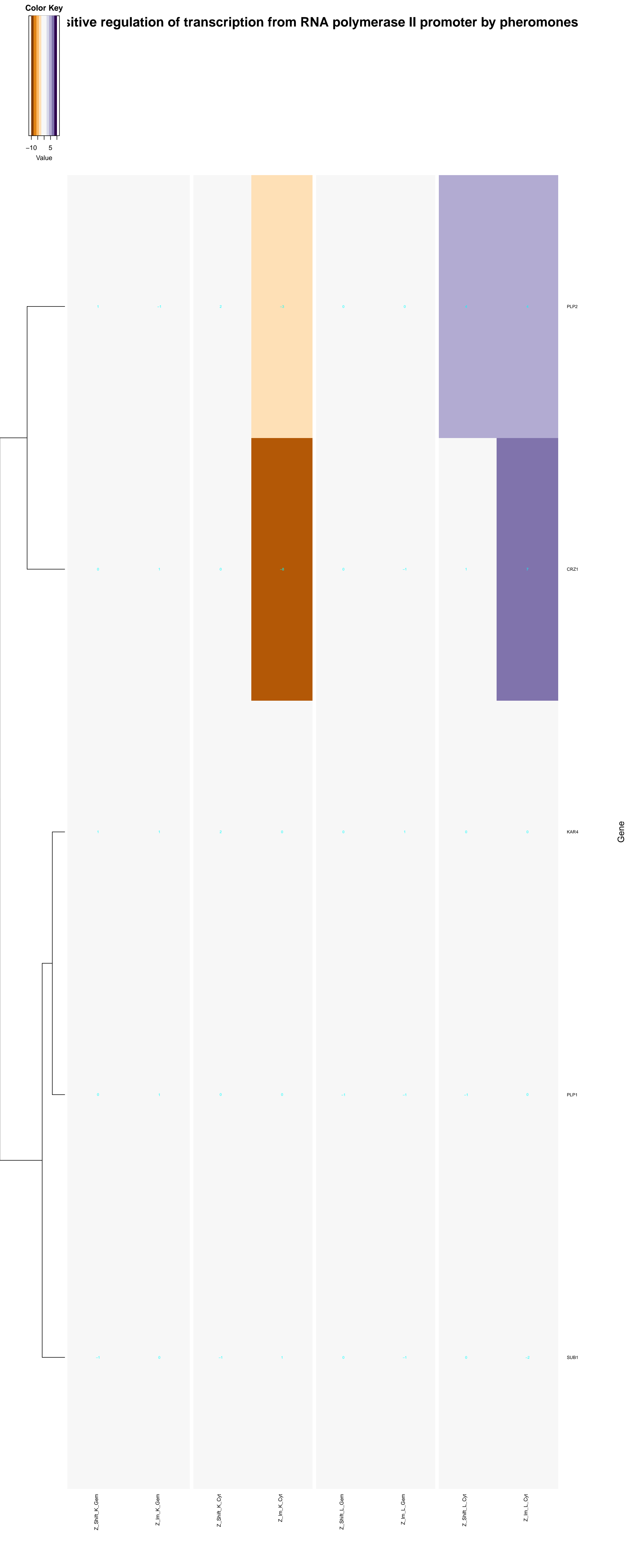
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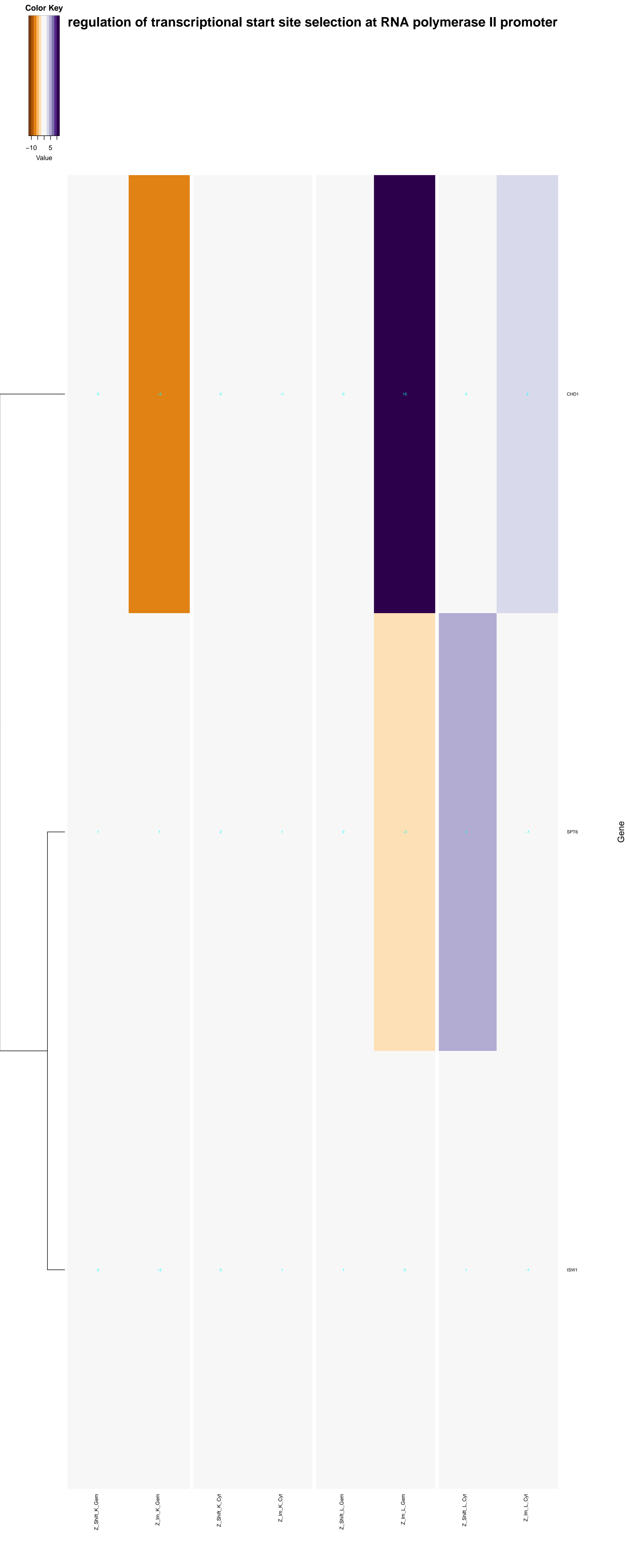
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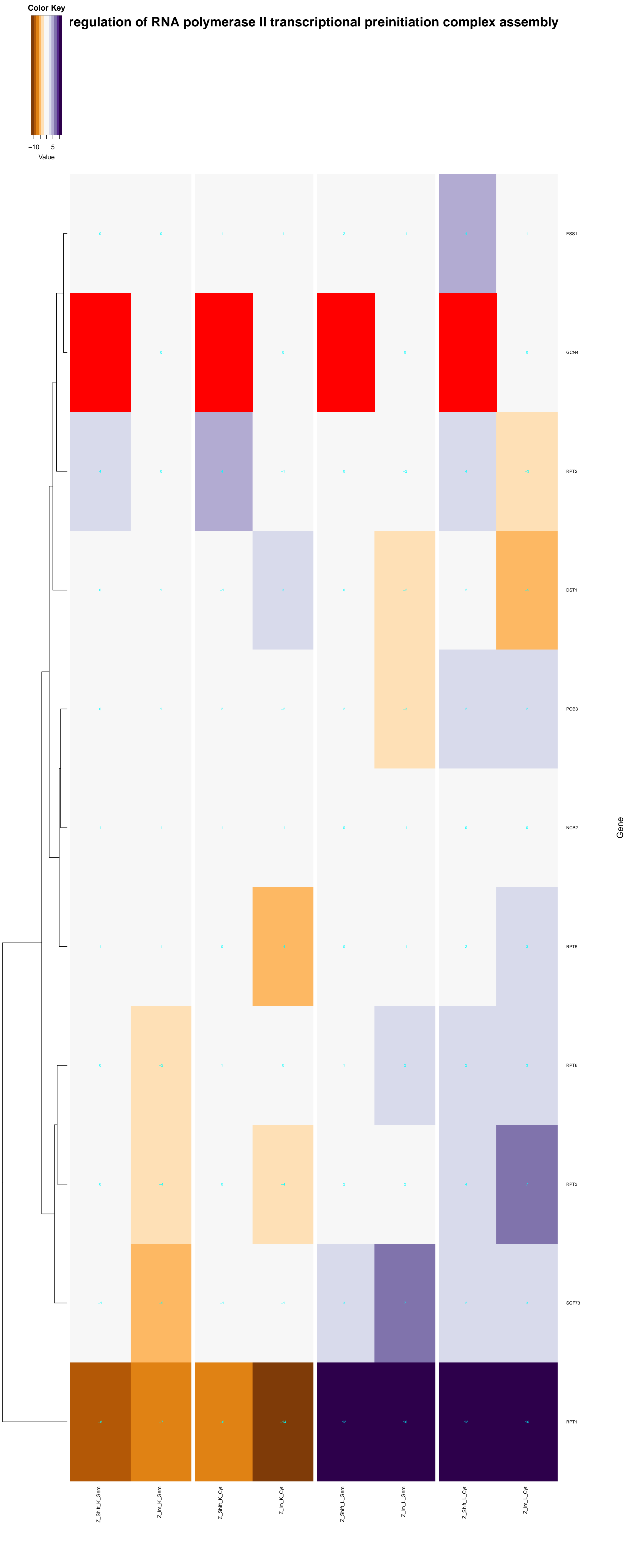




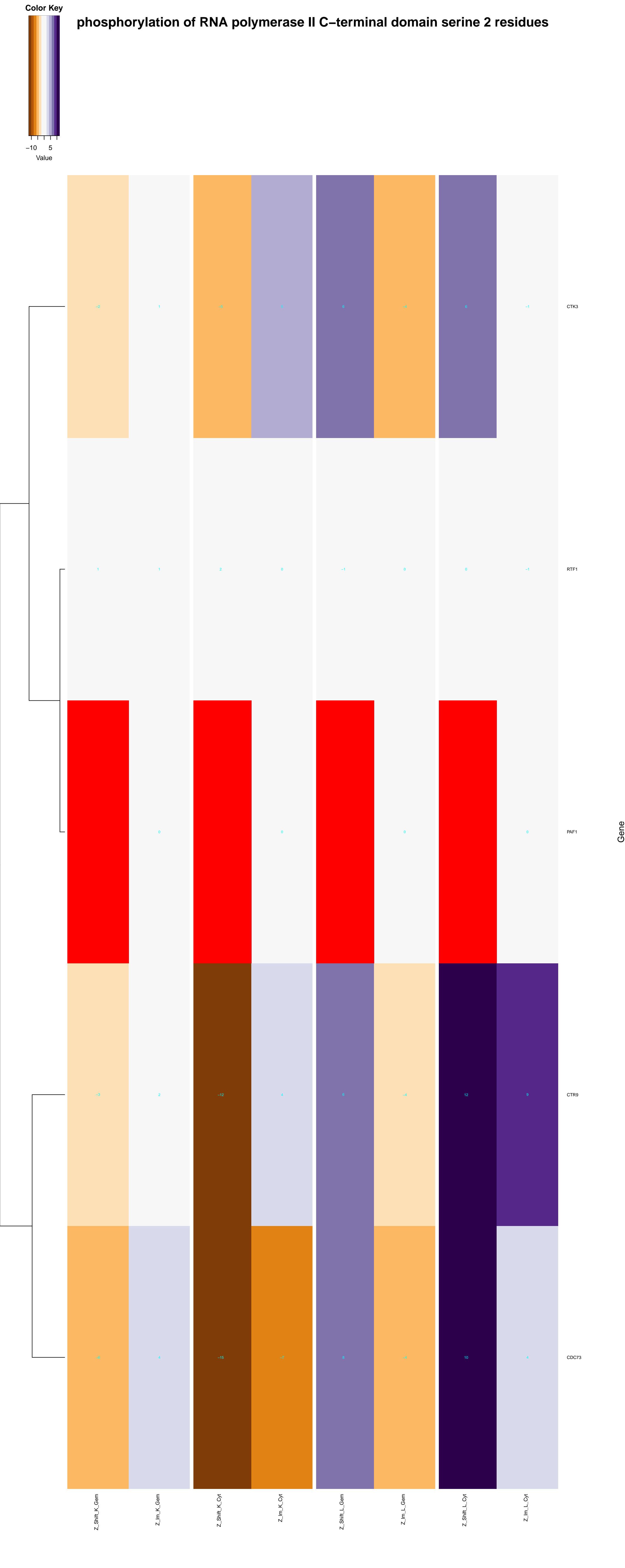


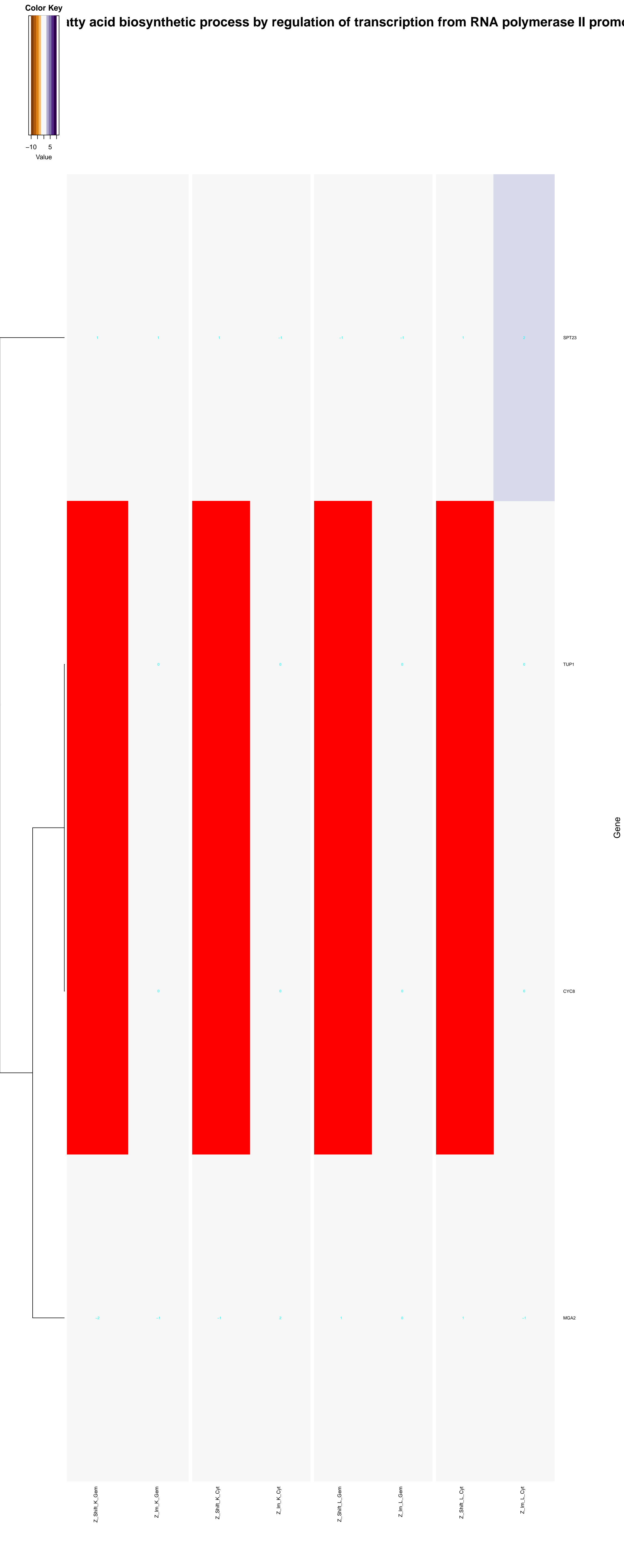


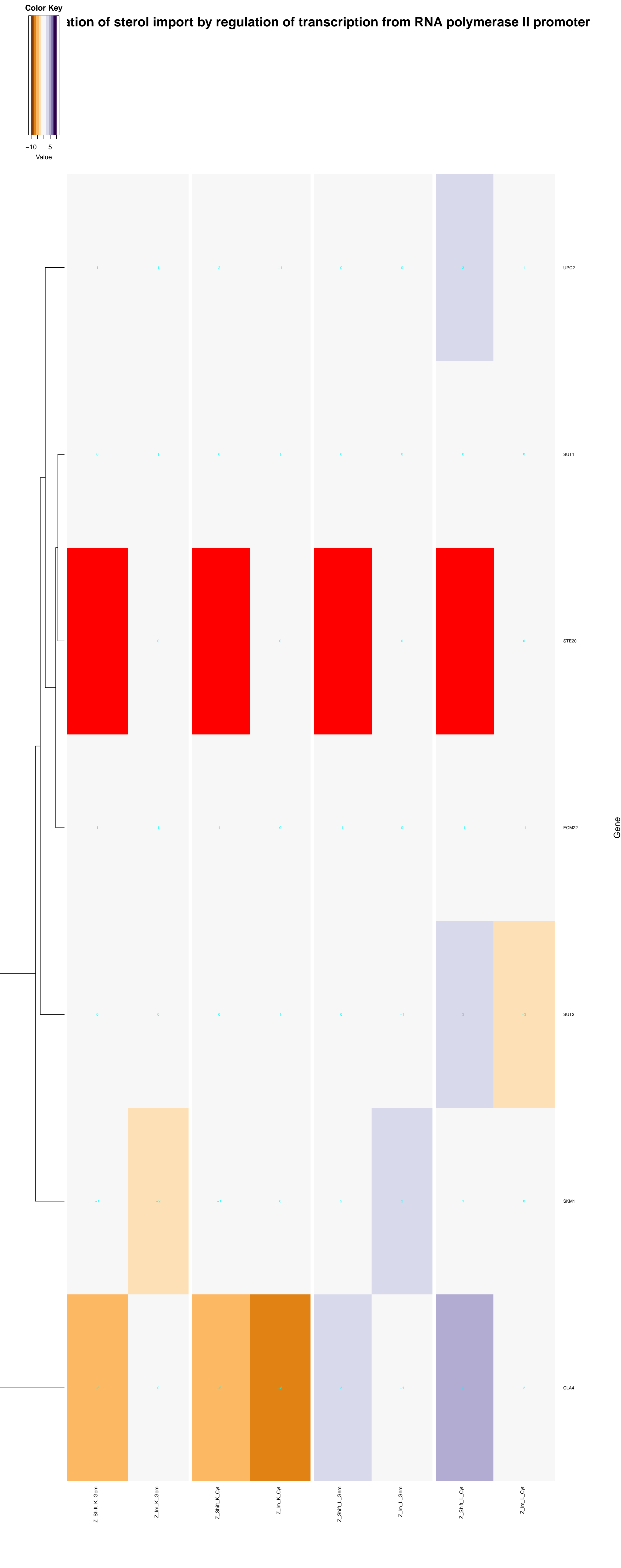


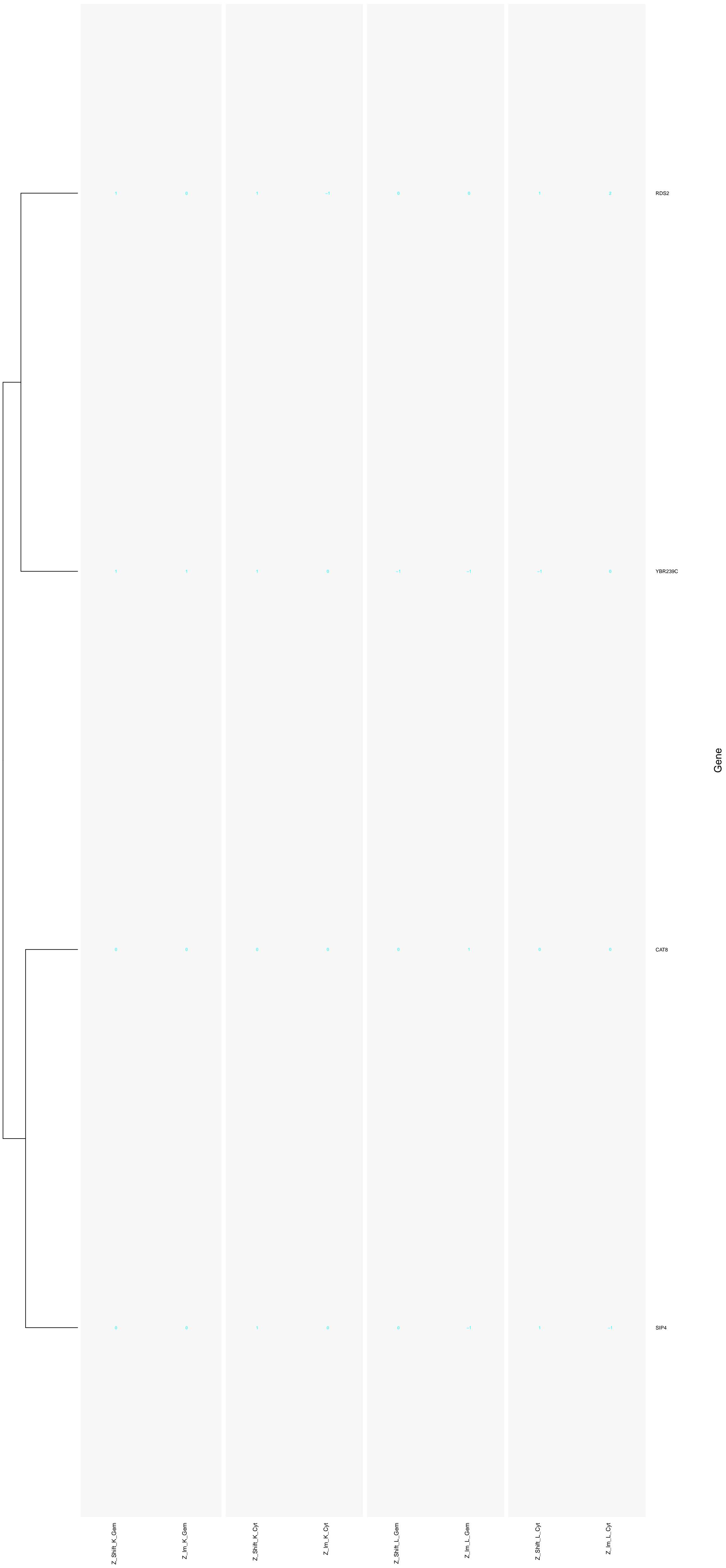
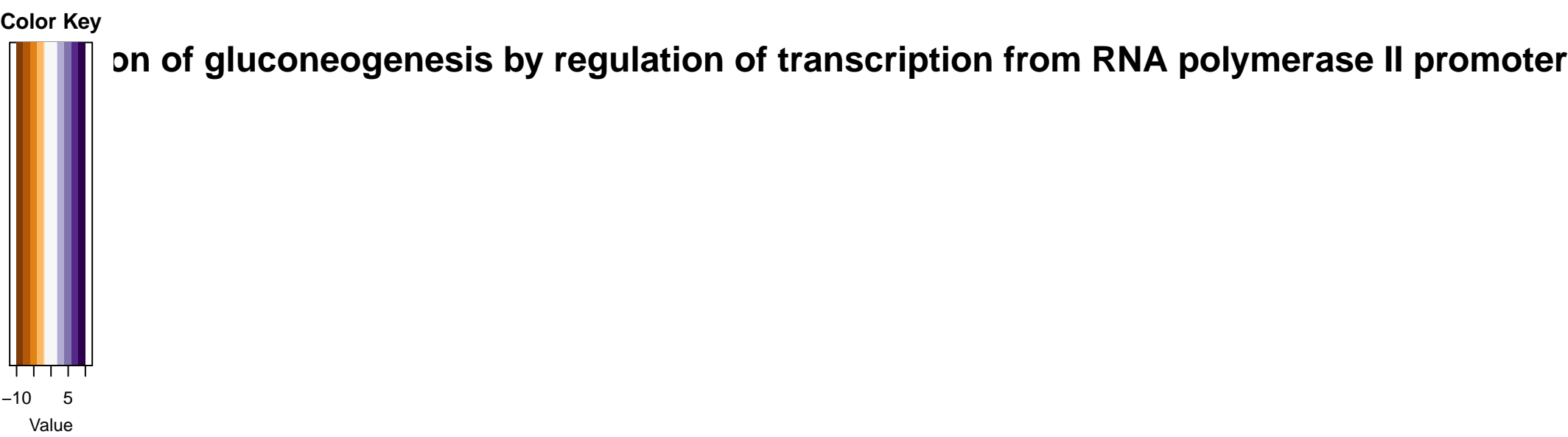


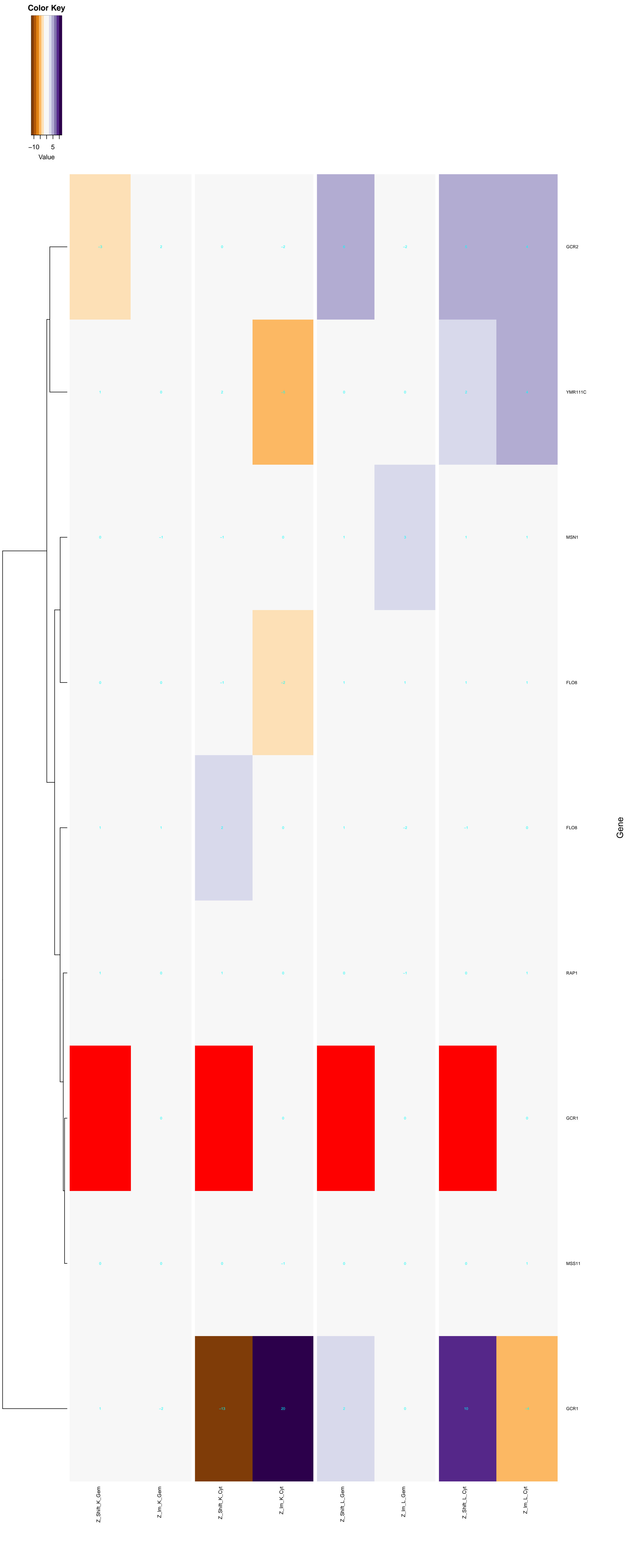


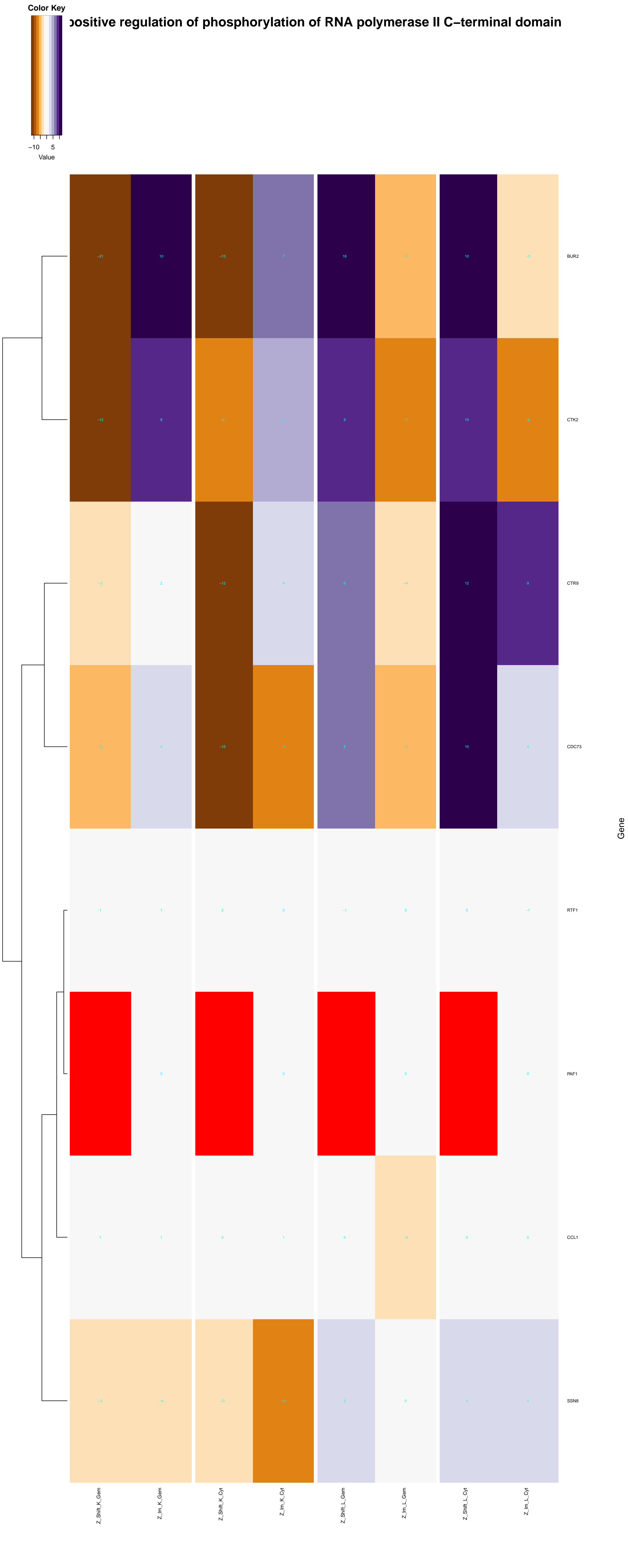


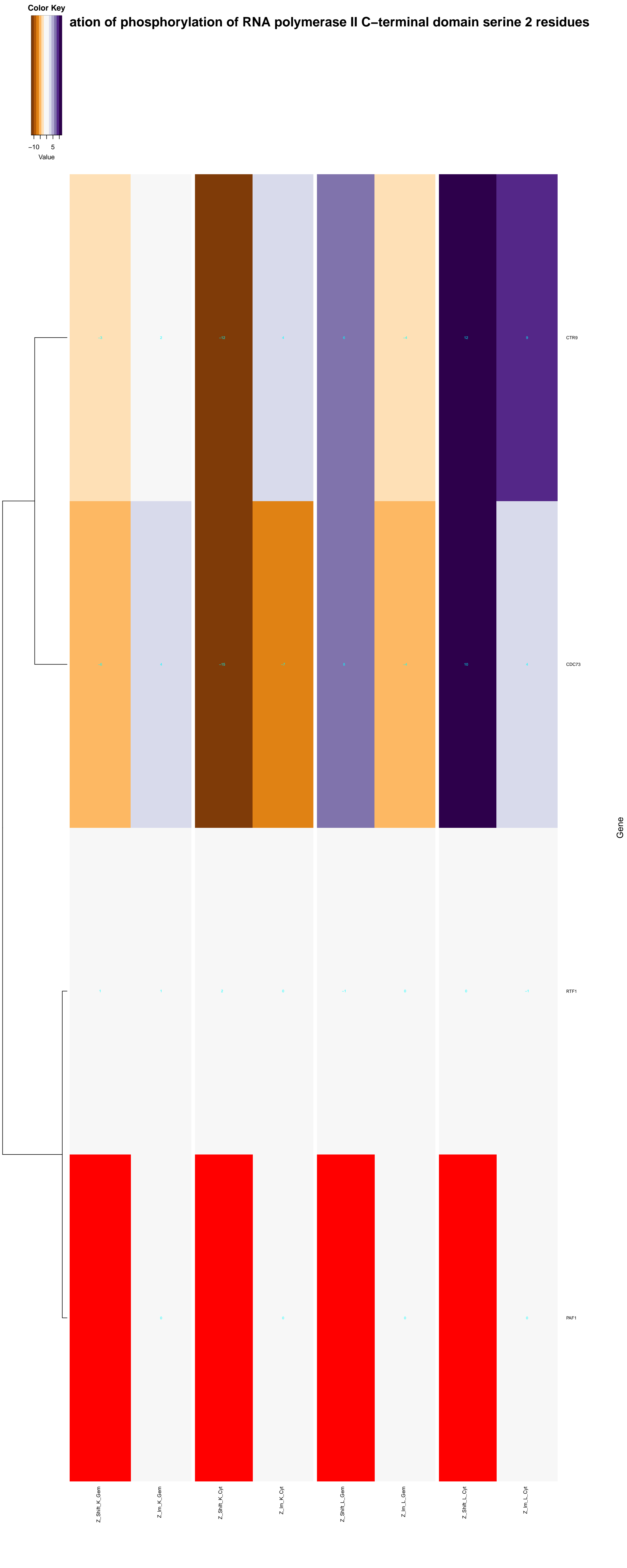






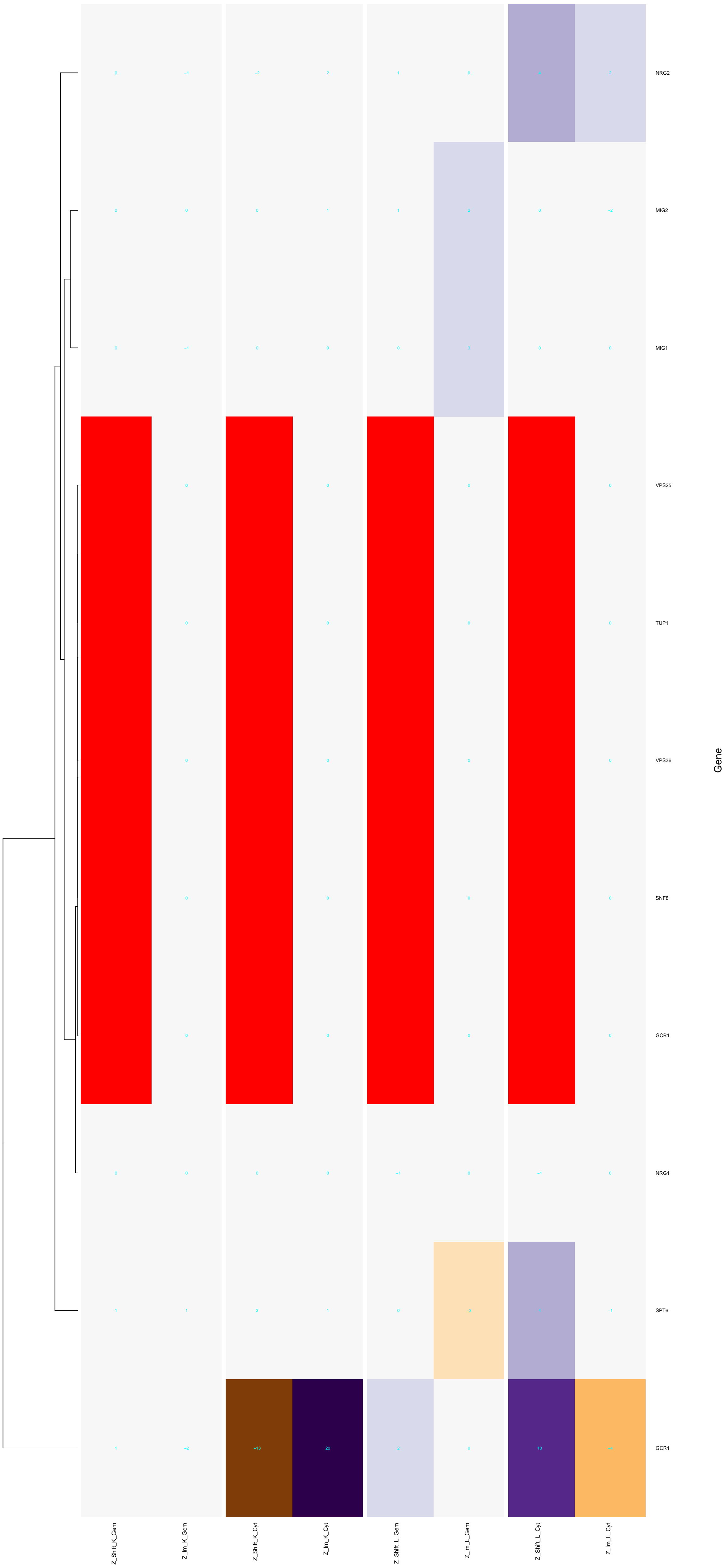




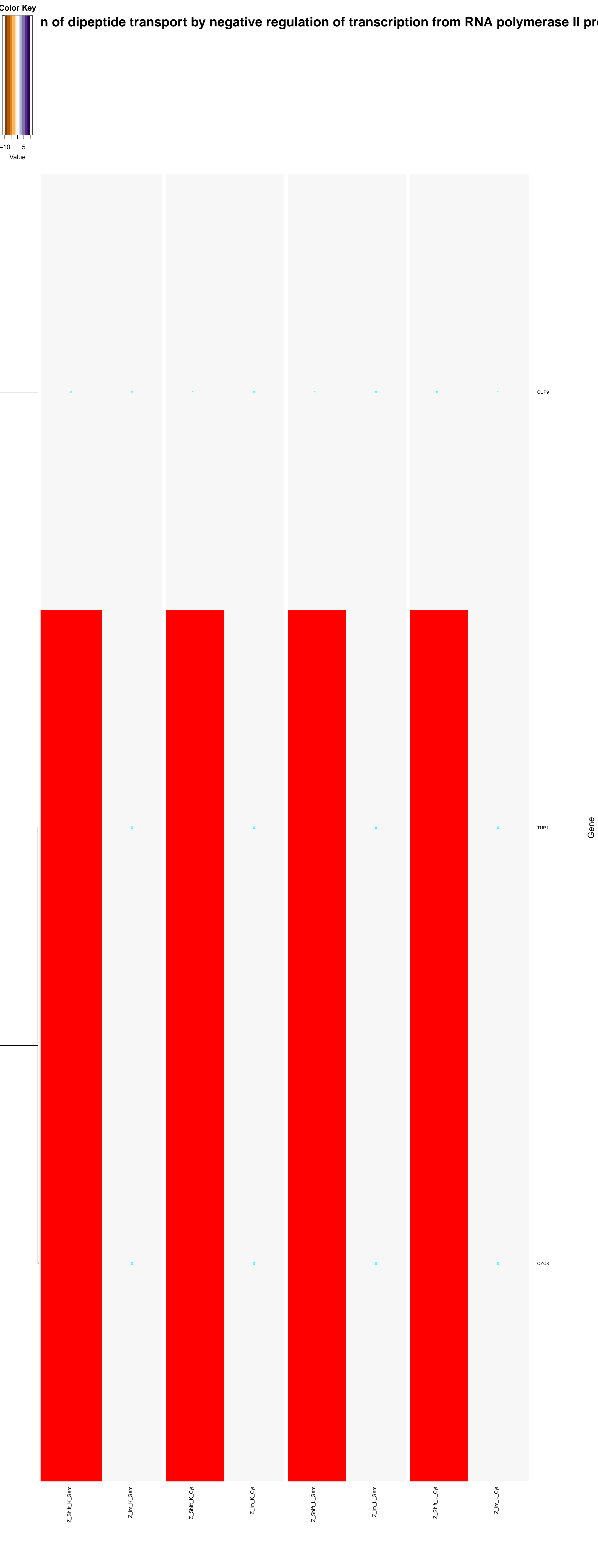


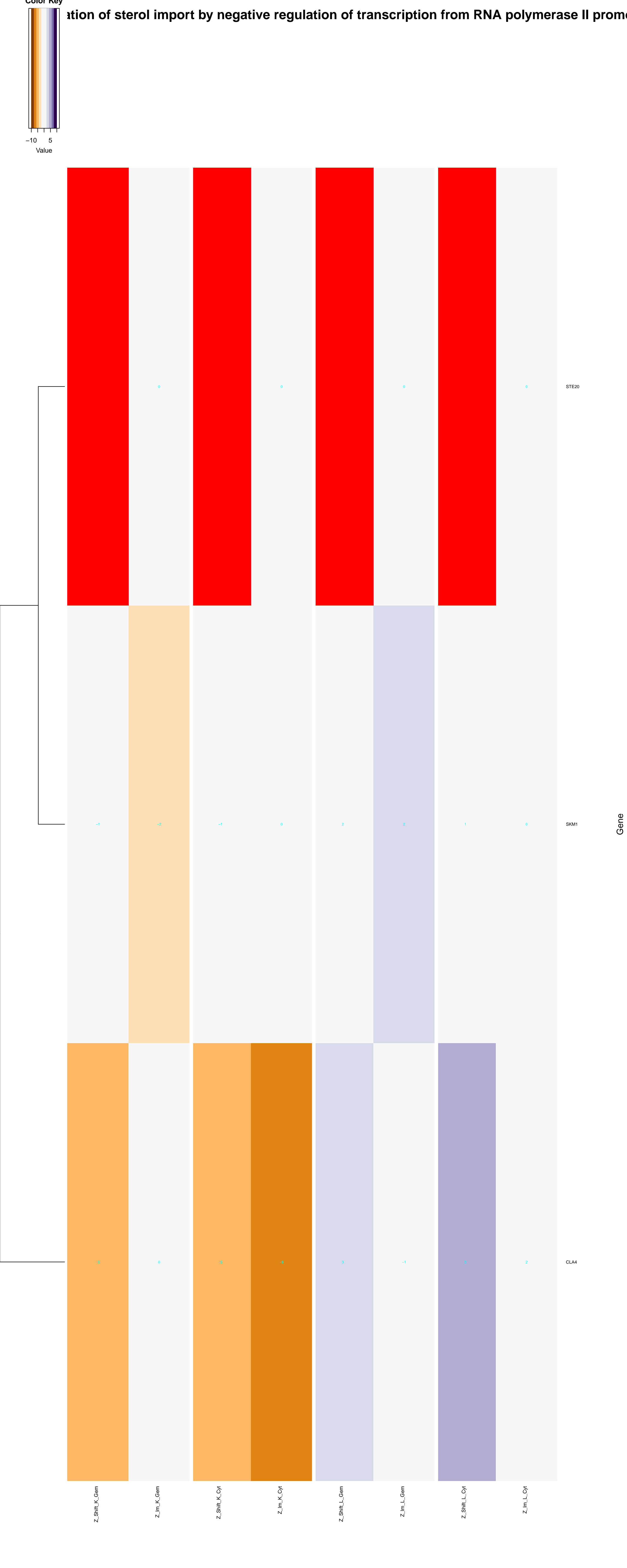


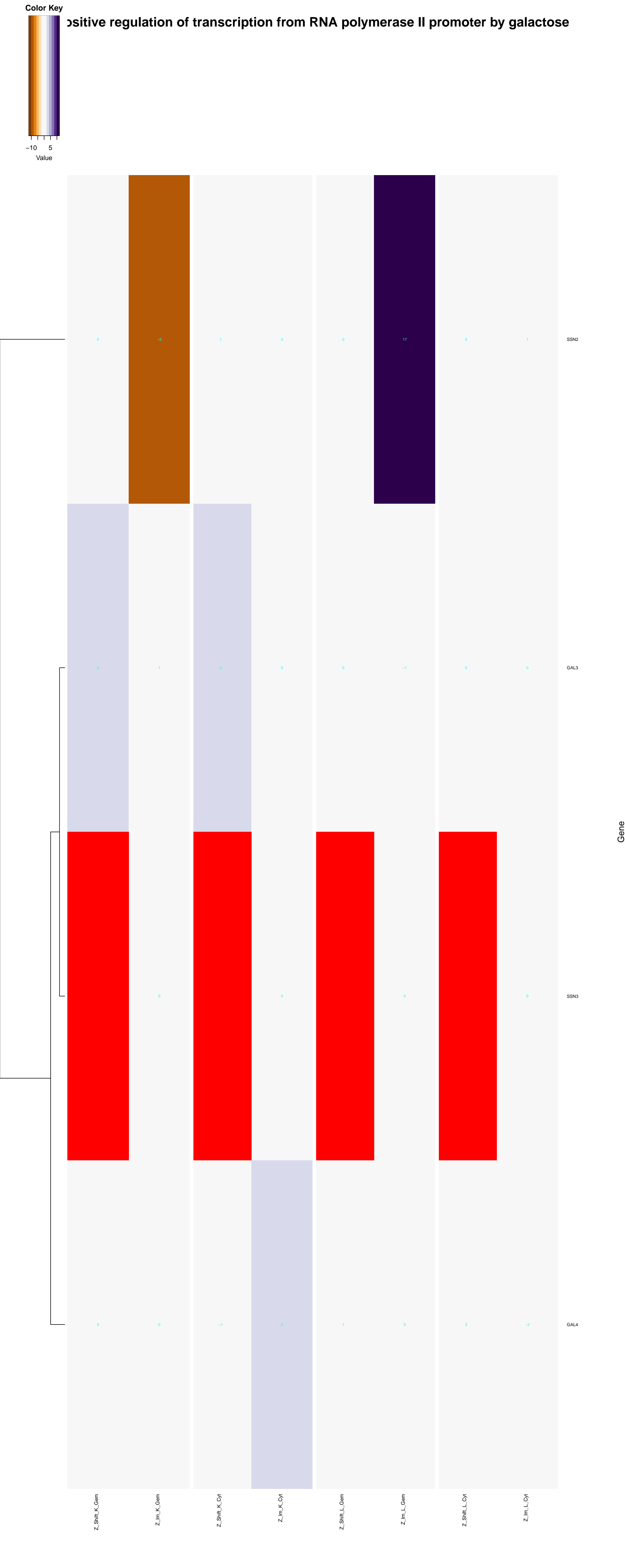
egative regulation of transcription from RNA polymerase II promoter by glucose









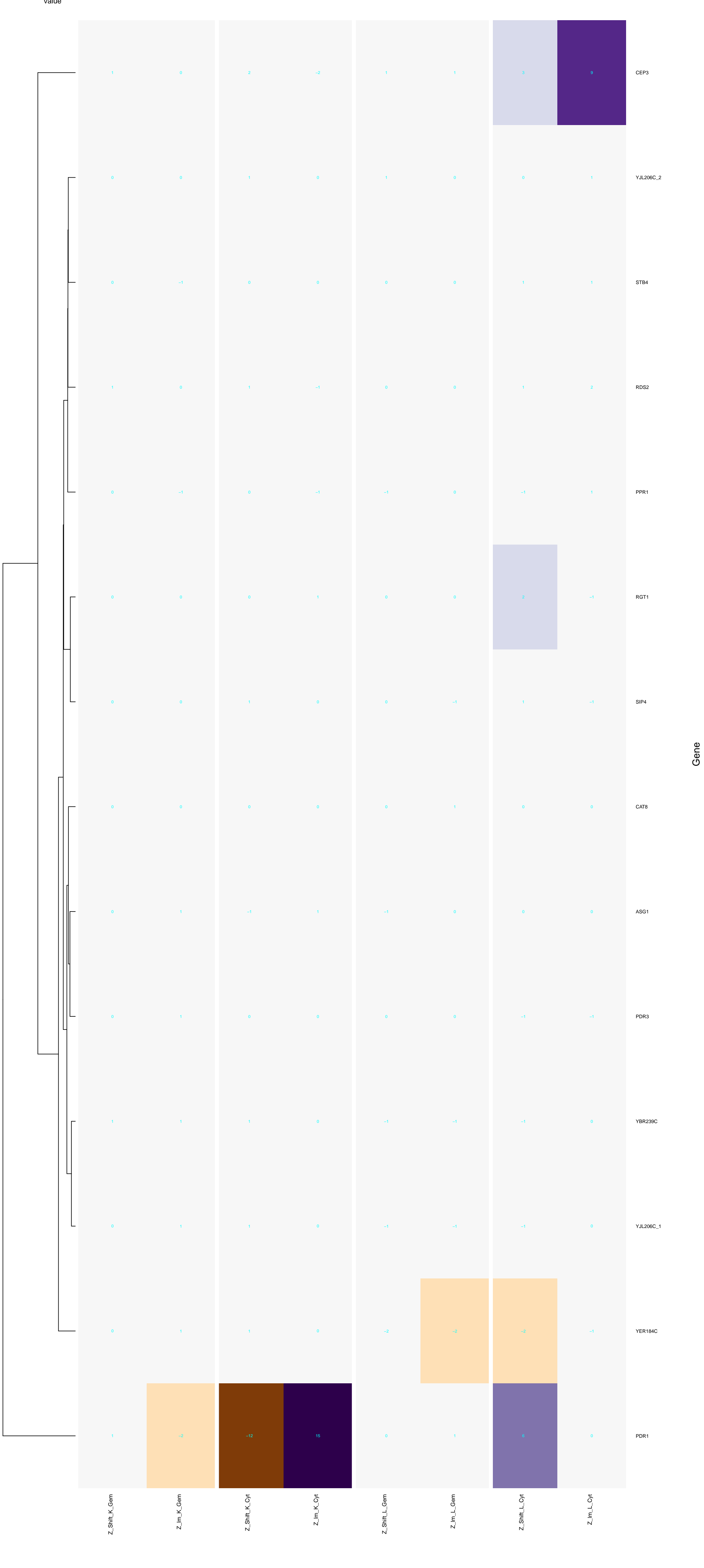


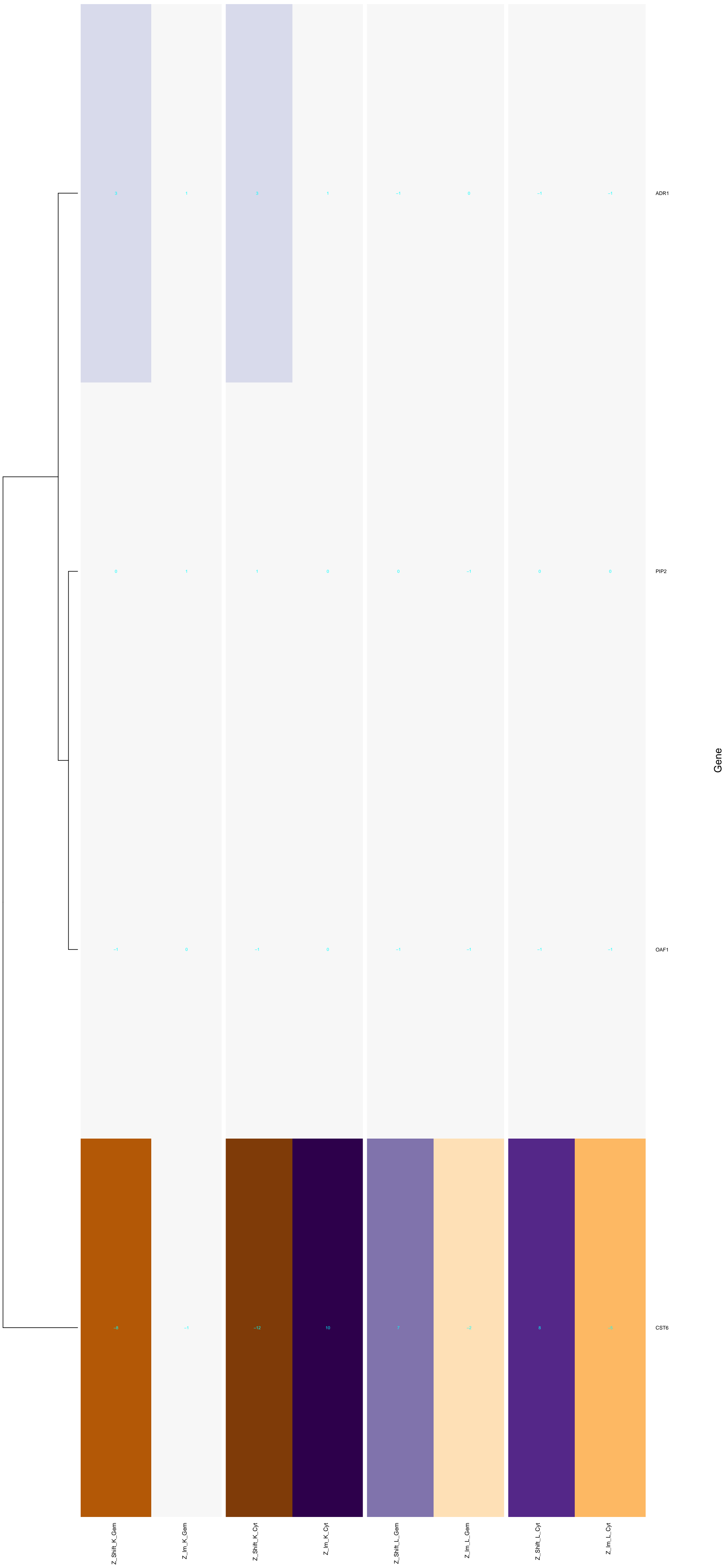
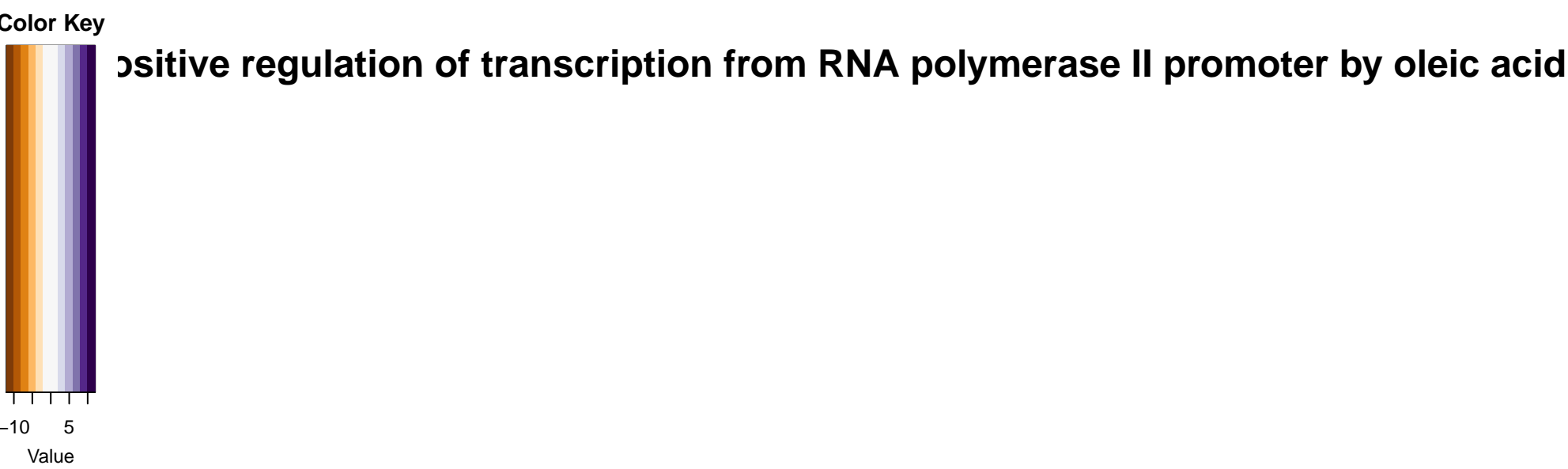
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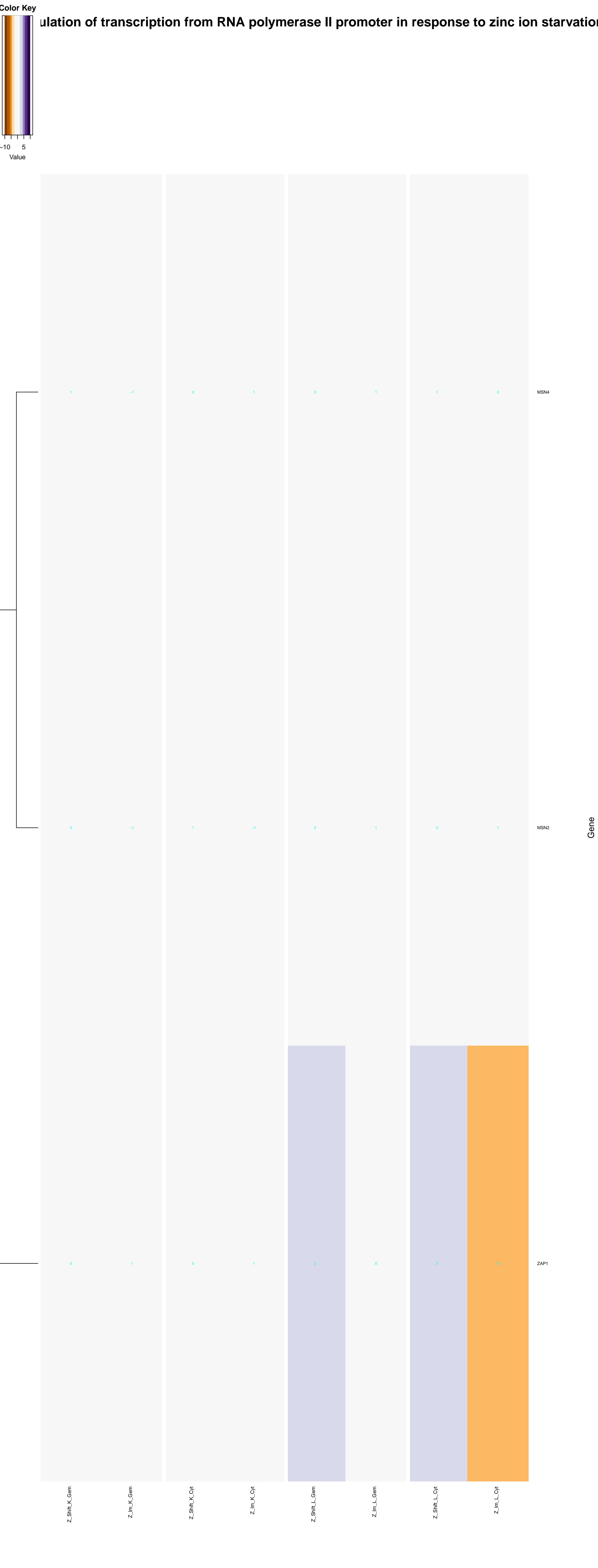
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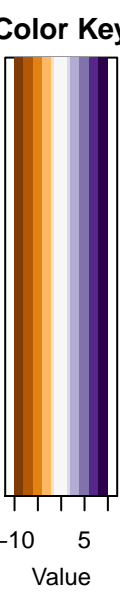
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lation of transcription from RNA polymerase II promoter by a nonfermentable carbon source

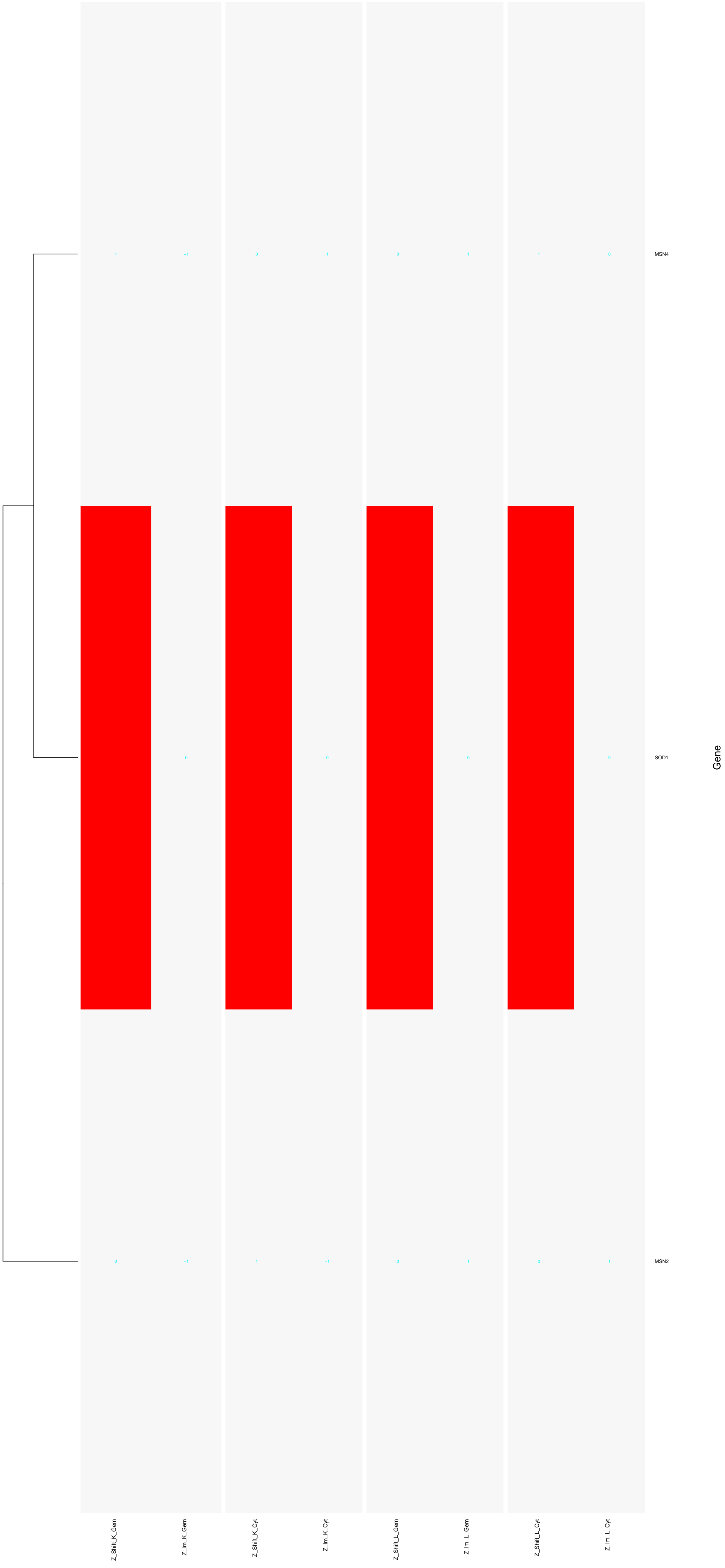








gulation of transcription from RNA polymerase II promoter in response to oxidative stress



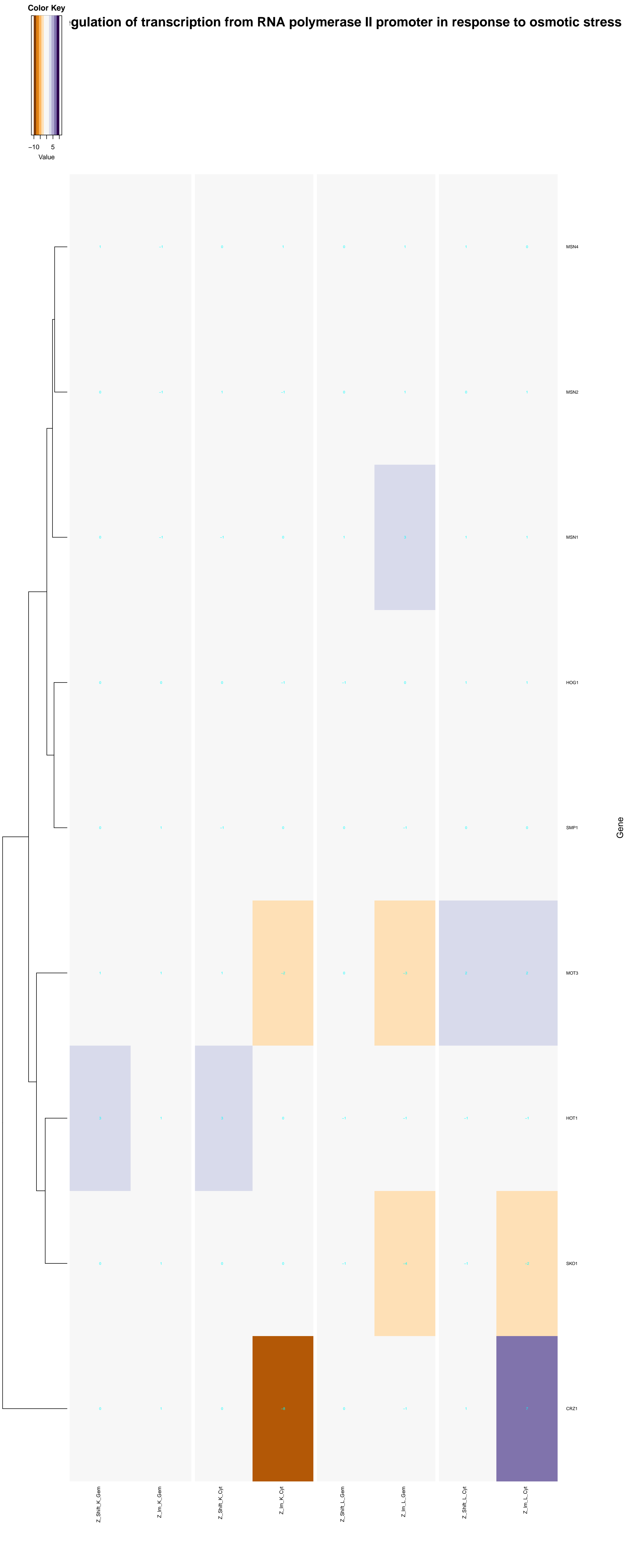
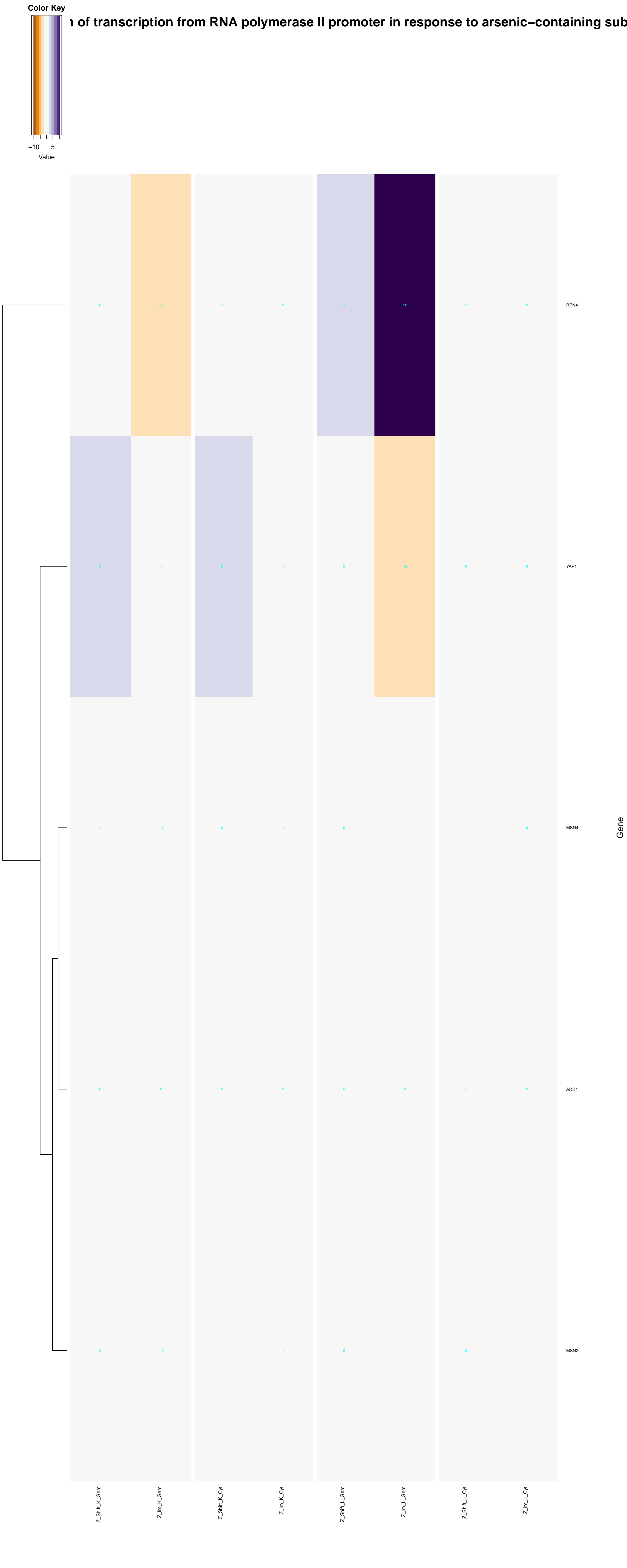
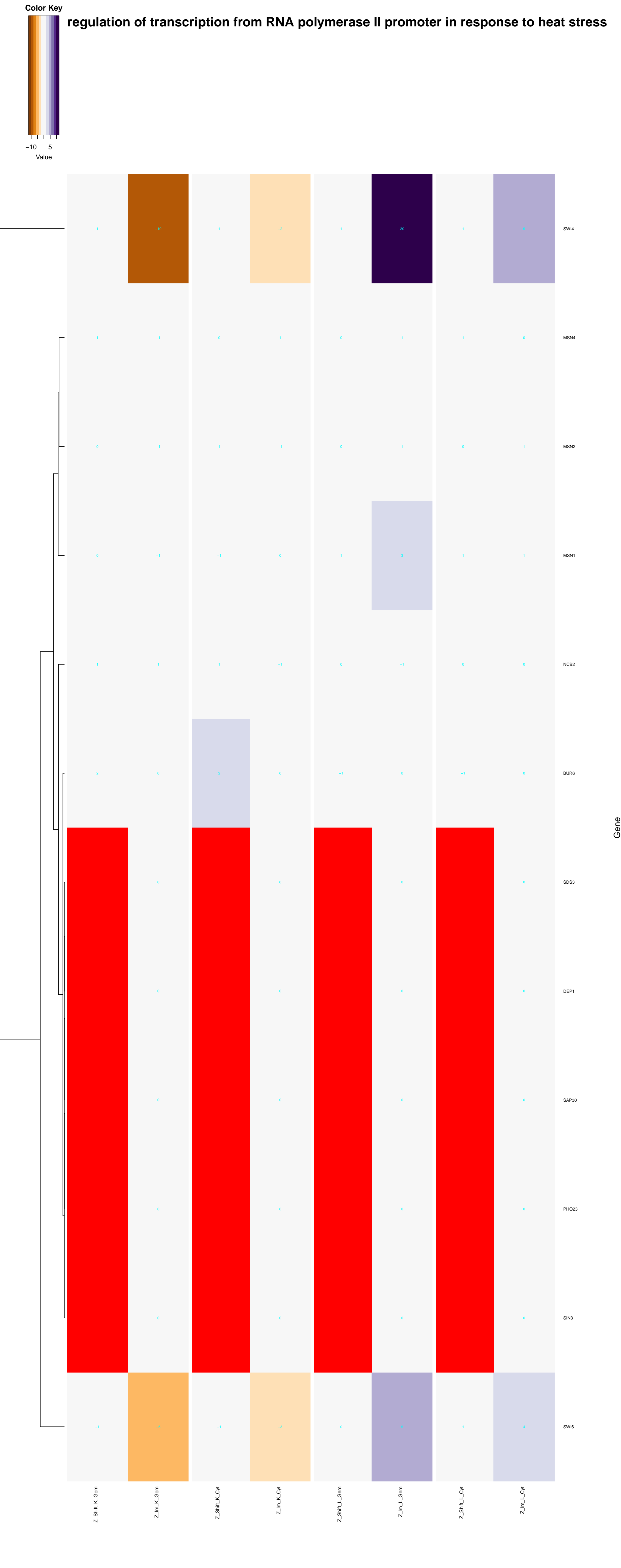
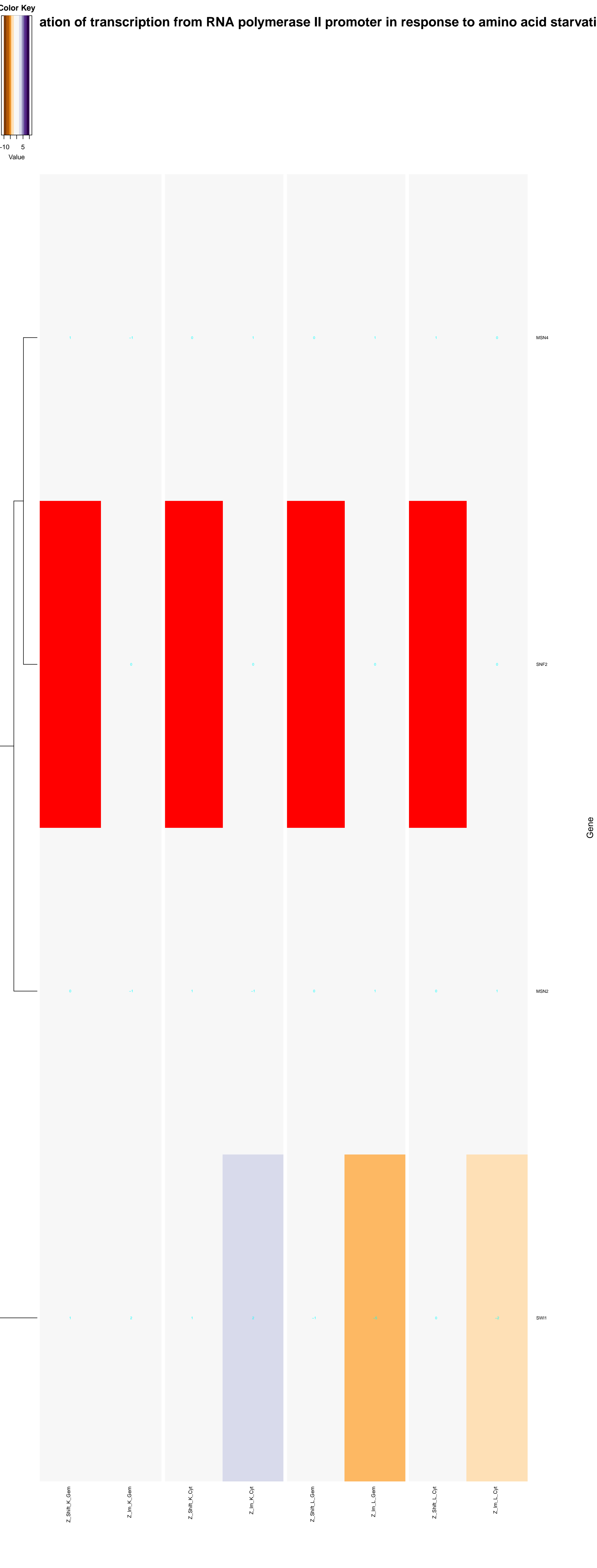


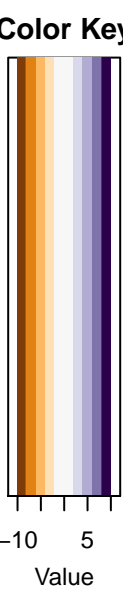


Figure 1: Heatmap showing the log2 fold change of transcription from RNA polymerase II promoter in response to arsenic-containing substances.

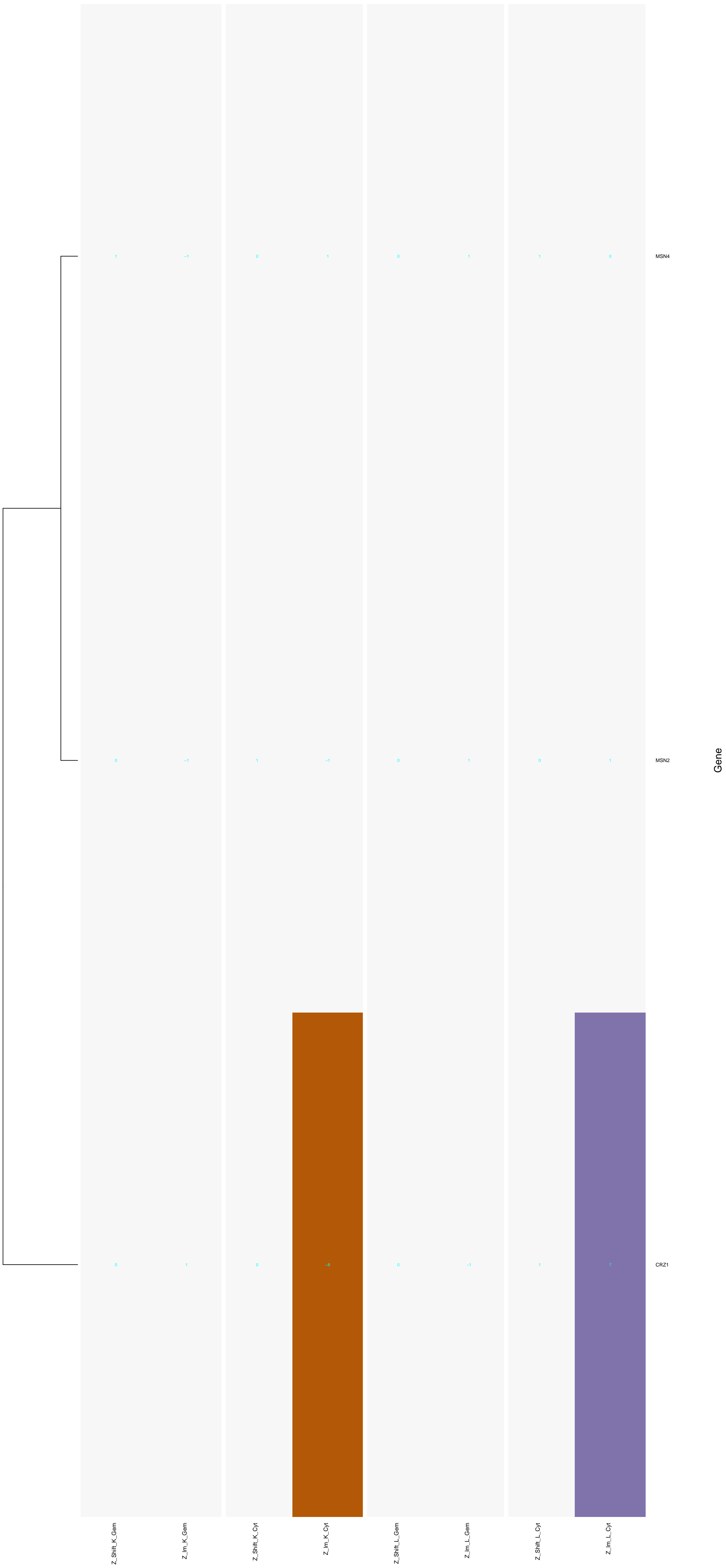








regulation of transcription from RNA polymerase II promoter in response to alkaline pH

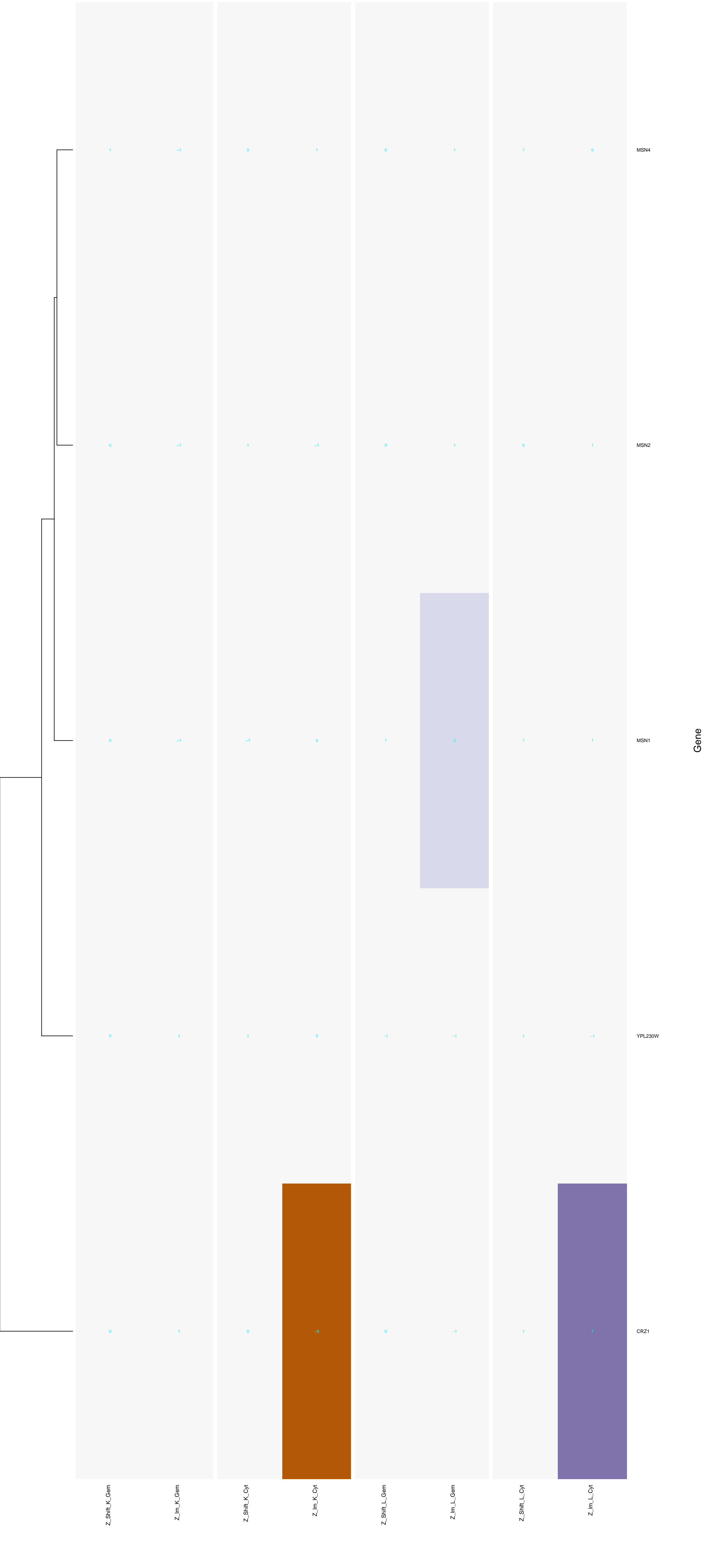


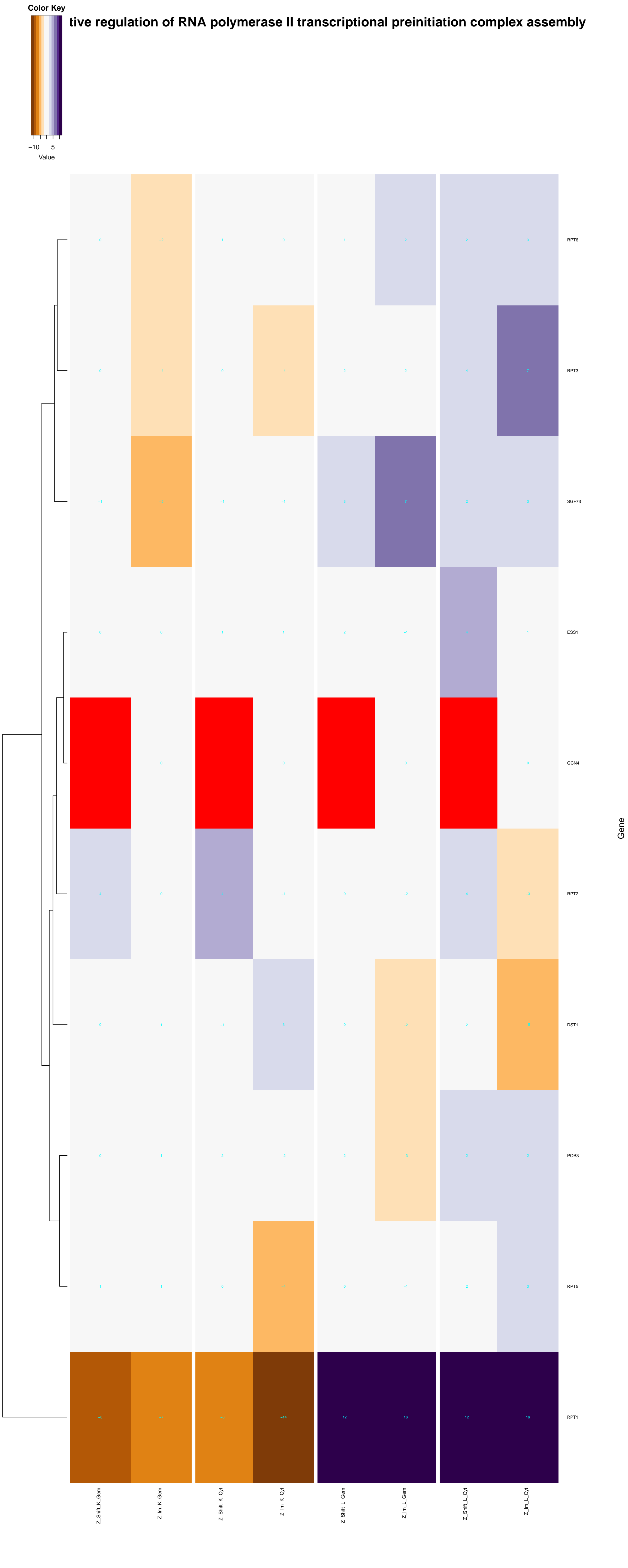
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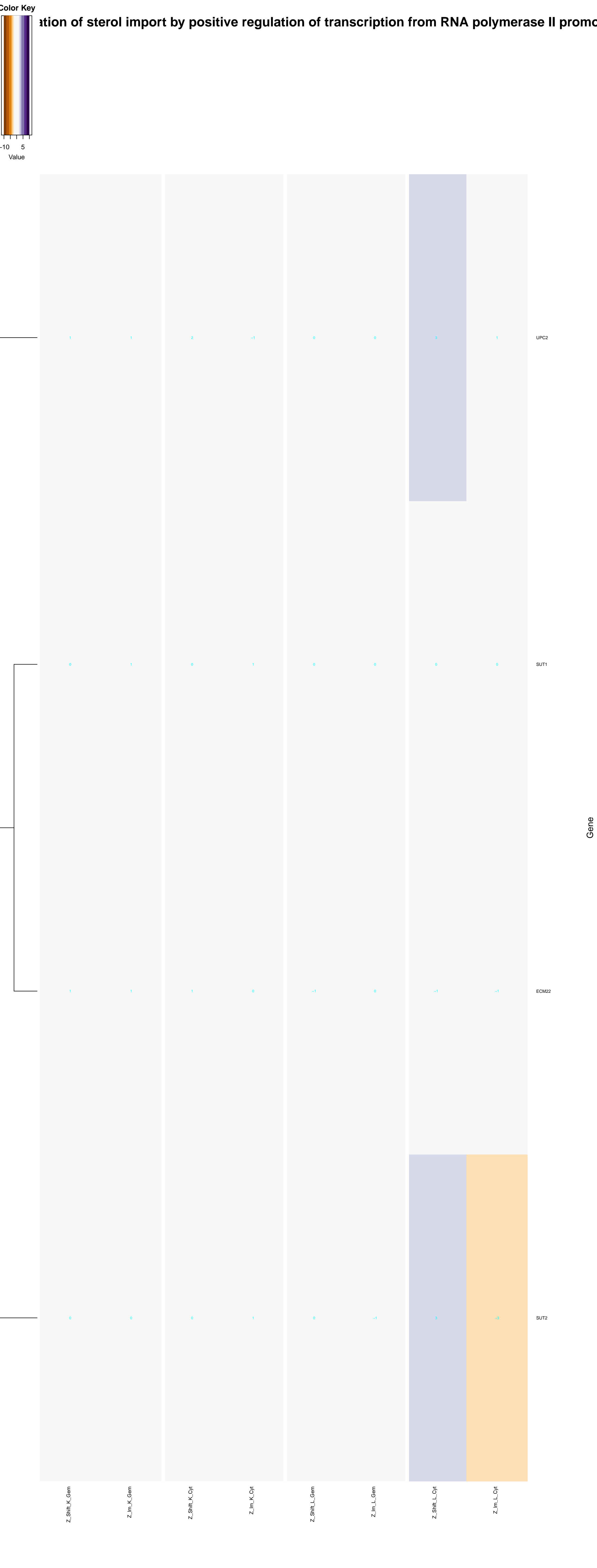
Correlation of transcription from RNA polymerase II promoter in response to salt stress

Value

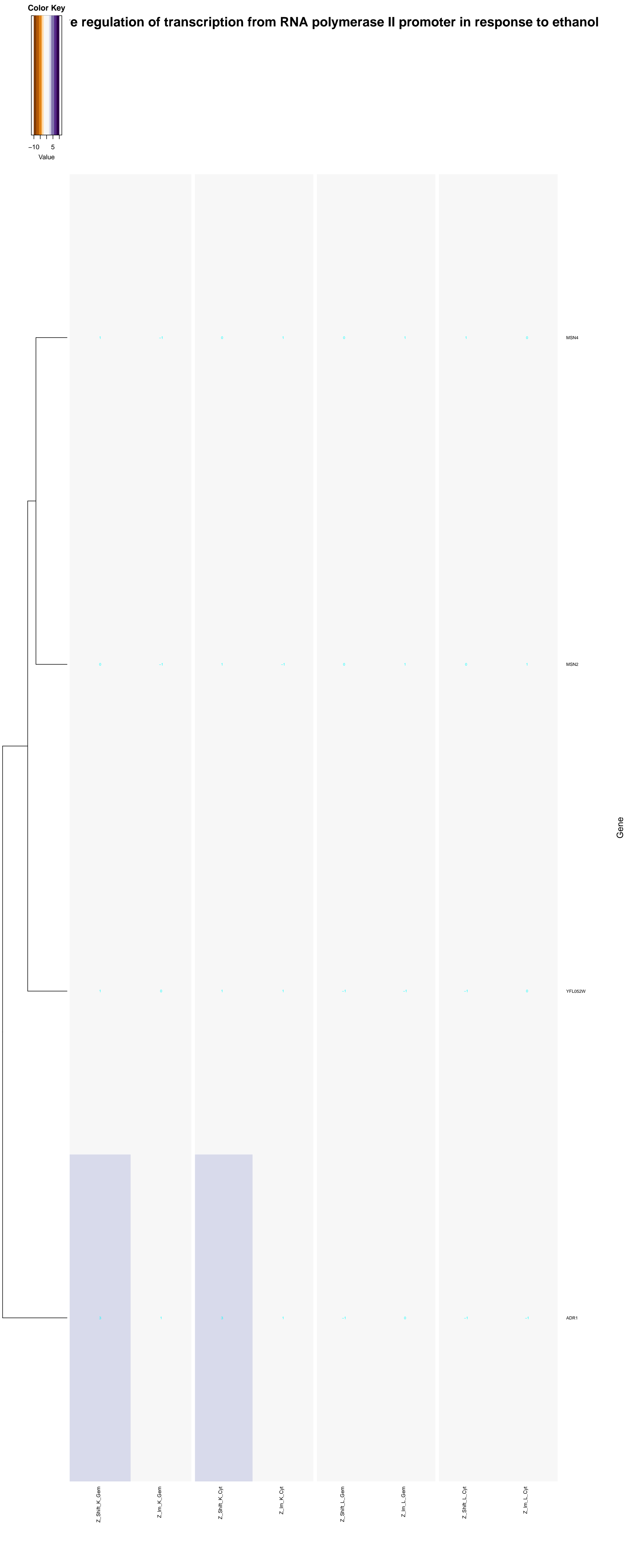
-10 5



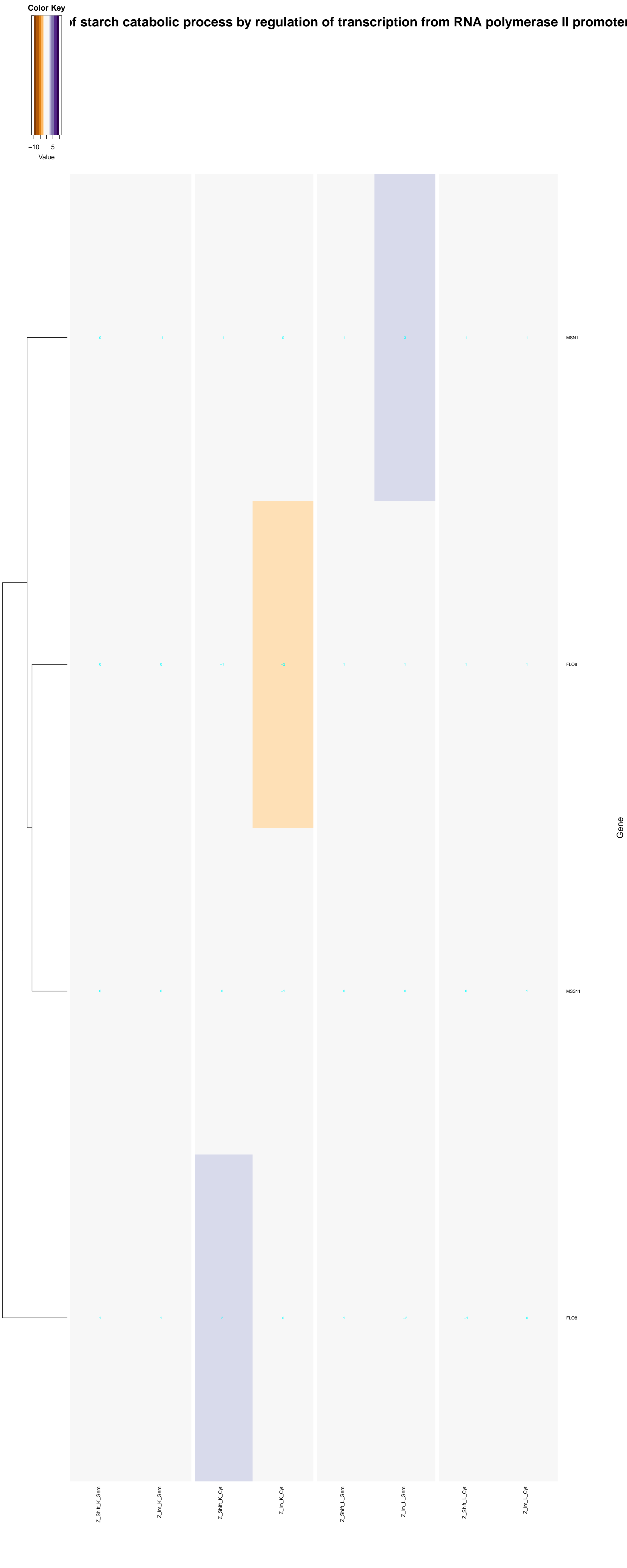


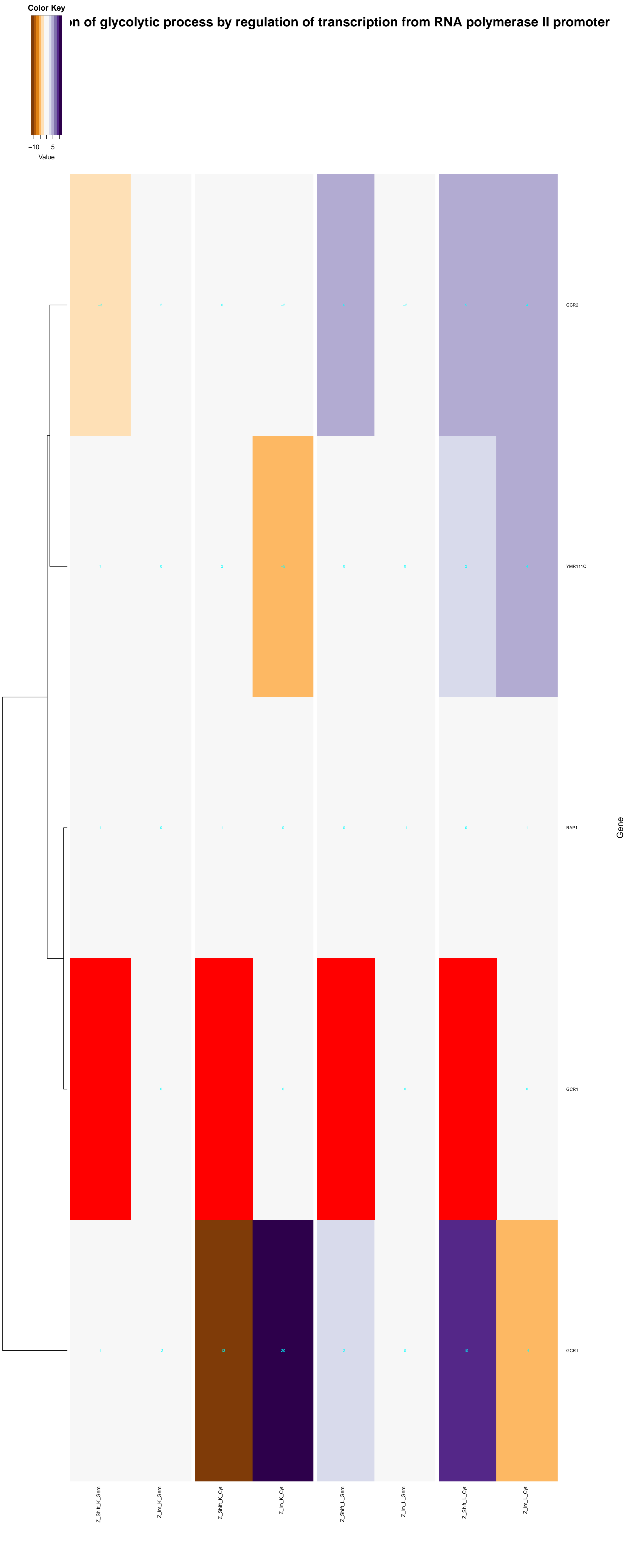


the regulation of transcription from RNA polymerase II promoter in response to ethanol



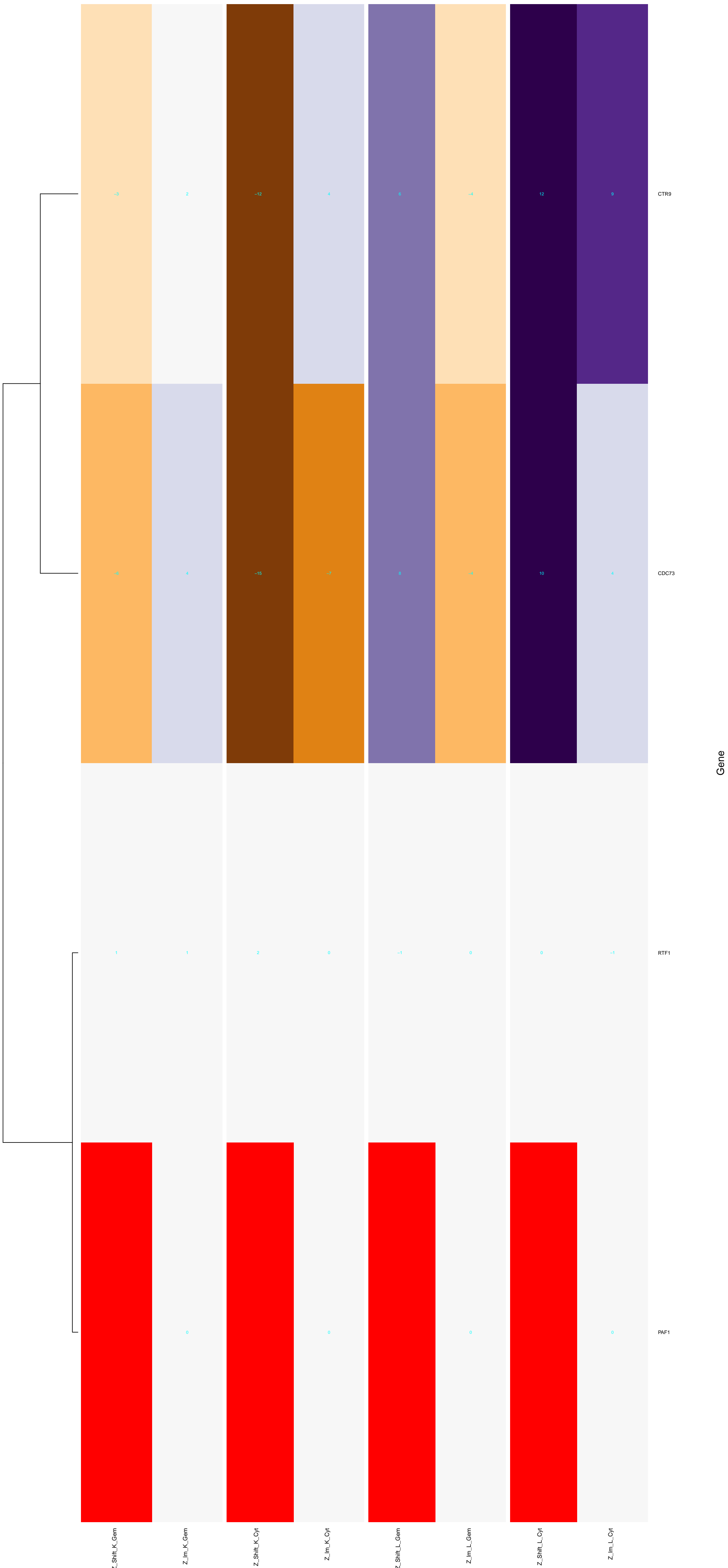


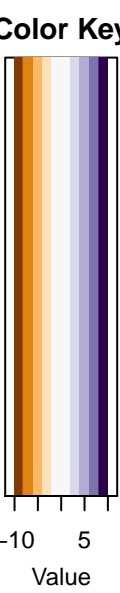






egulation of phosphorylation of RNA polymerase II C-terminal domain serine 2 residues





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

