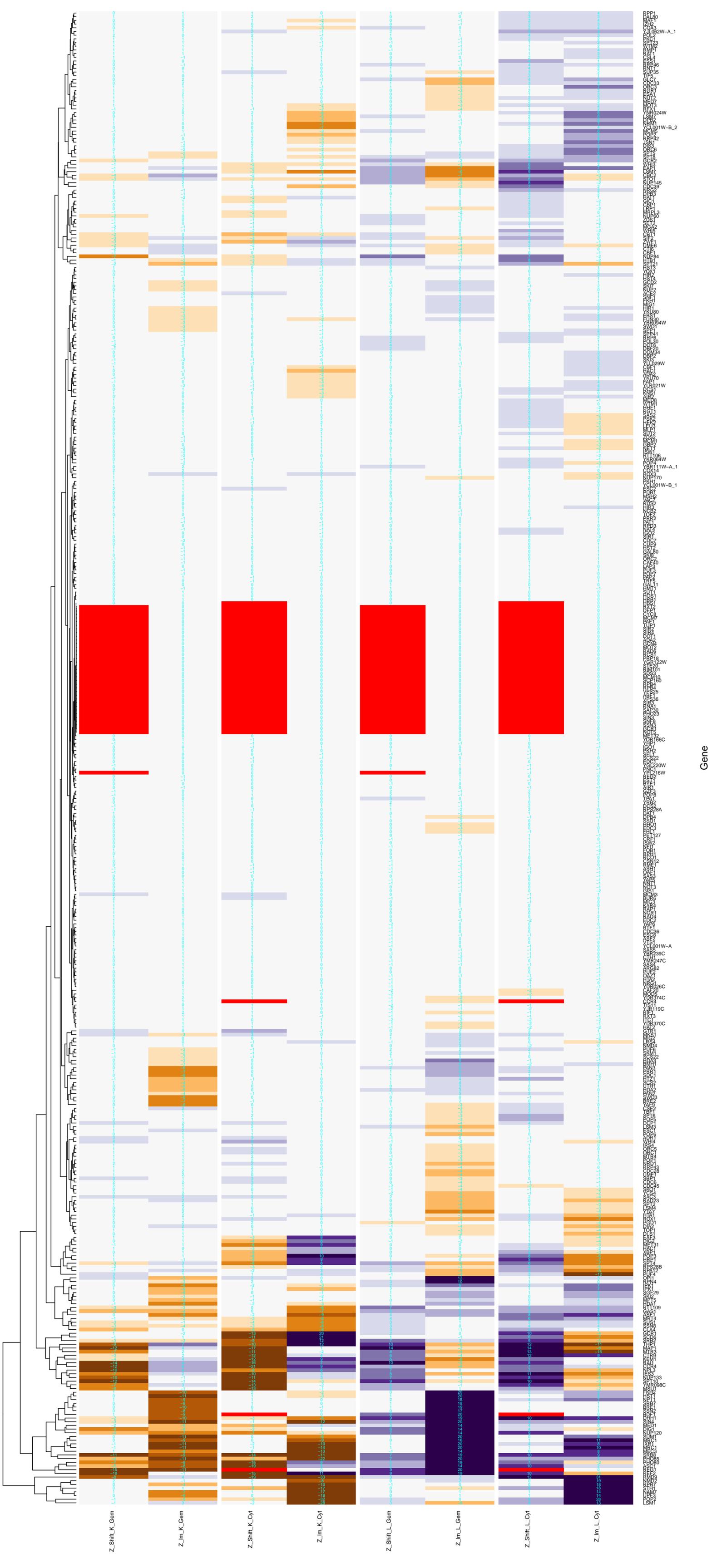
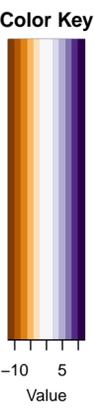
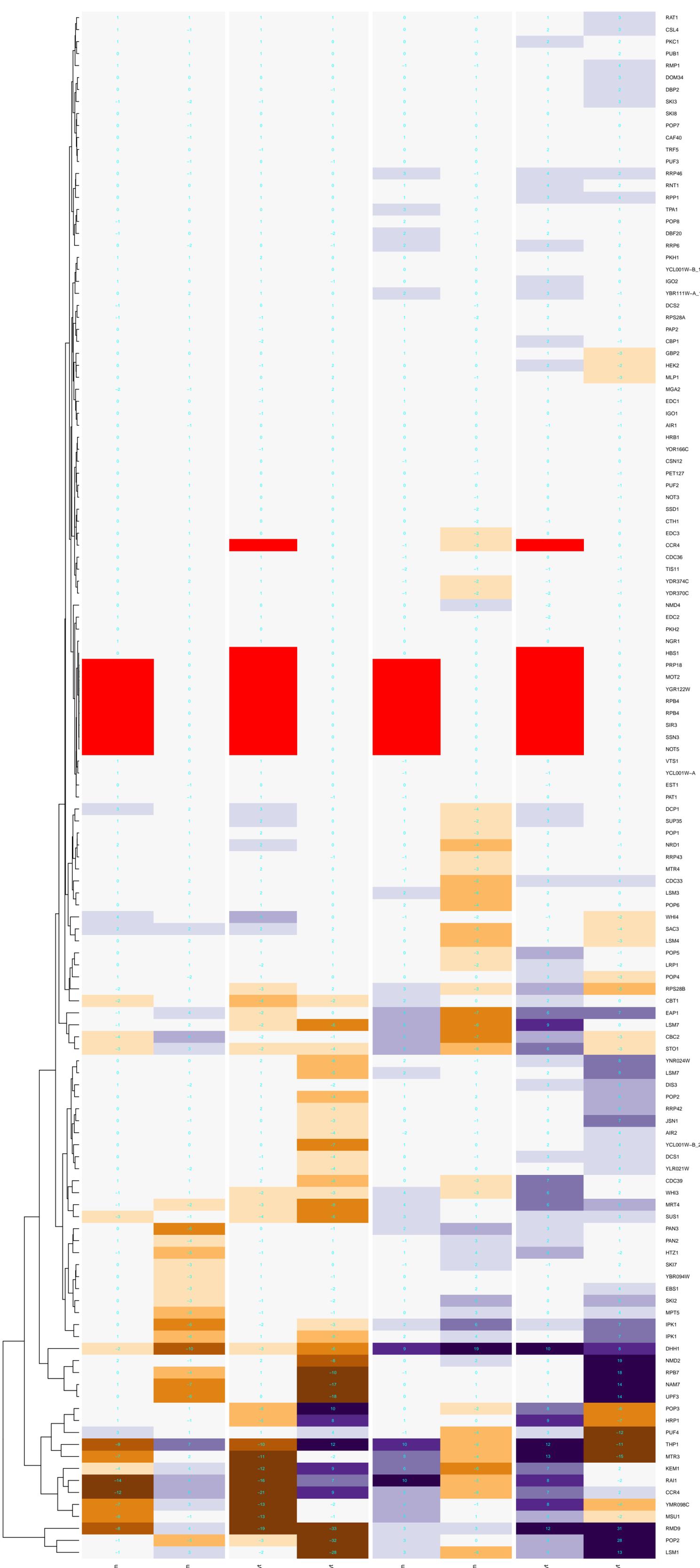
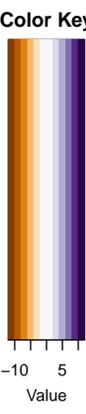


negative regulation of gene expression

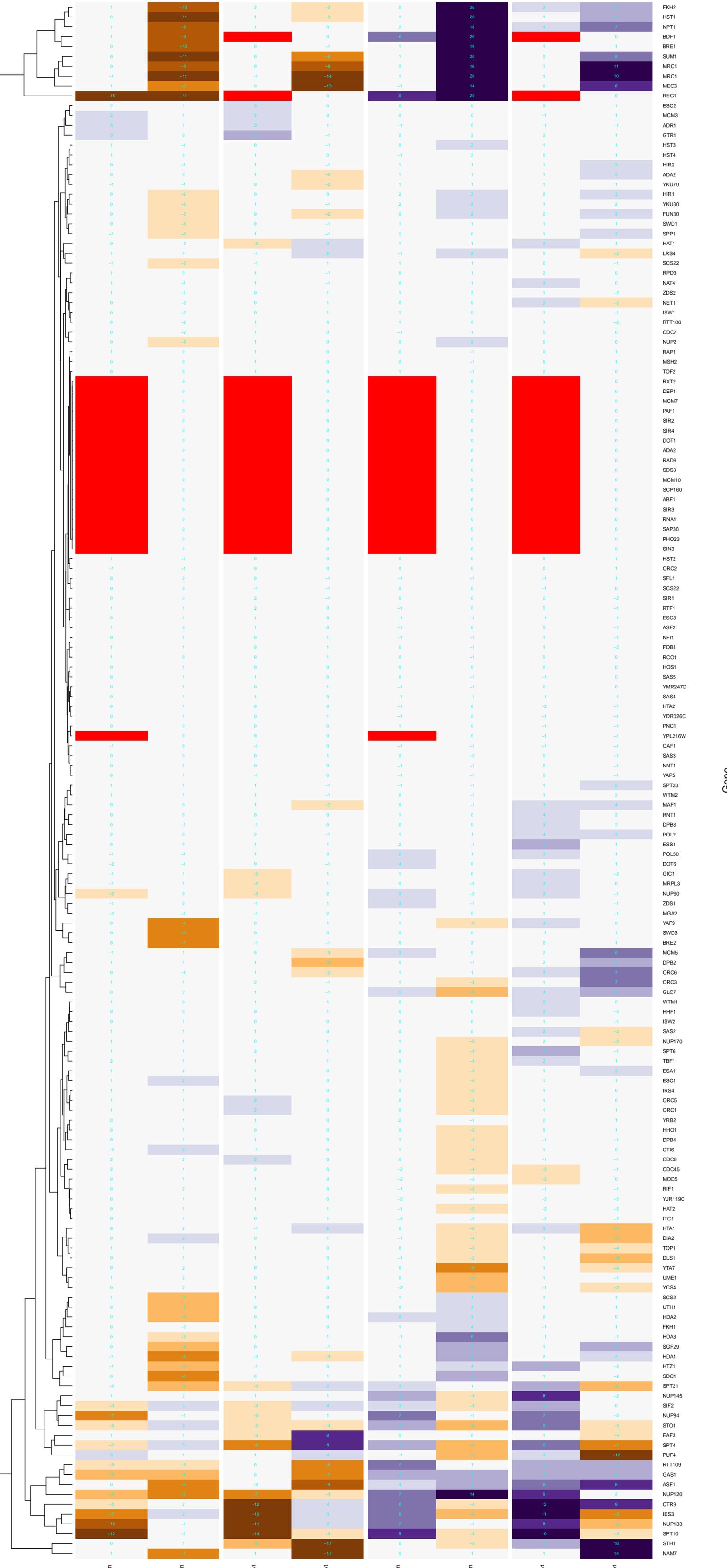
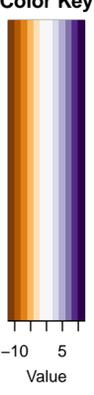


mRNA catabolic process



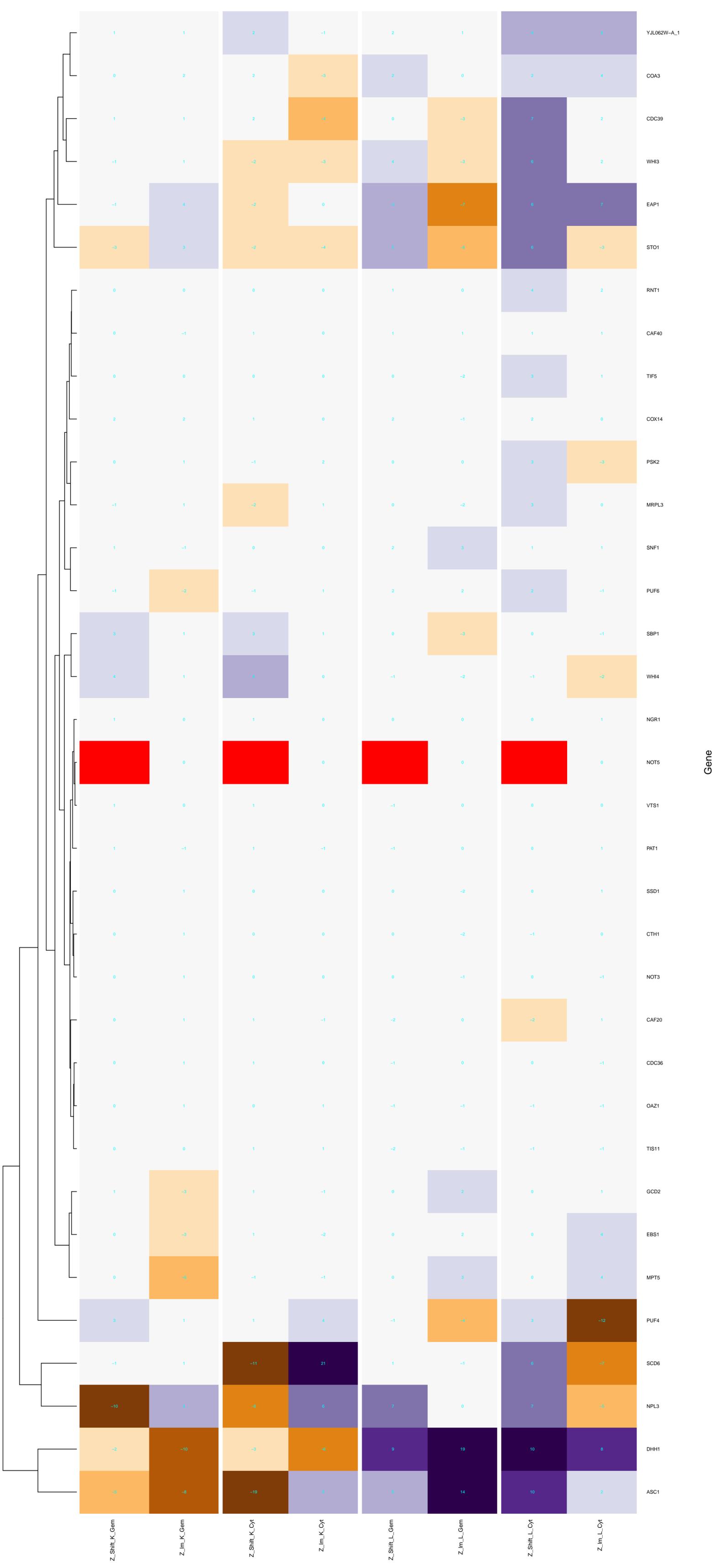
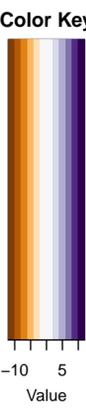
Gene

gene silencing

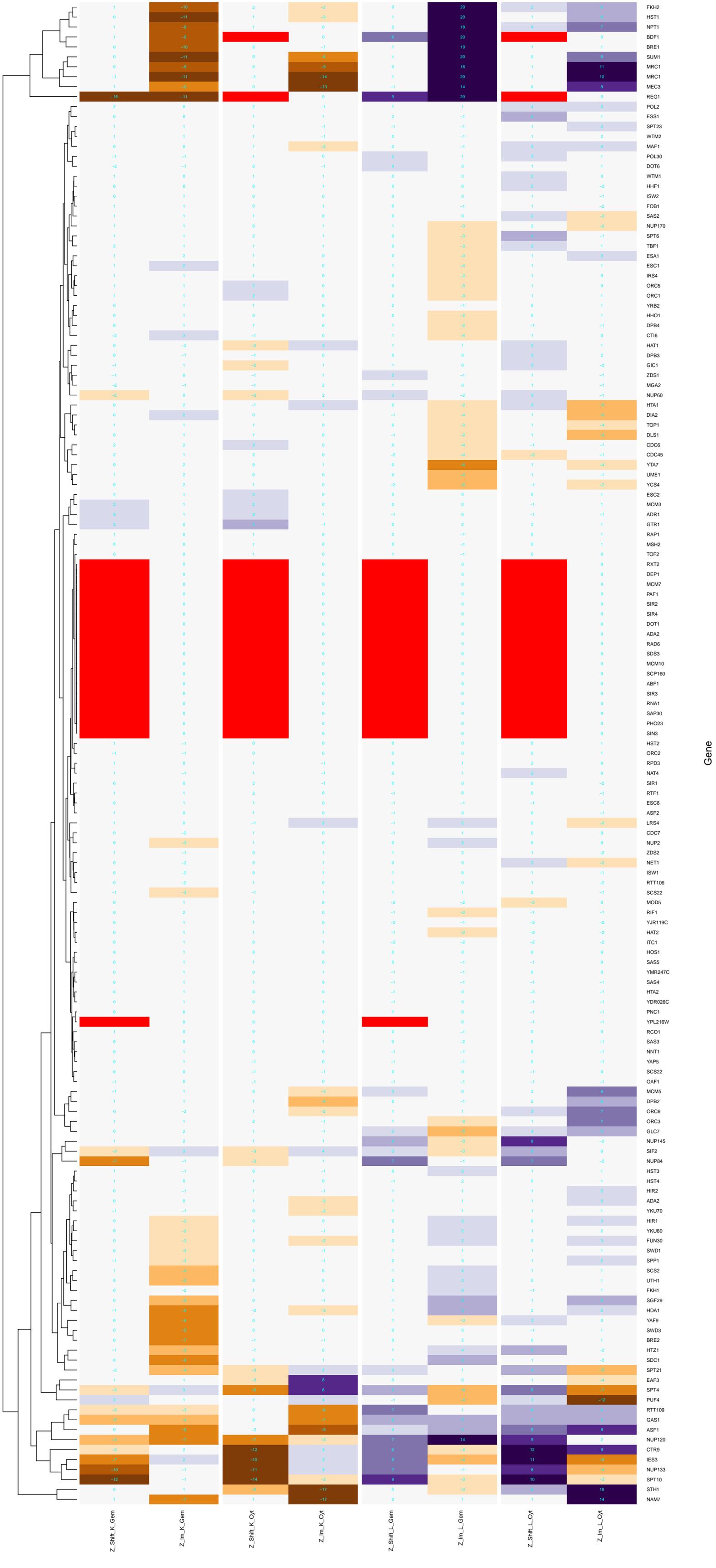


Gene

negative regulation of translation

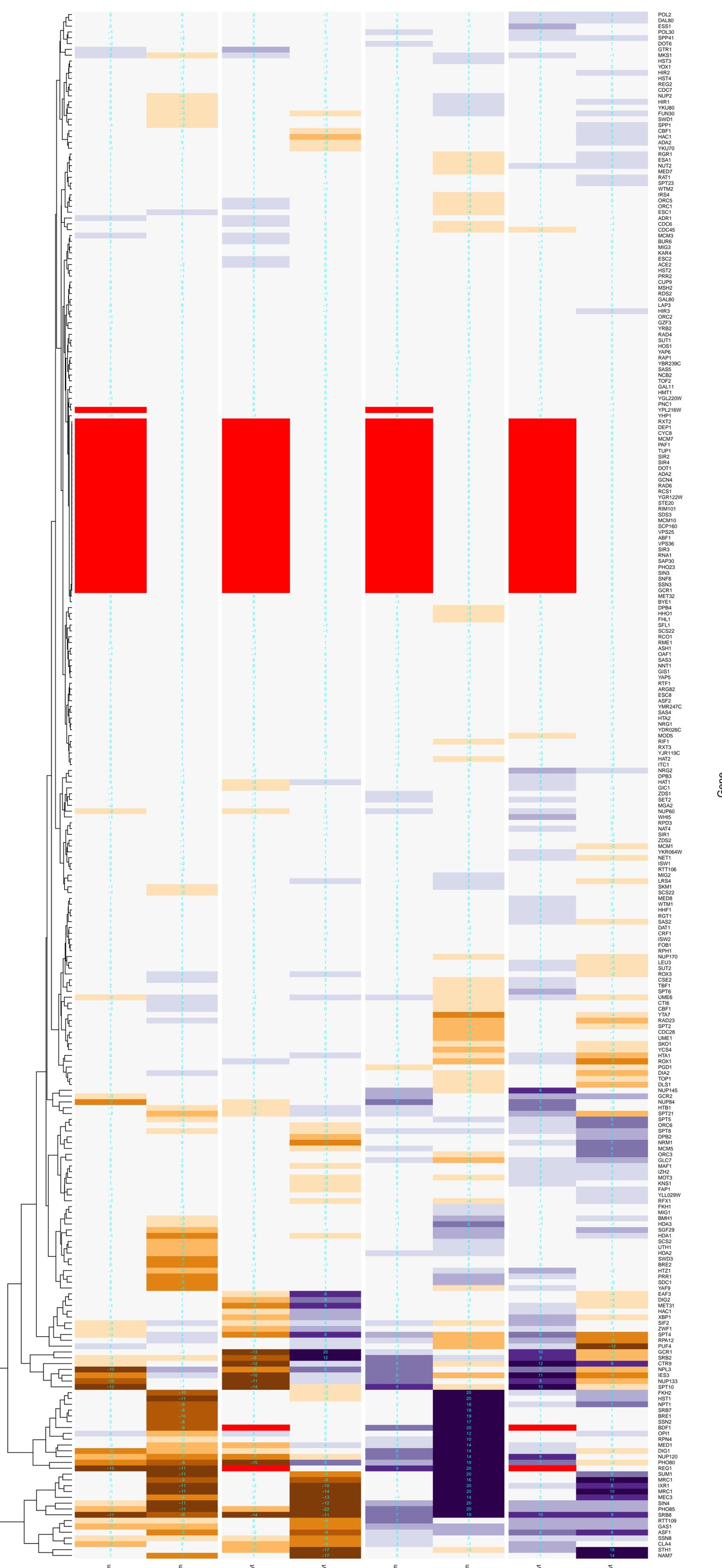
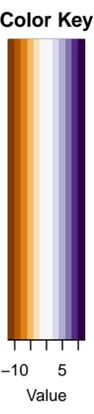


negative regulation of gene expression, epigenetic

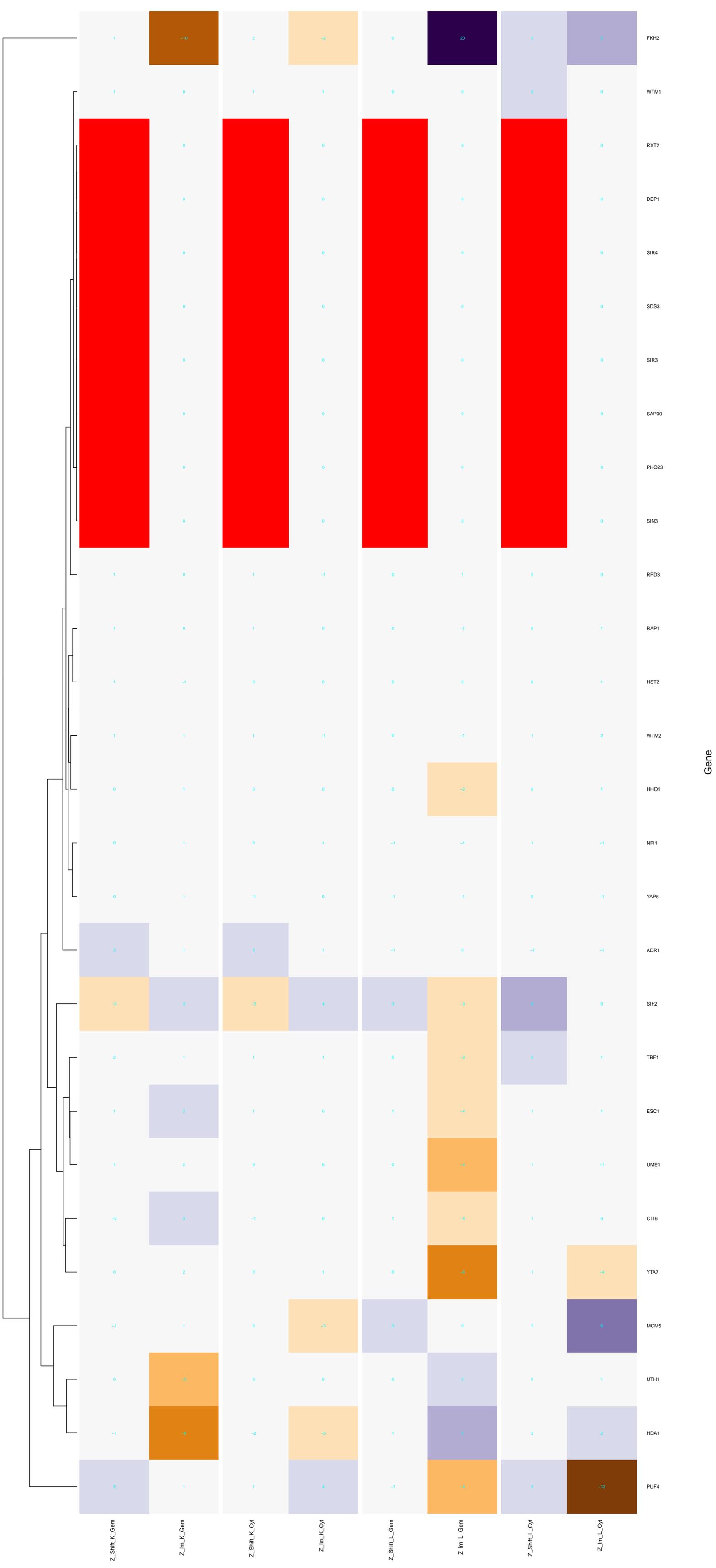
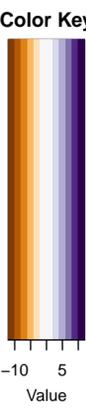


Gene

negative regulation of transcription, DNA-templated

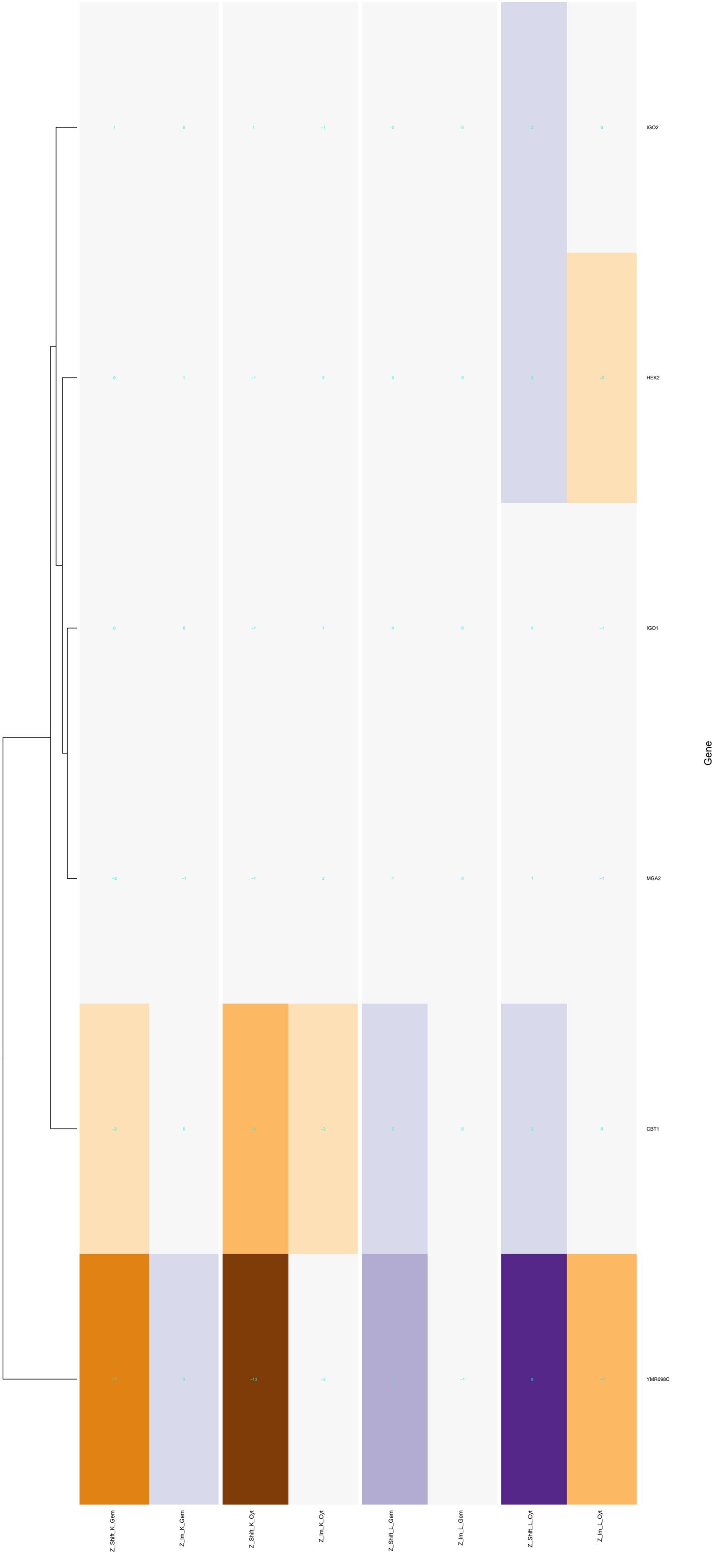
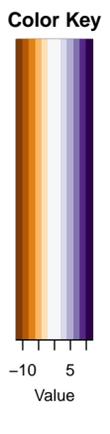


negative regulation of gene silencing

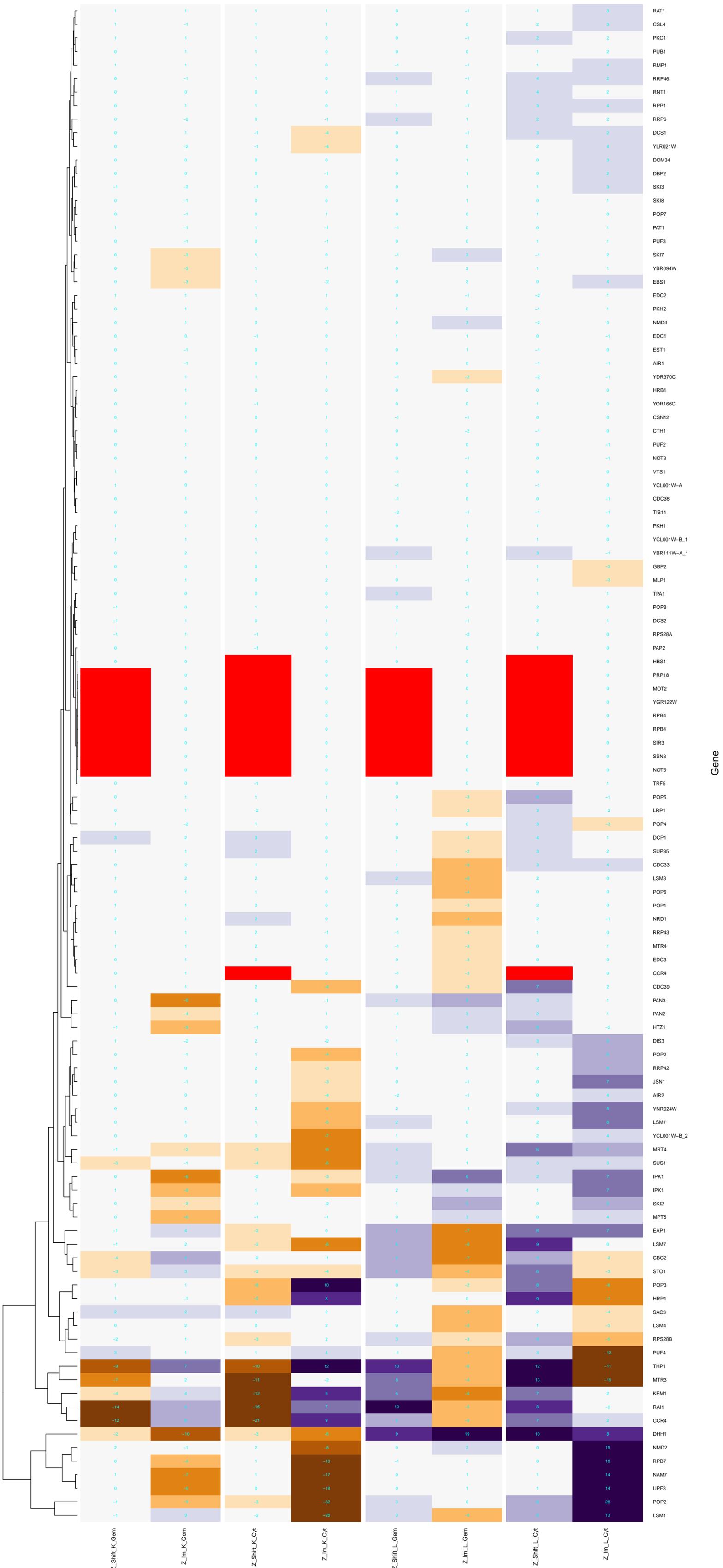
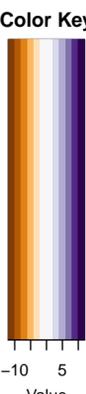


Gene

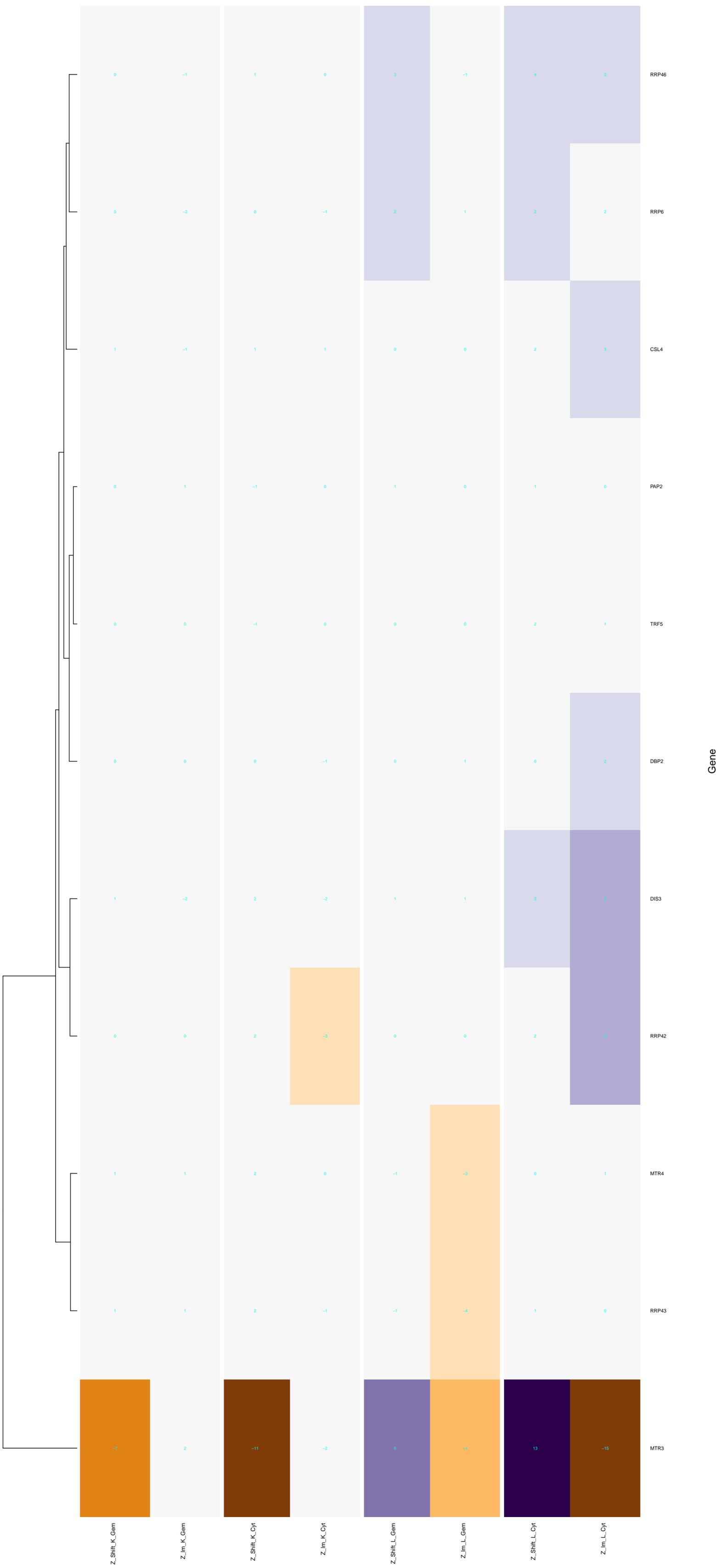
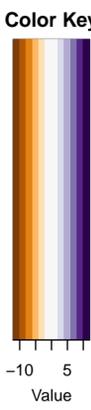
negative regulation of mRNA catabolic process



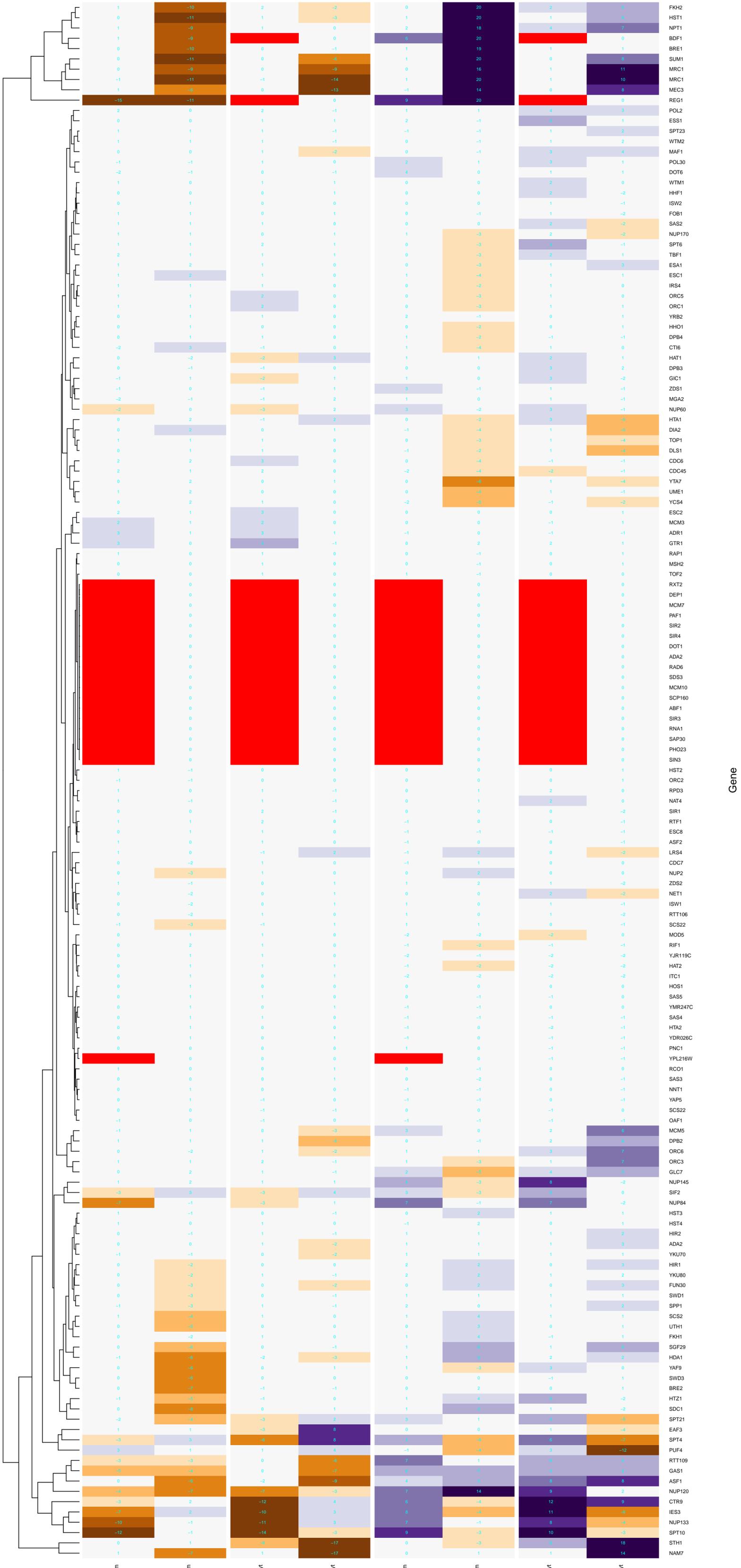
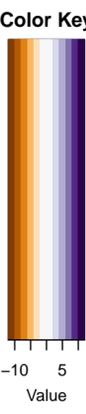
nuclear-transcribed mRNA catabolic process



polyadenylation-dependent mRNA catabolic process

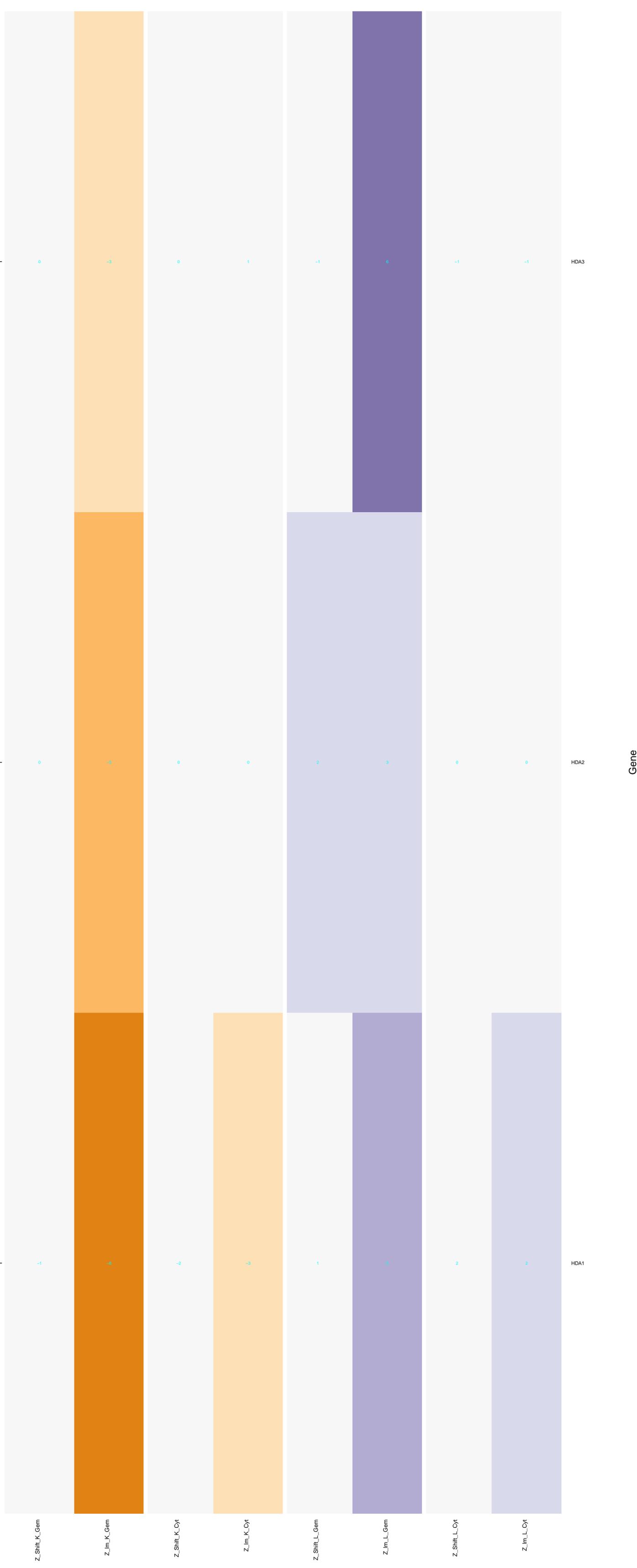
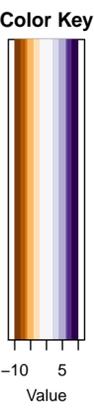


chromatin silencing

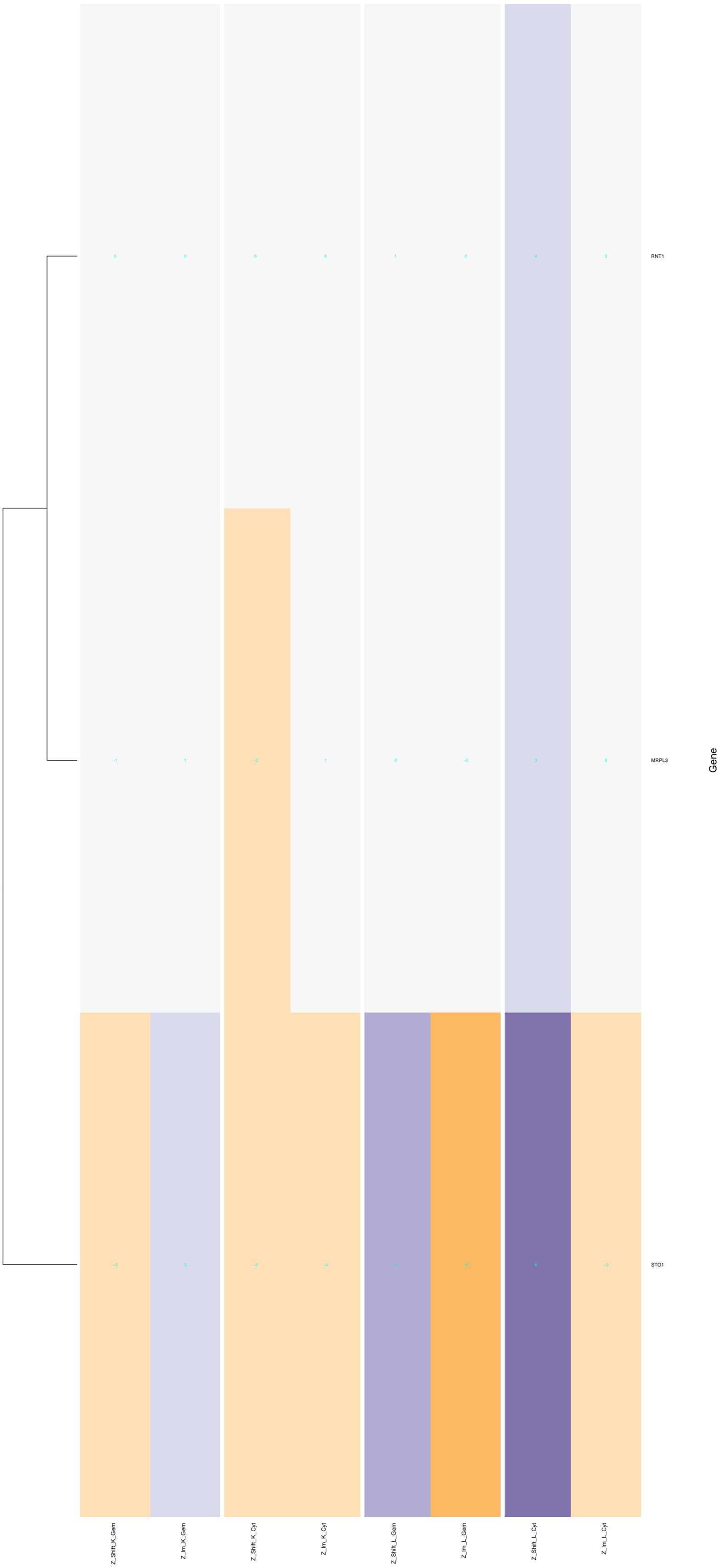
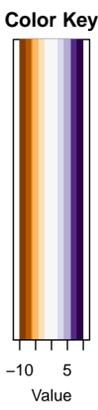


Gene

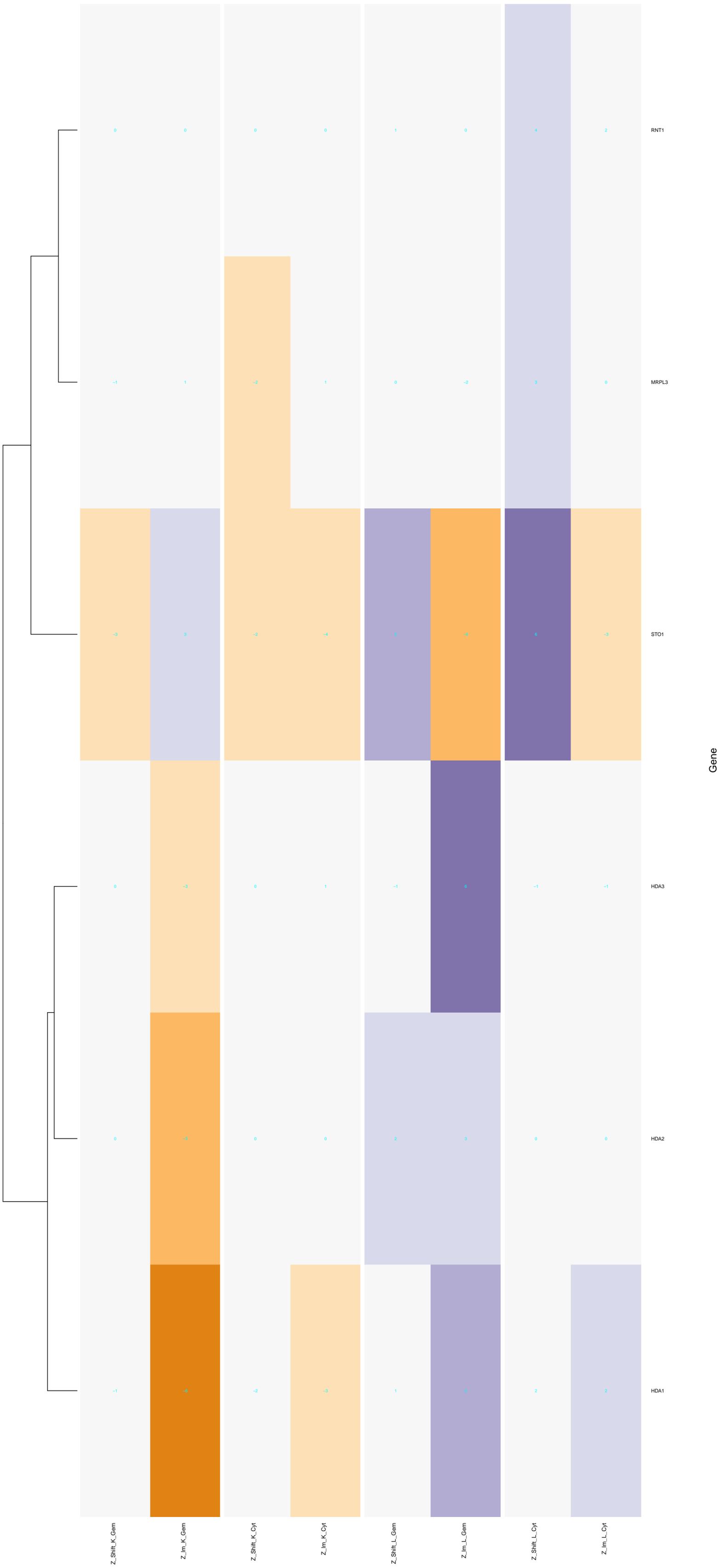
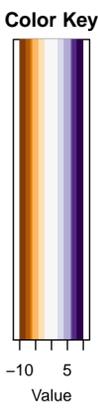
gene silencing involved in chronological cell aging



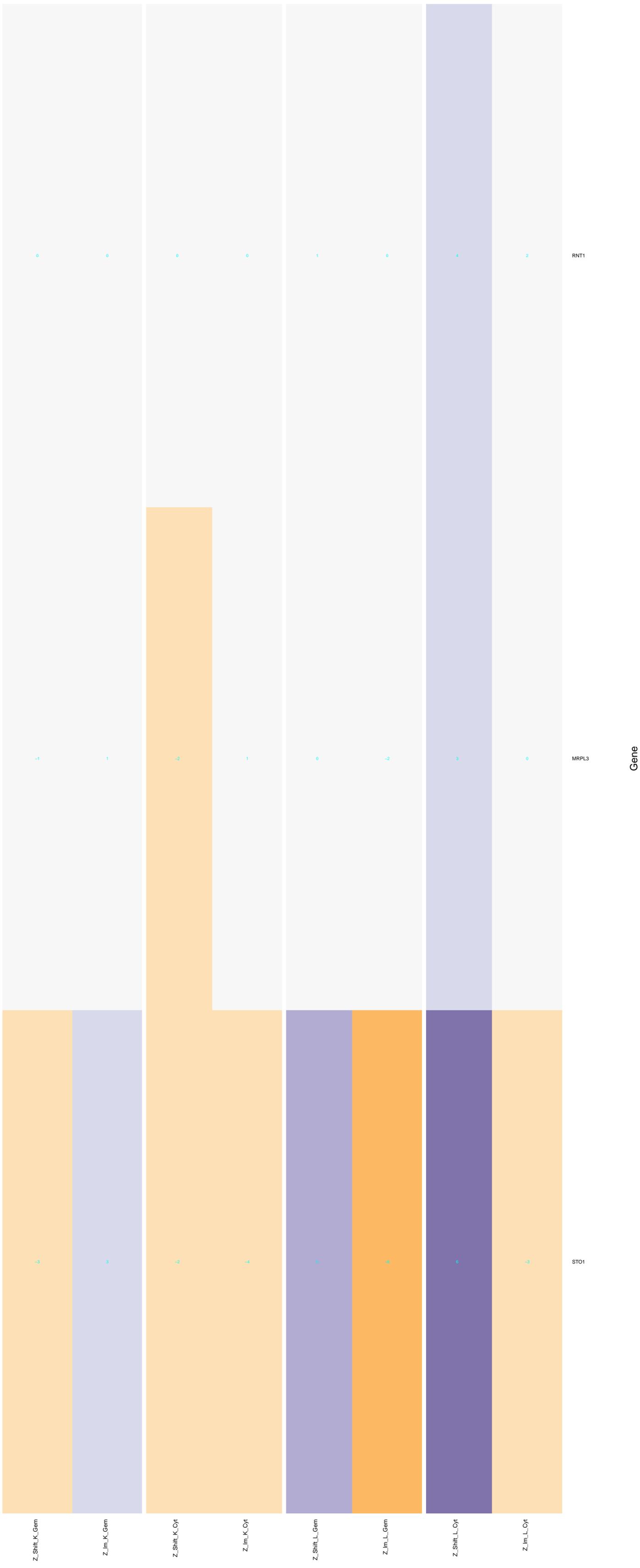
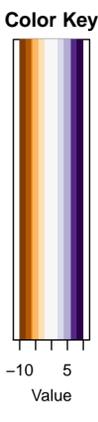
posttranscriptional gene silencing



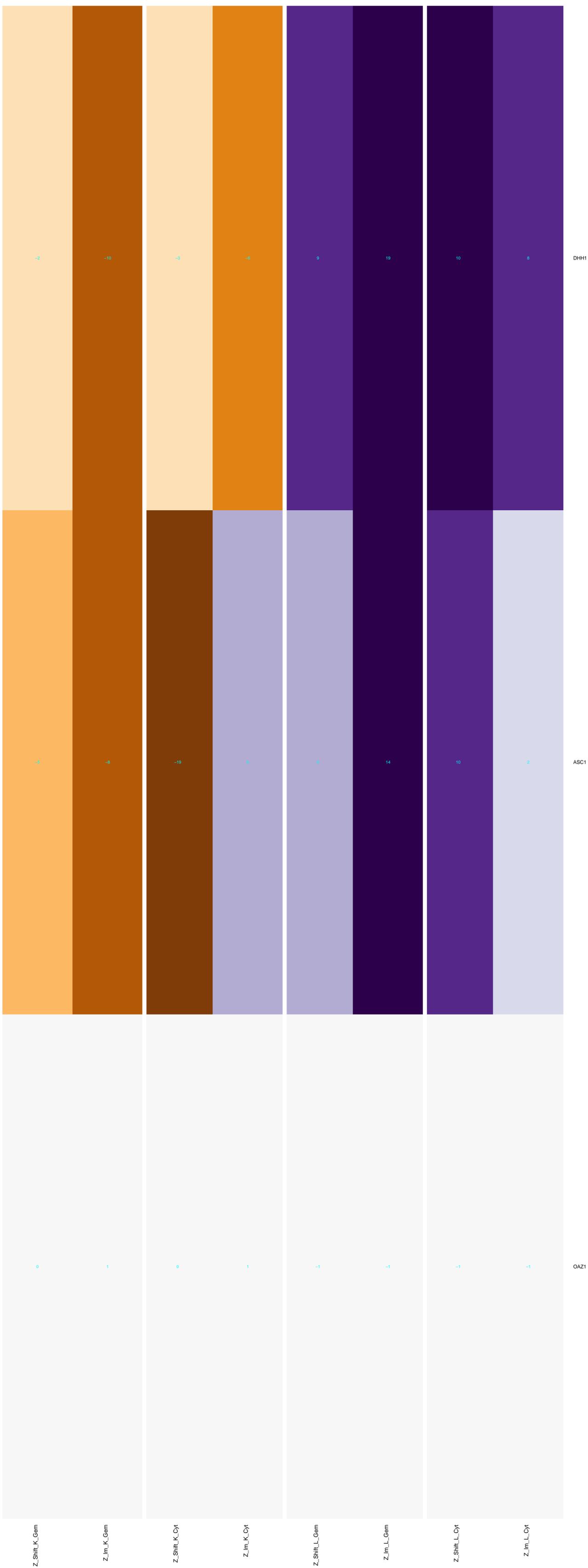
gene silencing by RNA



gene silencing by miRNA

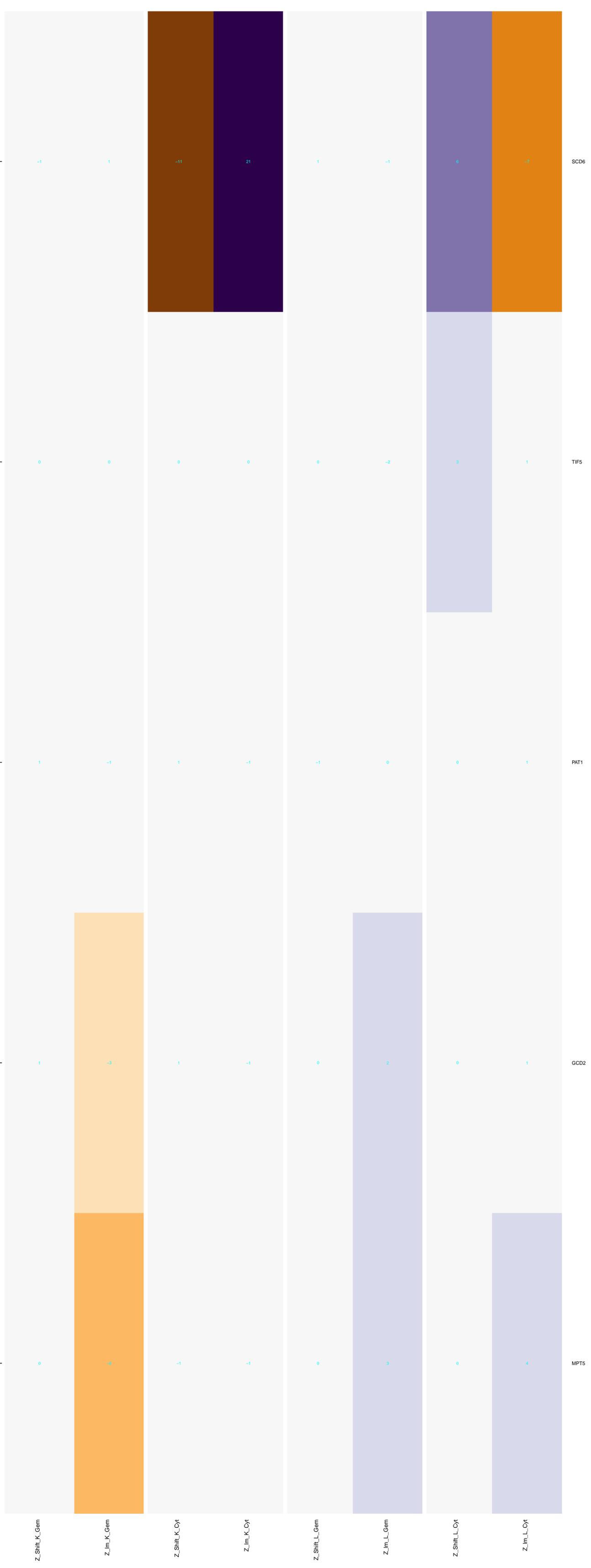
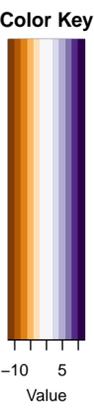


negative regulation of translational elongation



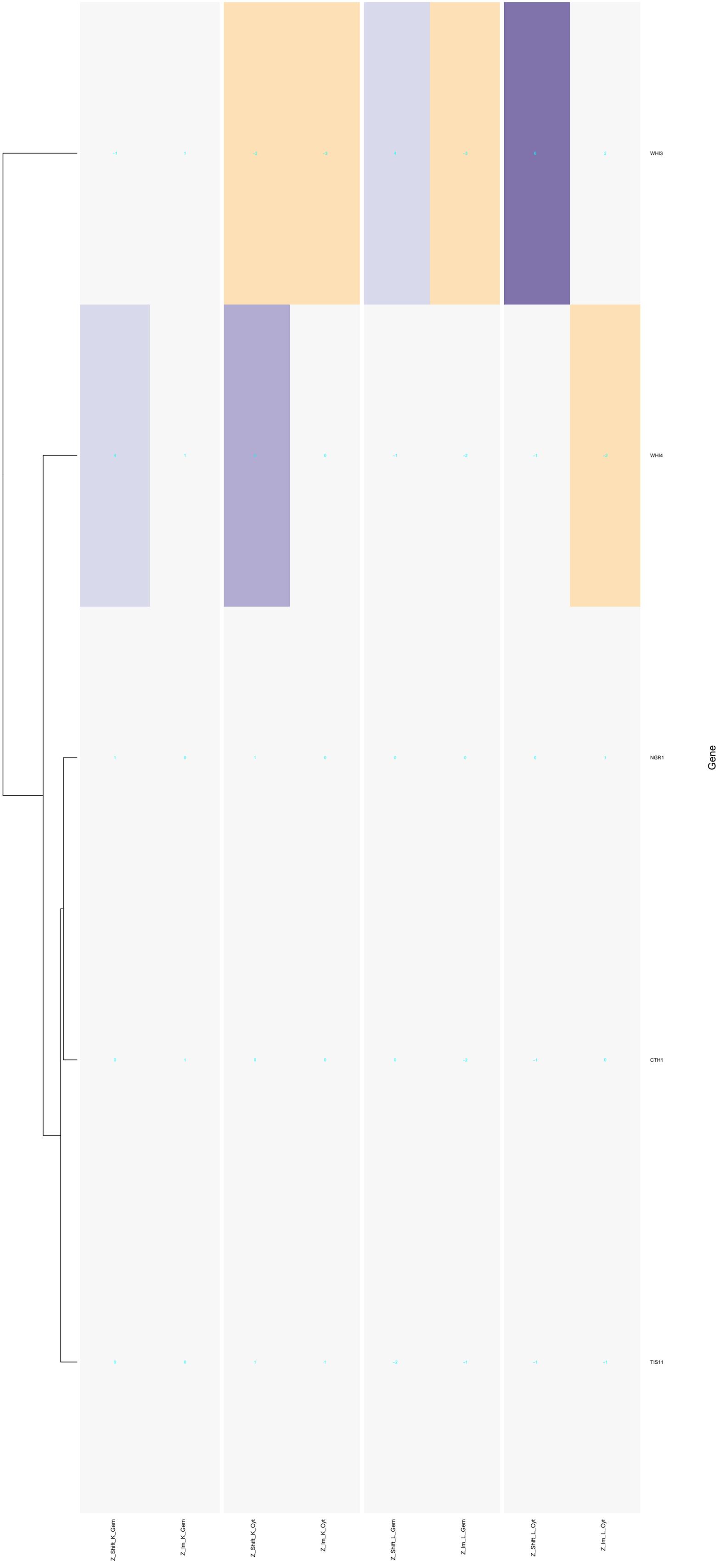
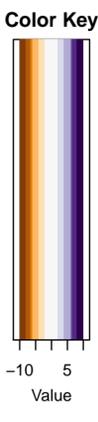
Gene

negative regulation of translational initiation

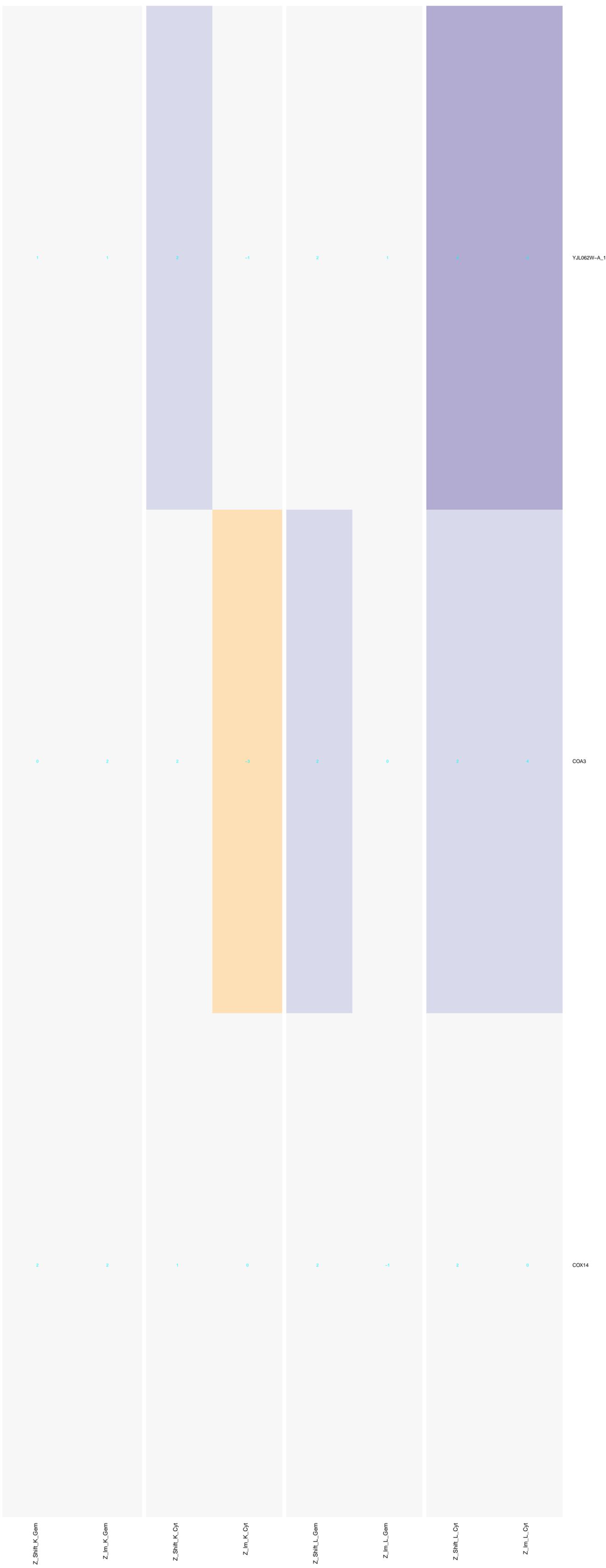
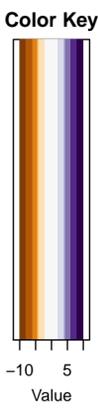


Gene

mRNA destabilization

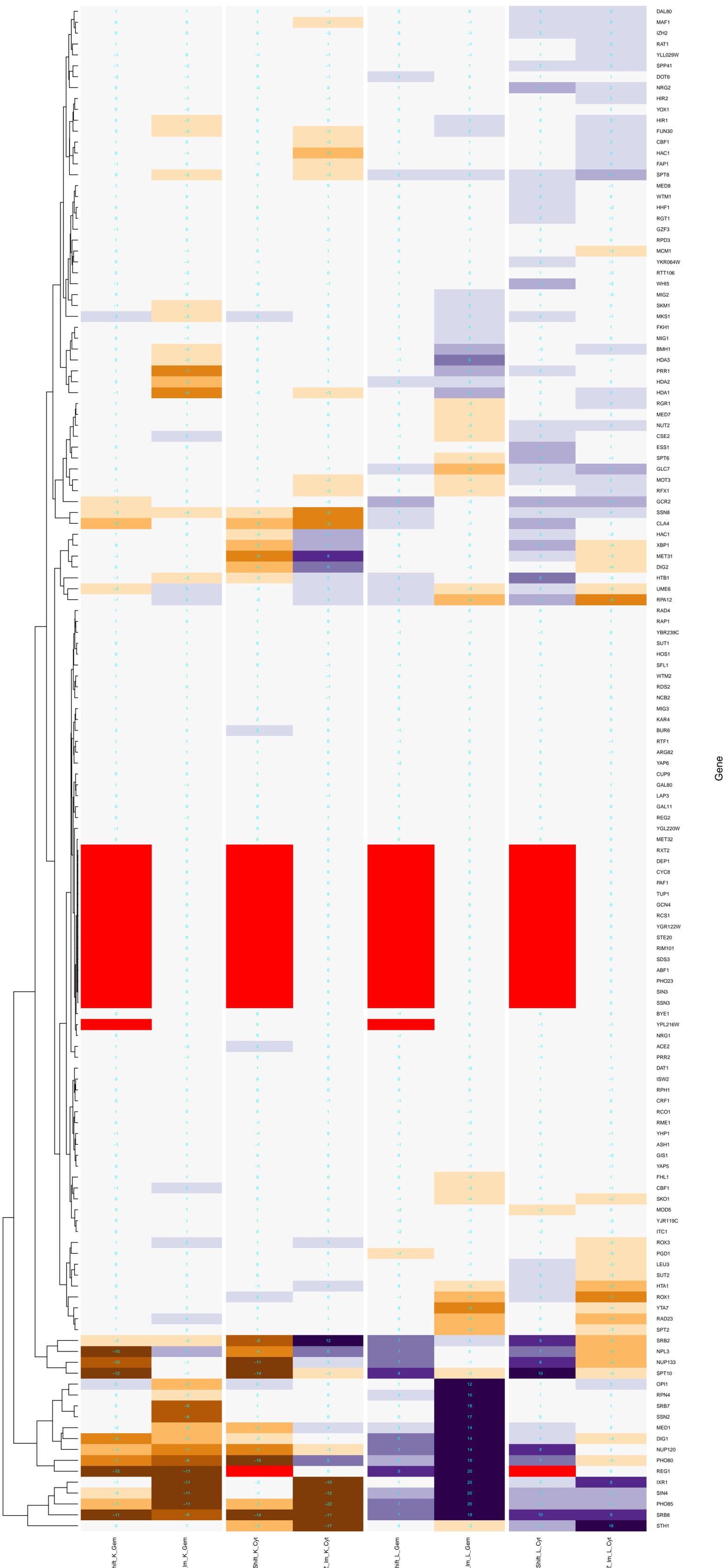
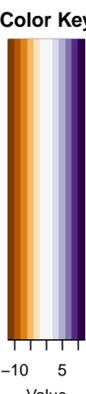


negative regulation of mitochondrial translation

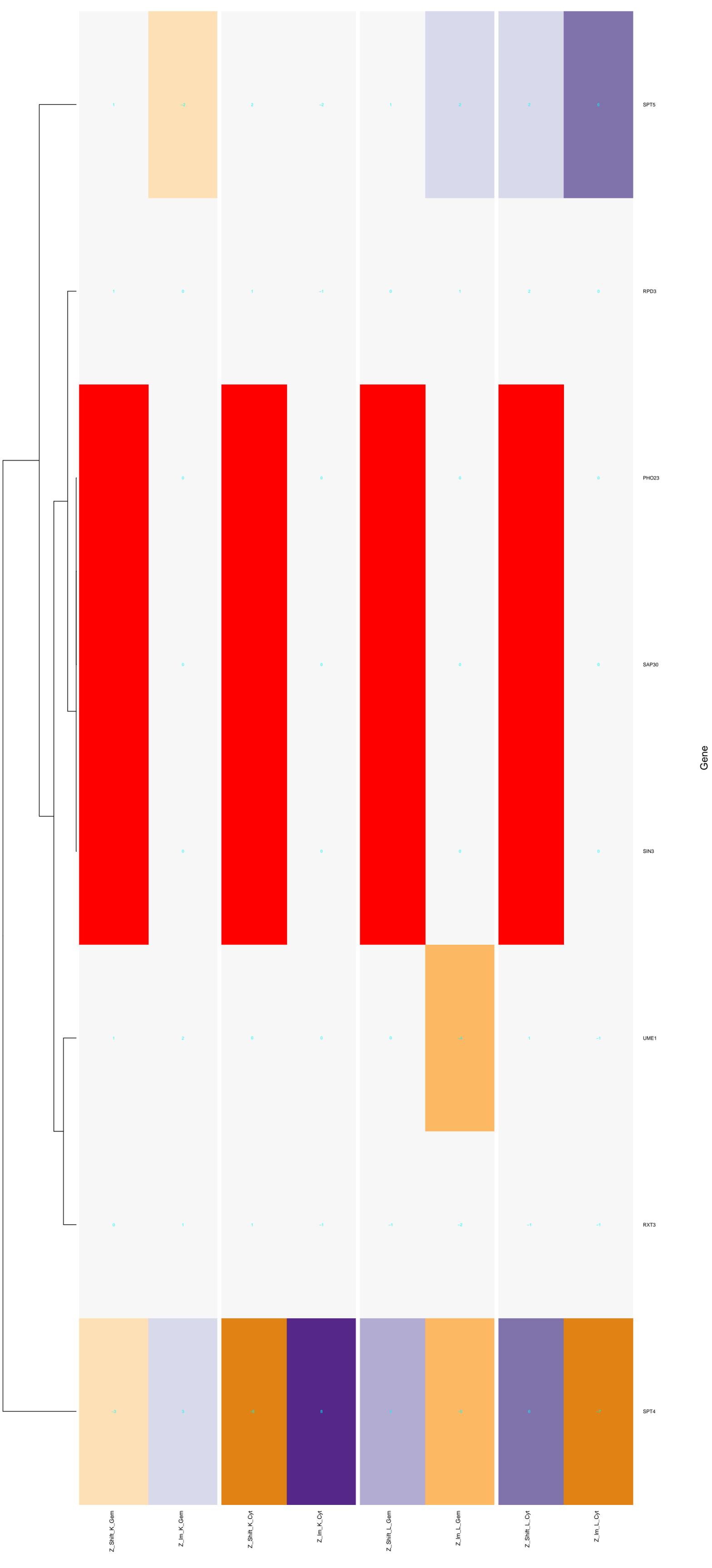
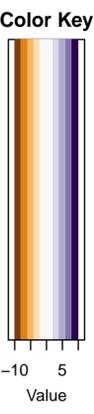


Gene

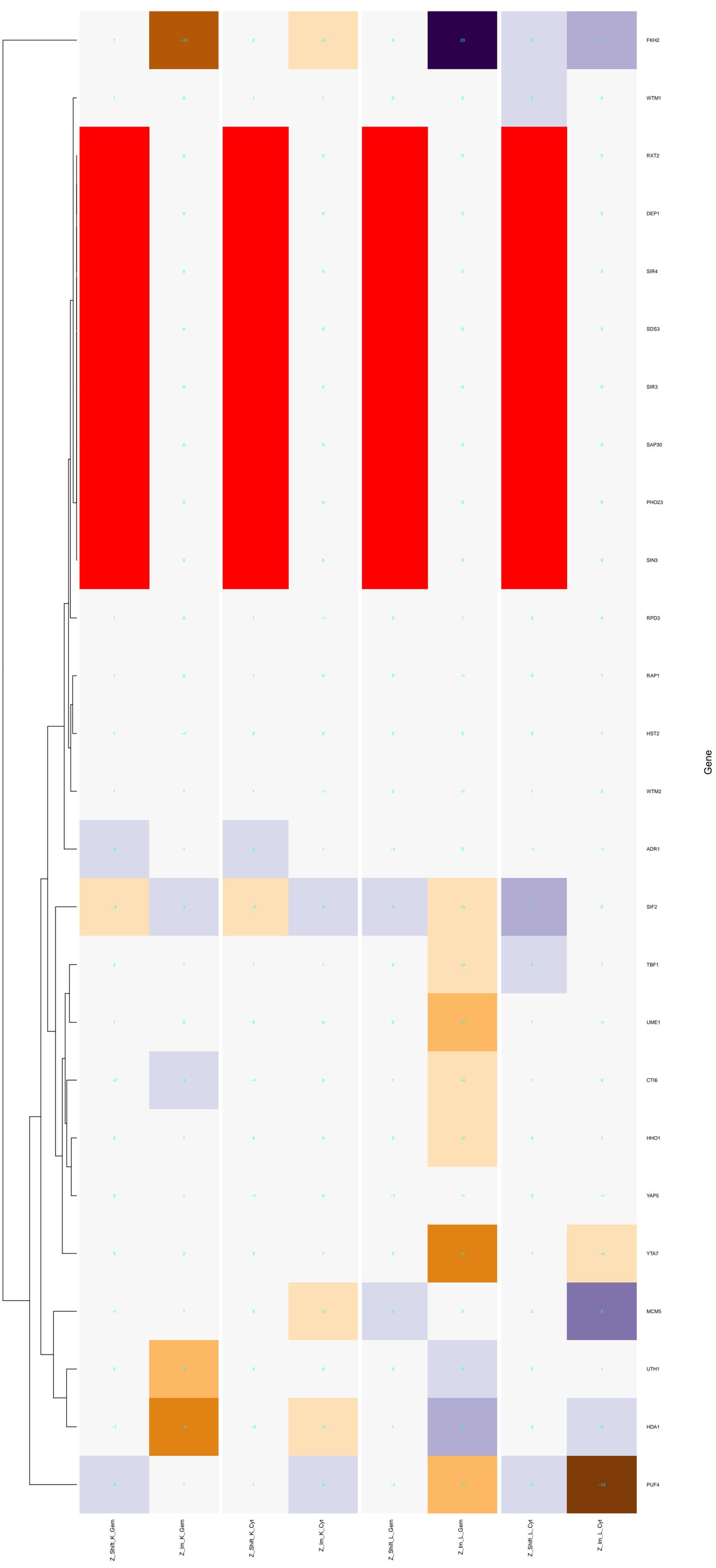
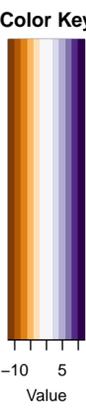
negative regulation of transcription from RNA polymerase II promoter



negative regulation of transcription from RNA polymerase I promoter

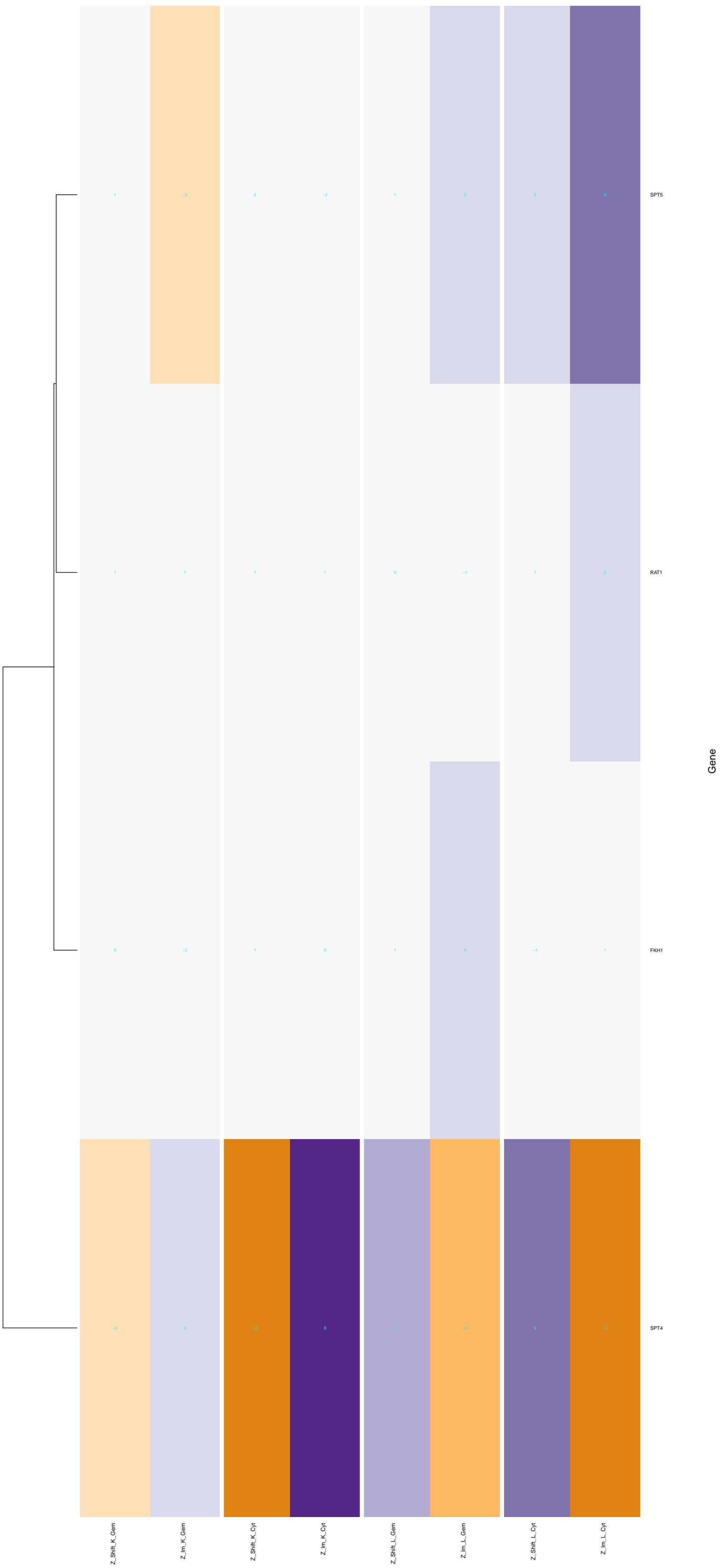
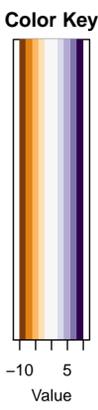


negative regulation of chromatin silencing

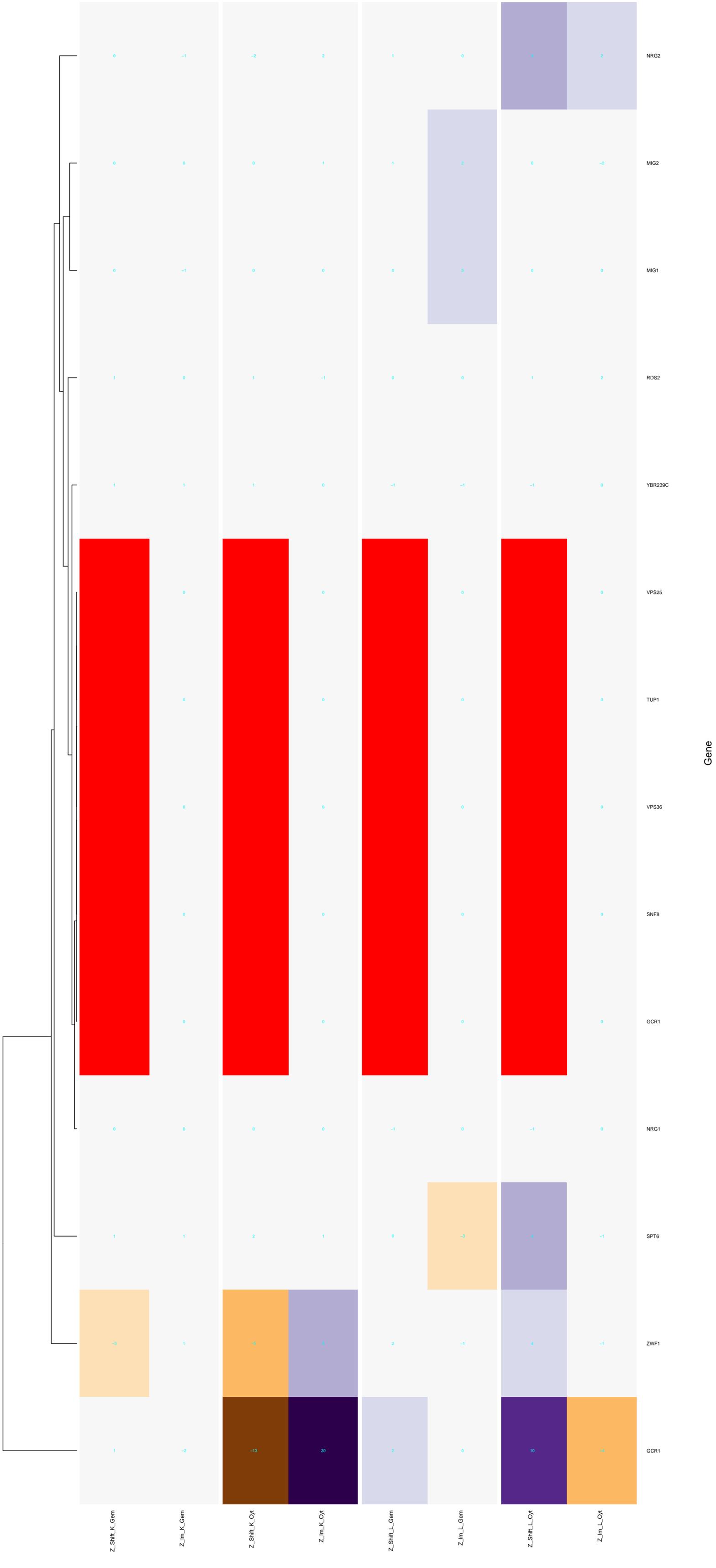
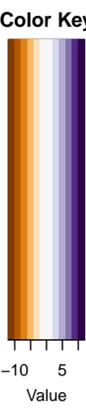


Gene

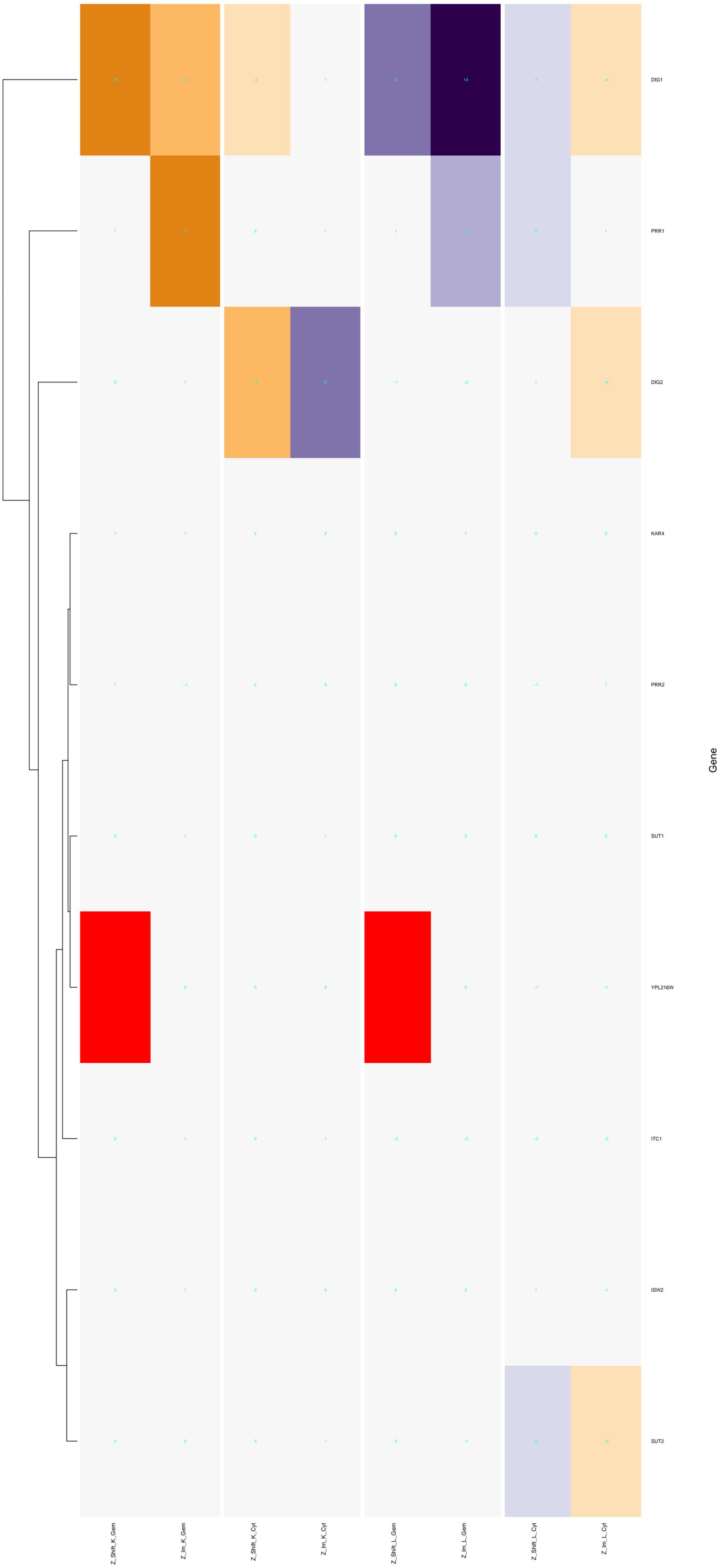
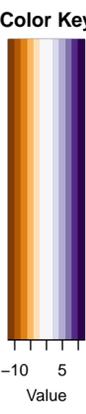
negative regulation of DNA-templated transcription, elongation



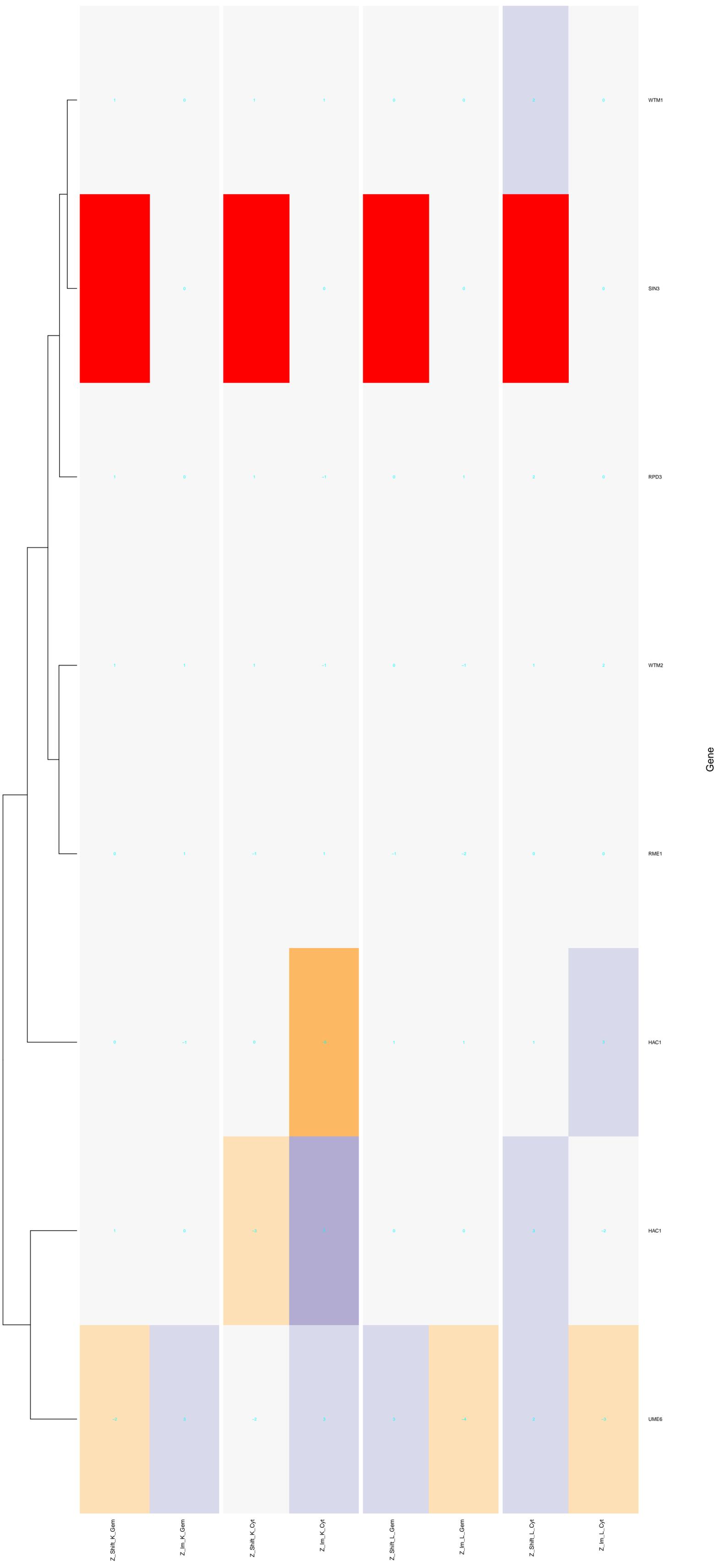
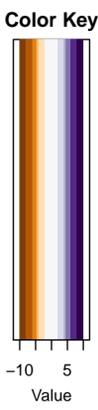
carbon catabolite repression of transcription



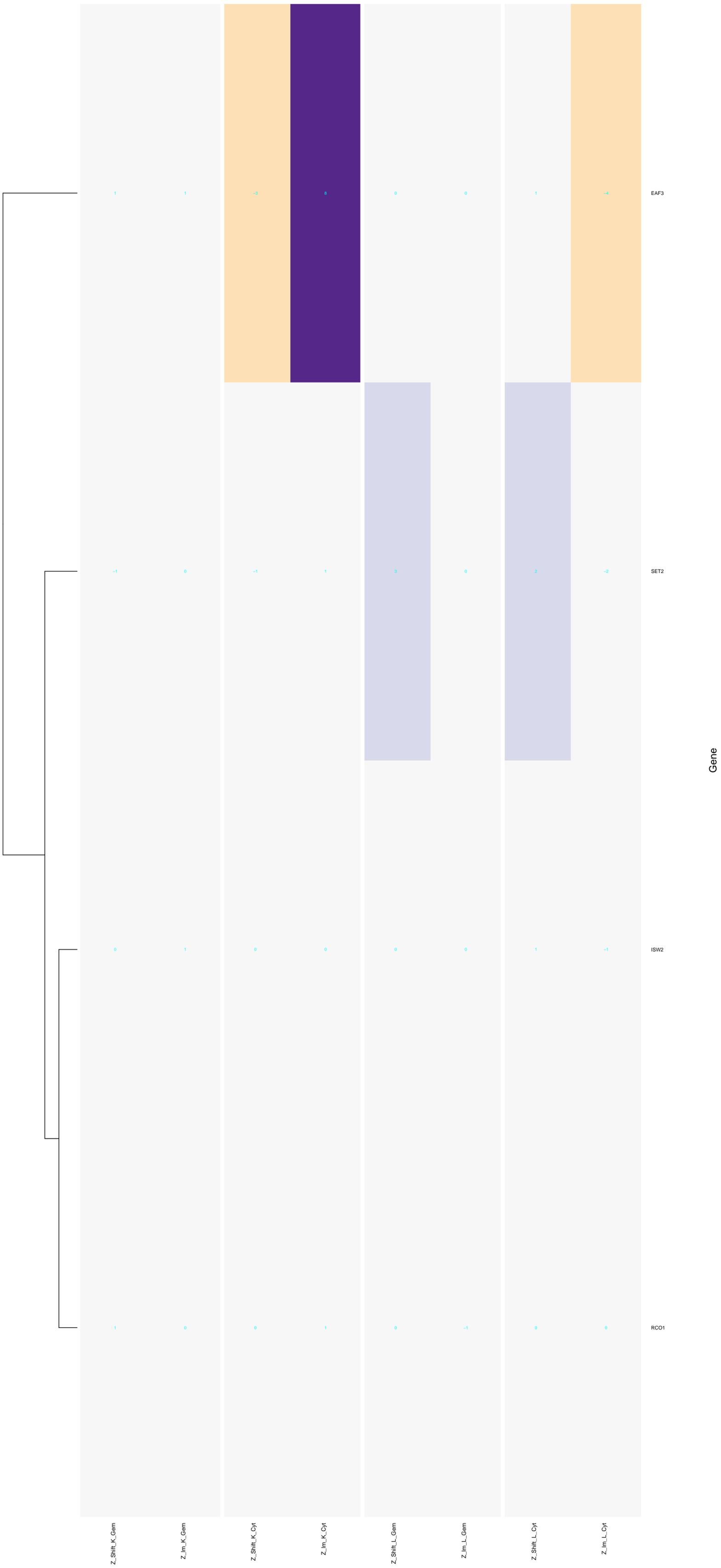
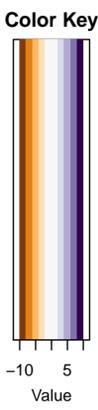
negative regulation of transcription by pheromones



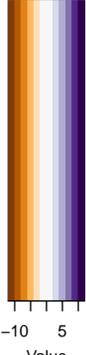
negative regulation of transcription involved in meiotic cell cycle



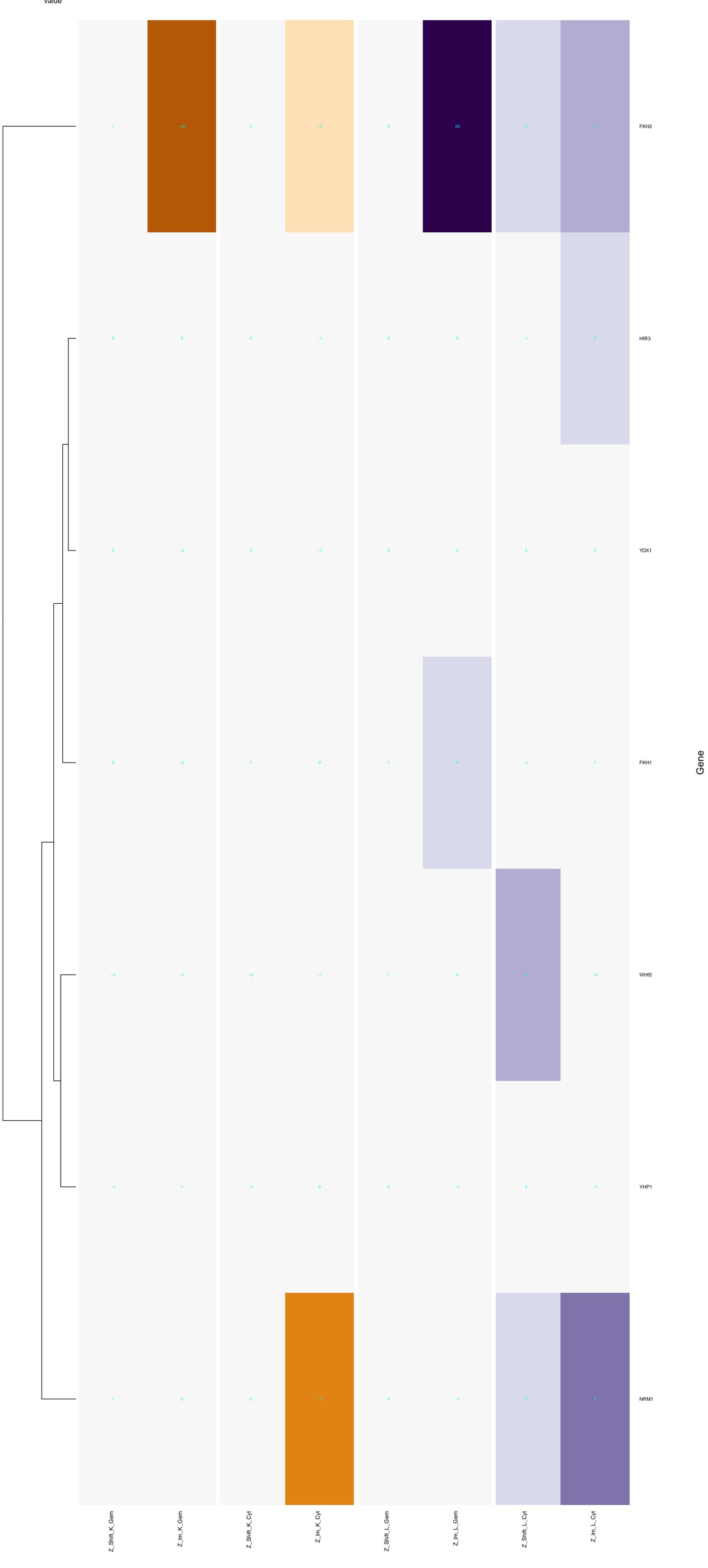
negative regulation of antisense RNA transcription



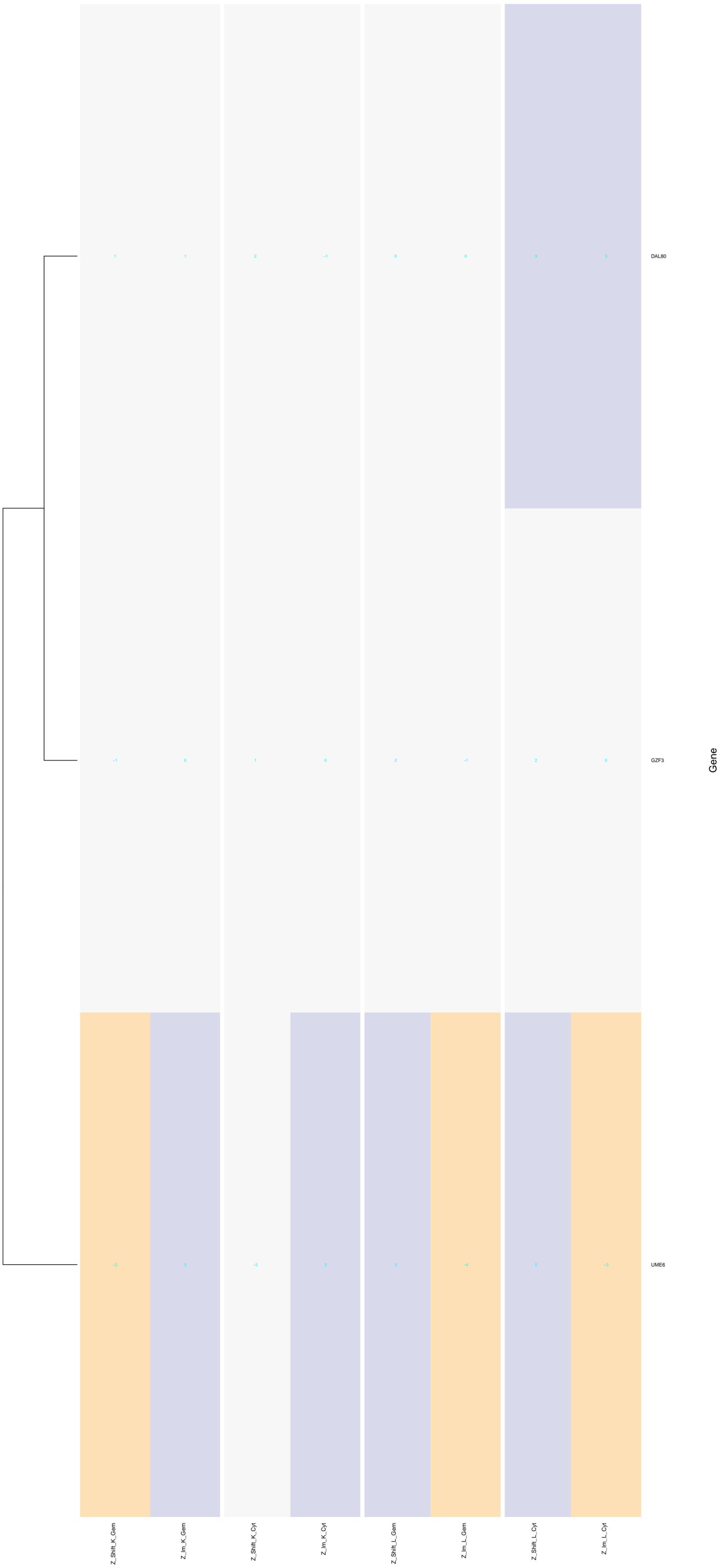
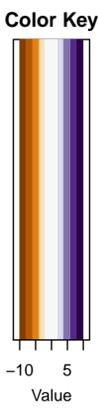
Color Key

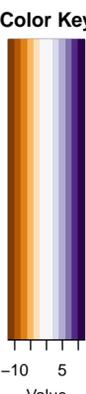


Negative regulation of transcription involved in G1/S transition of mitotic cell cycle

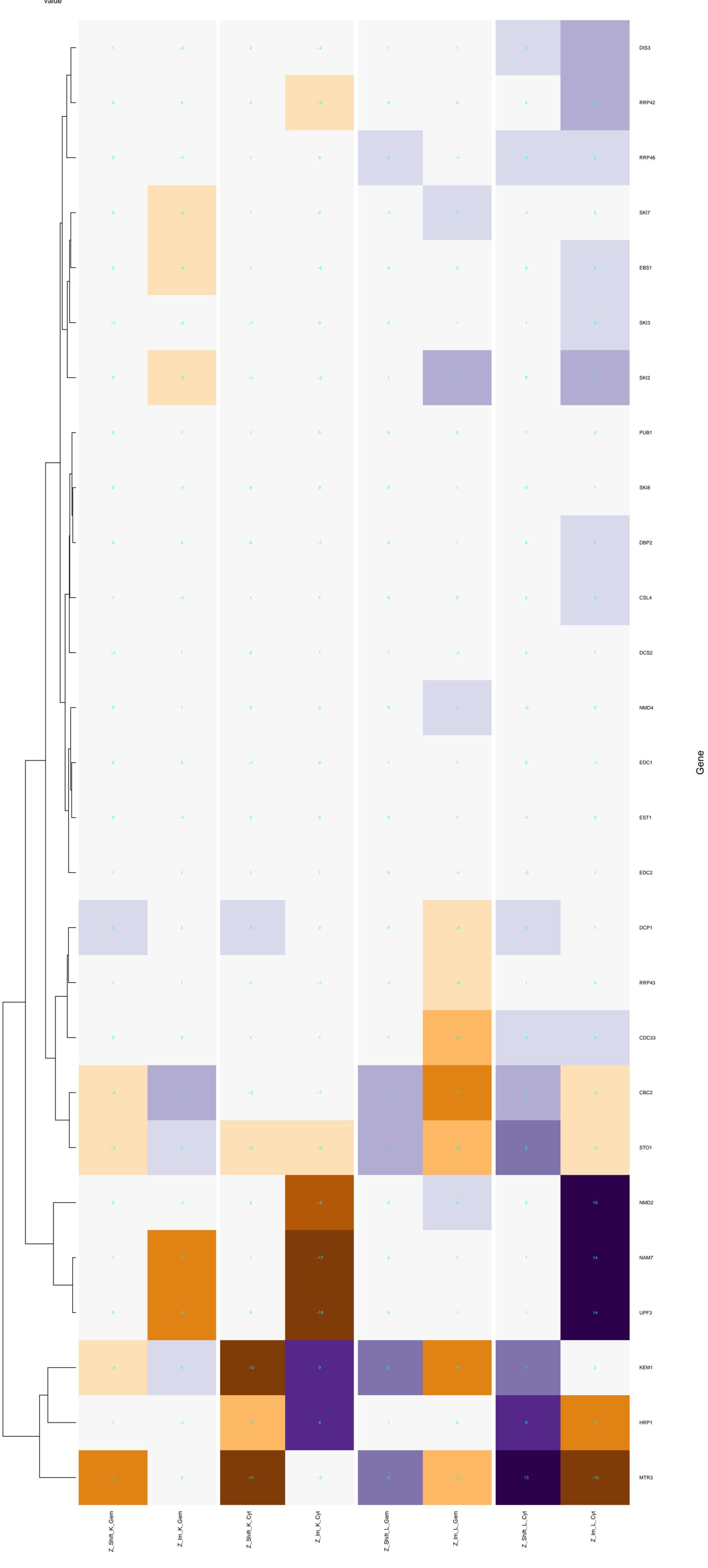


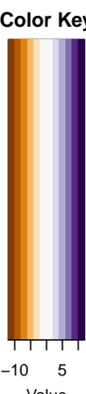
nitrogen catabolite repression of transcription



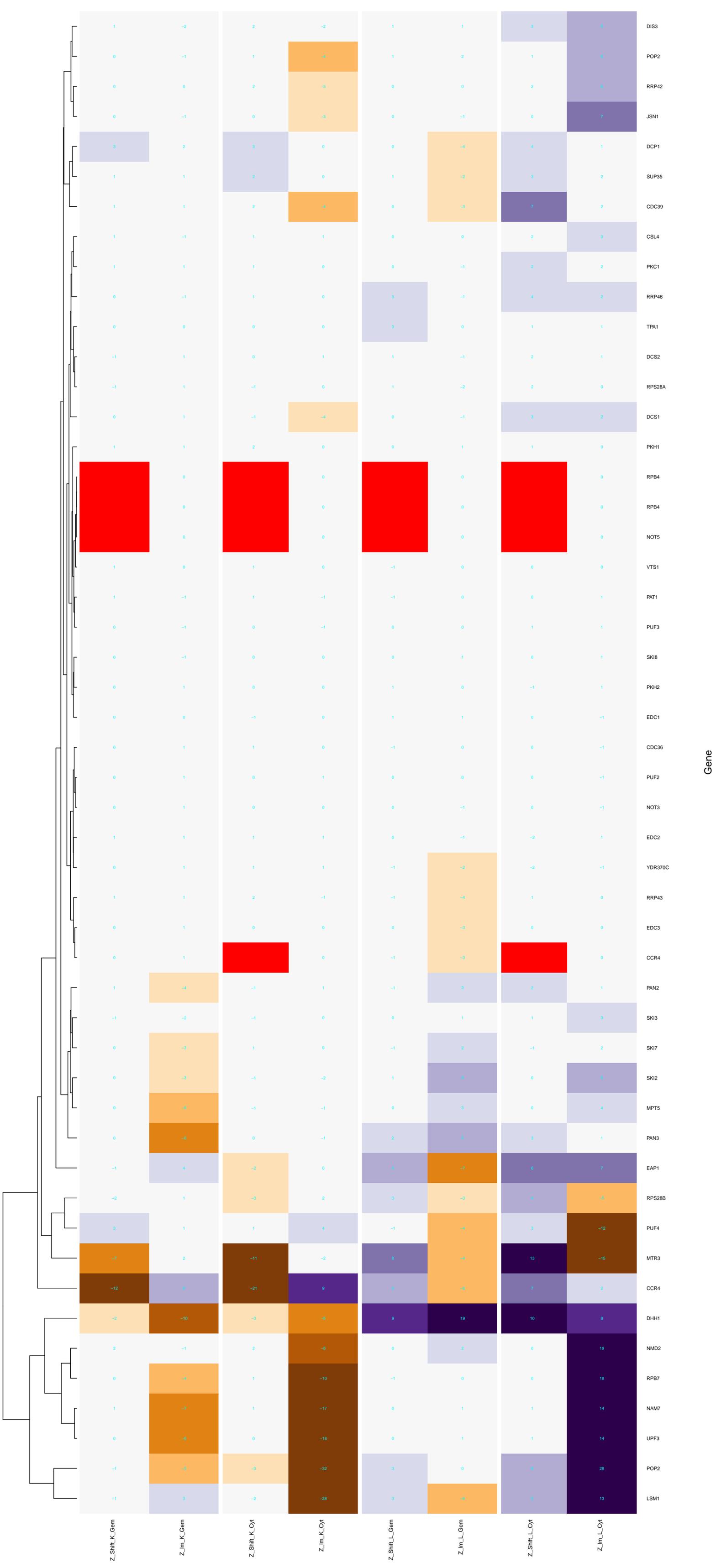


nuclear-transcribed mRNA catabolic process, nonsense-mediated decay

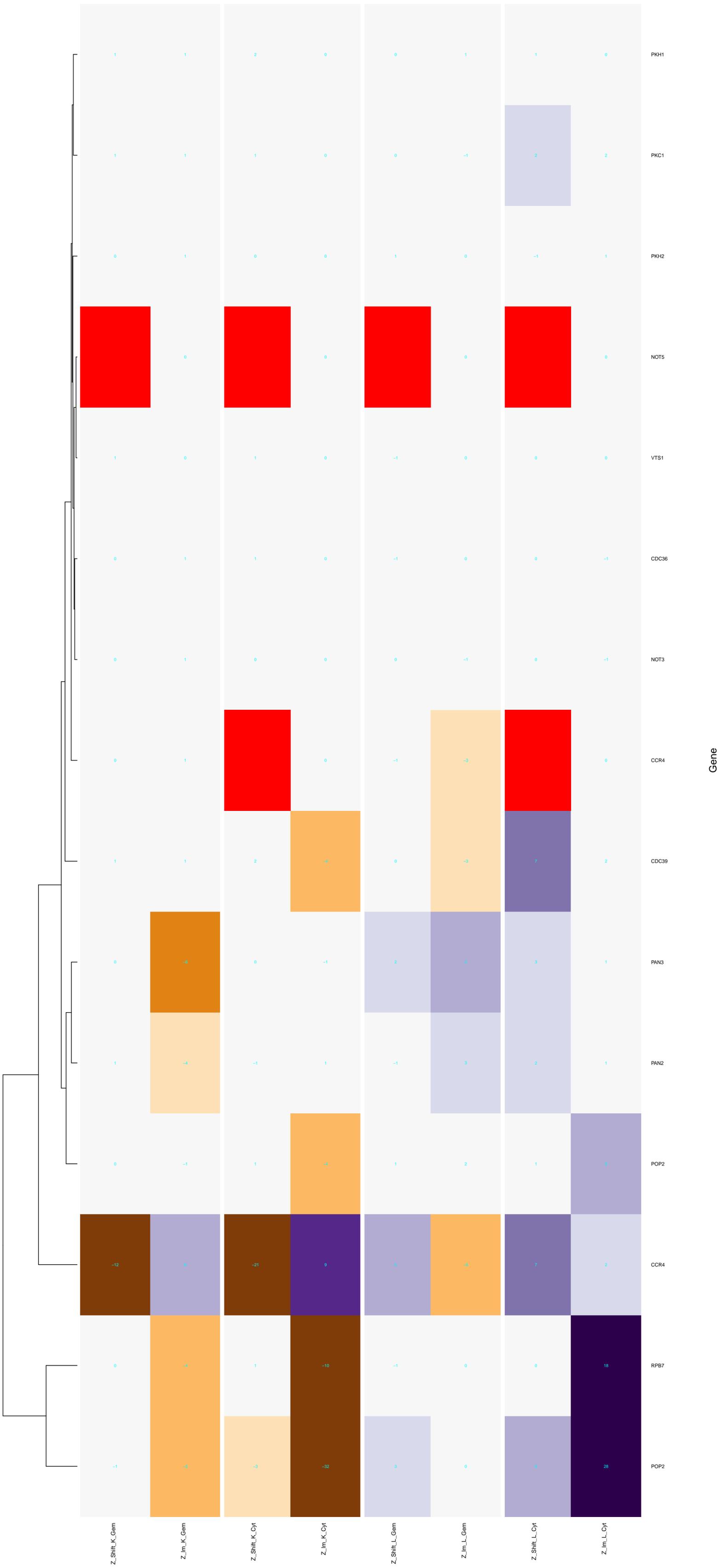
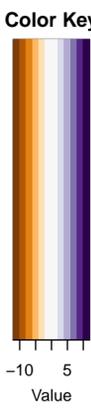




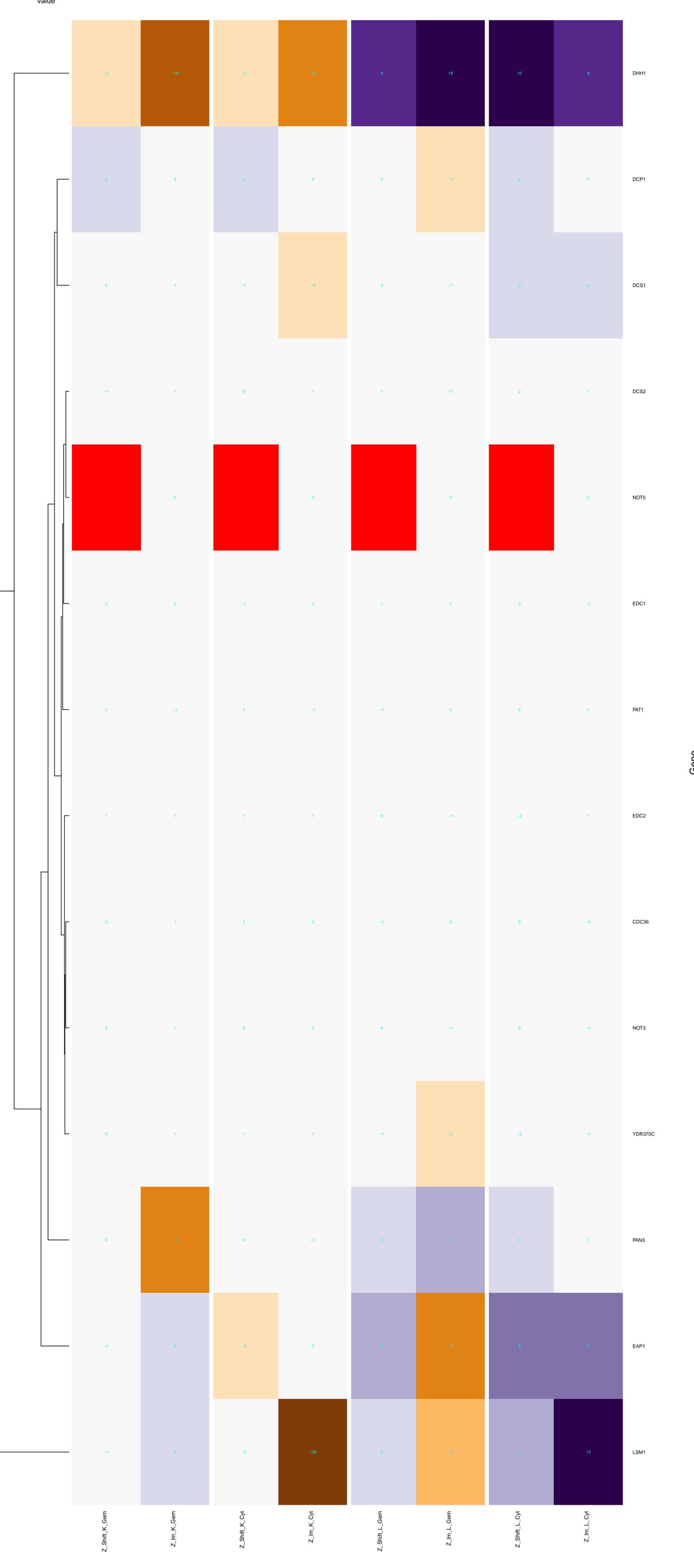
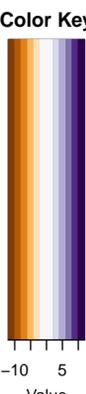
nuclear-transcribed mRNA catabolic process, deadenylation-dependent decay



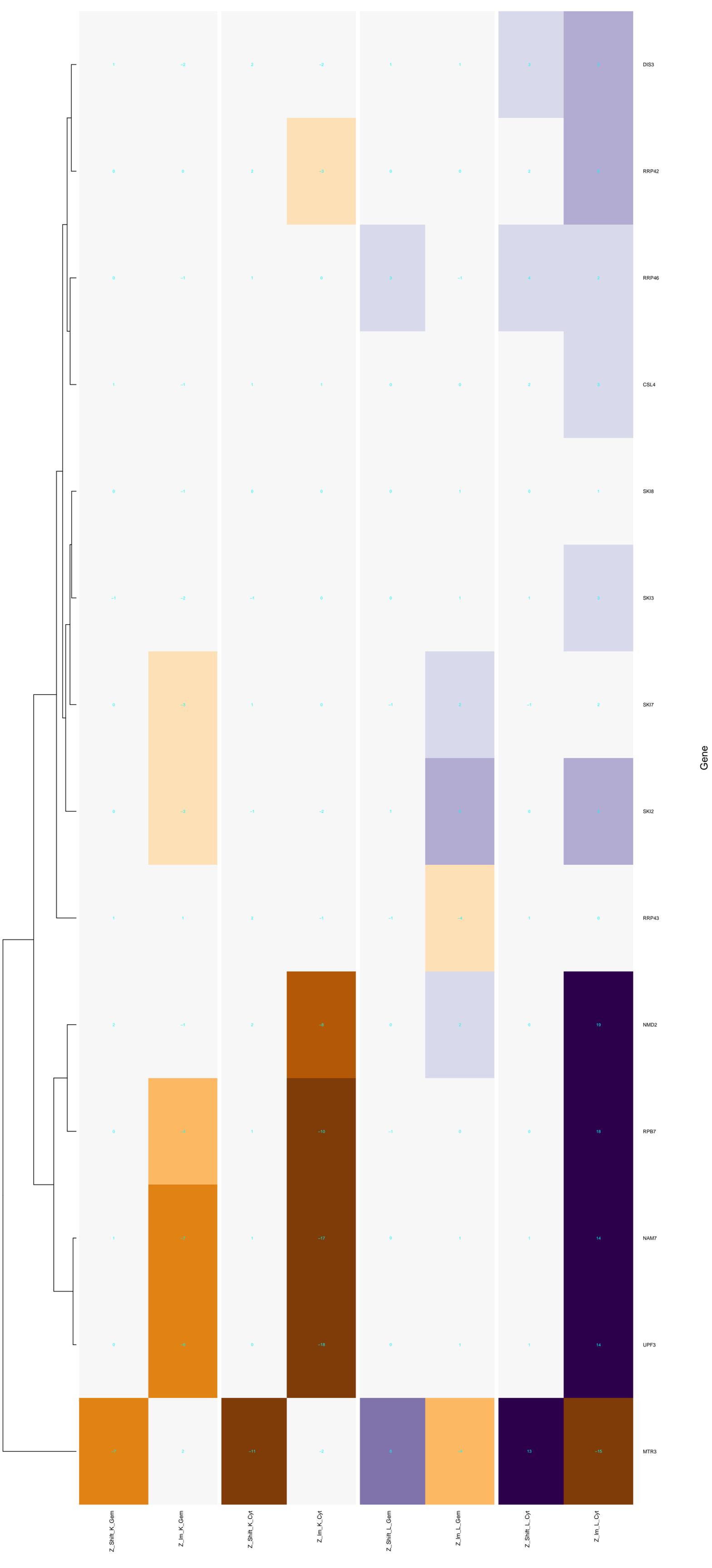
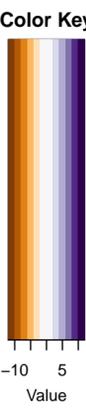
nuclear-transcribed mRNA poly(A) tail shortening

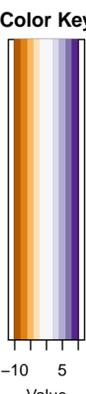


deadenylation-dependent decapping of nuclear-transcribed mRNA

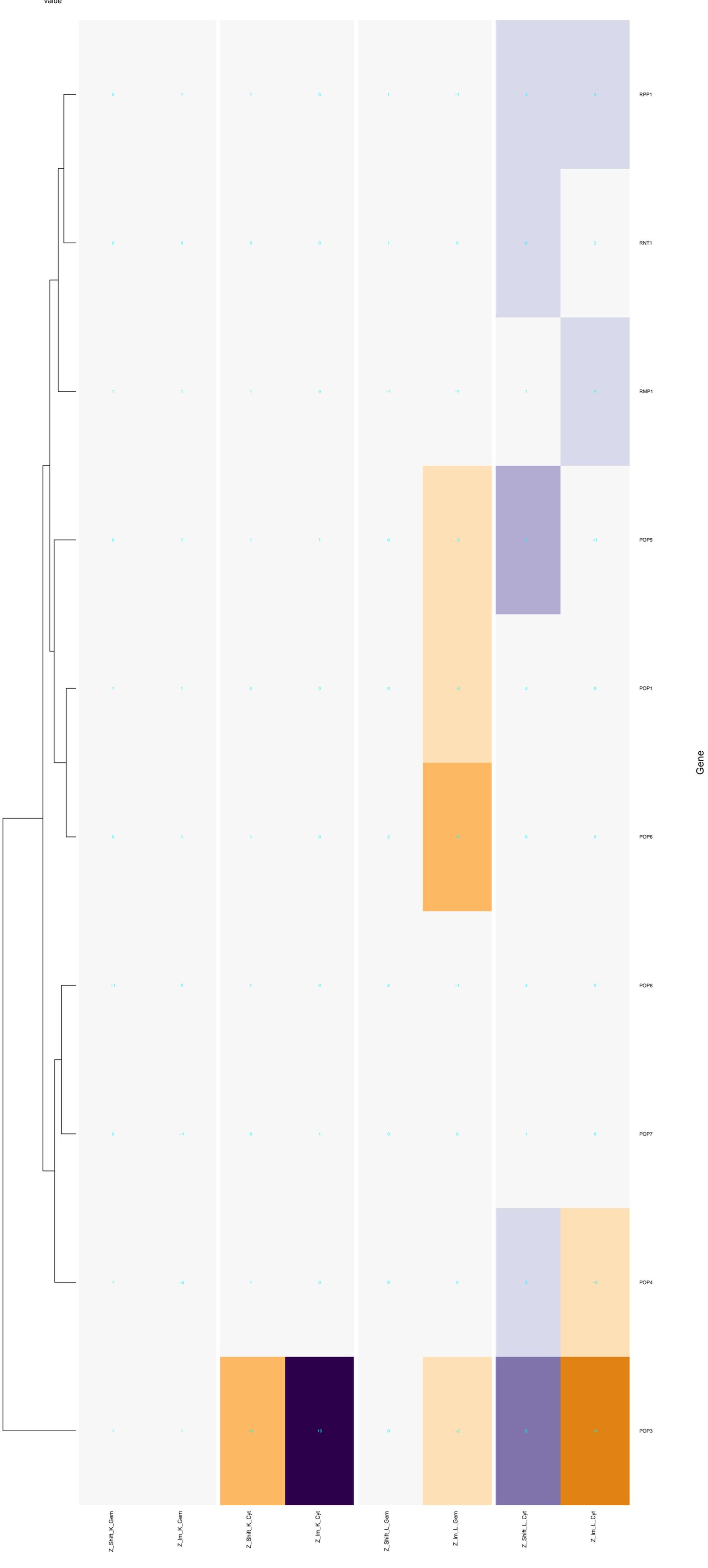


nuclear-transcribed mRNA catabolic process, exonucleolytic



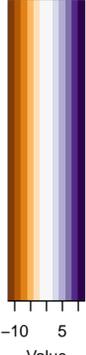


ar-transcribed mRNA catabolic process, endonucleolytic cleavage-dependent decay

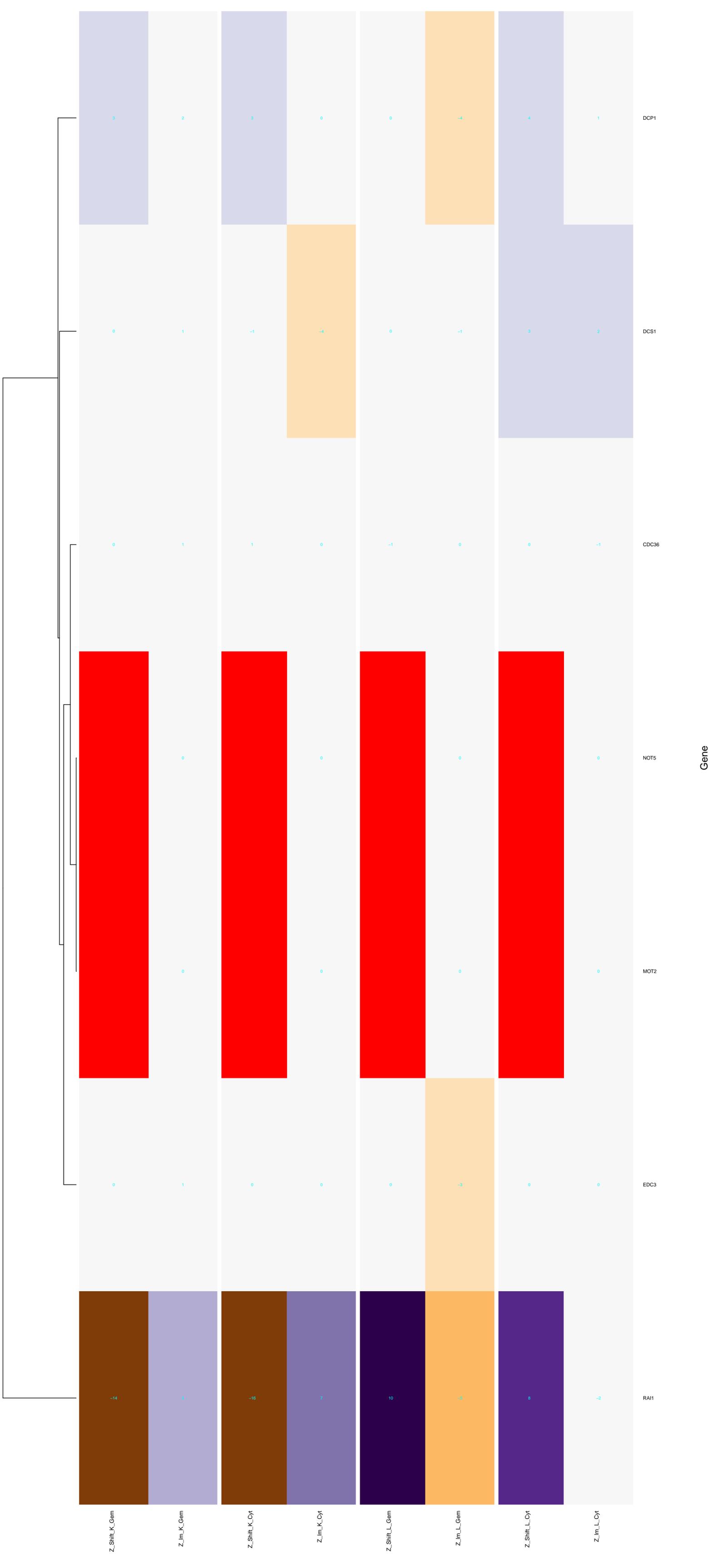


Gene

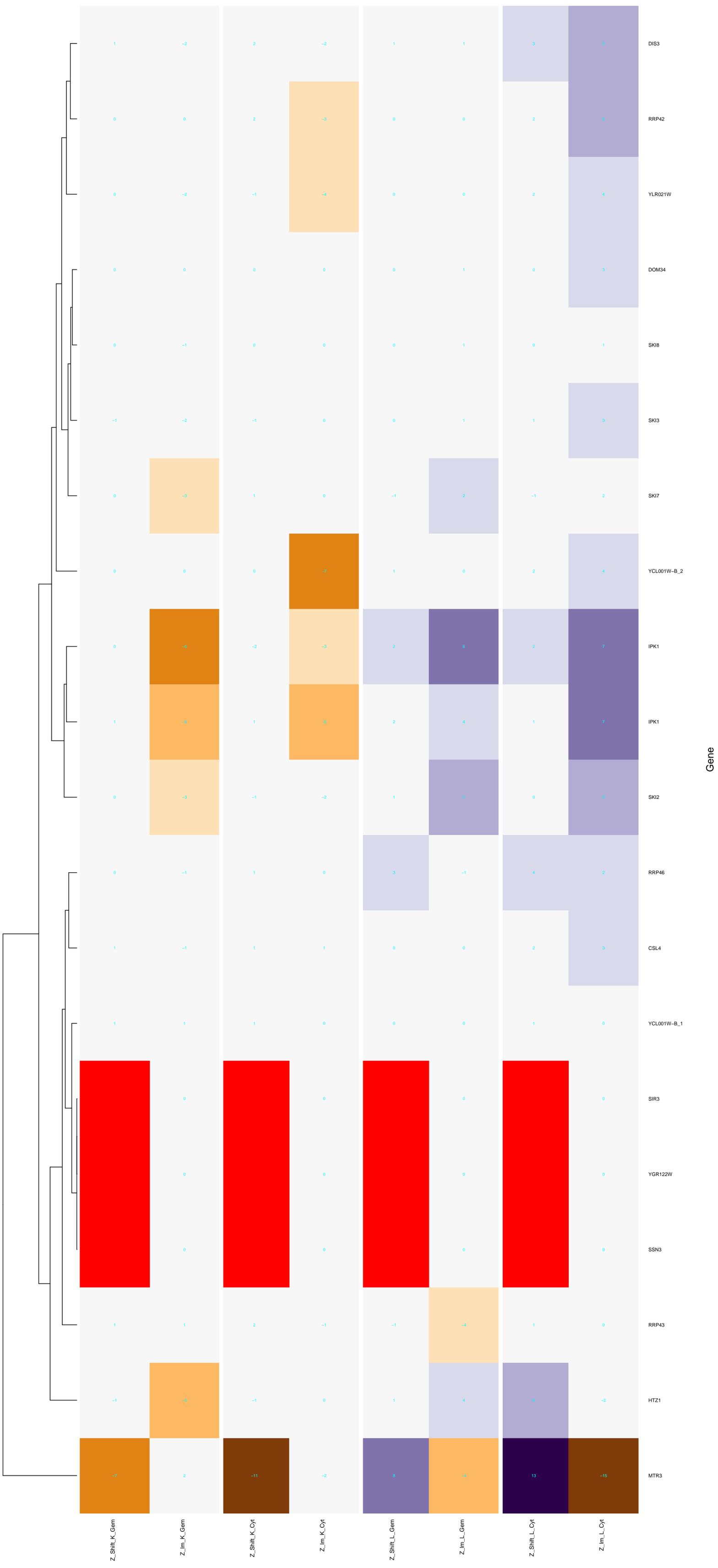
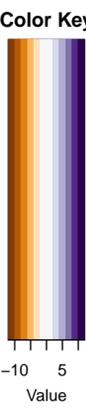
Color Key



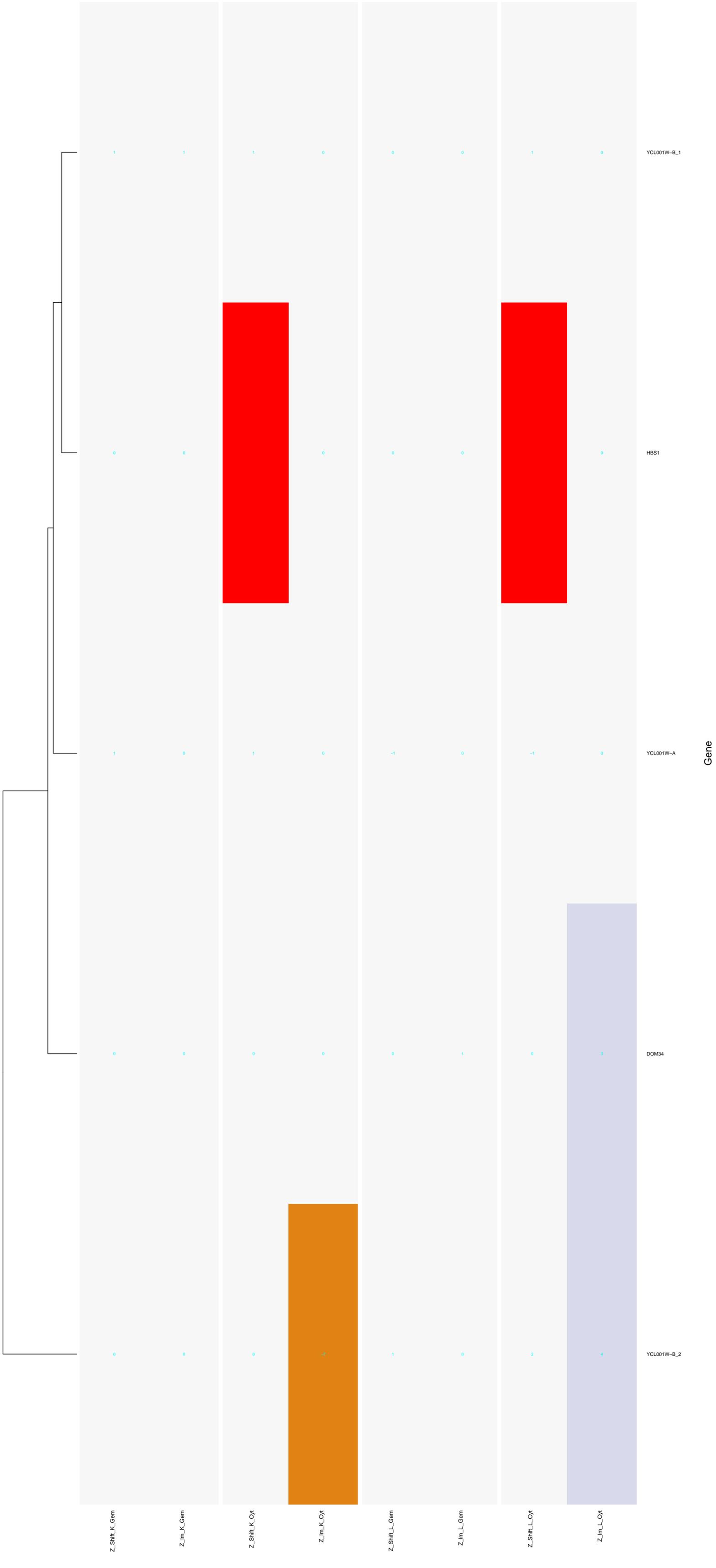
uclear-transcribed mRNA catabolic process, deadenylation-independent decay



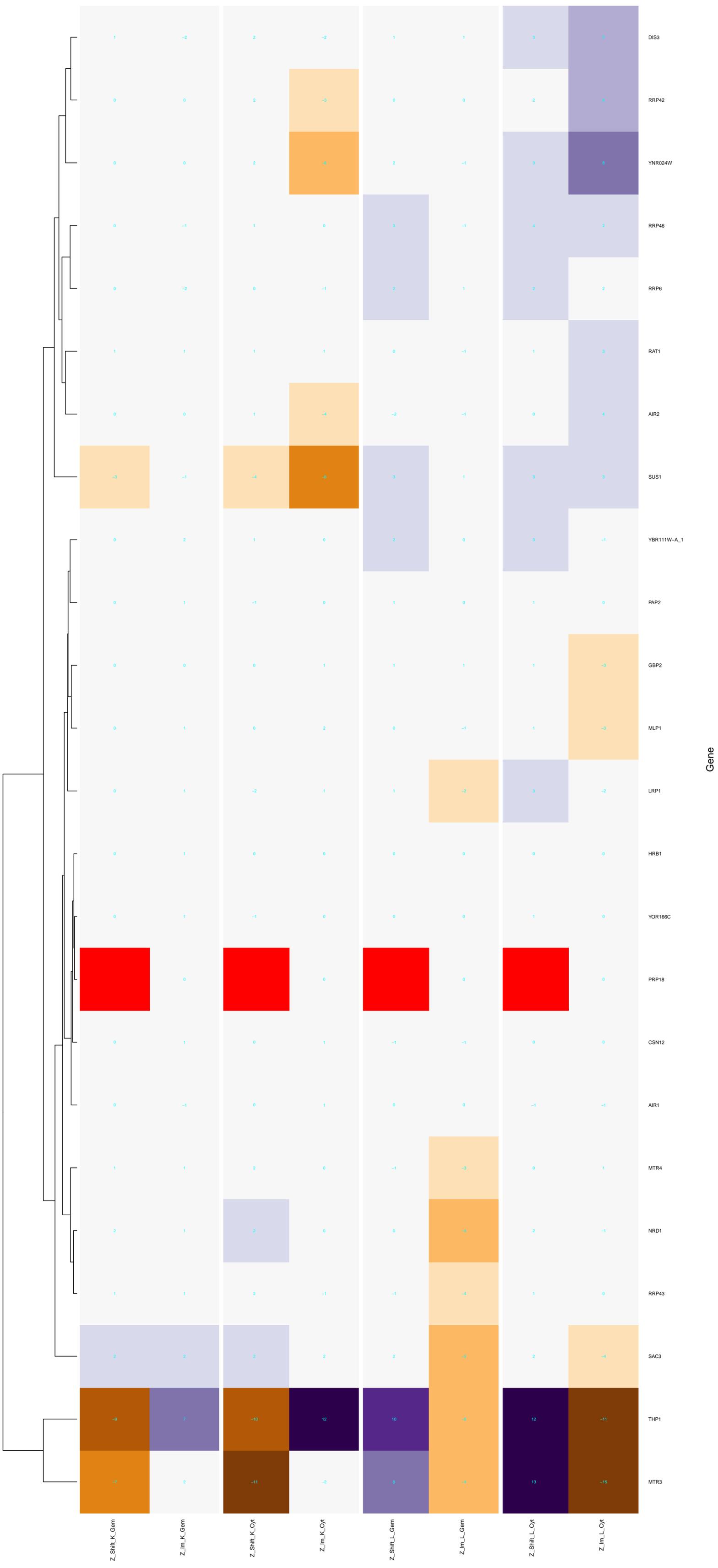
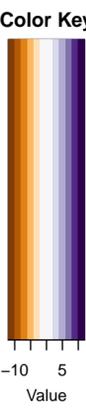
nuclear-transcribed mRNA catabolic process, non-stop decay



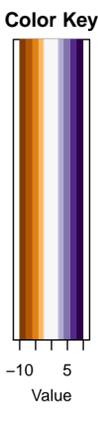
nuclear-transcribed mRNA catabolic process, no-go decay



nuclear mRNA surveillance

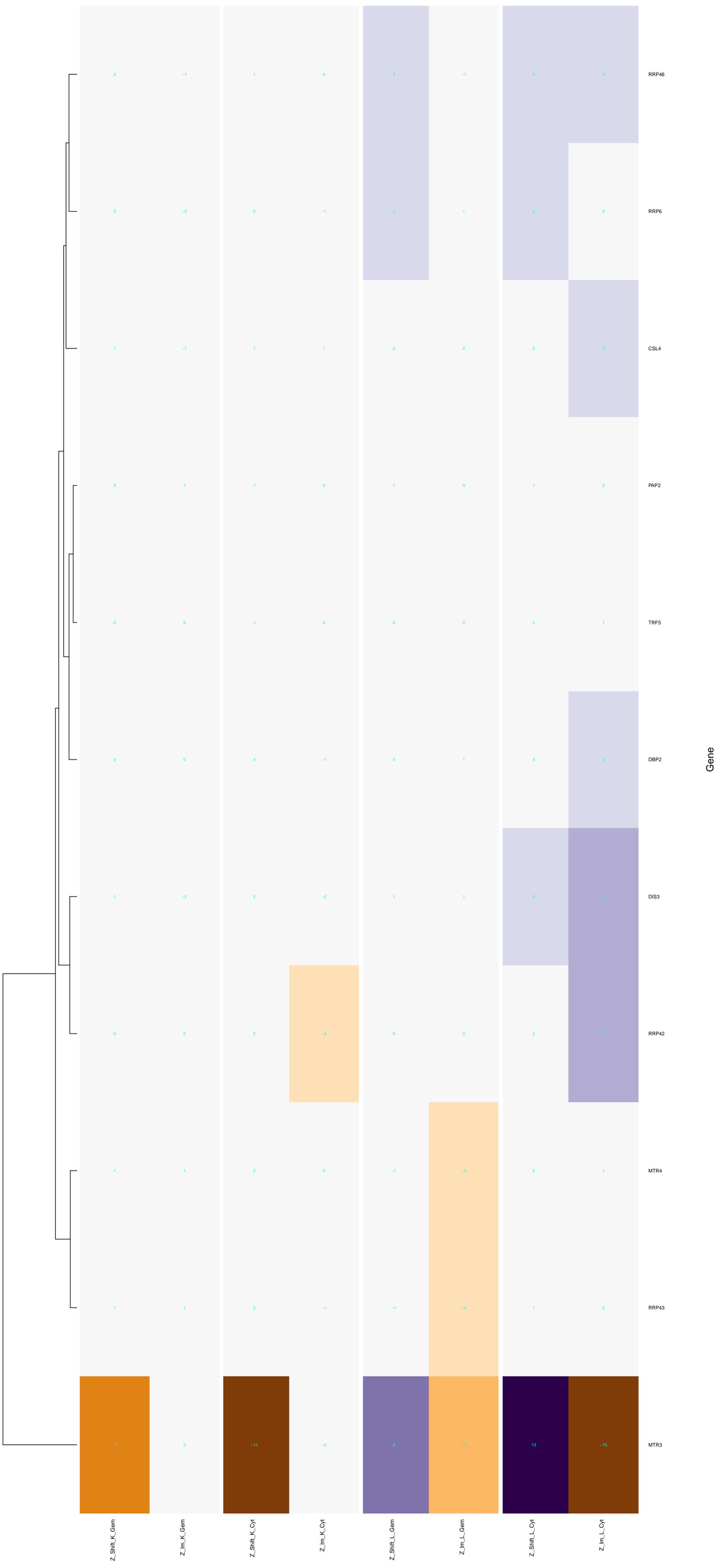
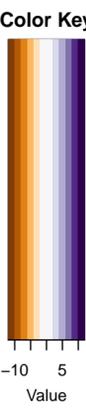


histone mRNA catabolic process

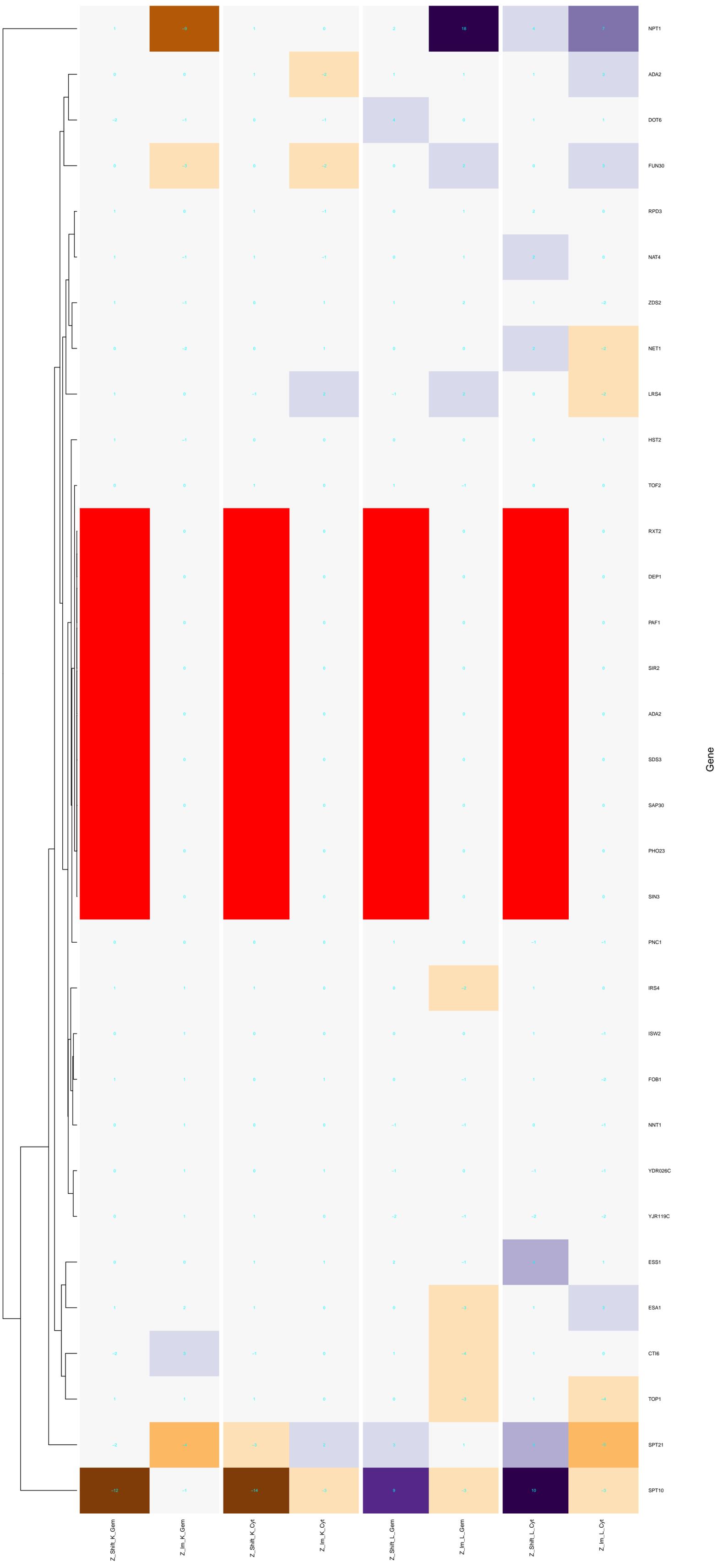
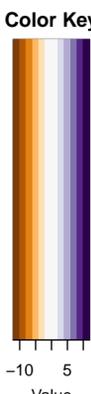


Gene

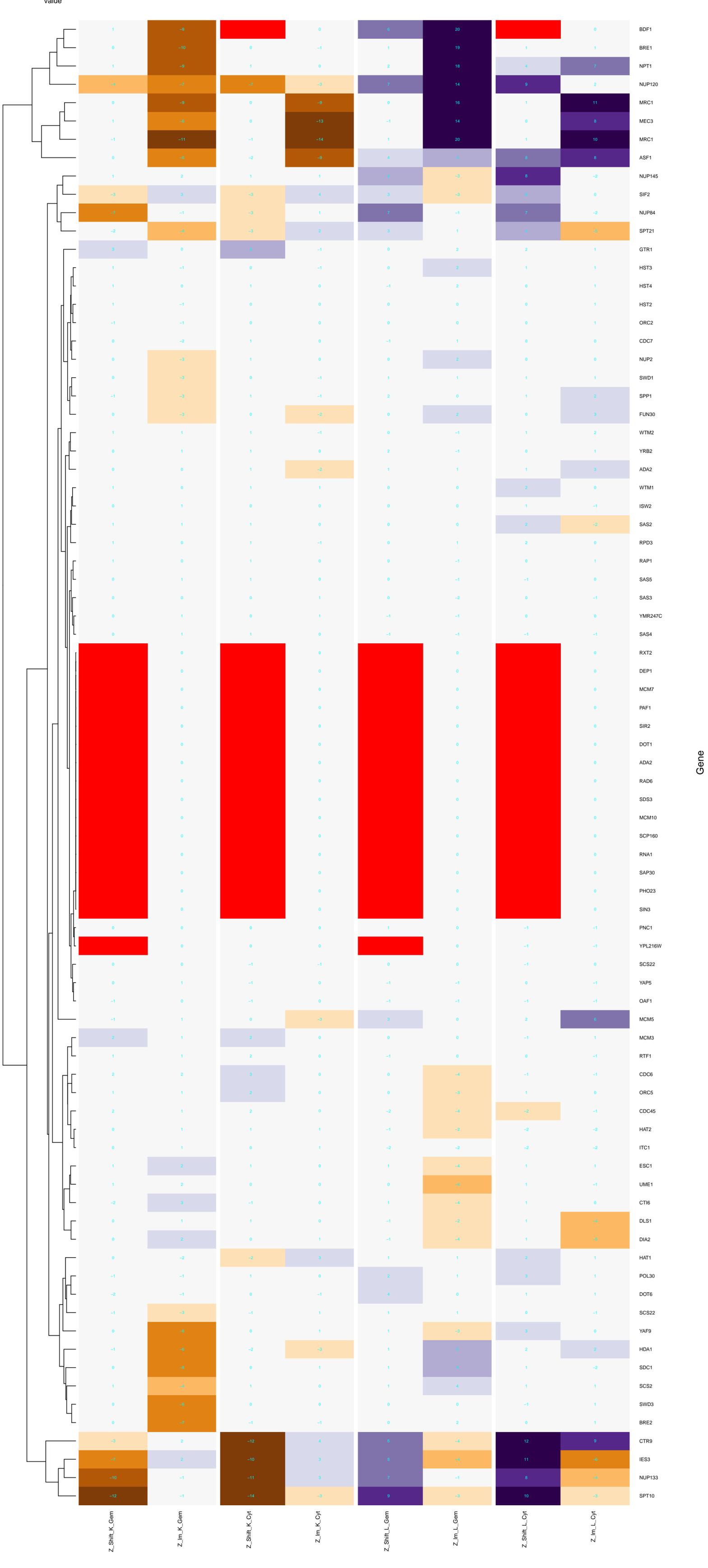
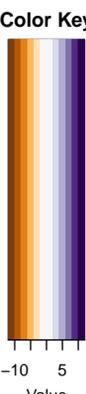
nuclear polyadenylation–dependent mRNA catabolic process



chromatin silencing at rDNA

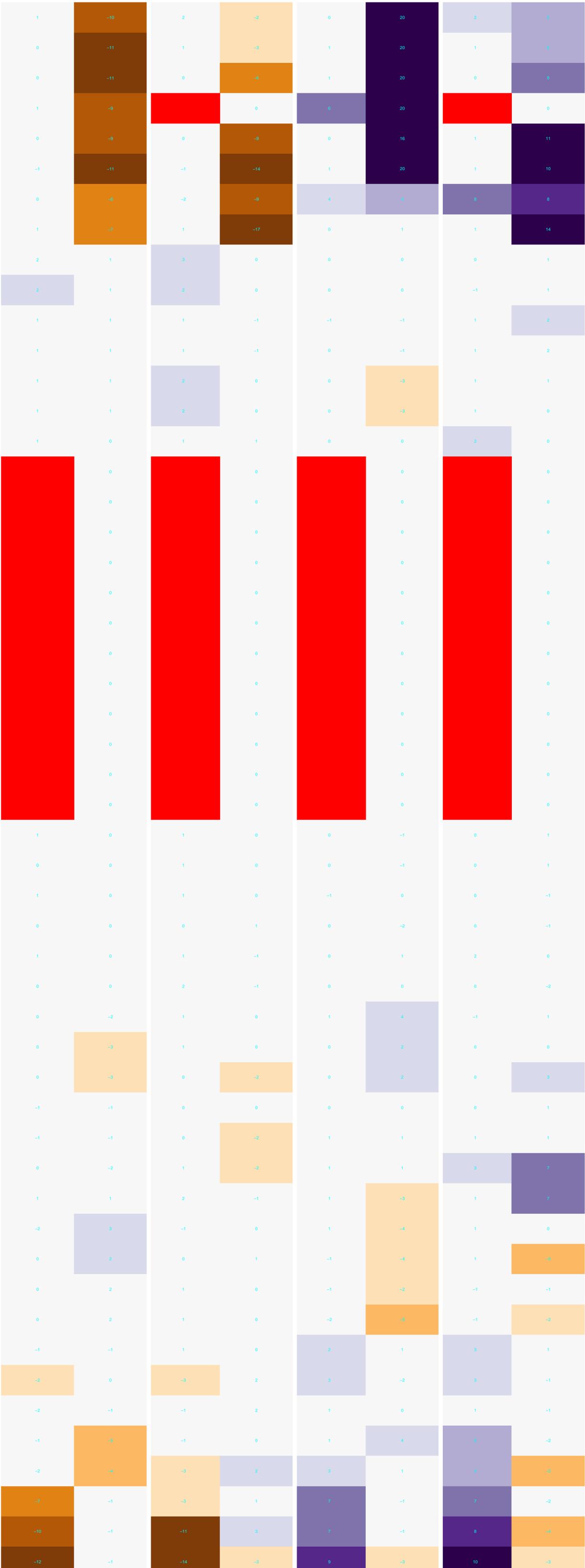
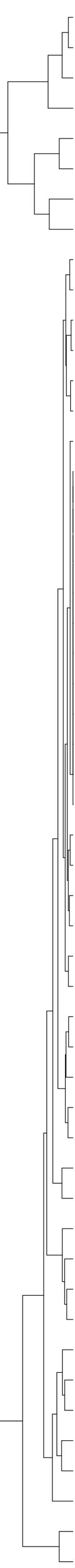
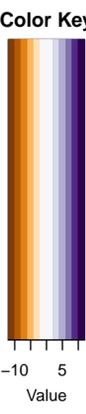


chromatin silencing at telomere



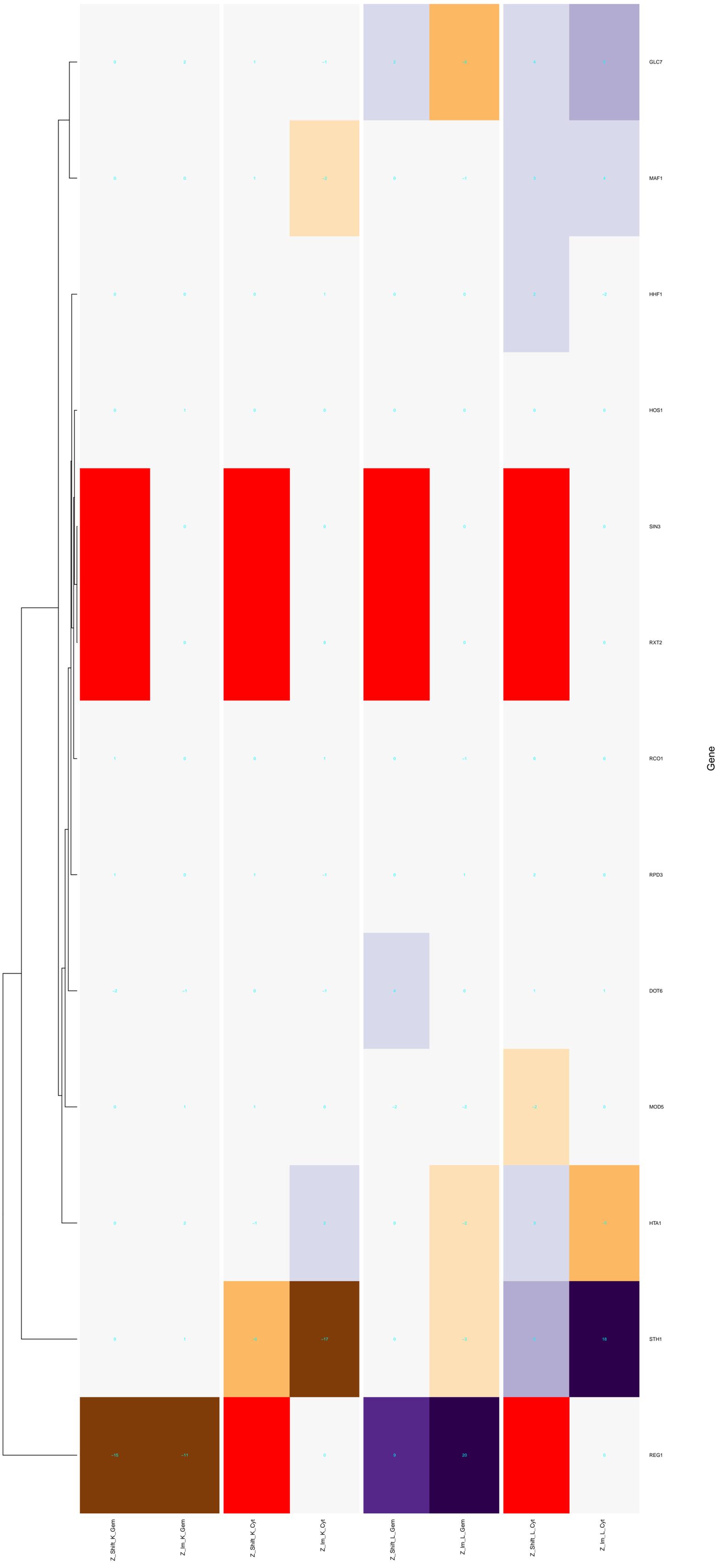
Gene

chromatin silencing at silent mating-type cassette

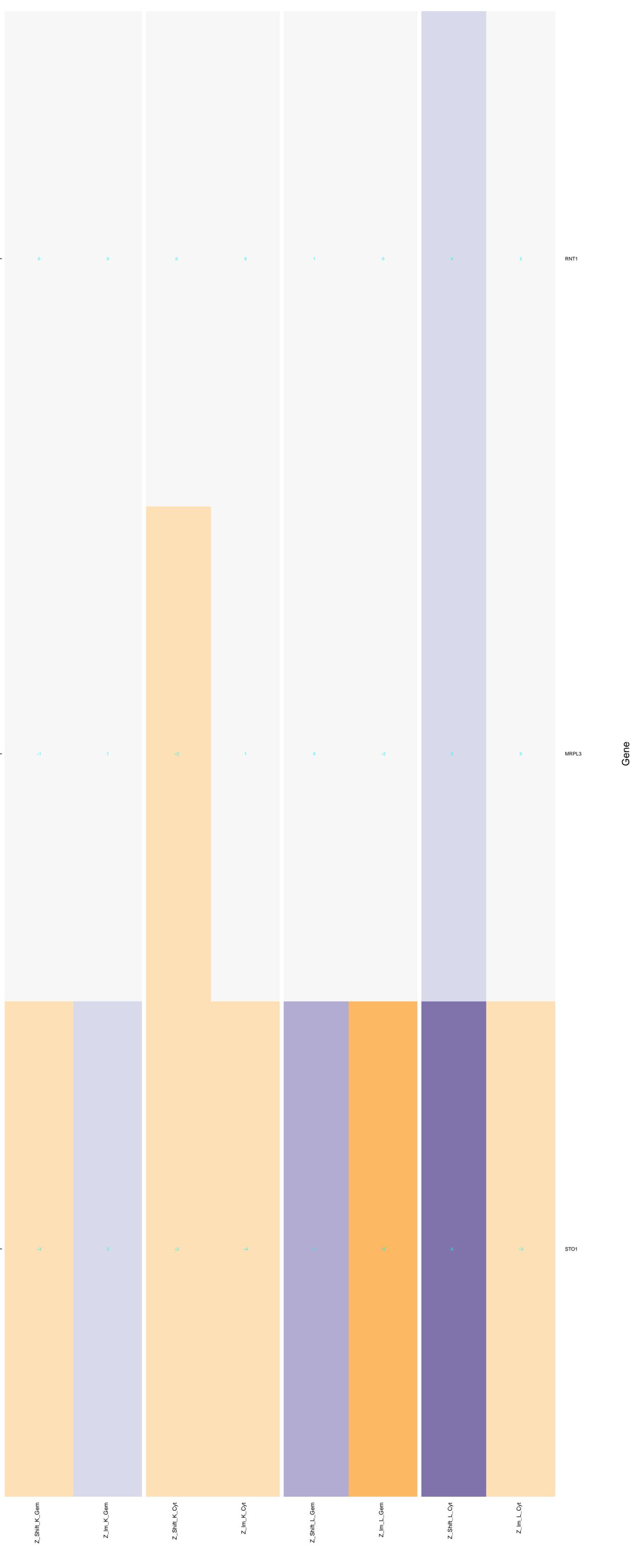
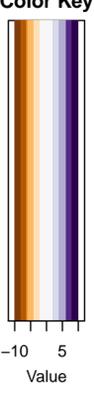


Gene

transfer RNA gene-mediated silencing



posttranscriptional gene silencing by RNA

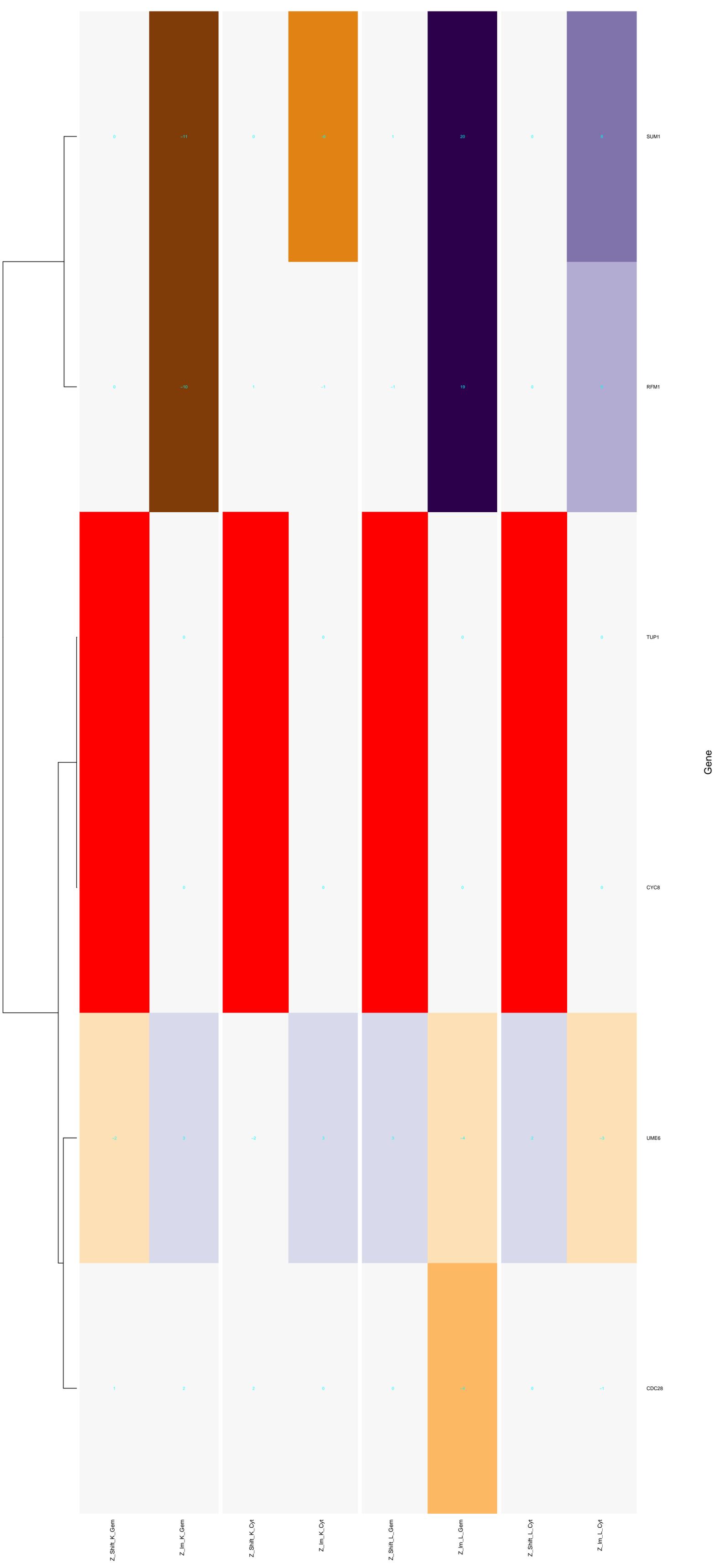
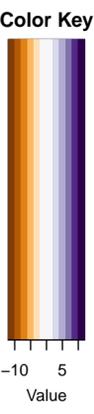


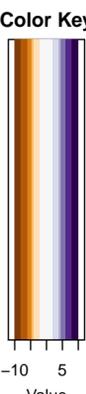
3'-UTR-mediated mRNA destabilization



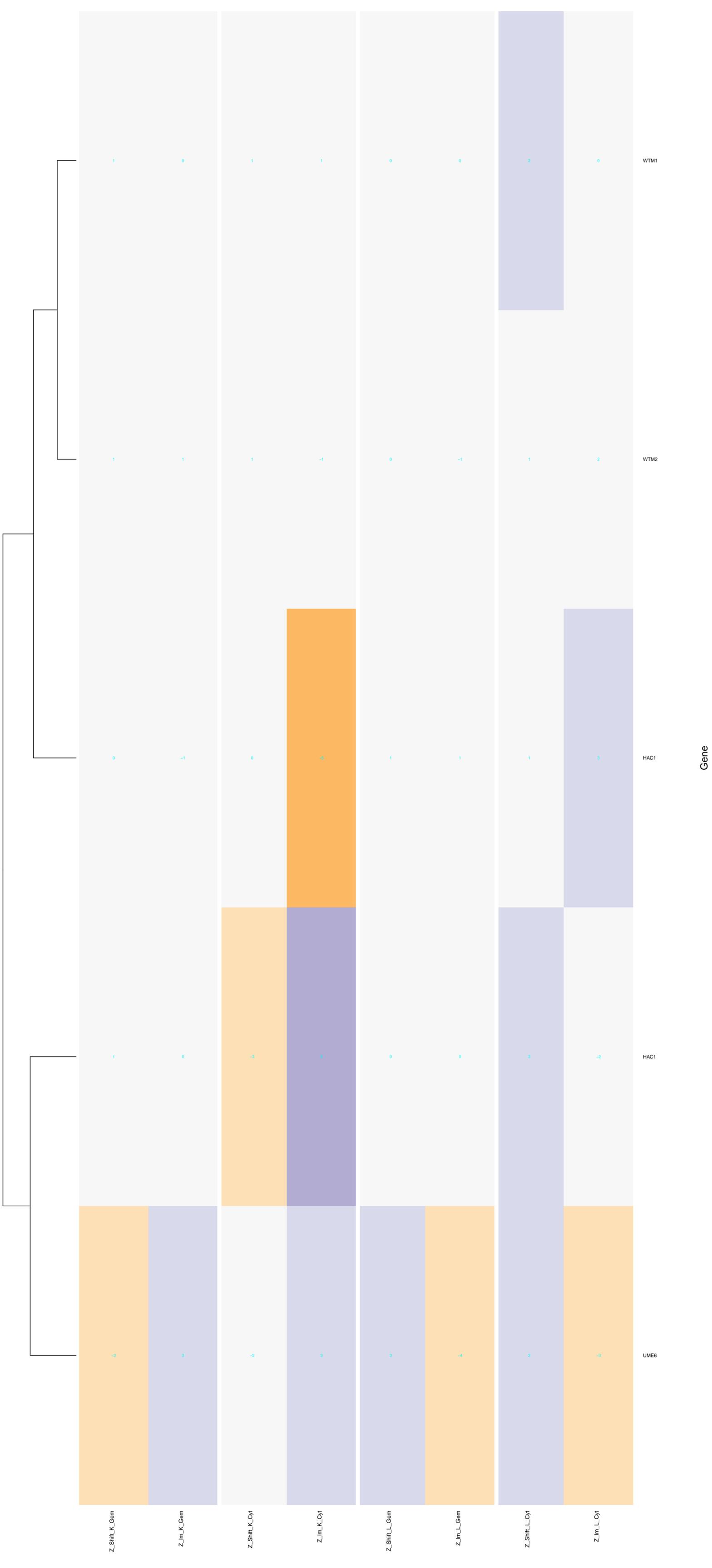
Gene

Gene regulation of transcription from RNA polymerase II promoter during mitotic cell cycle

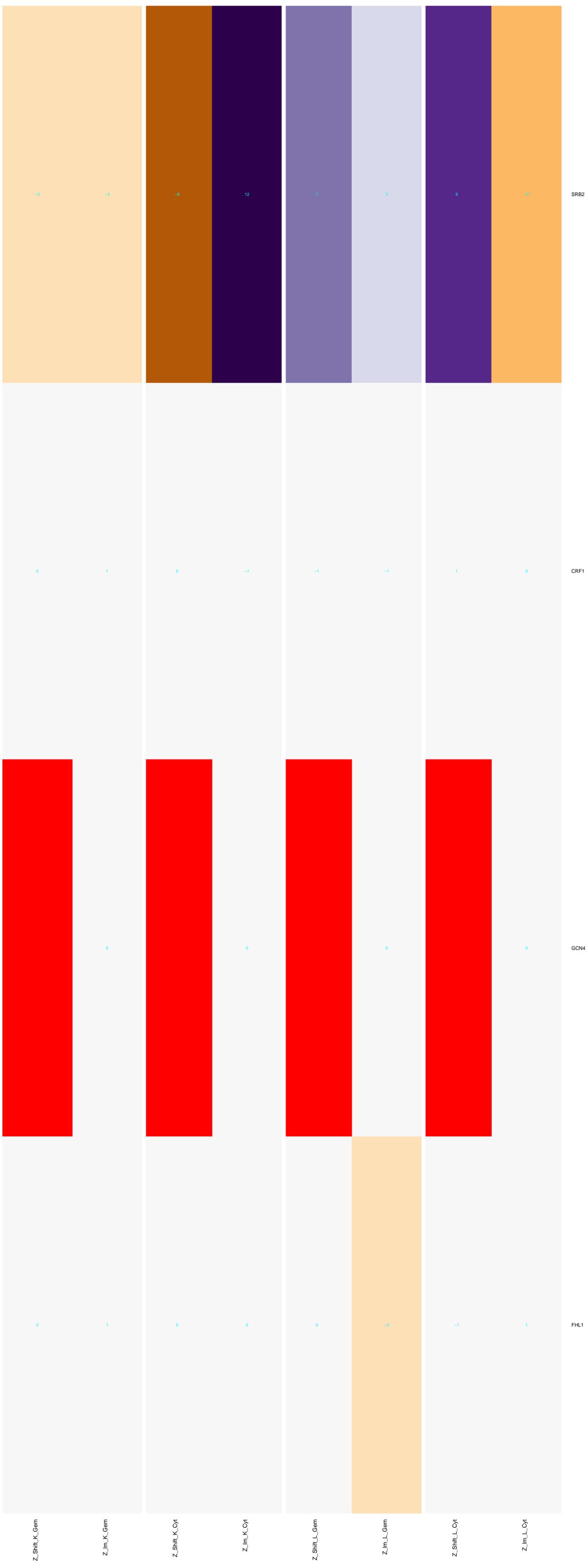
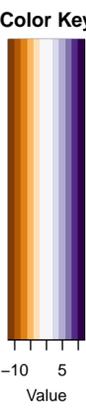




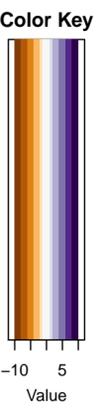
regulation of transcription from RNA polymerase II promoter involved in meiotic cell cycle



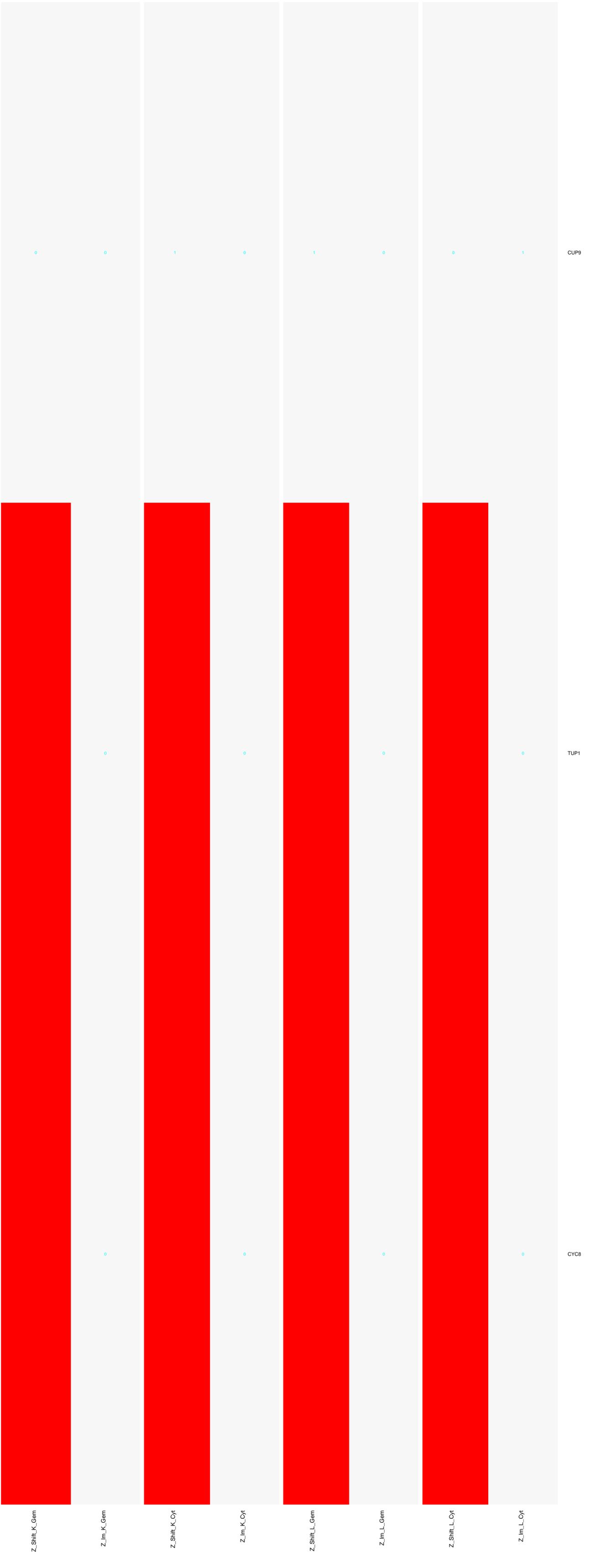
e regulation of ribosomal protein gene transcription from RNA polymerase II promoter



Gene

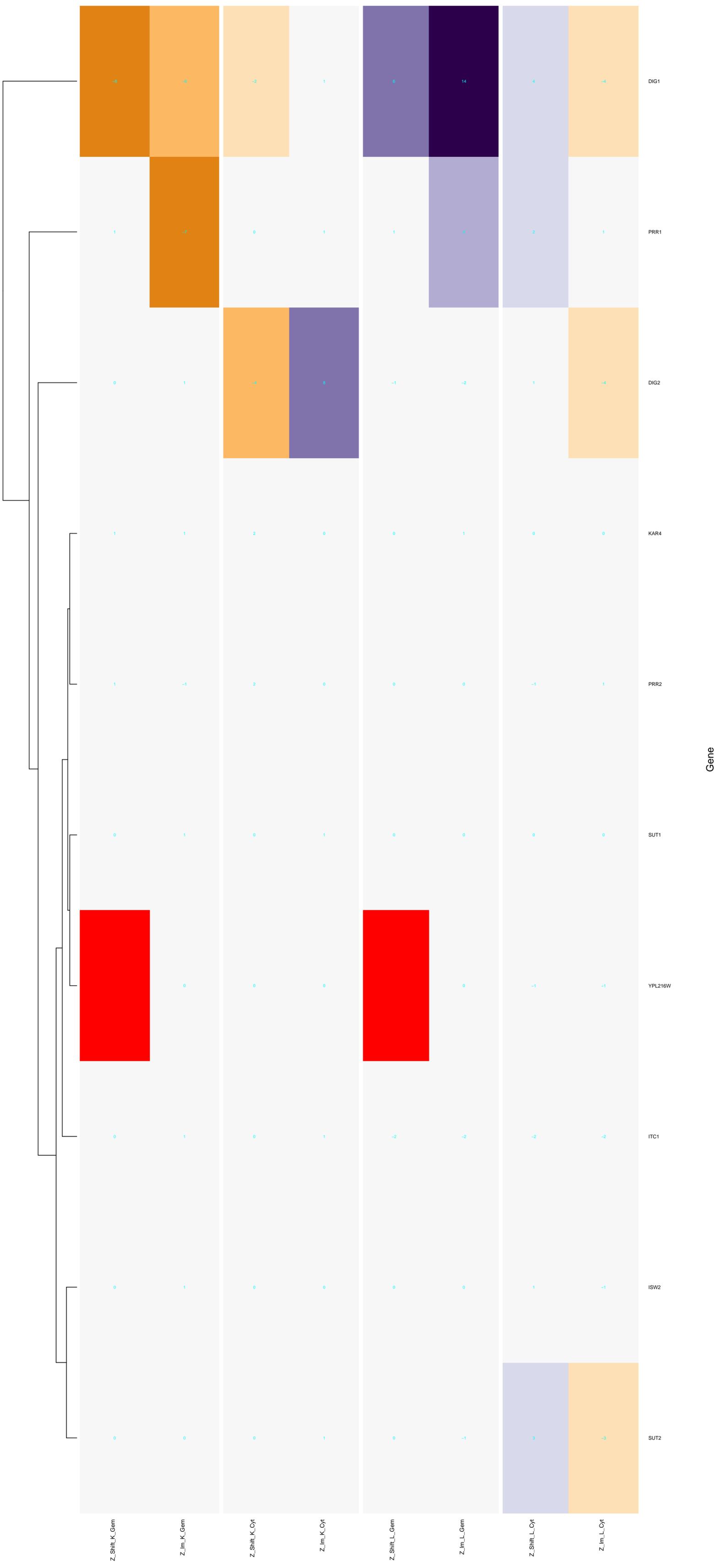
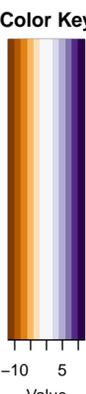


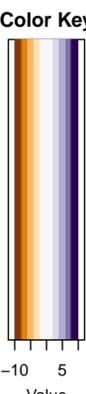
of oligopeptide transport by negative regulation of transcription from RNA polymerase II p



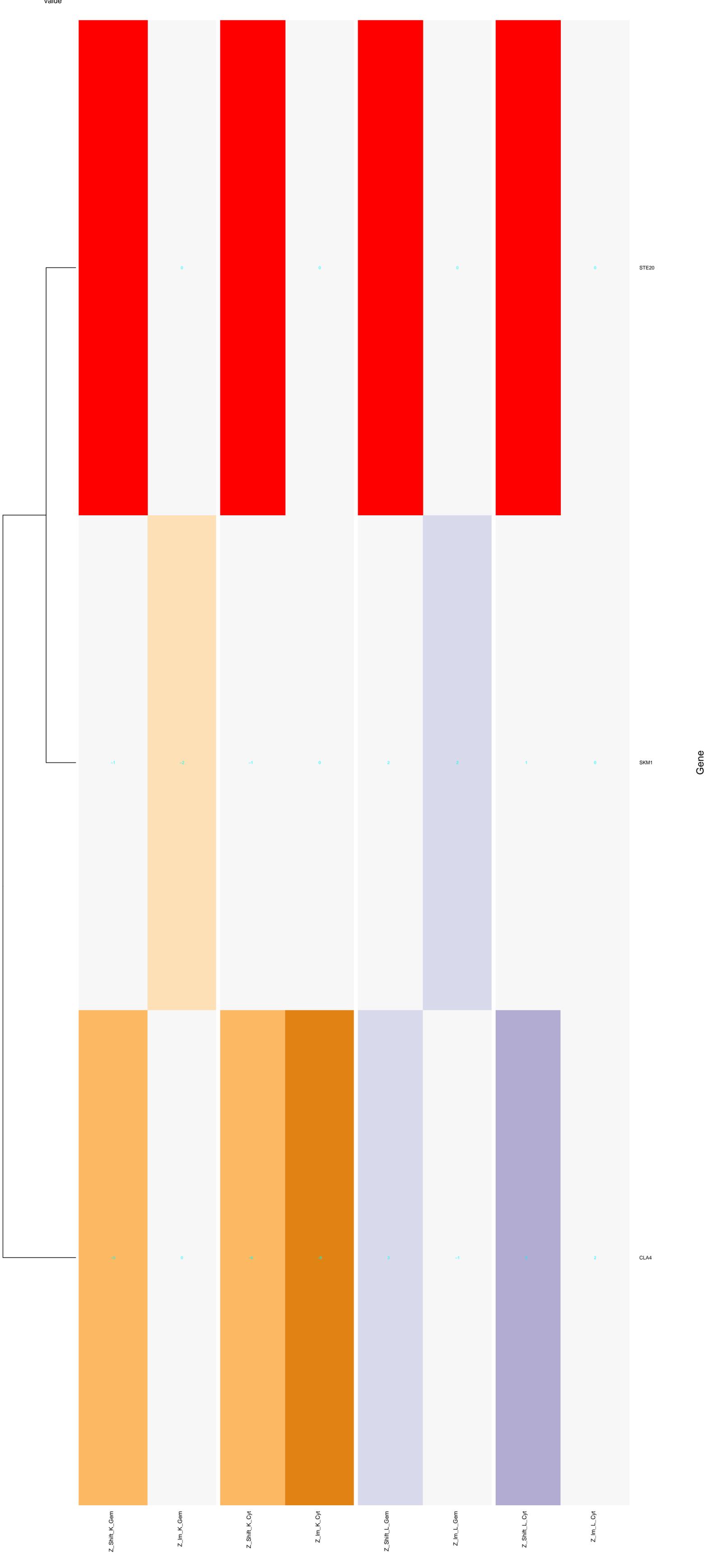
Gene

Active regulation of transcription from RNA polymerase II promoter by pheromones

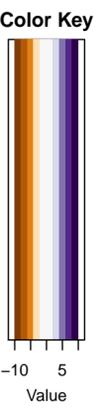




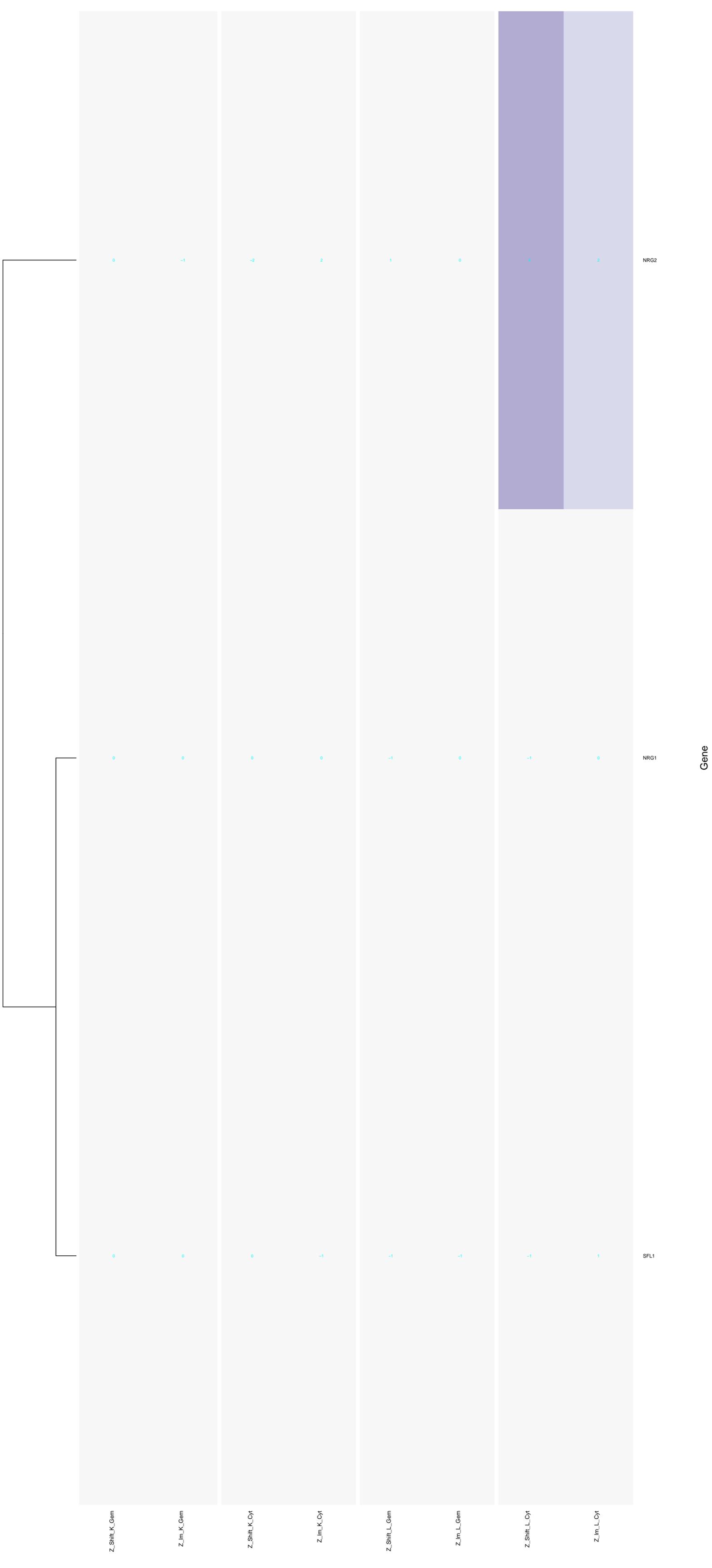
of lipid transport by negative regulation of transcription from RNA polymerase II promoter



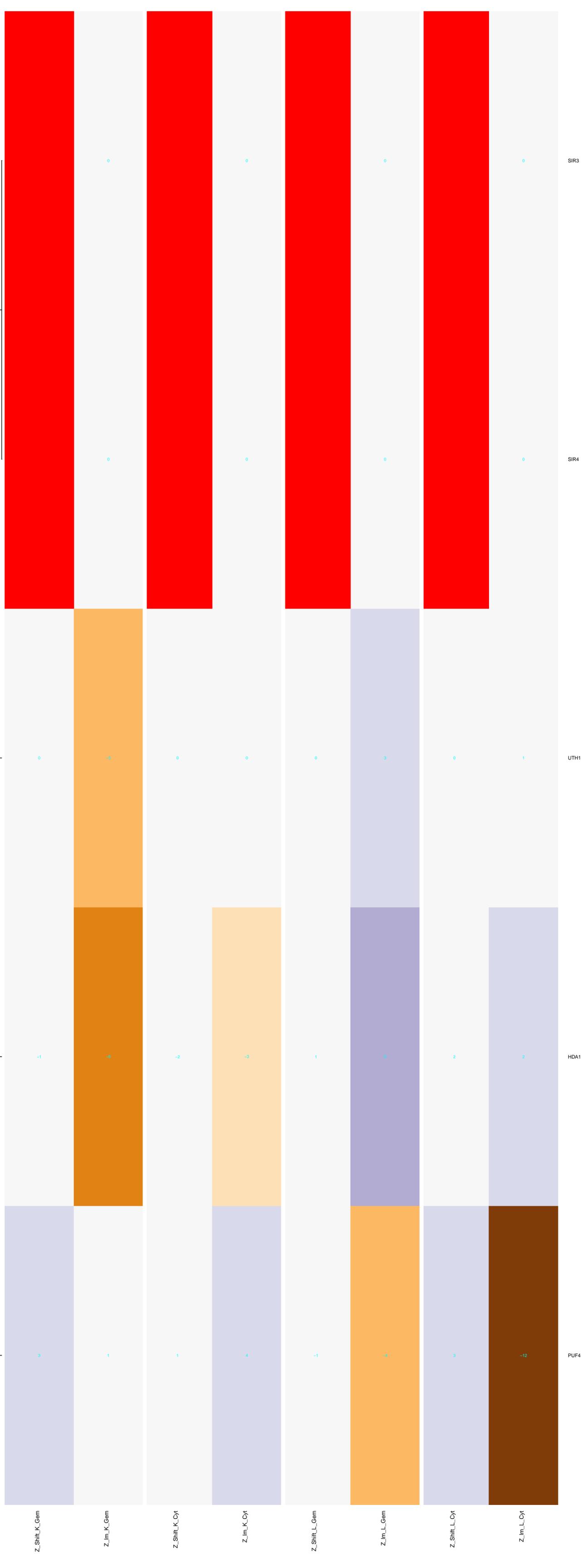
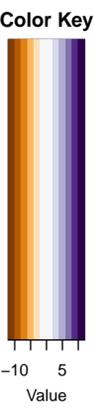
Gene



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

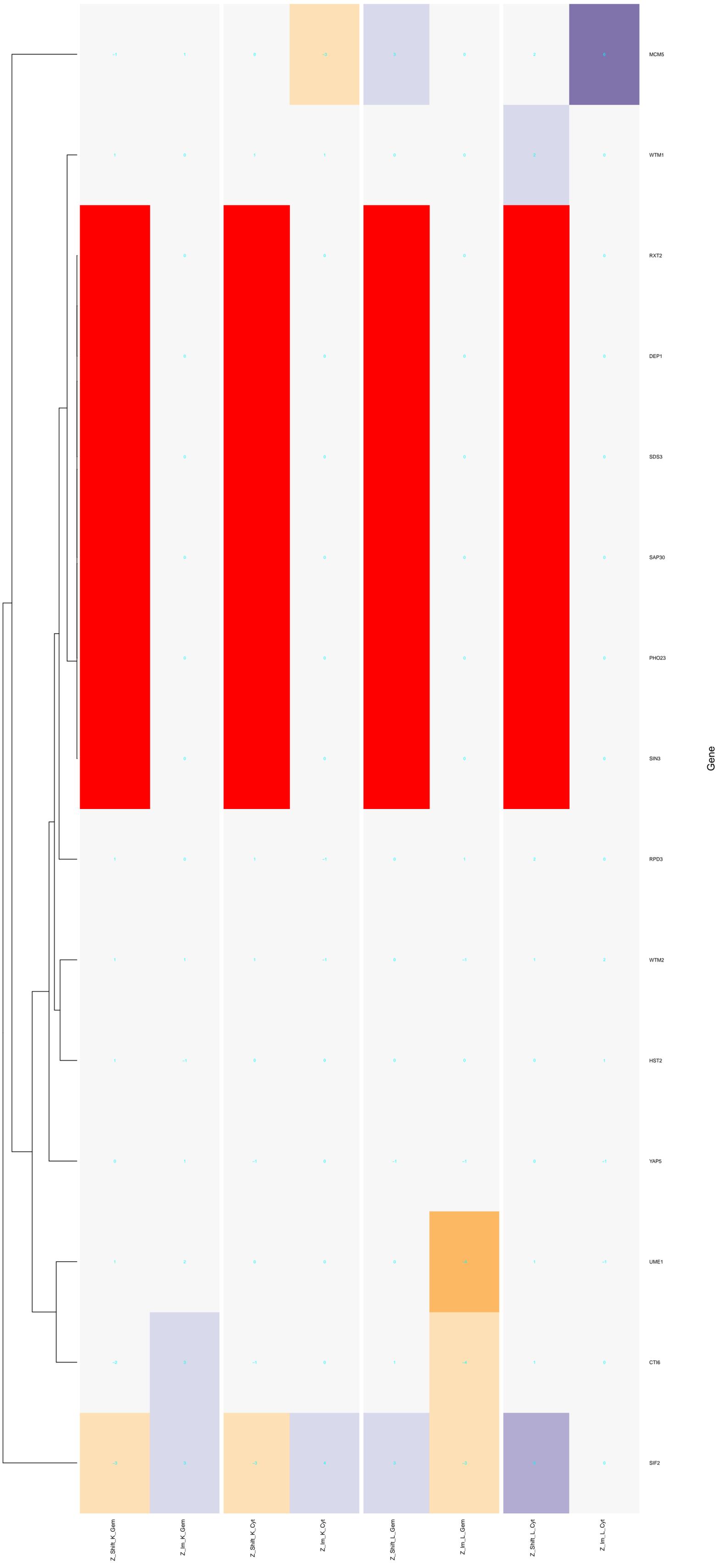
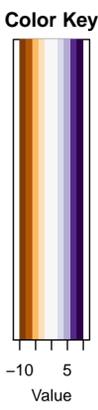


negative regulation of chromatin silencing involved in replicative cell aging

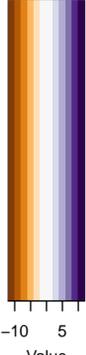


Gene

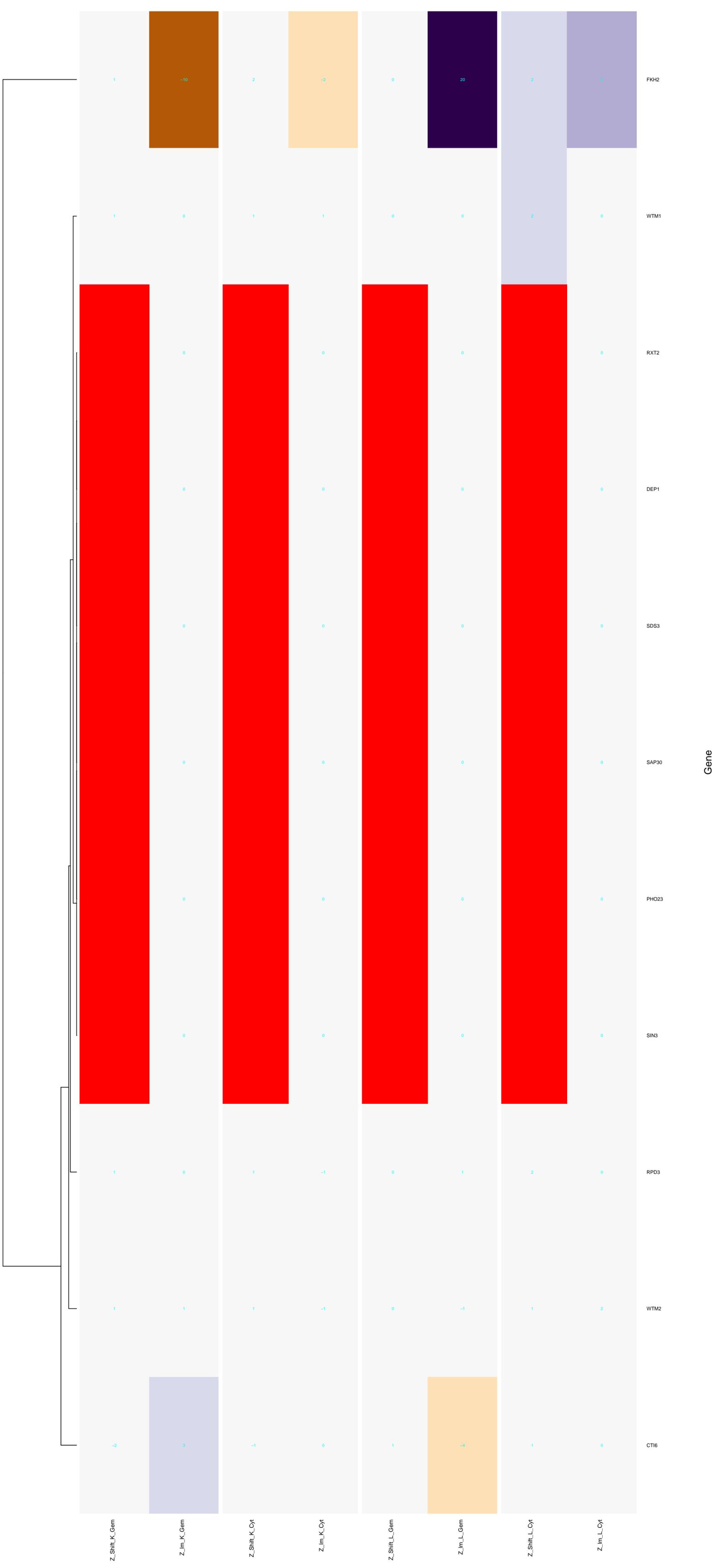
negative regulation of chromatin silencing at telomere



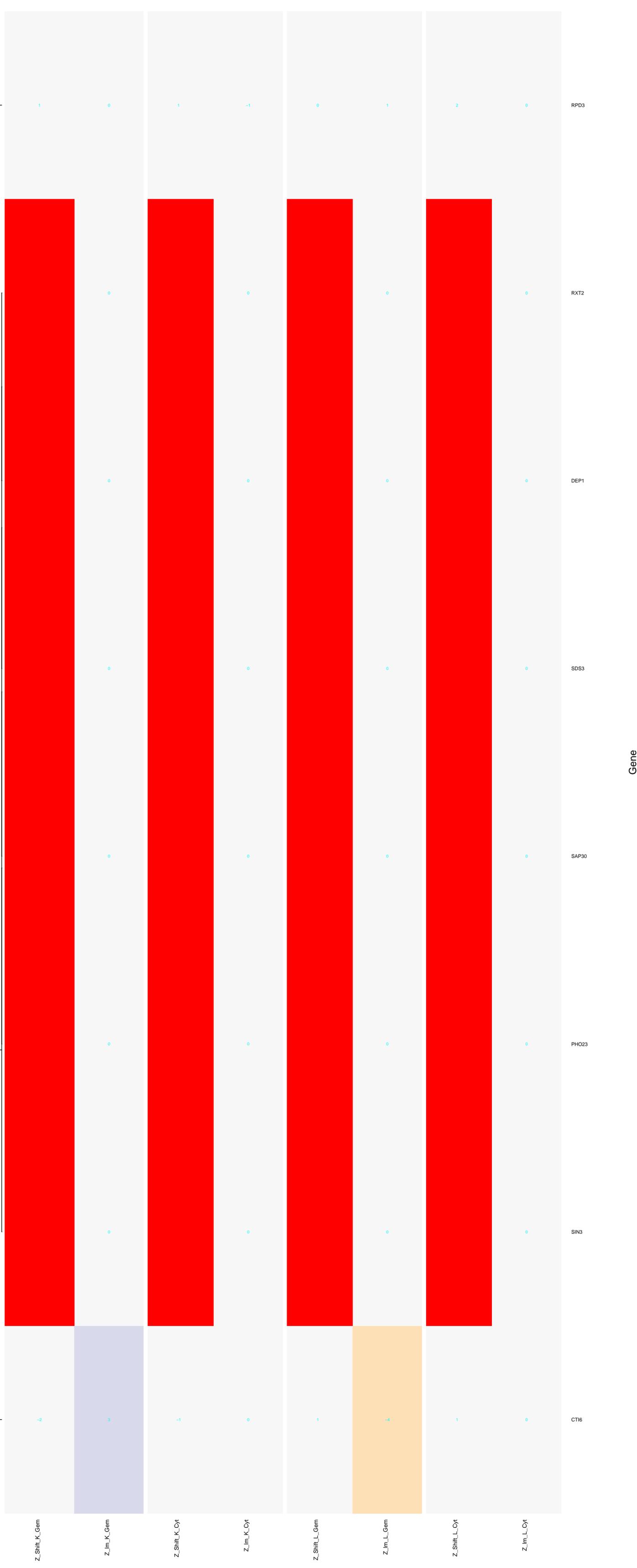
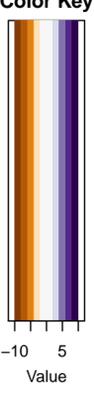
Color Key



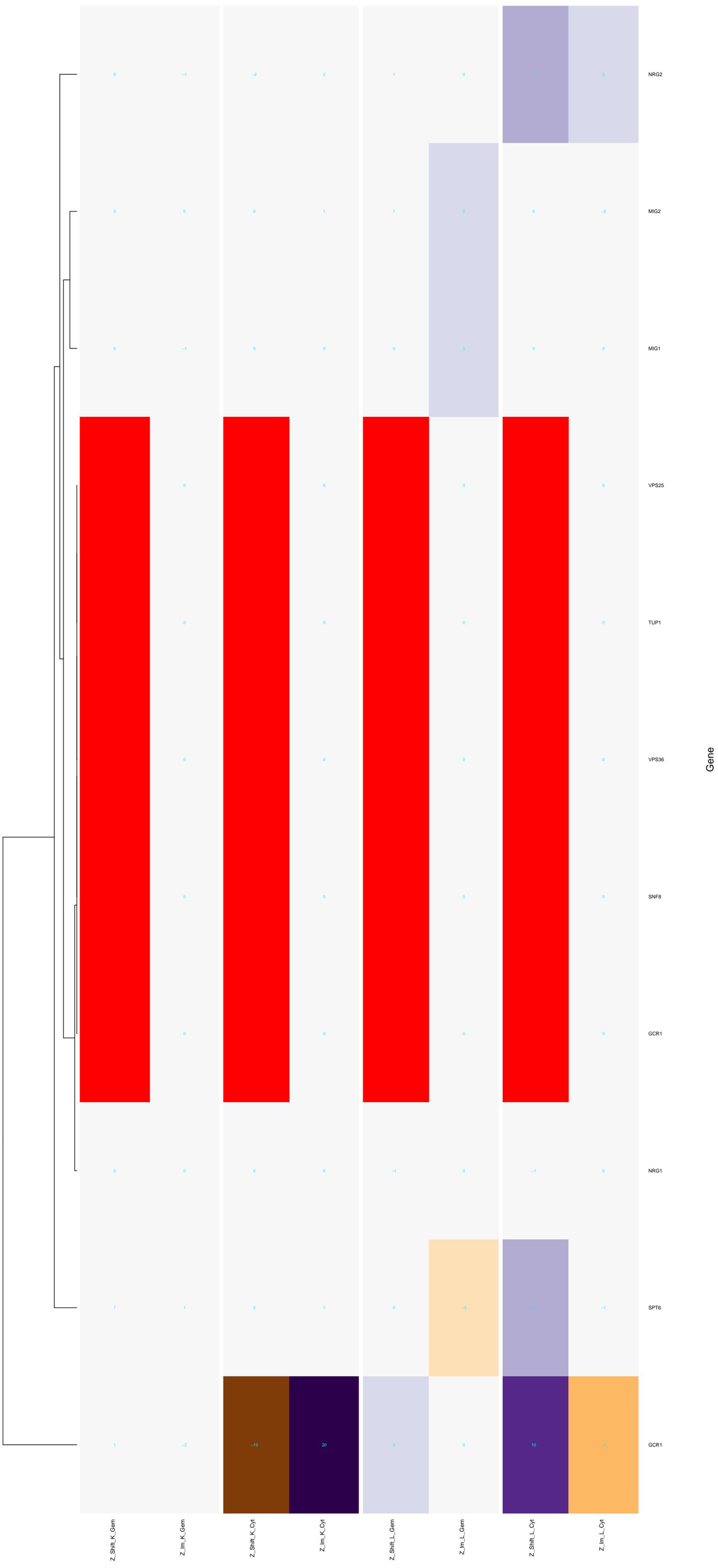
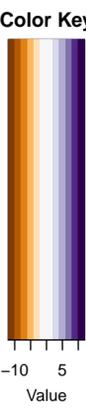
negative regulation of chromatin silencing at silent mating-type cassette

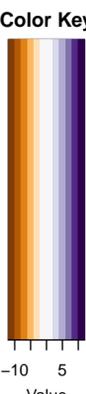


negative regulation of chromatin silencing at rDNA

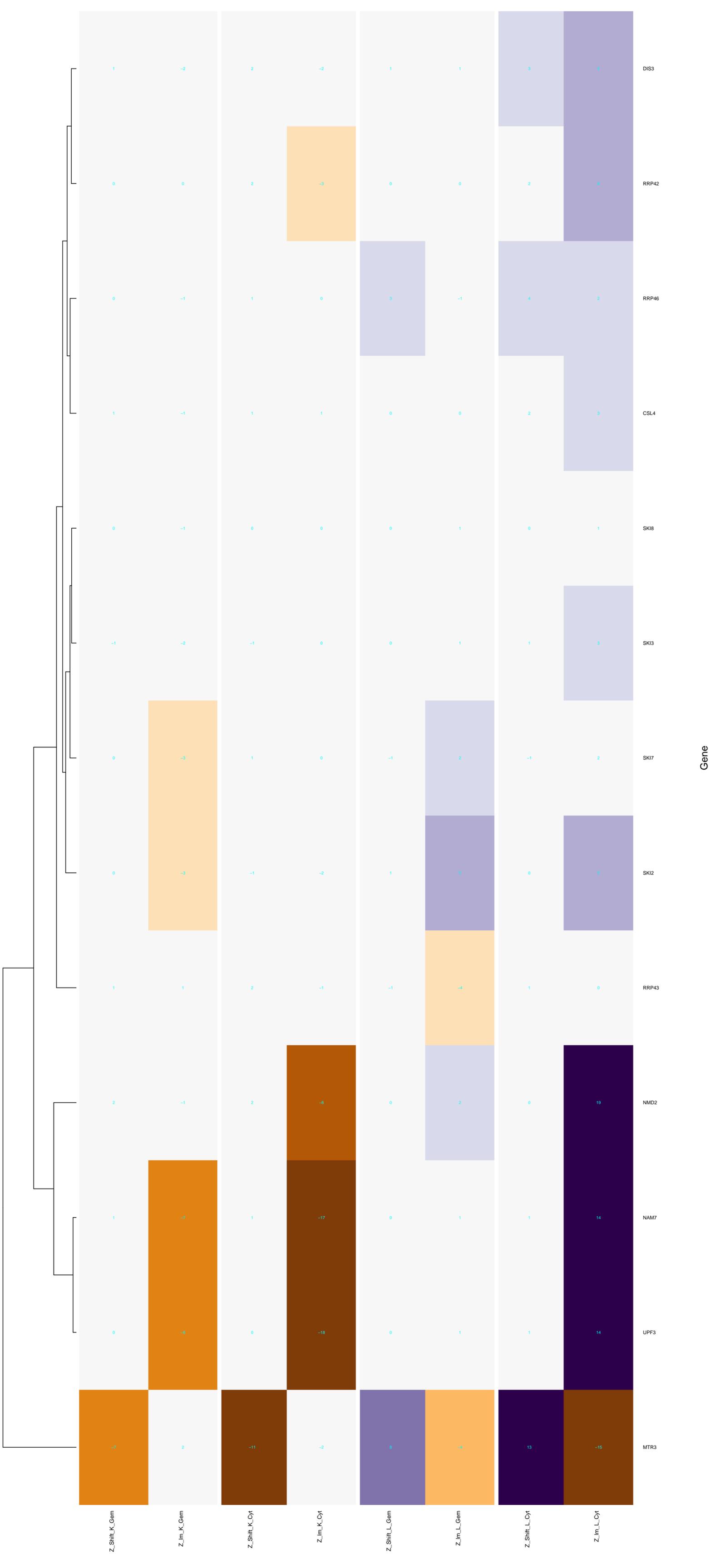


negative regulation of transcription by glucose

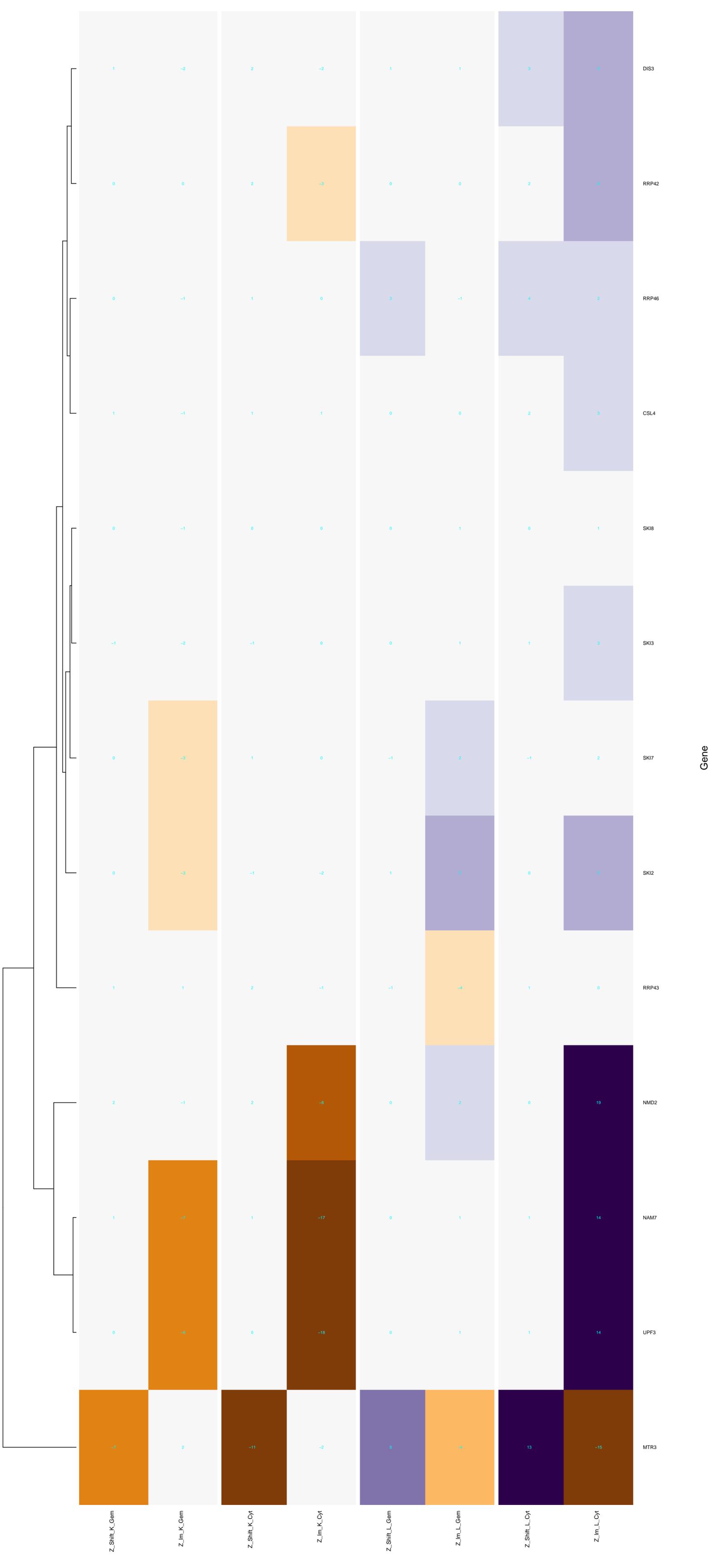
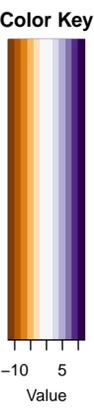


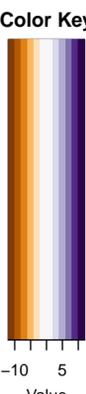


-transcribed mRNA catabolic process, 3'-5' exonucleolytic nonsense-mediated decay

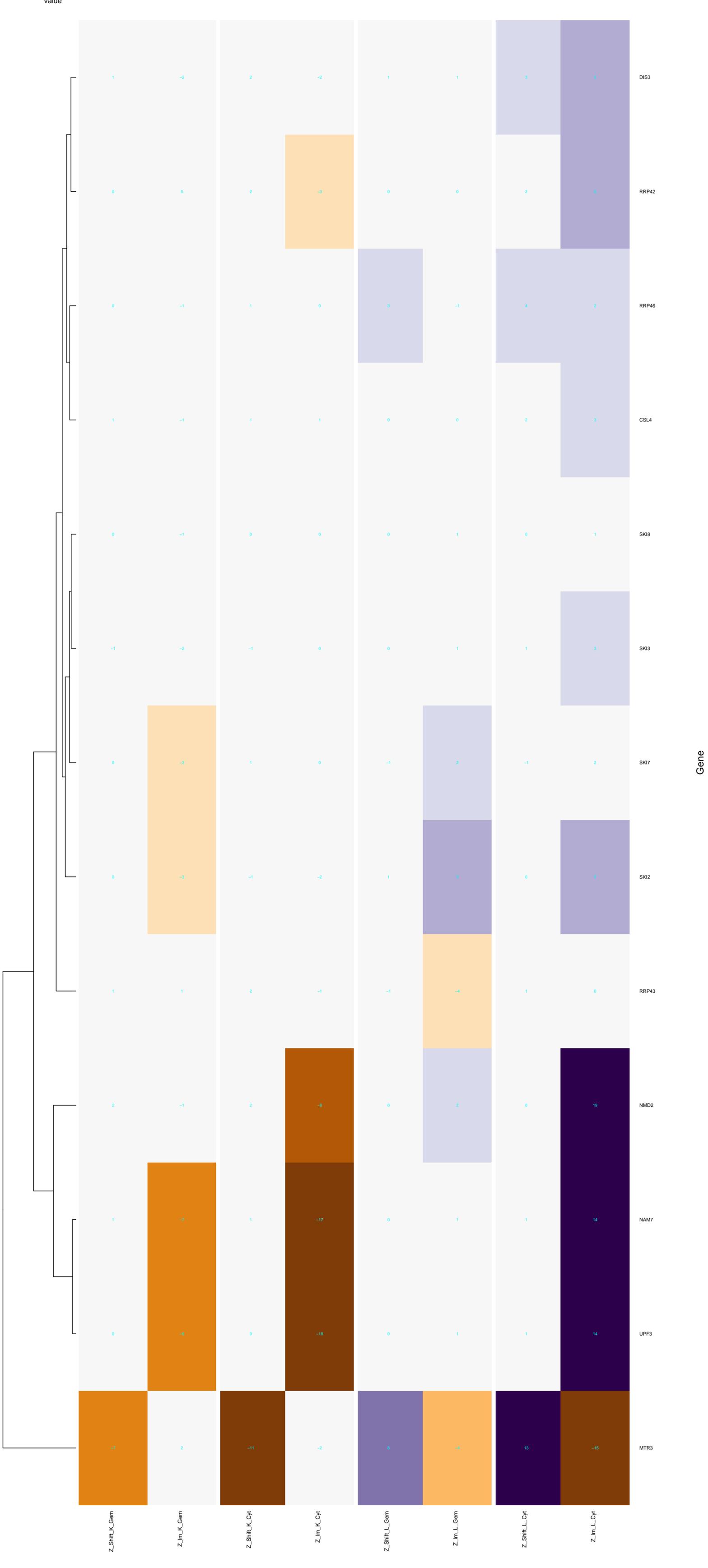


nuclear-transcribed mRNA catabolic process, exonucleolytic, 3'-5'

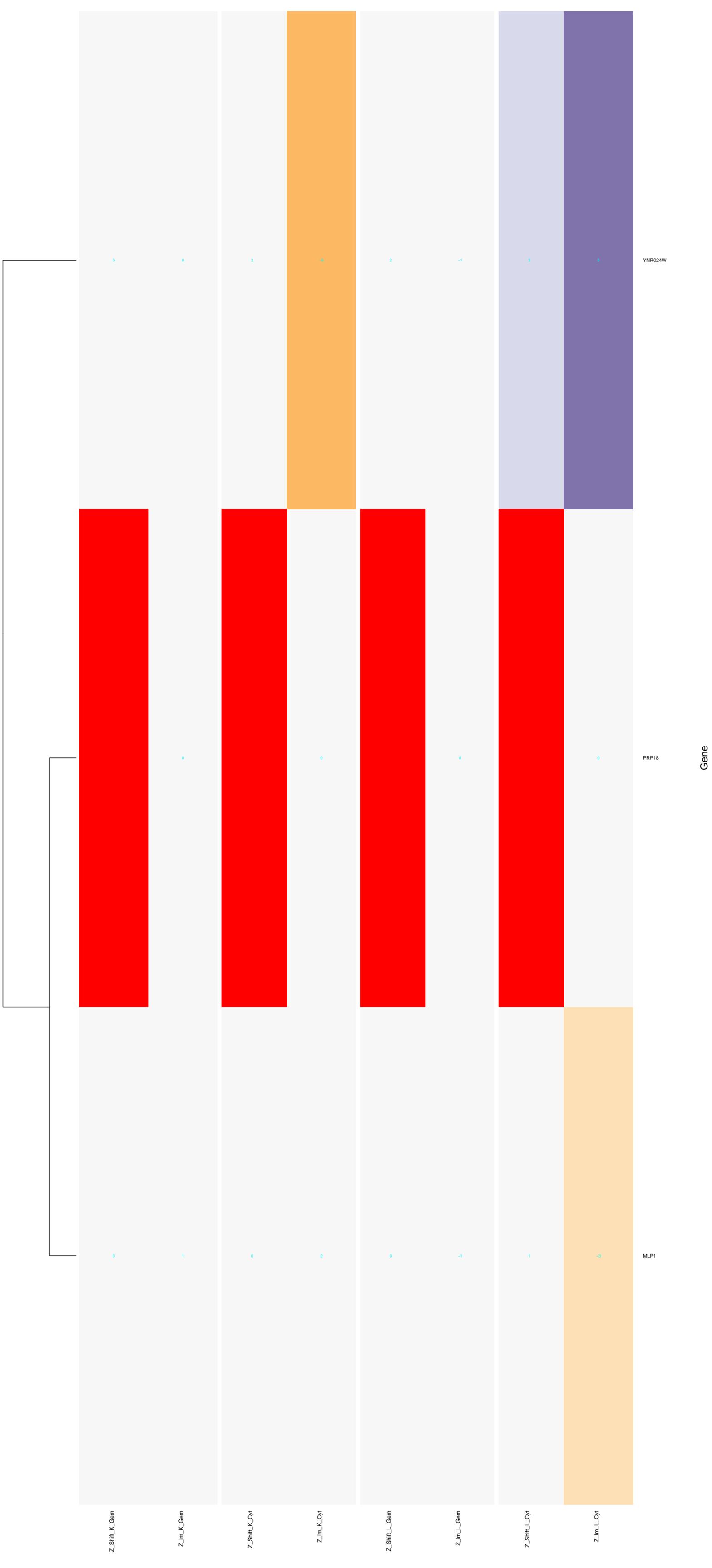
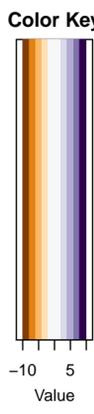




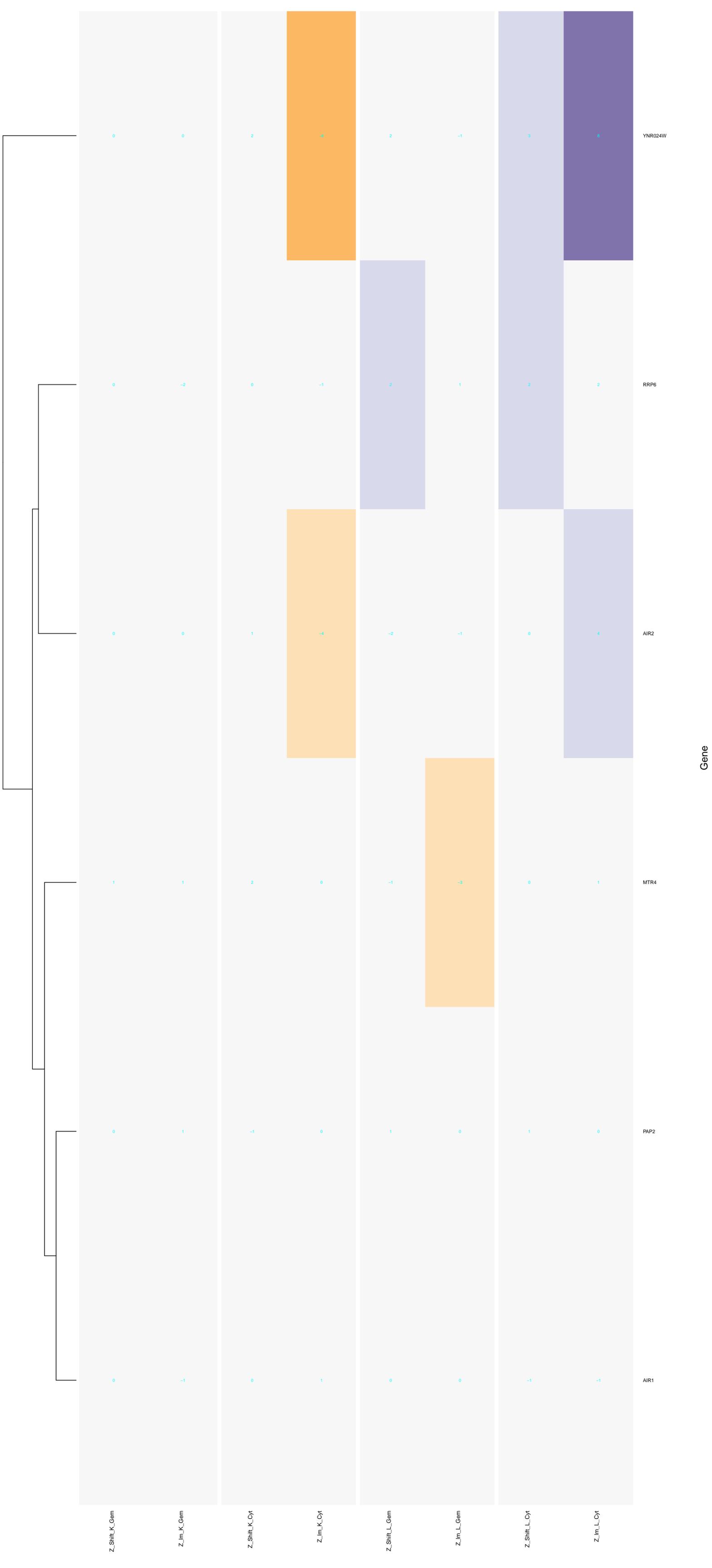
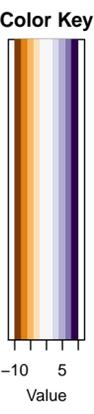
c nuclear-transcribed mRNA catabolic process involved in deadenylation-dependent decap



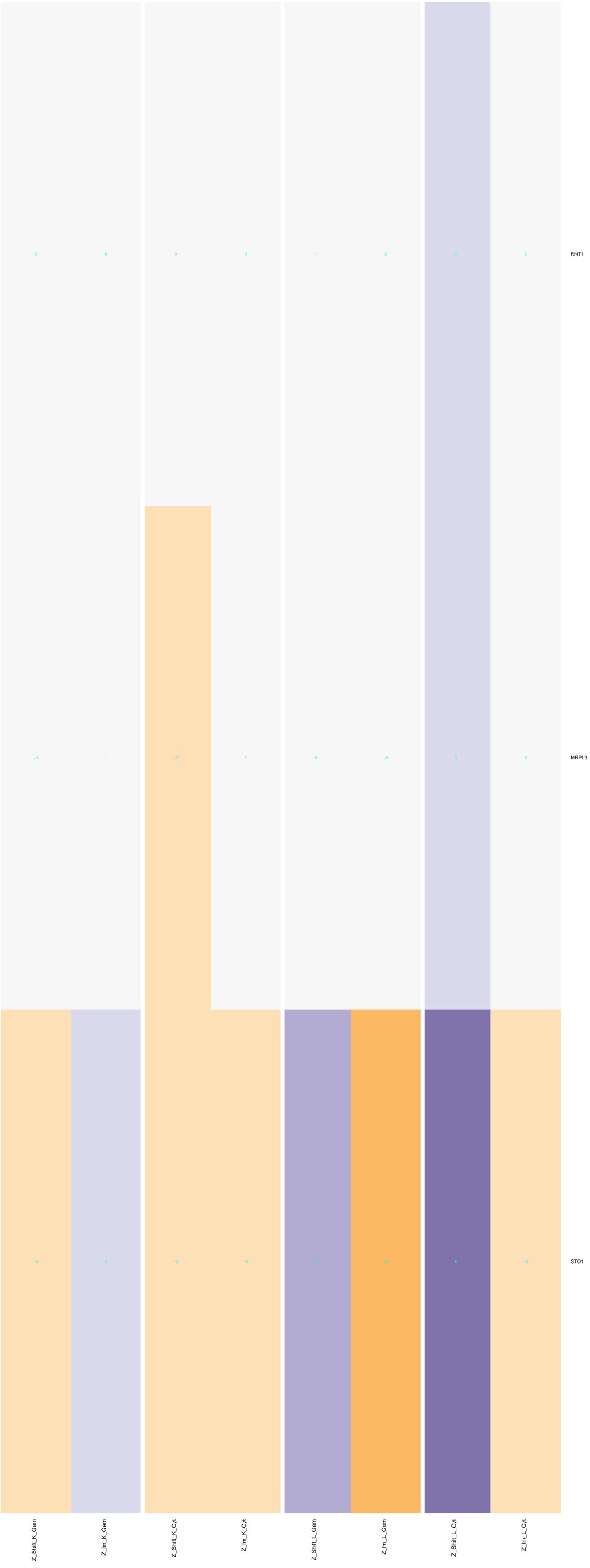
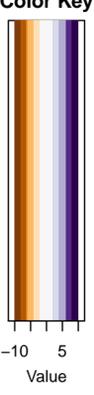
nuclear mRNA surveillance of spliceosomal pre-mRNA splicing



nuclear mRNA surveillance of mRNA 3'-end processing

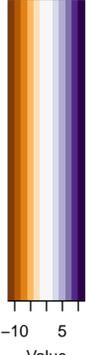


gene silencing by miRNA



Gene

Color Key



egative regulation of transcription from RNA polymerase II promoter by glucose

