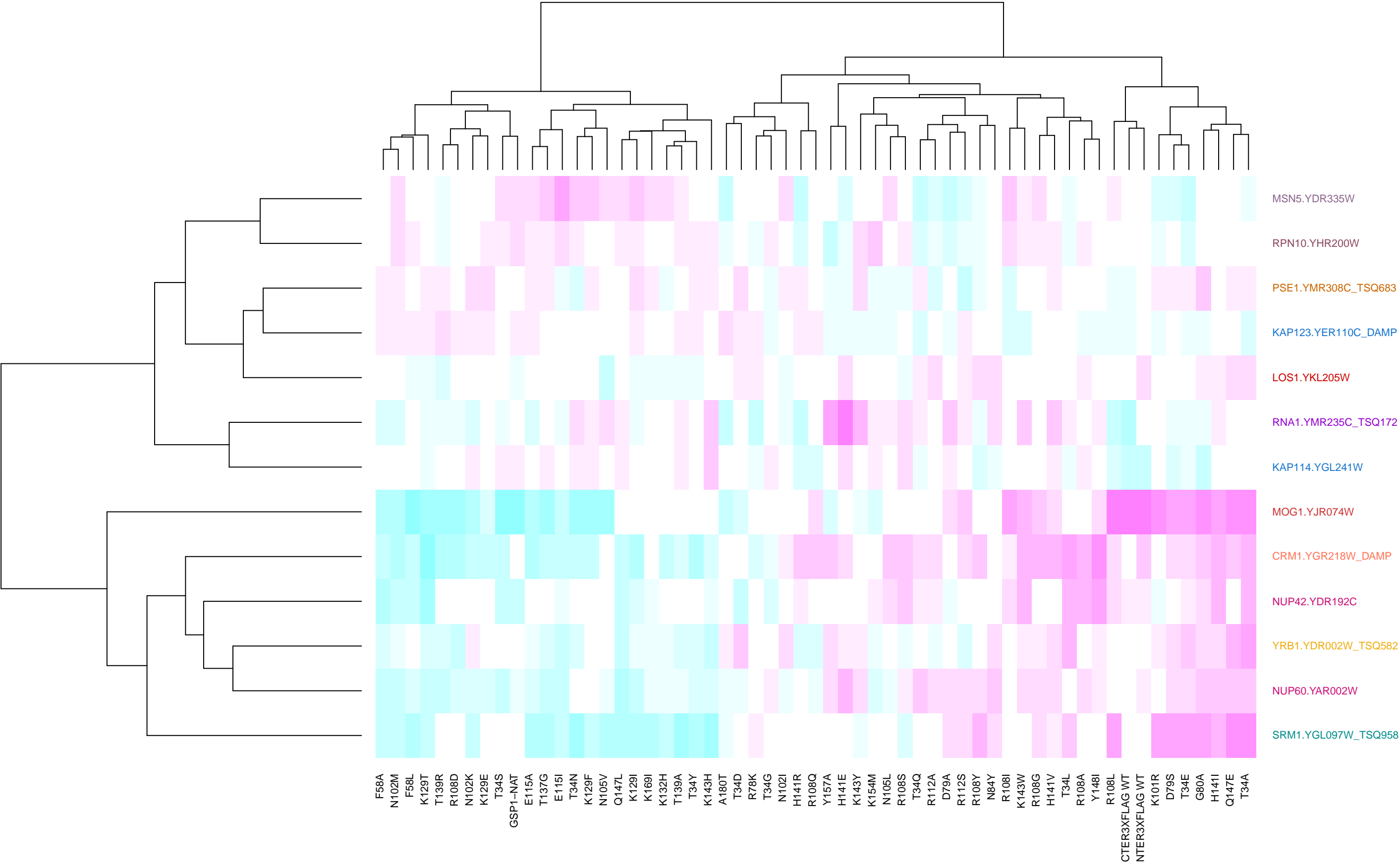
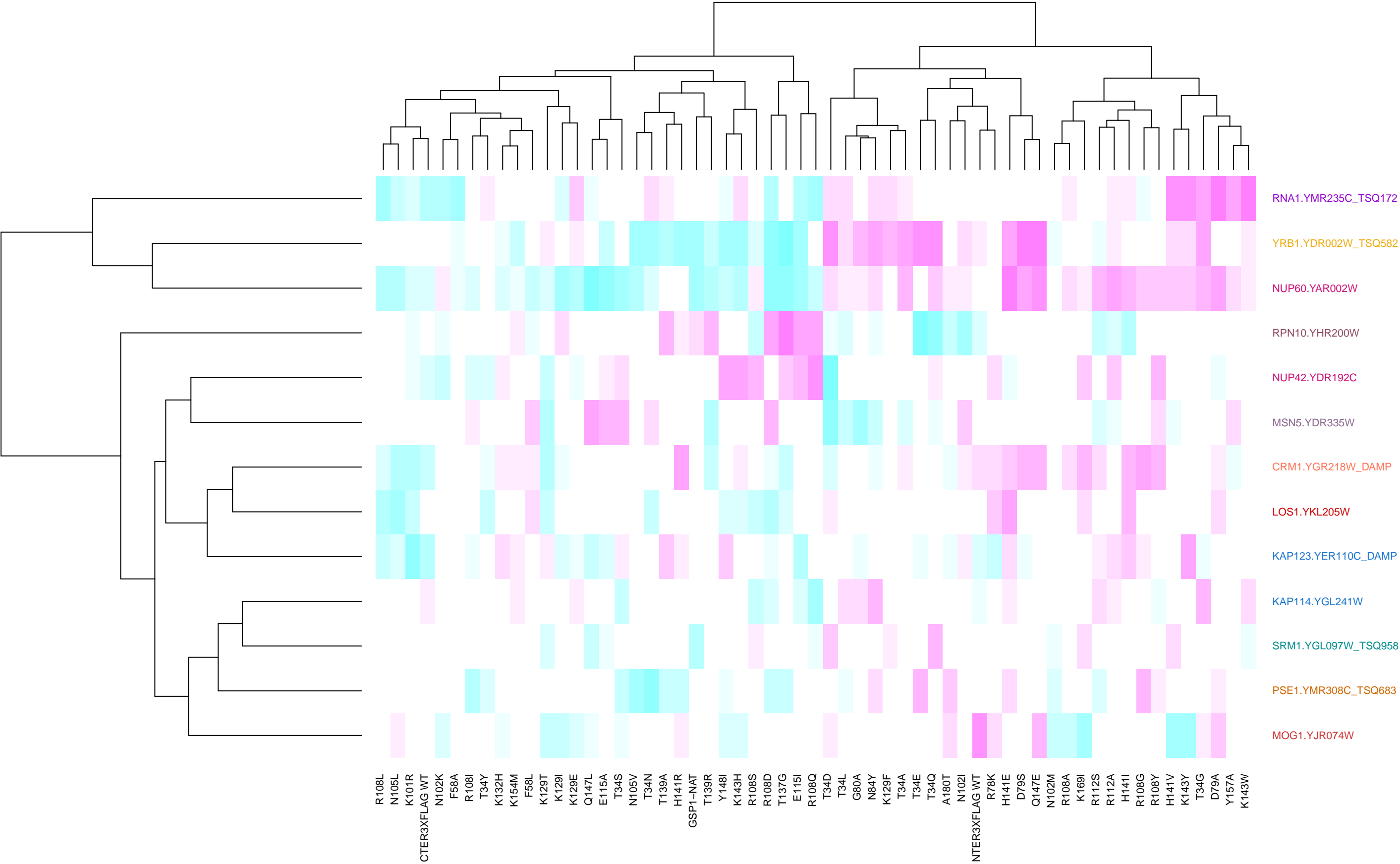


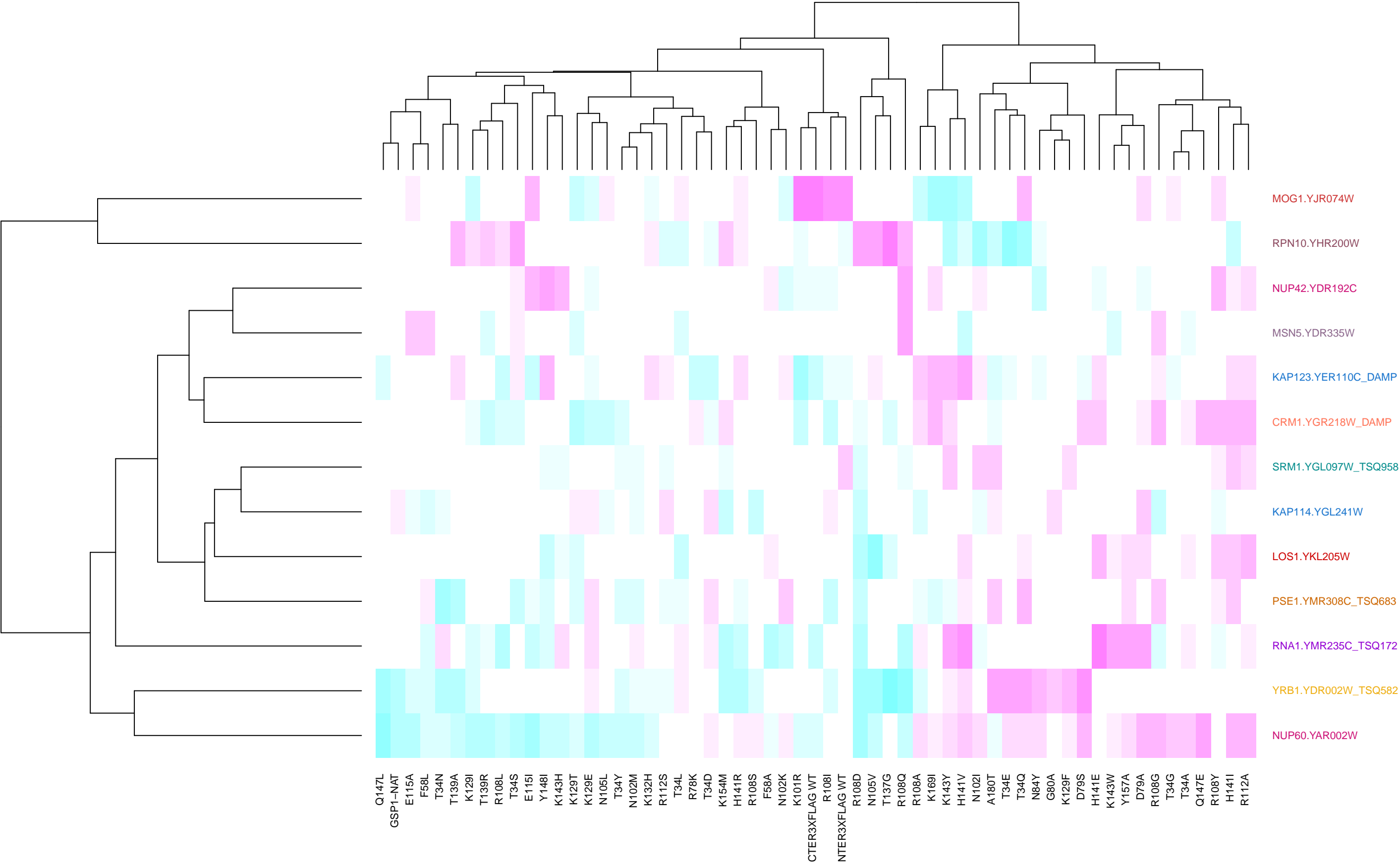
chromatin



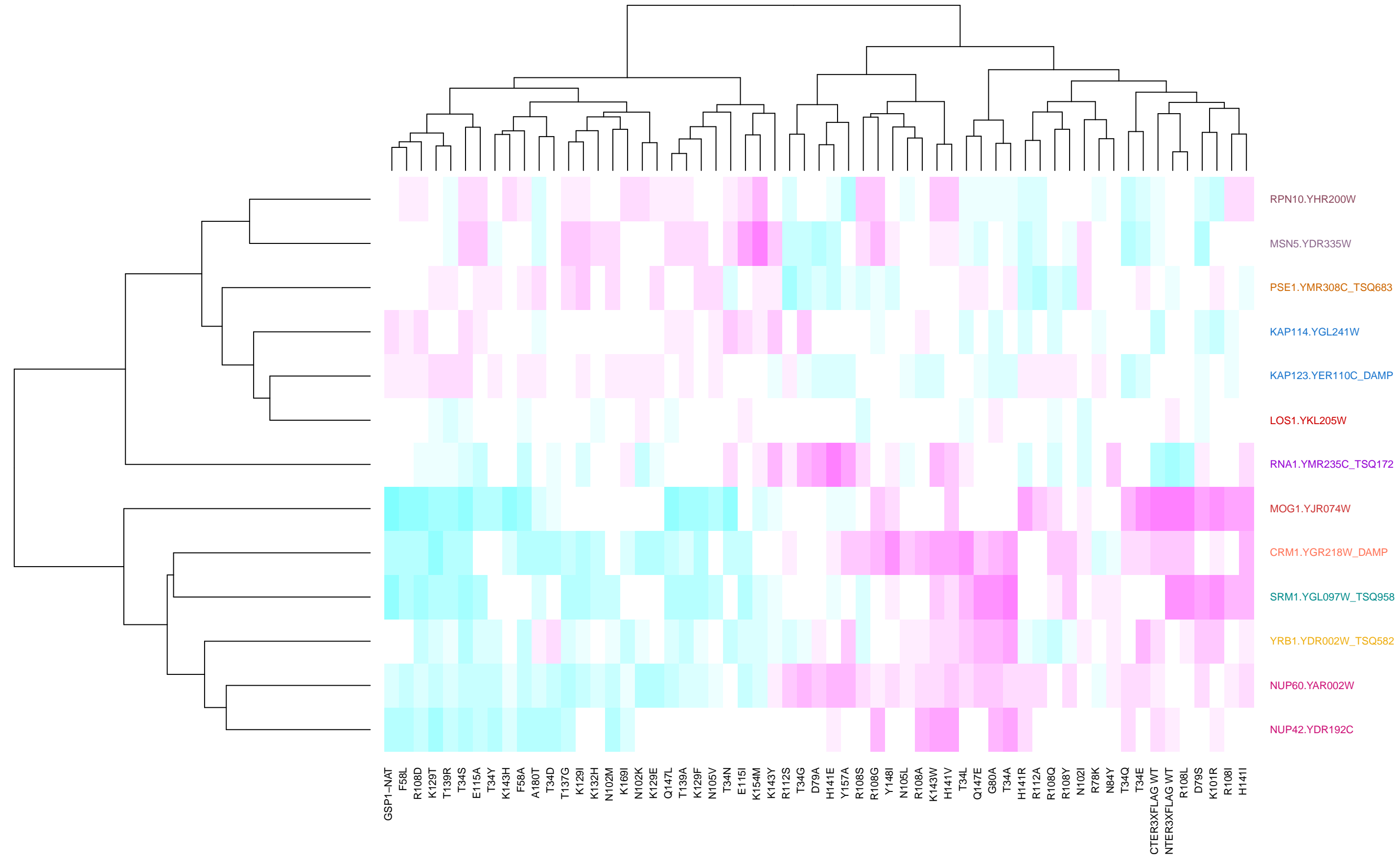
chromatin binding



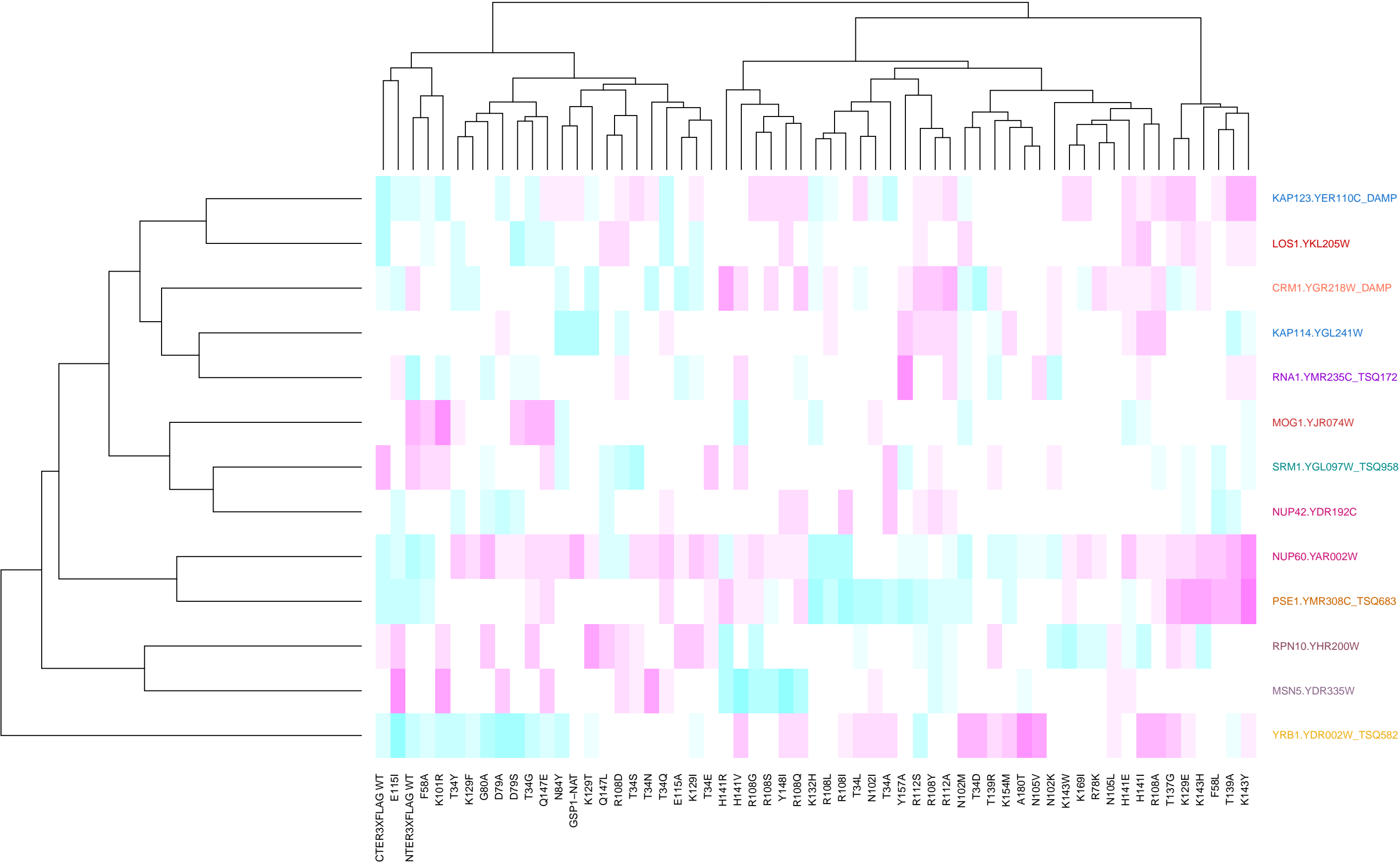
chromatin binding_GO_1



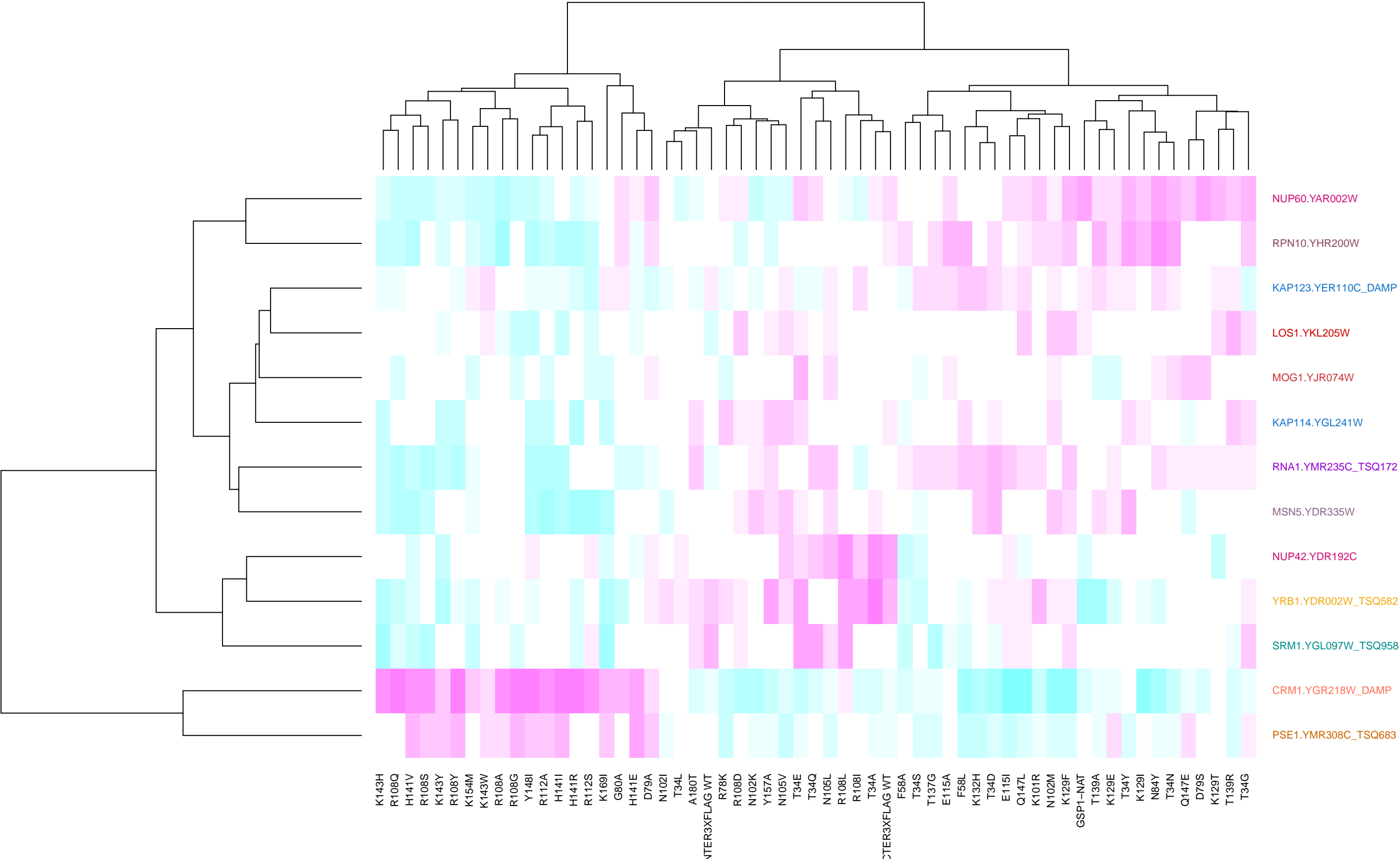
chromatin organization



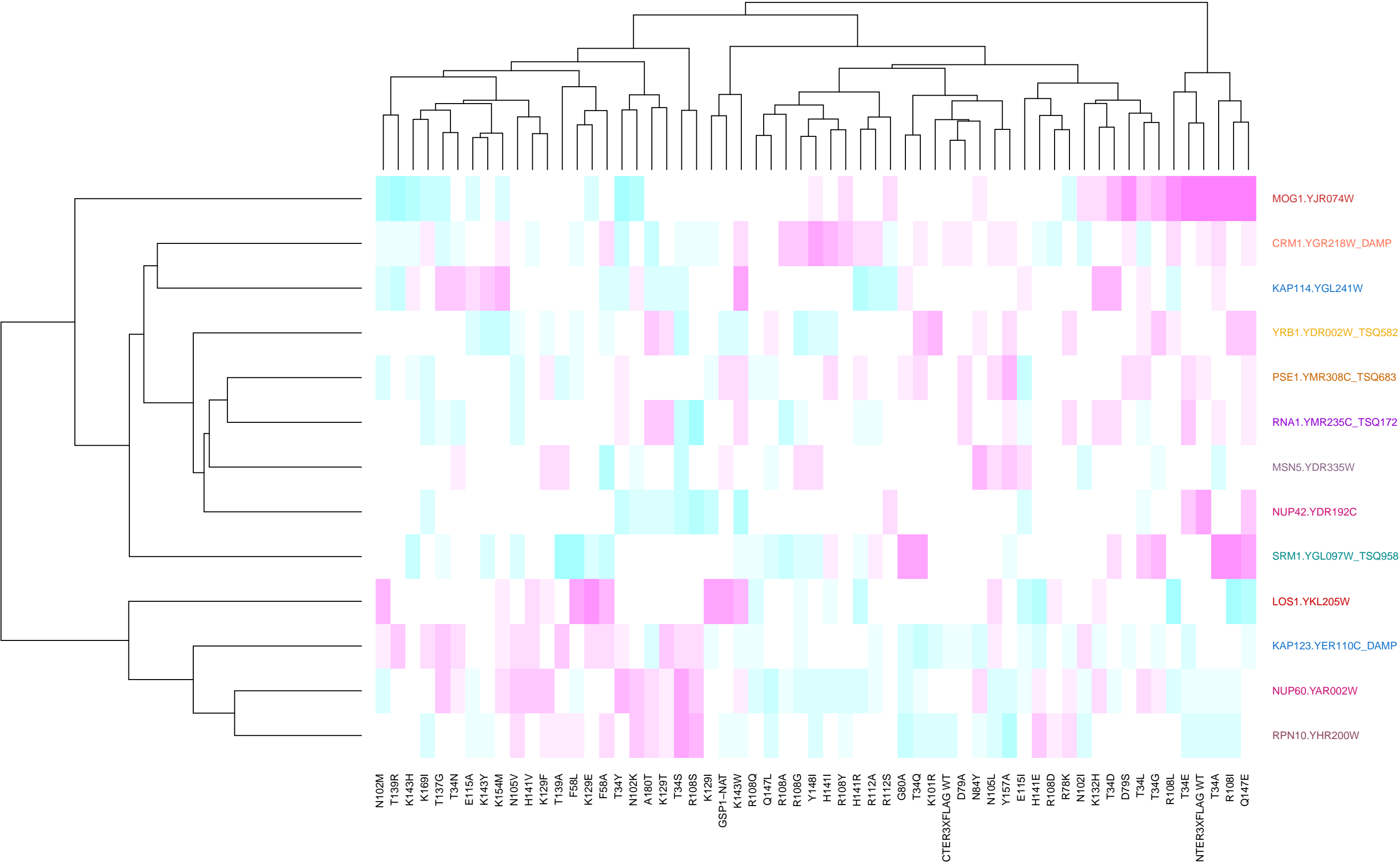
chromatin_GO_2



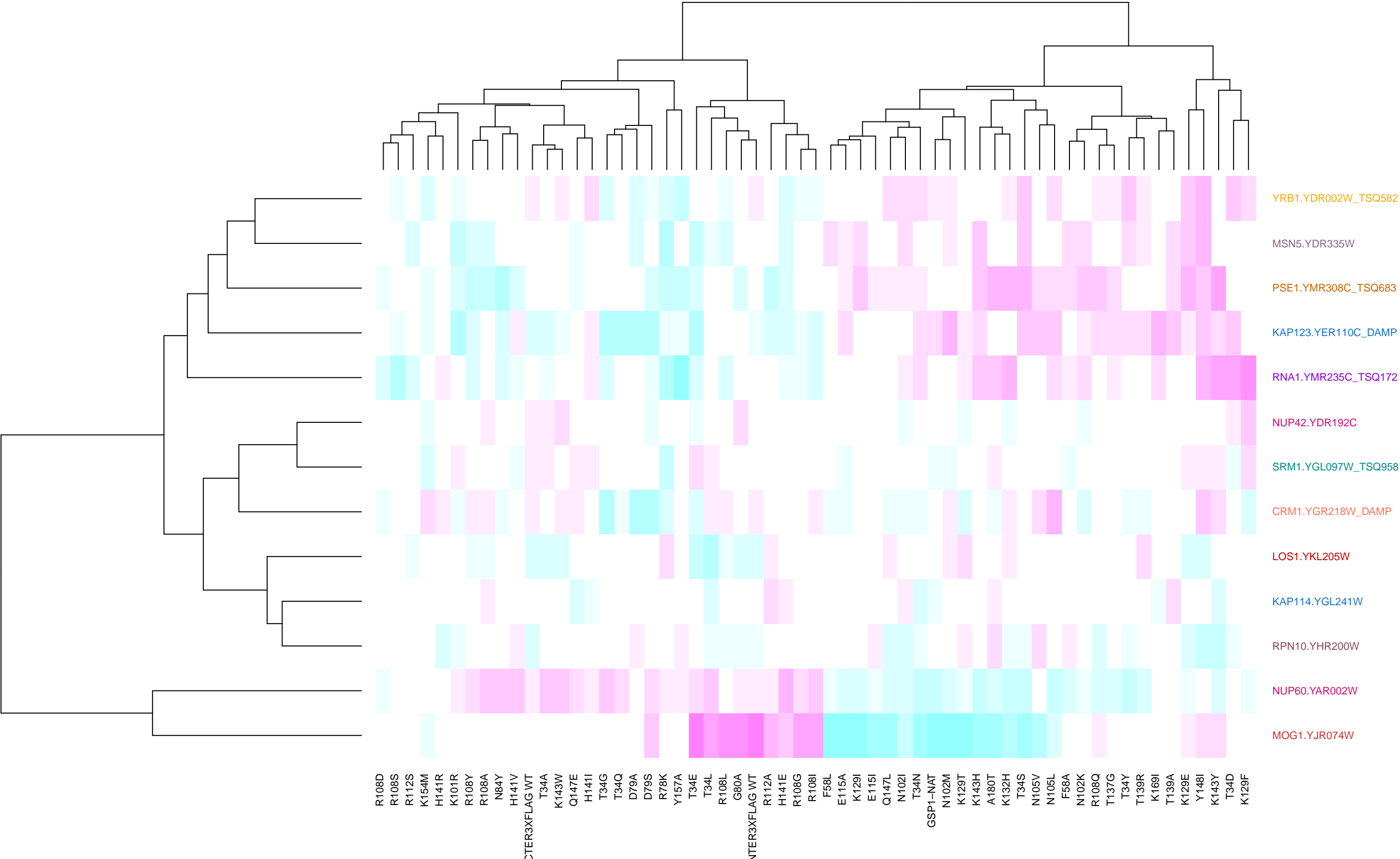
cytoskeleton organization



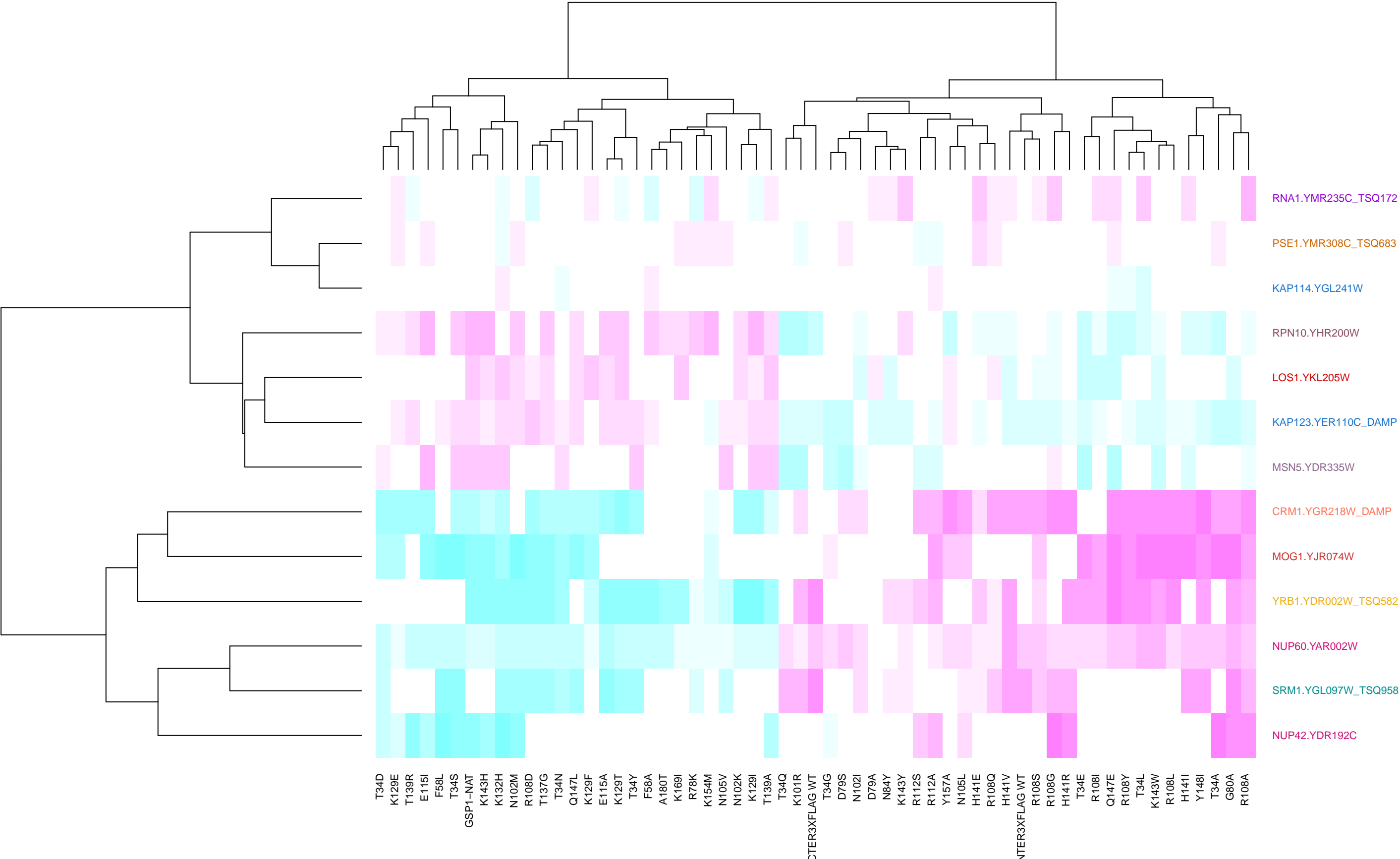
DNA binding_GO_2



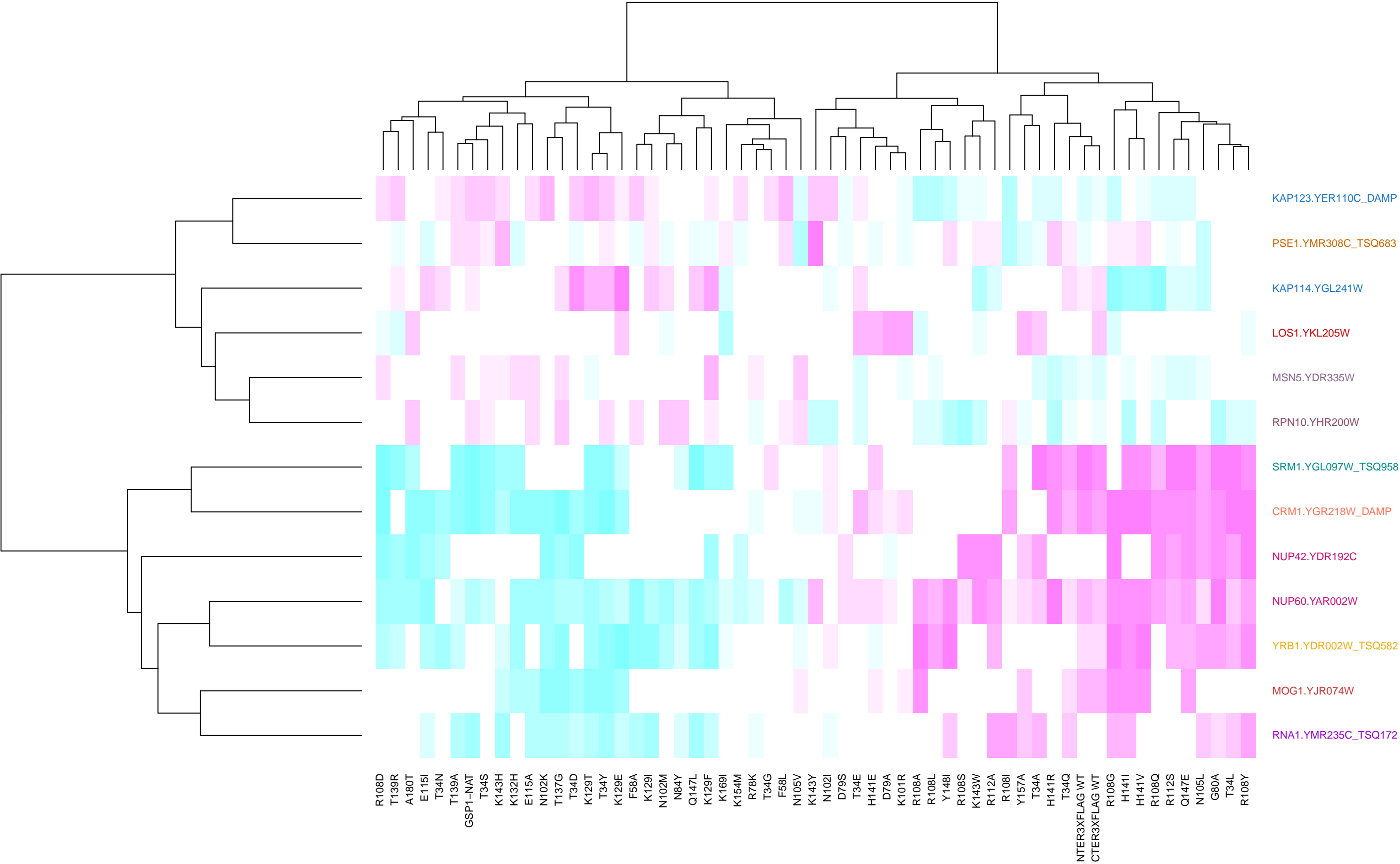
DNA recombination



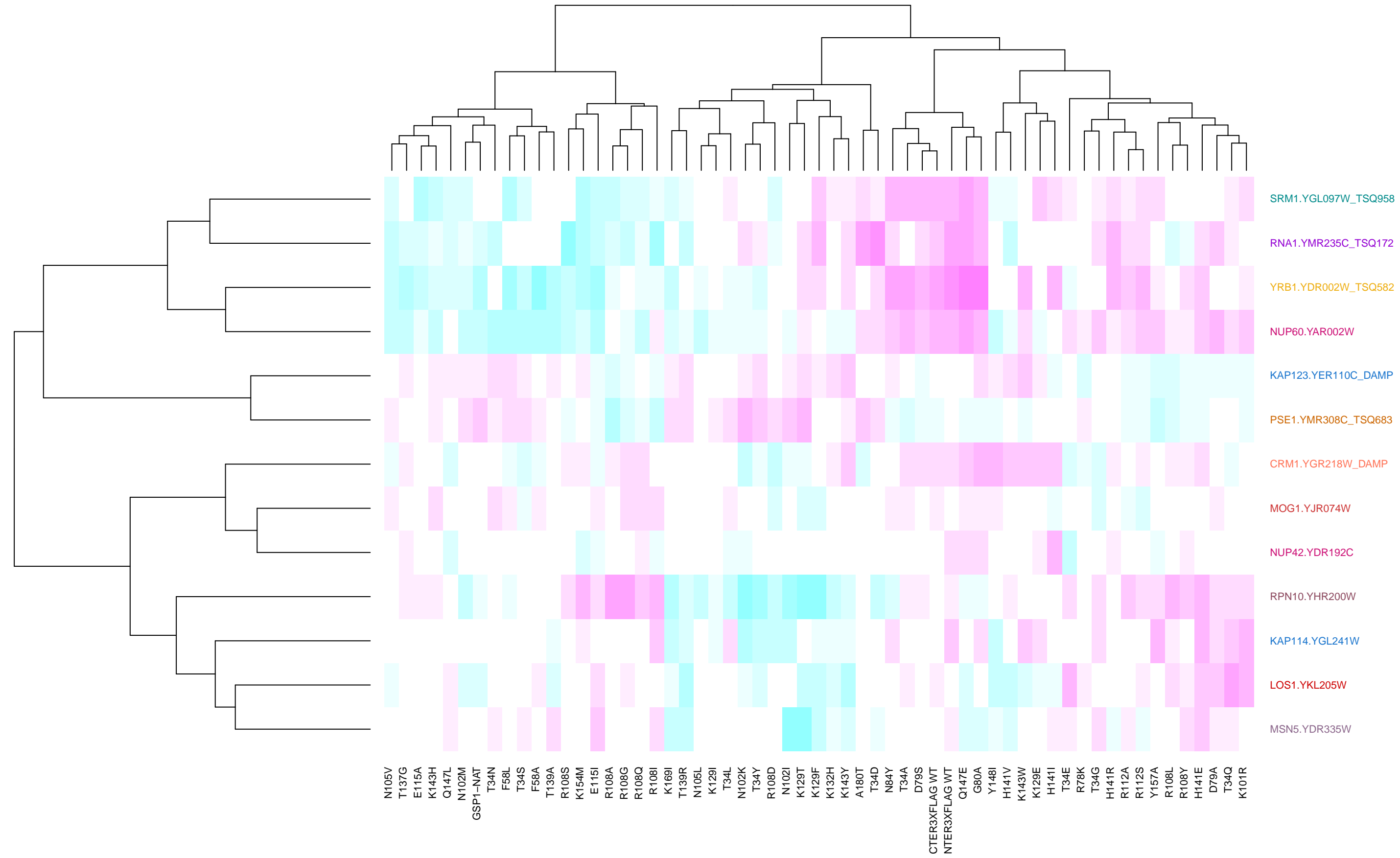
DNA repair



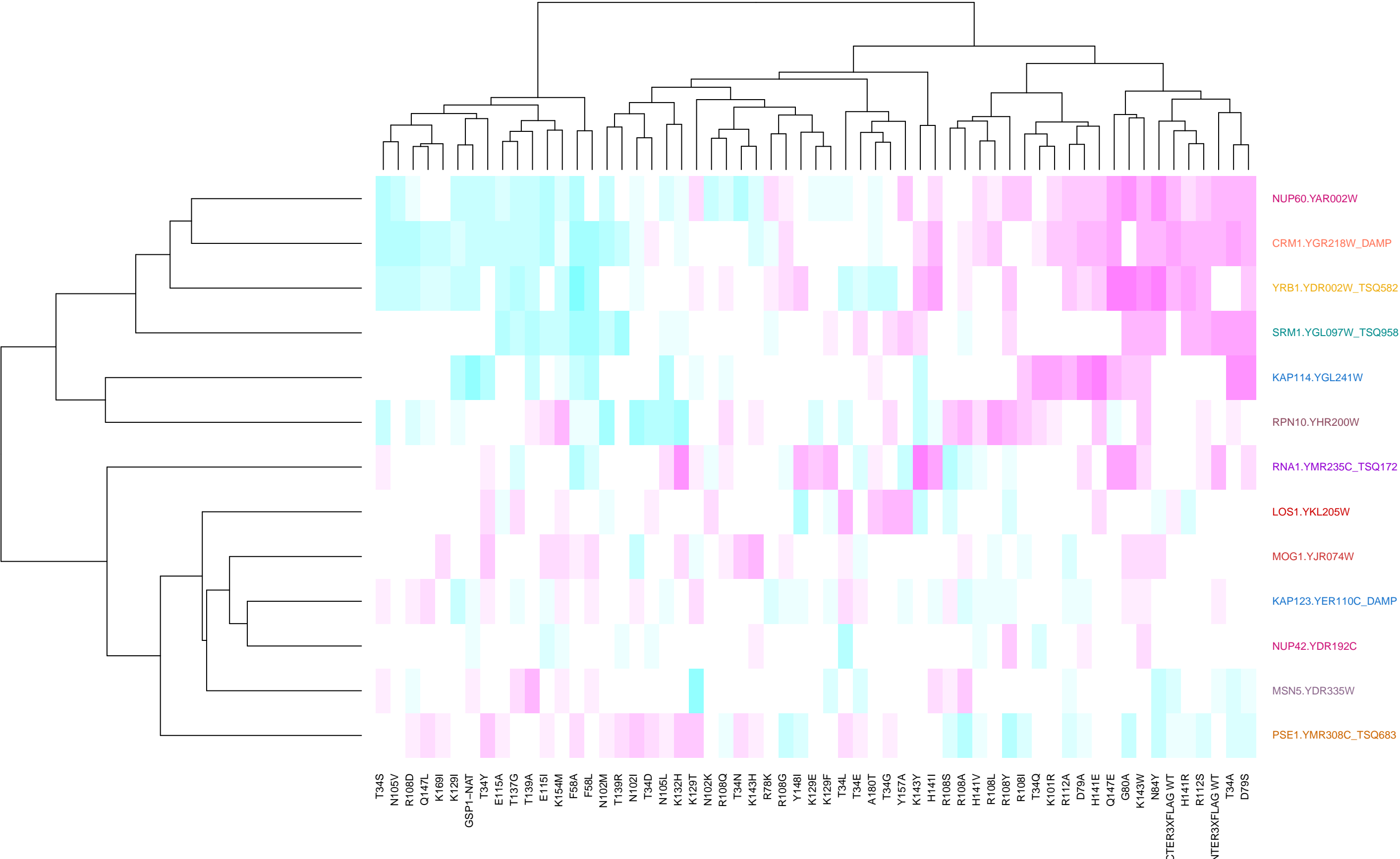
endomembrane system_GO_4



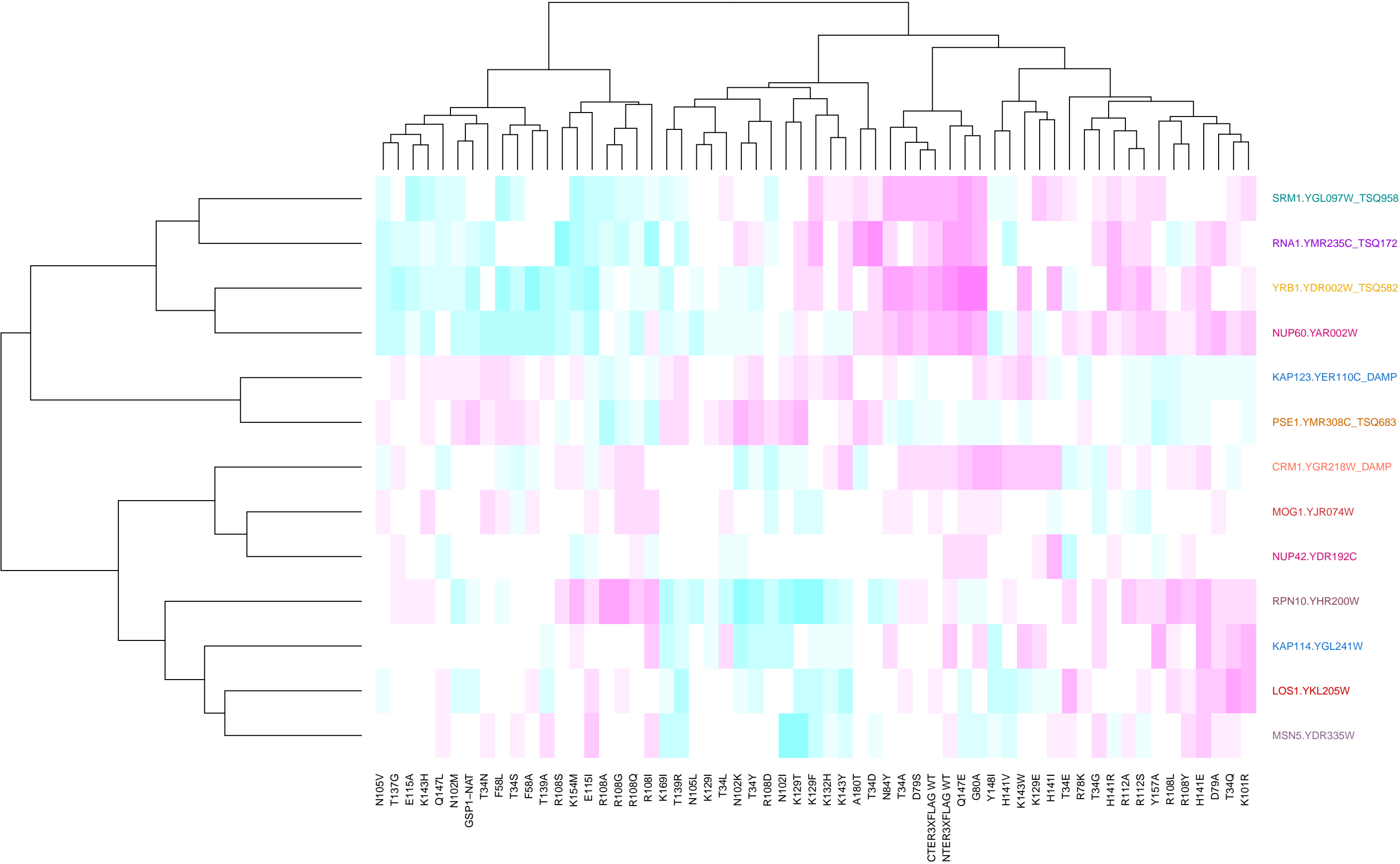
endoplasmic reticulum



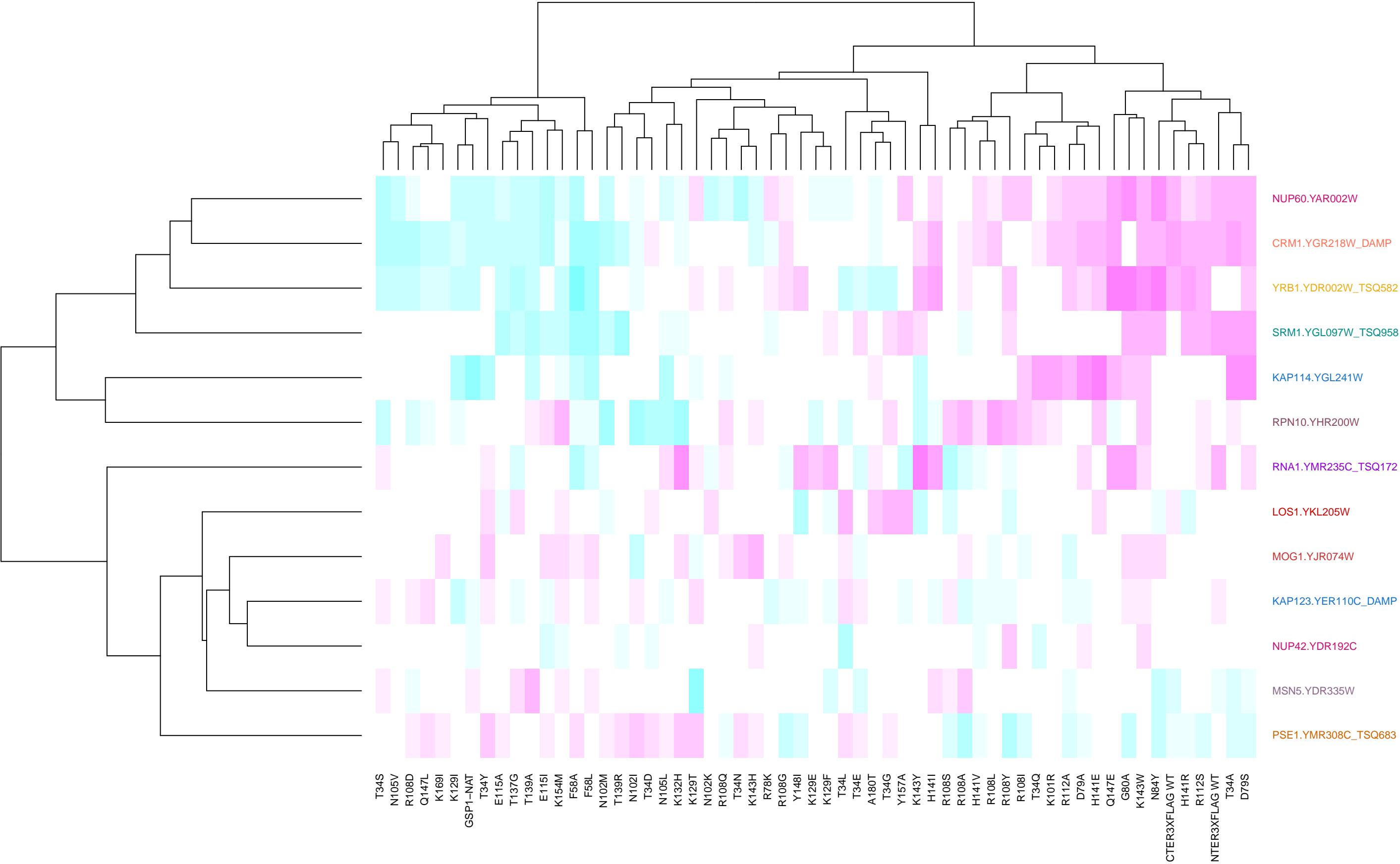
endoplasmic reticulum_GO_1



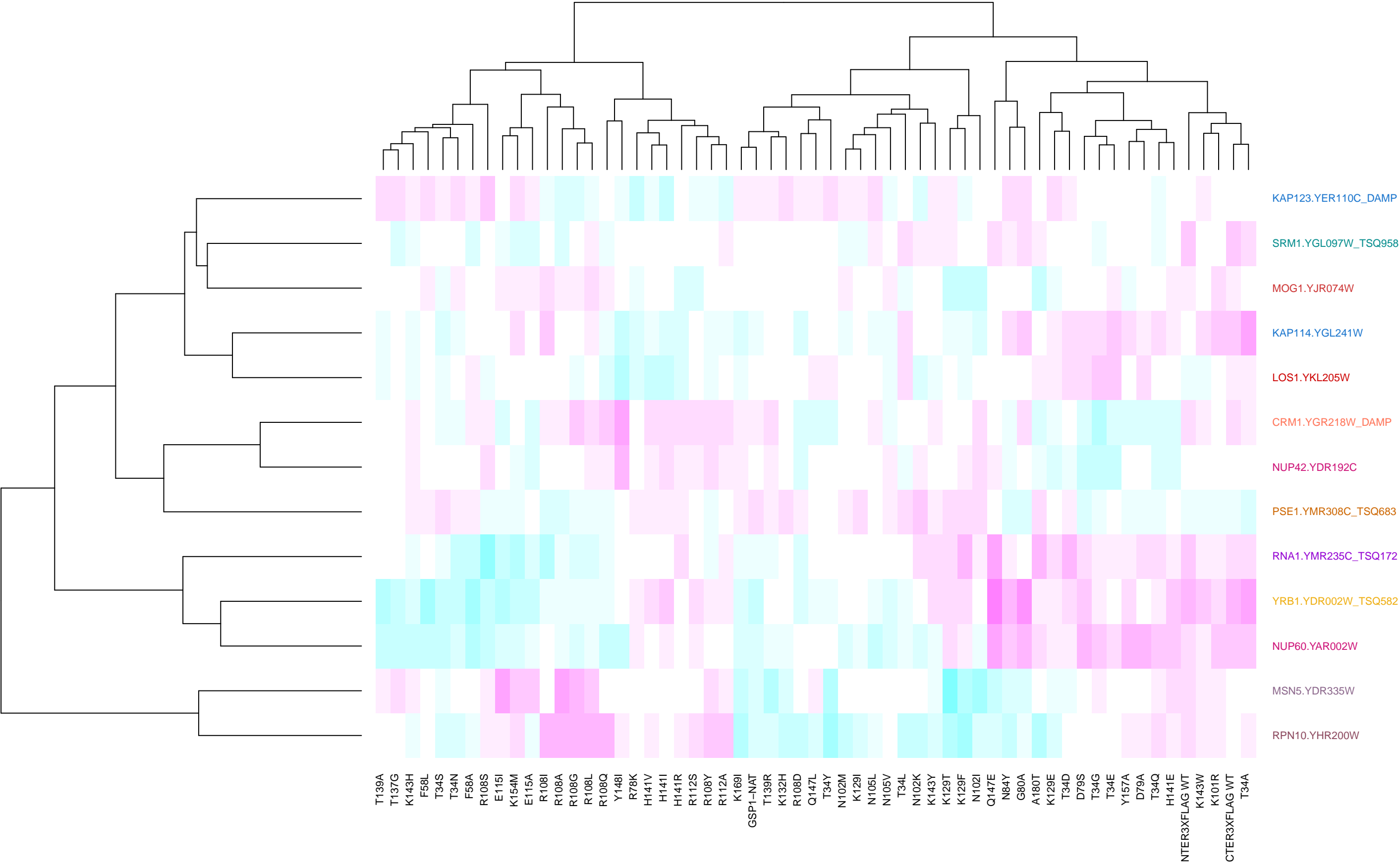
ER



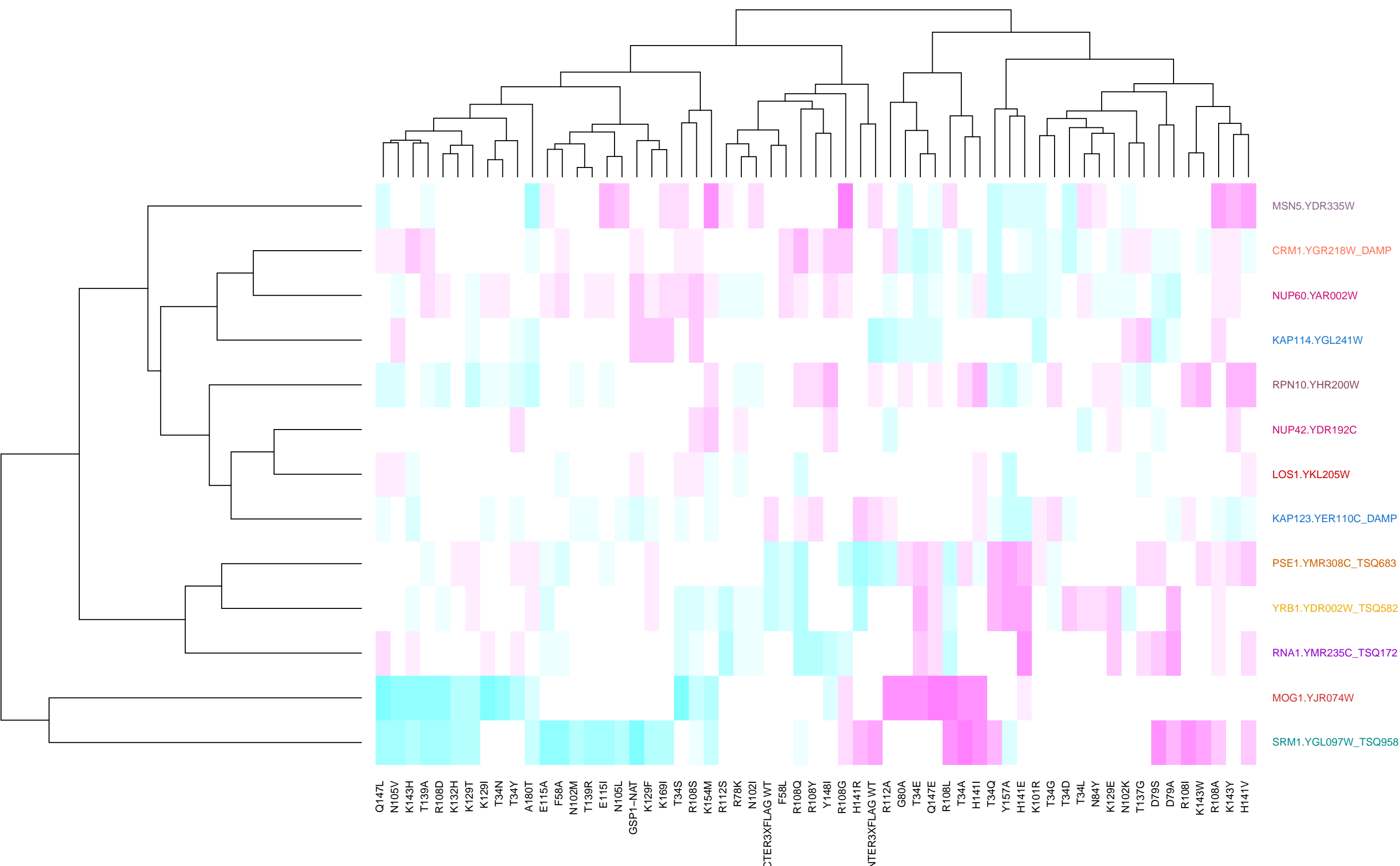
ER_GO_1



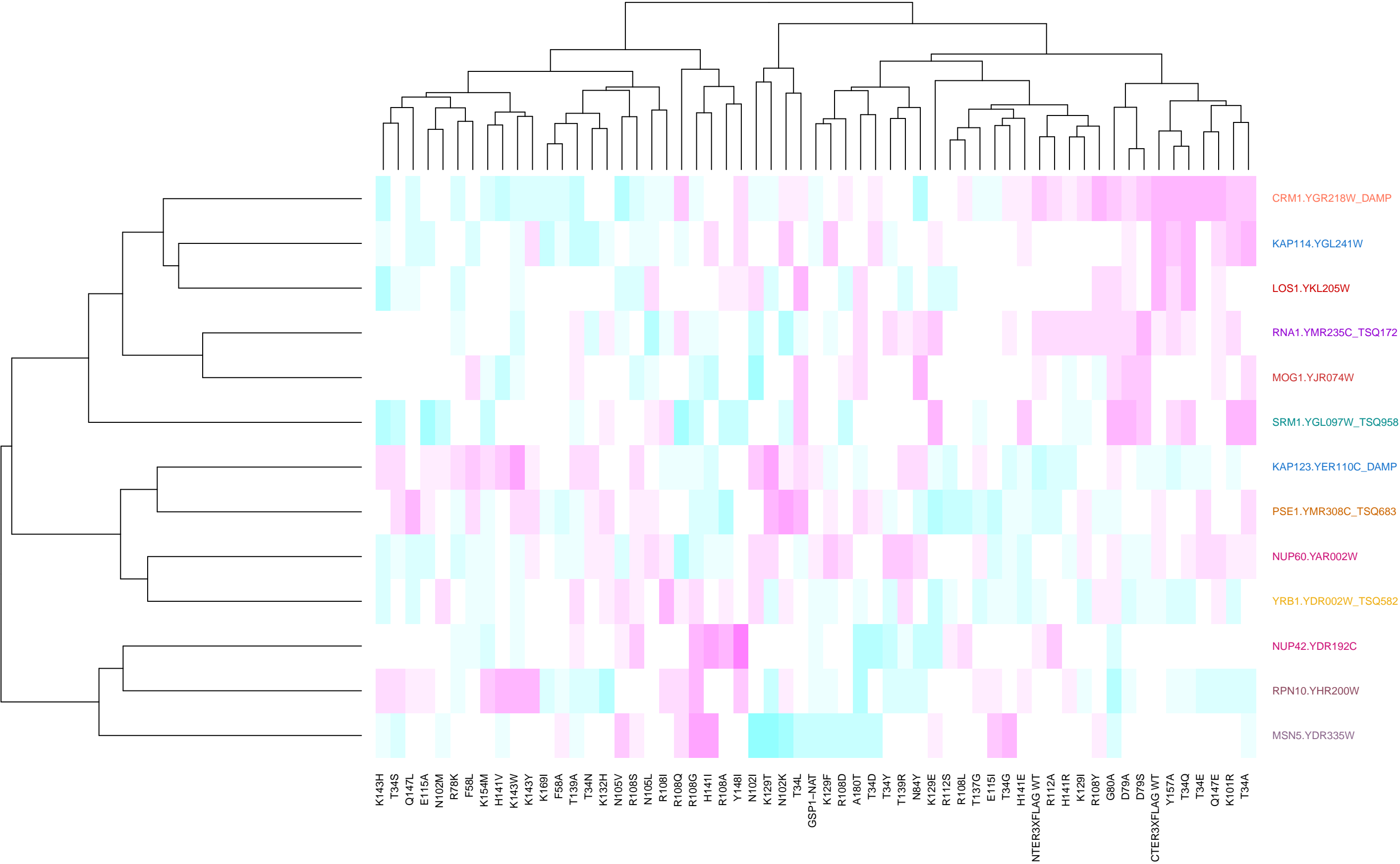
Golgi and ER



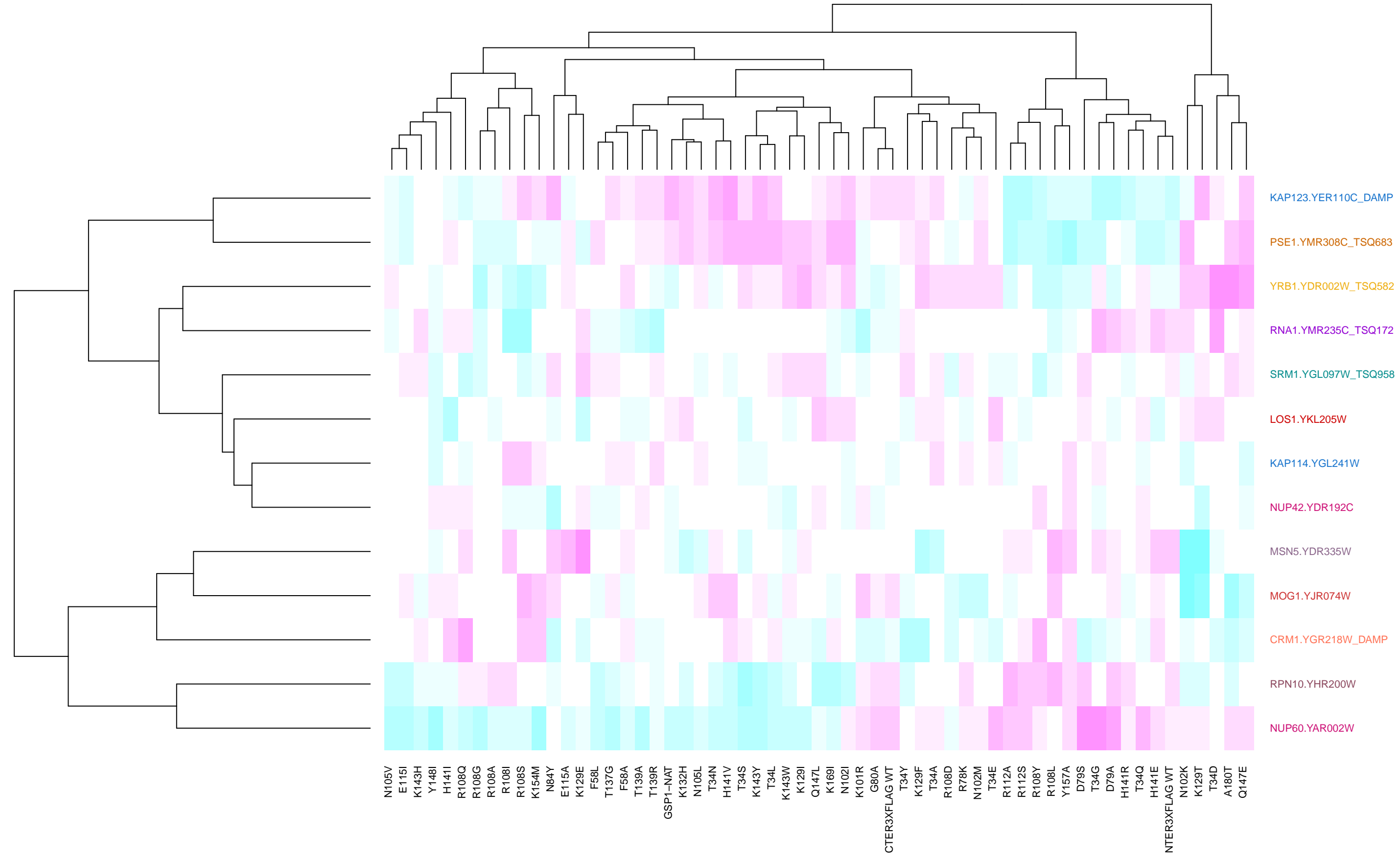
histone modification



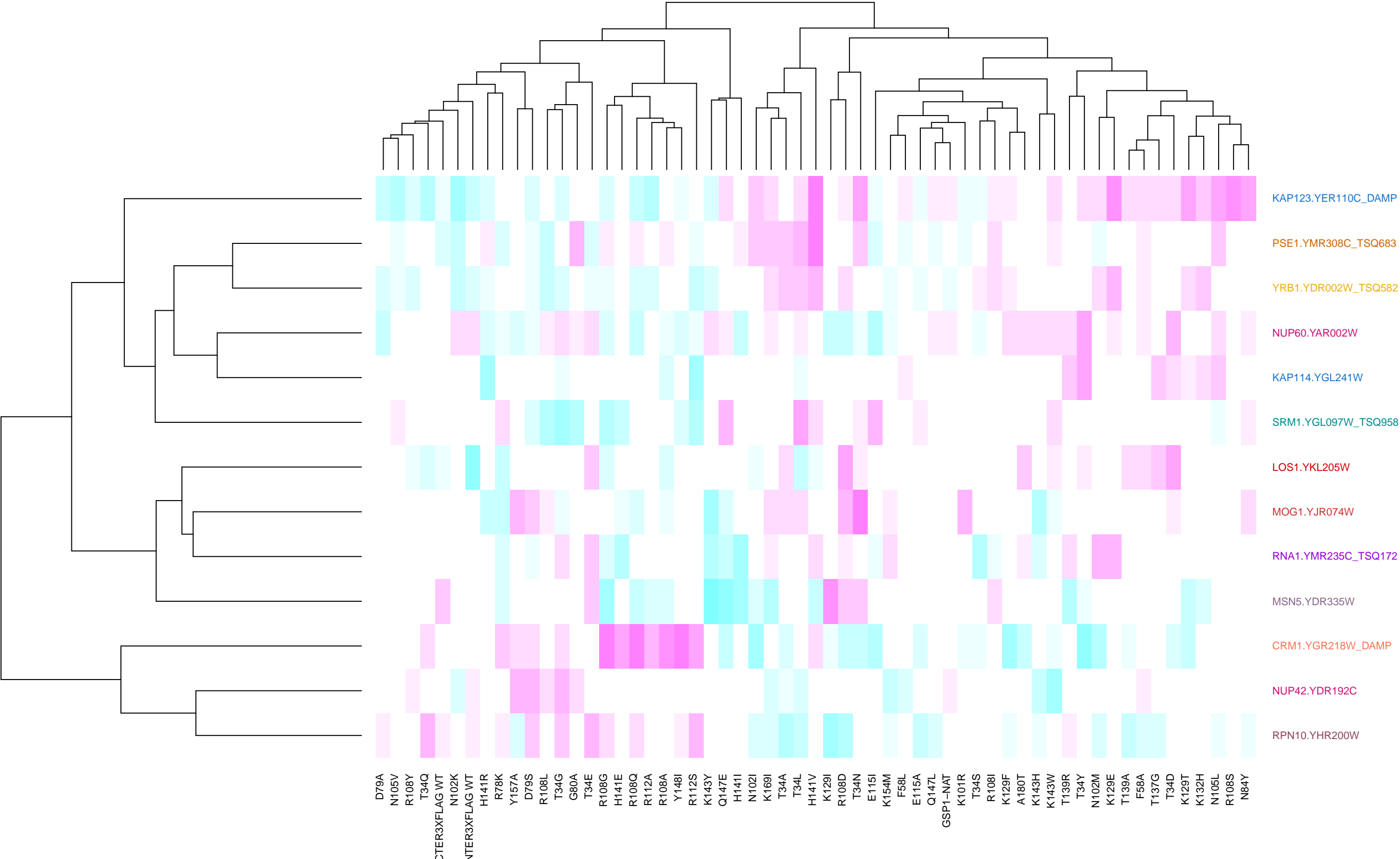
lipids



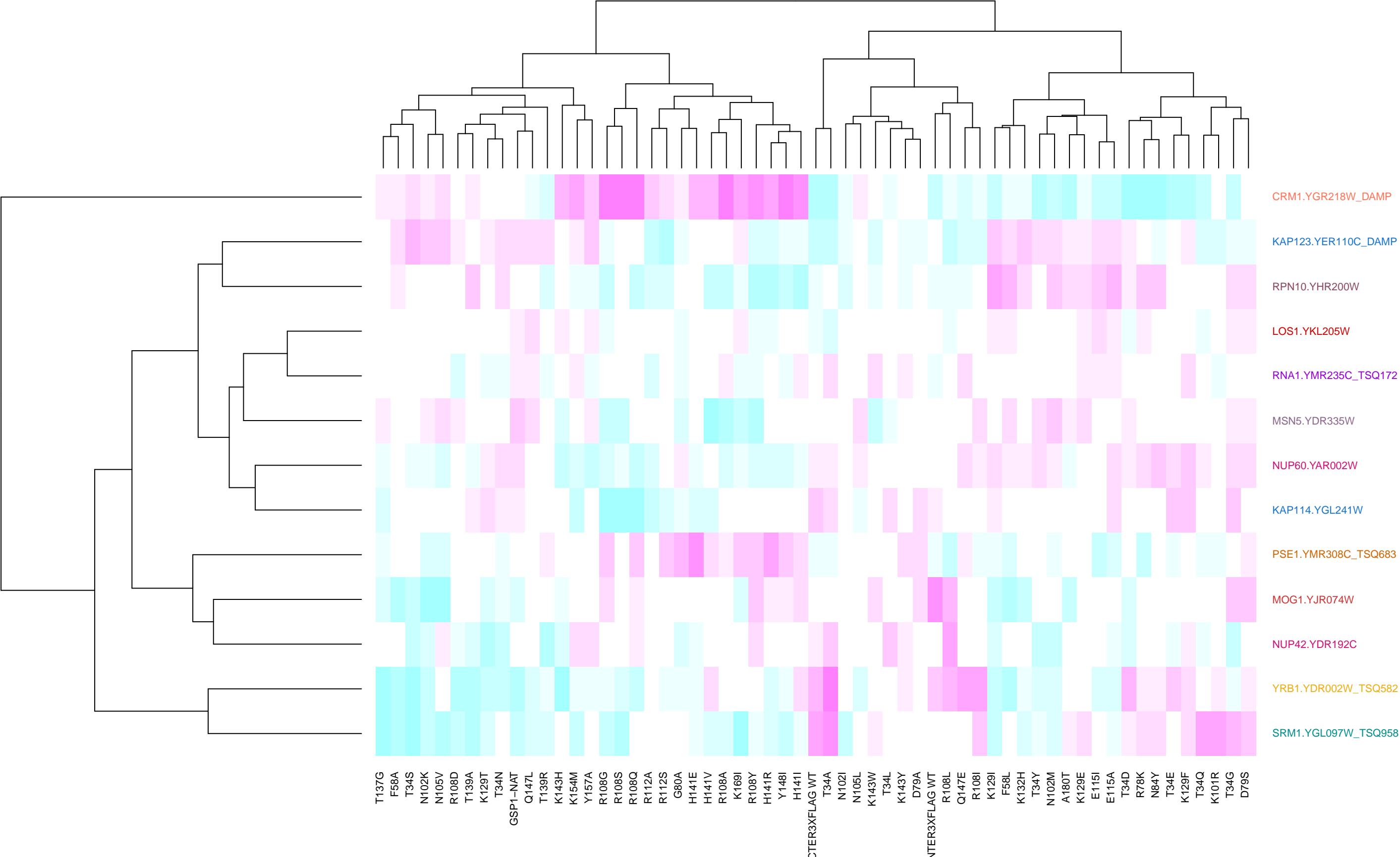
mitochondria



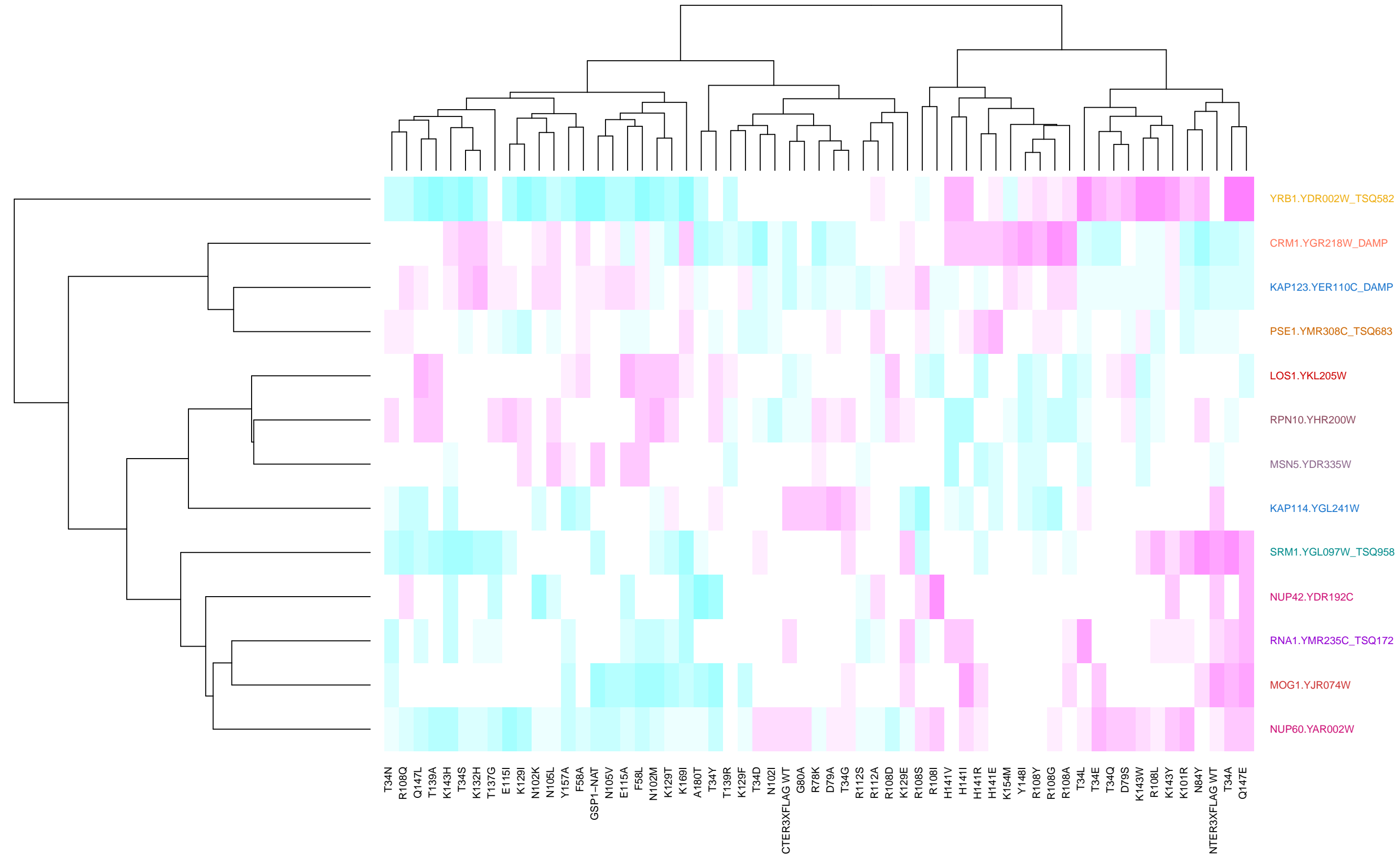
mitochondrion organization



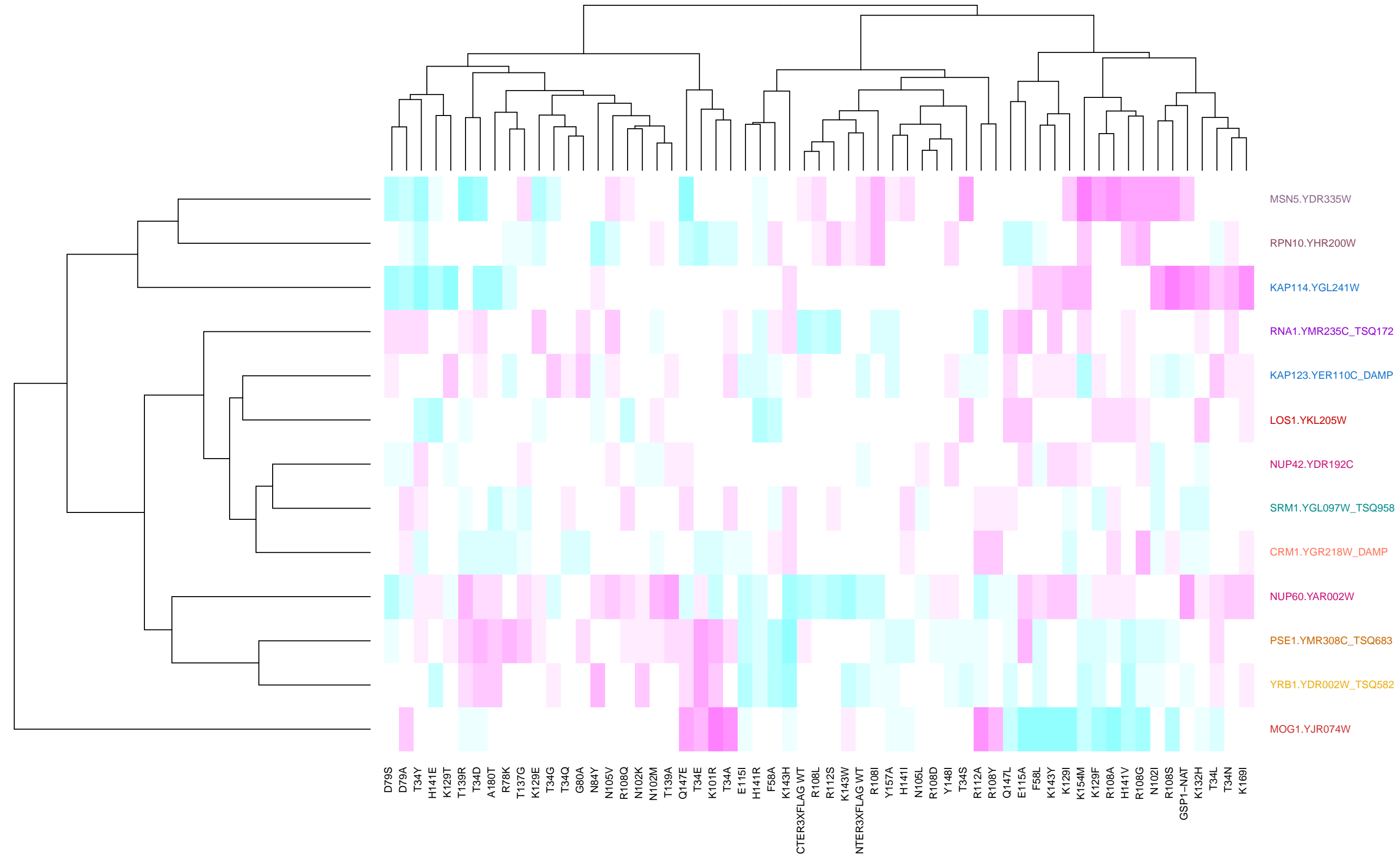
mitotic cell cycle



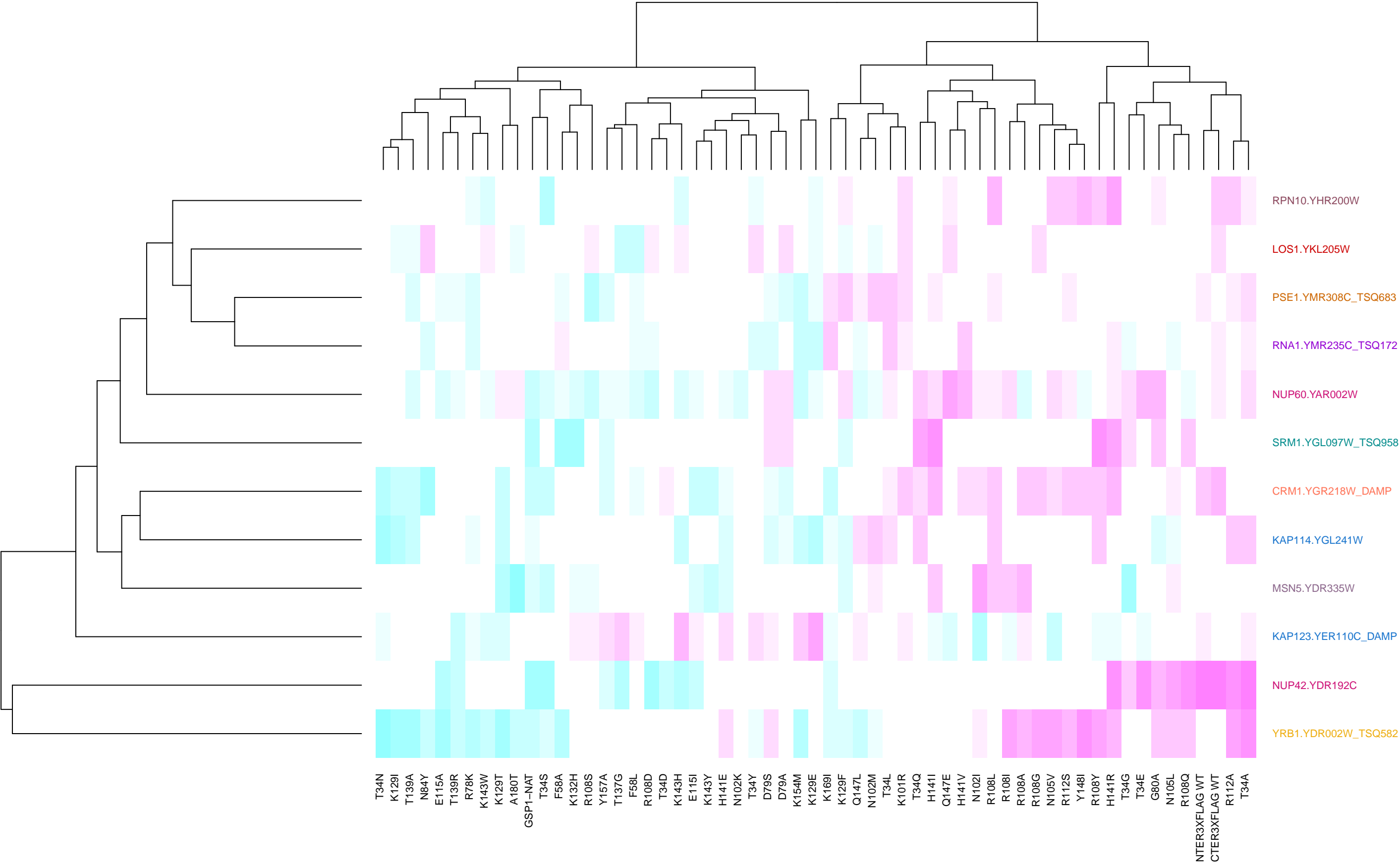
mitotic cell cycle_GO_1



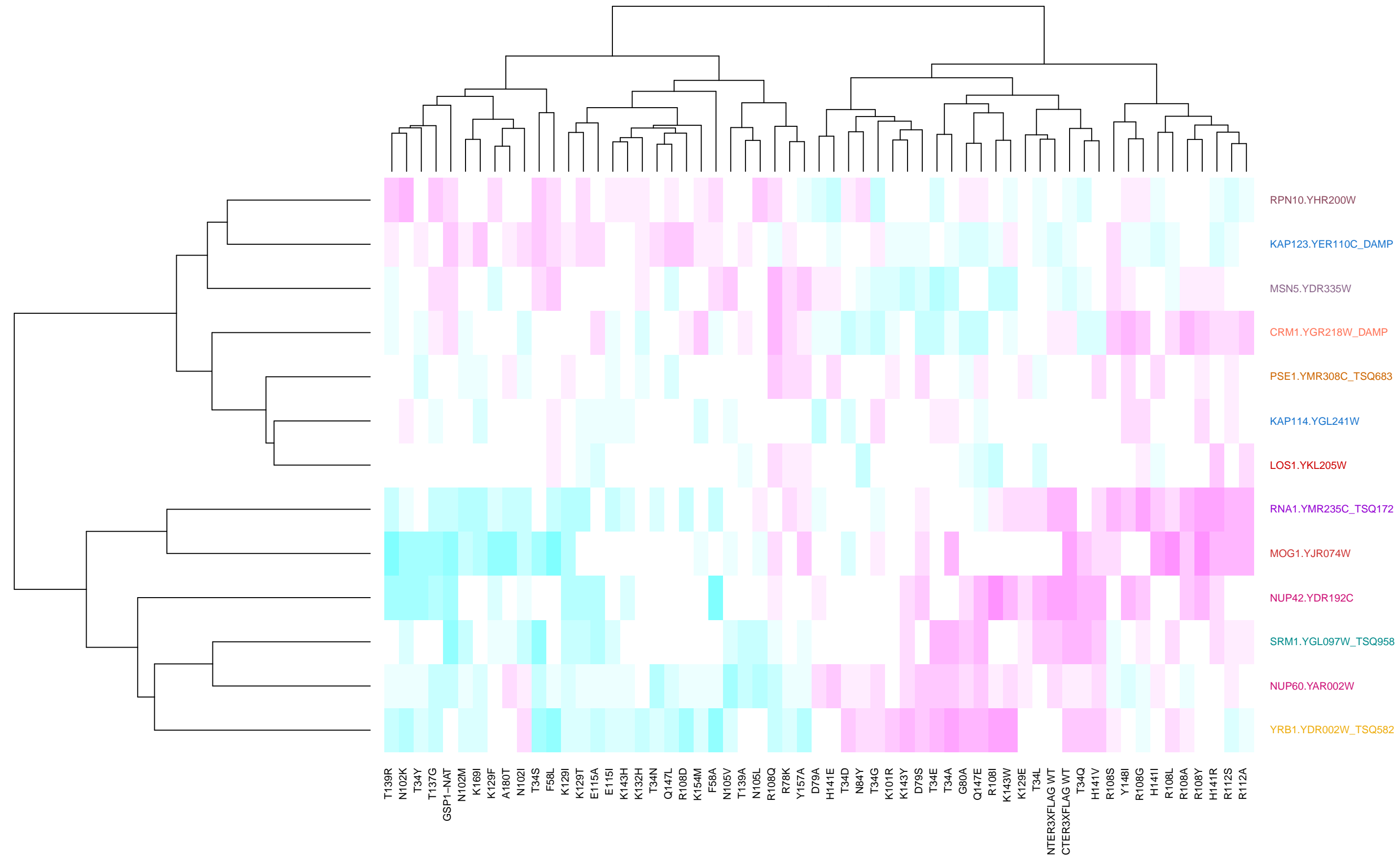
peptidyl-amino acid modification



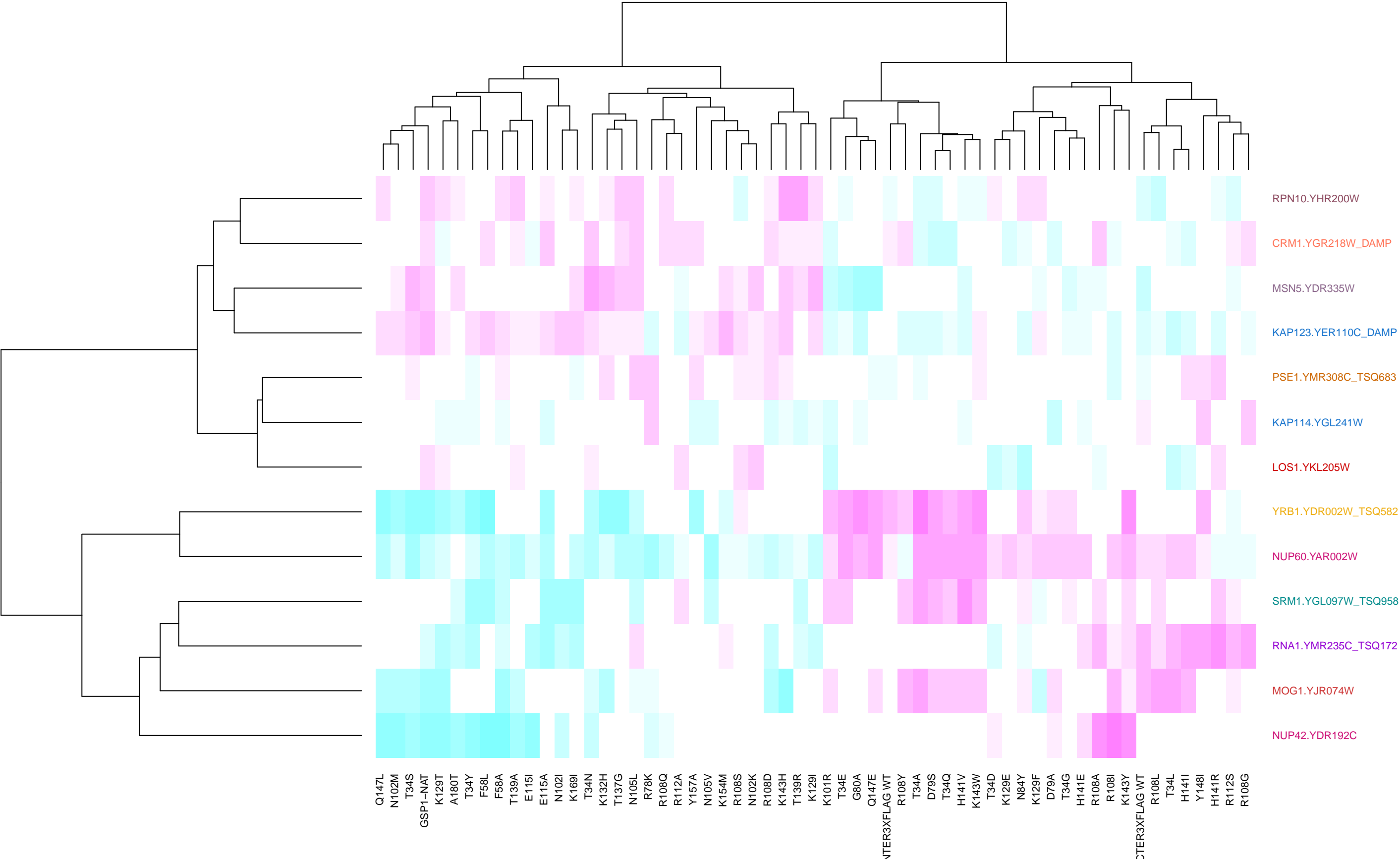
protein targeting_GO_2



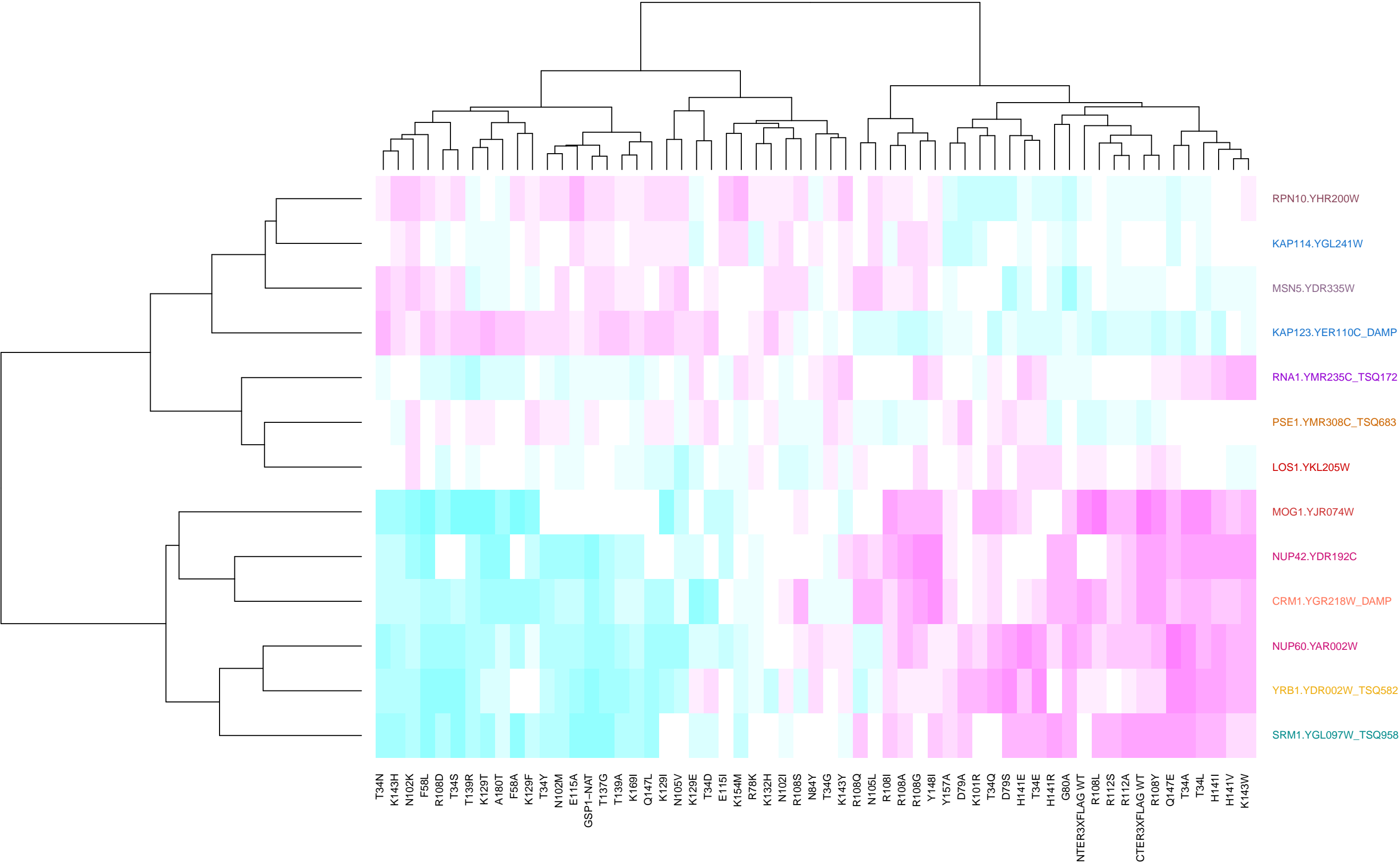
ribosomes and translation



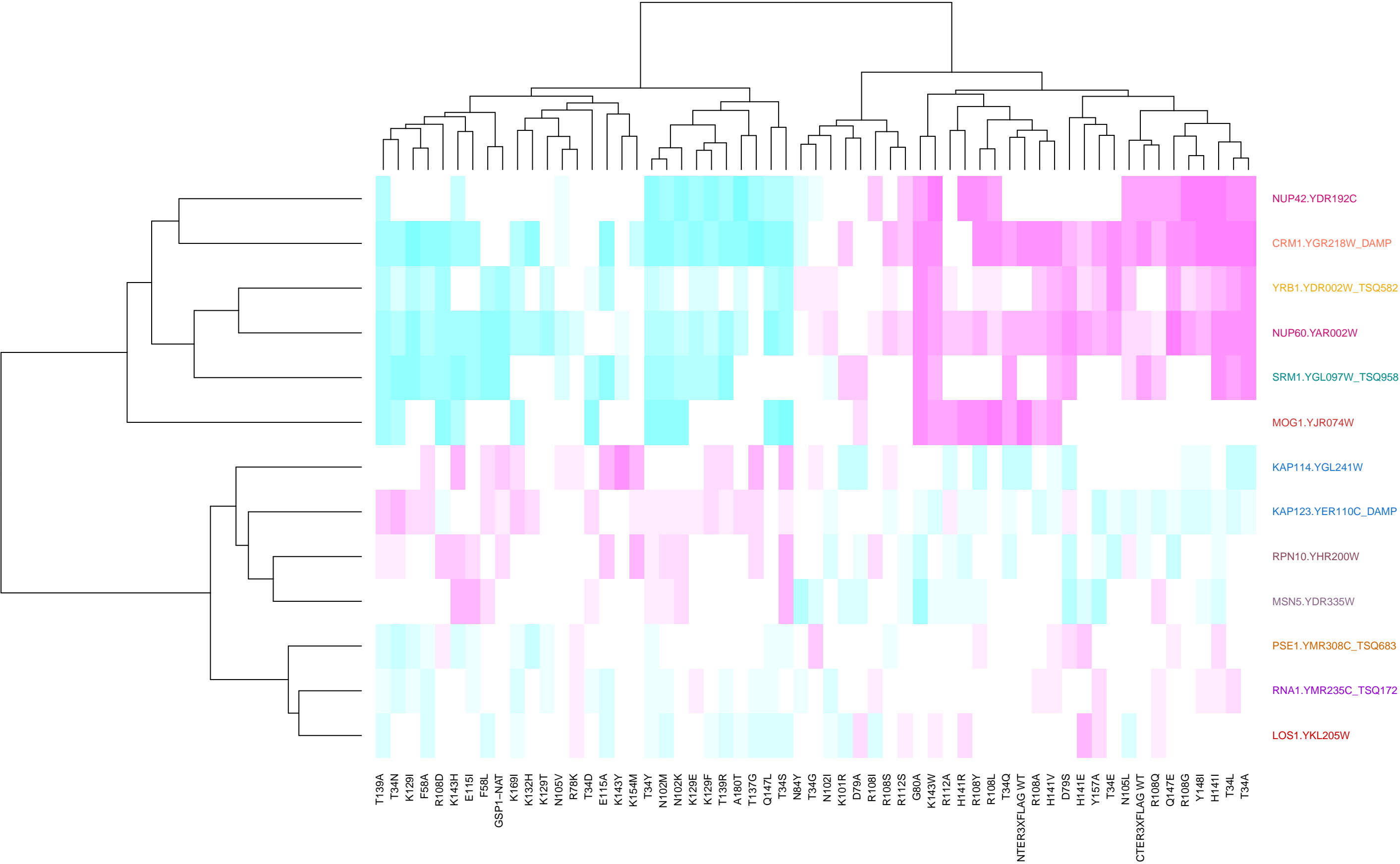
ribosomes and translation_GO_1



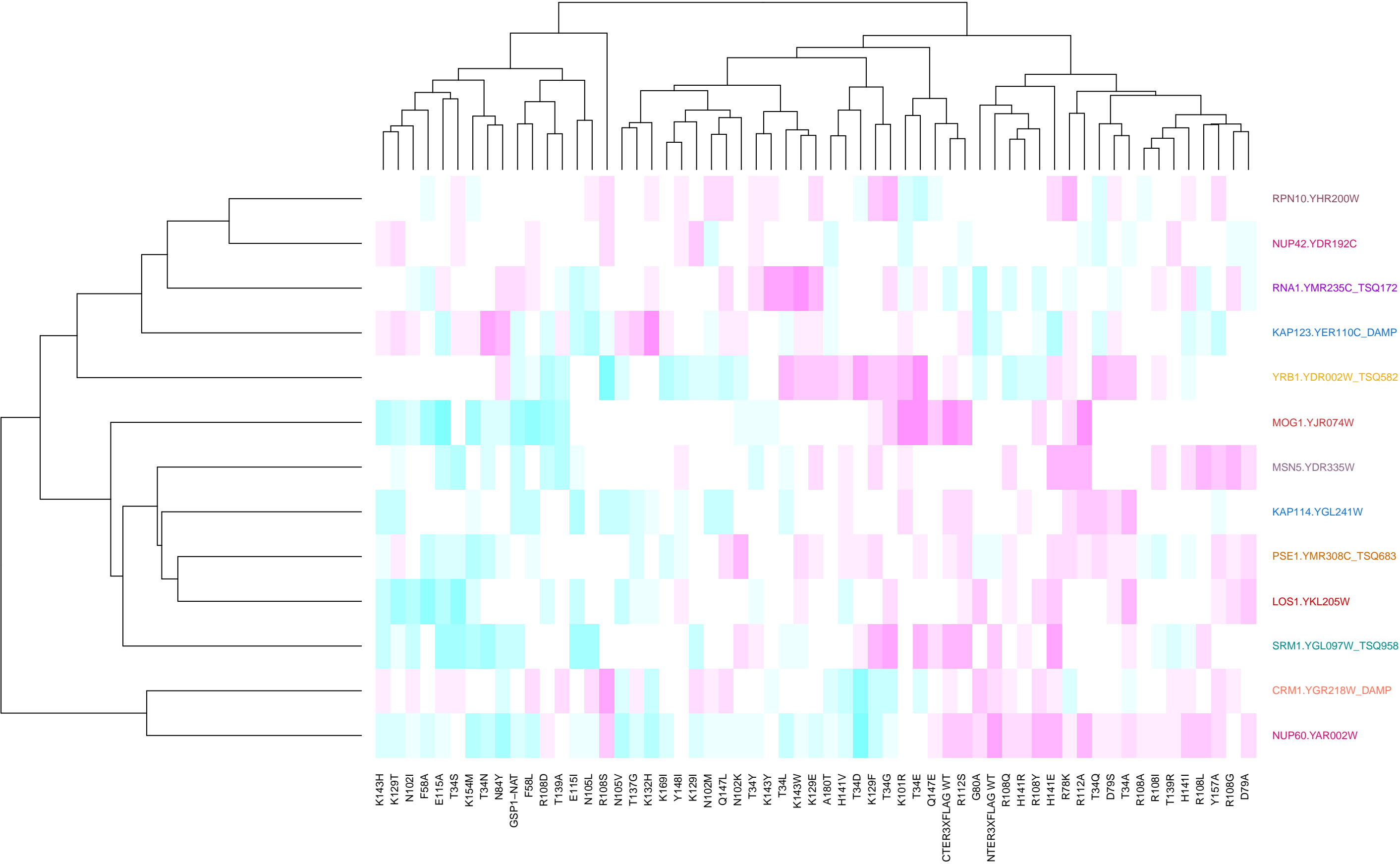
transcription and mRNA processing



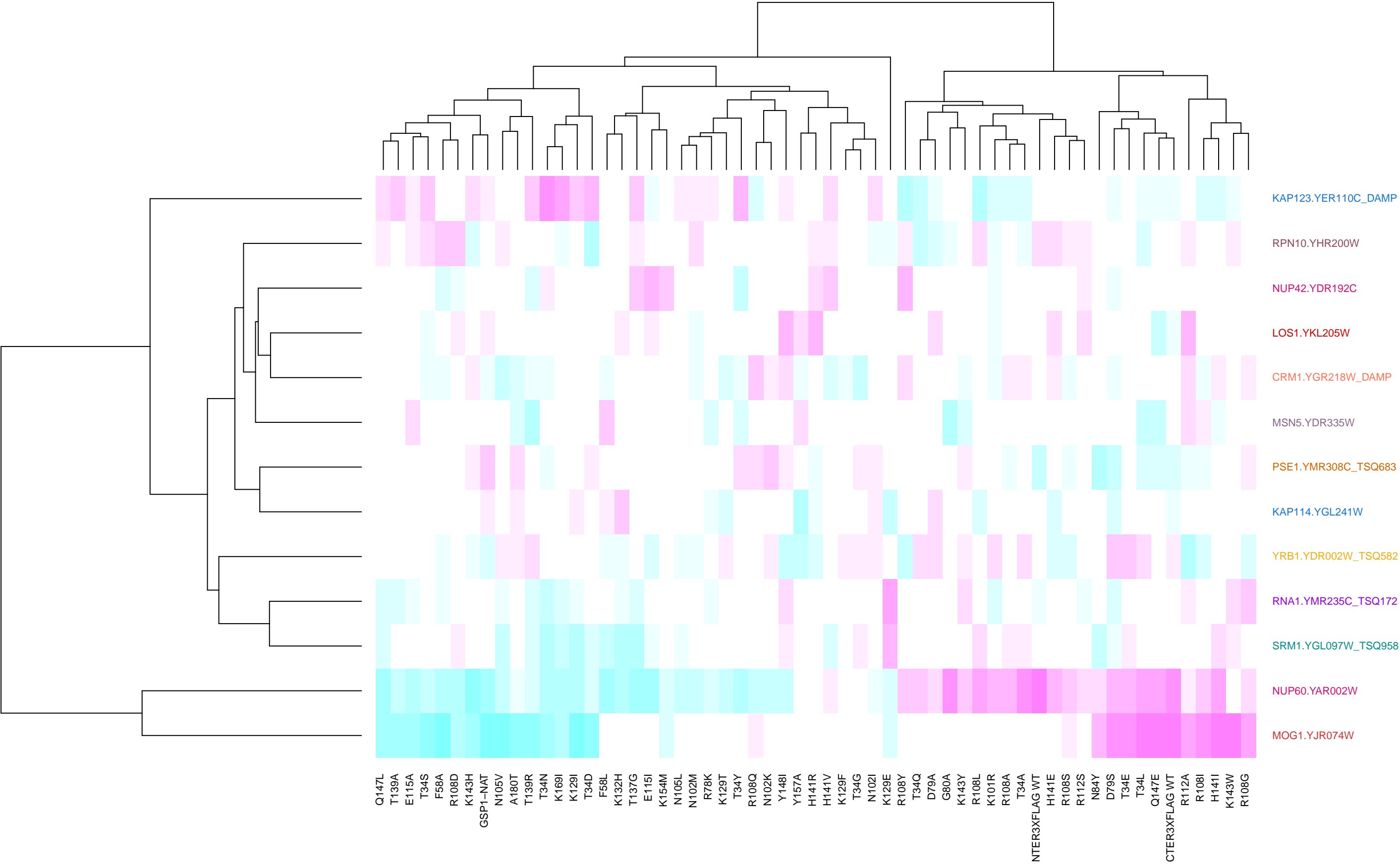
transcription and mRNA processing_GO_1



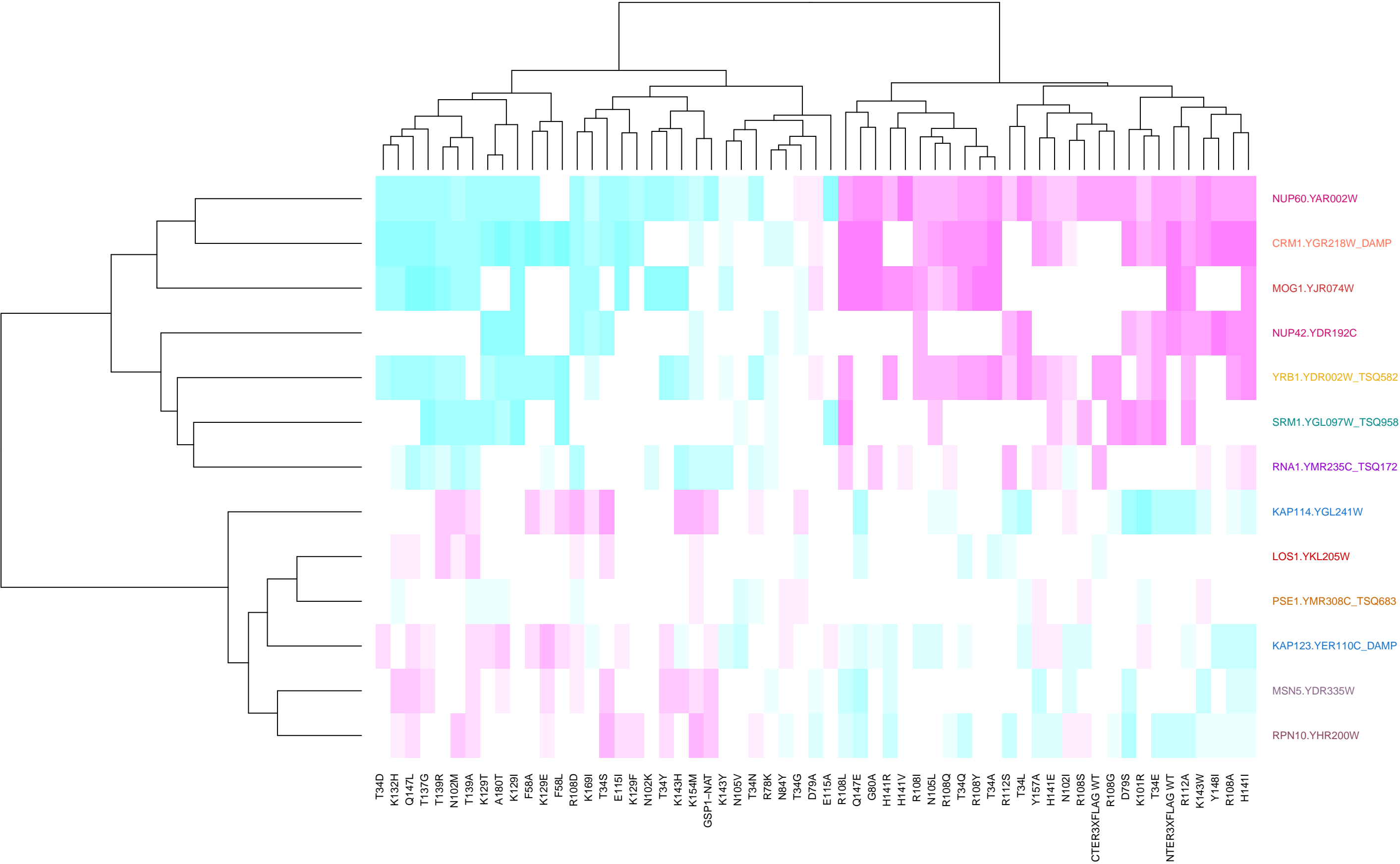
transcription from RNA polymerase II promoter_GO_1



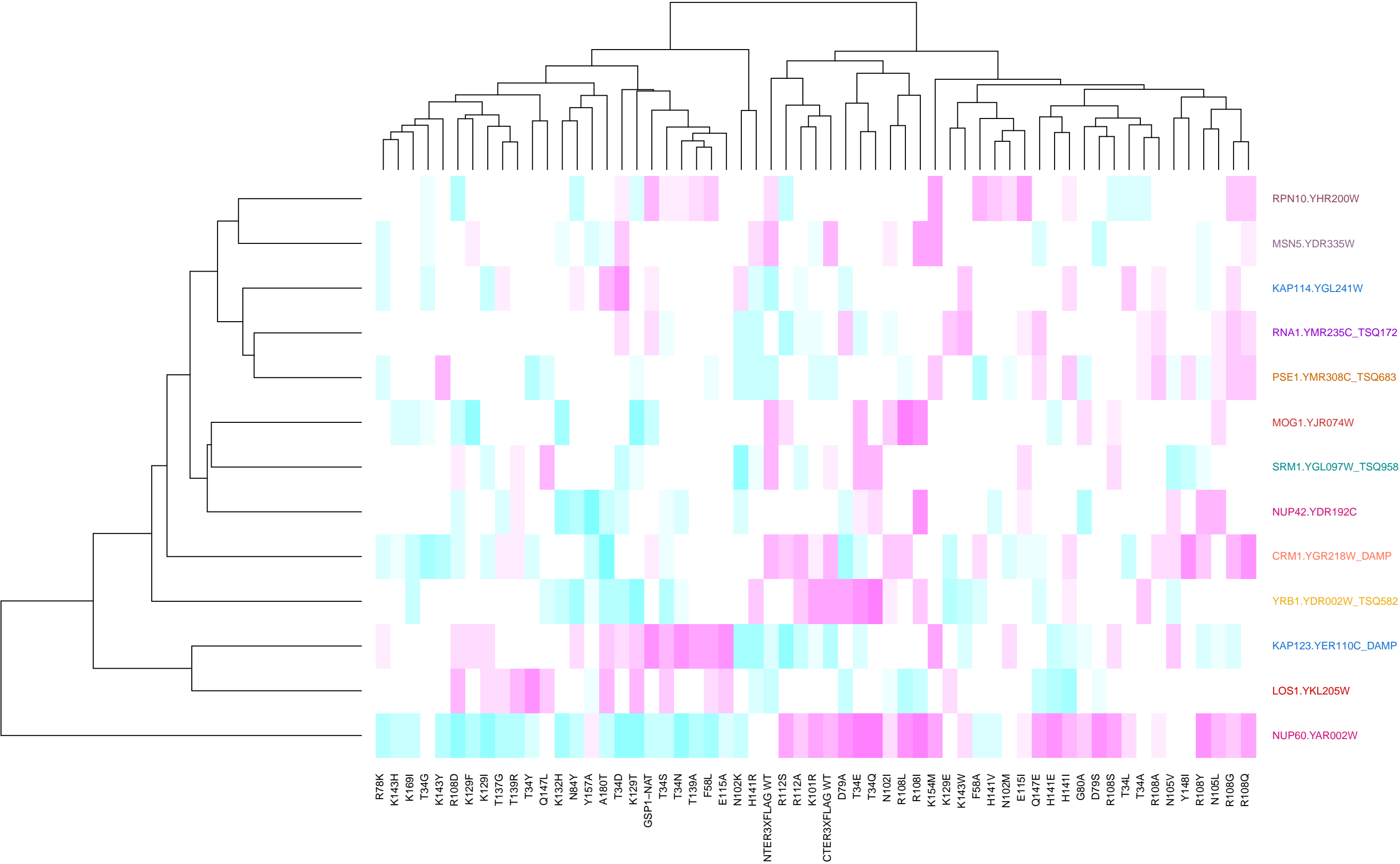
transcription_GO_2



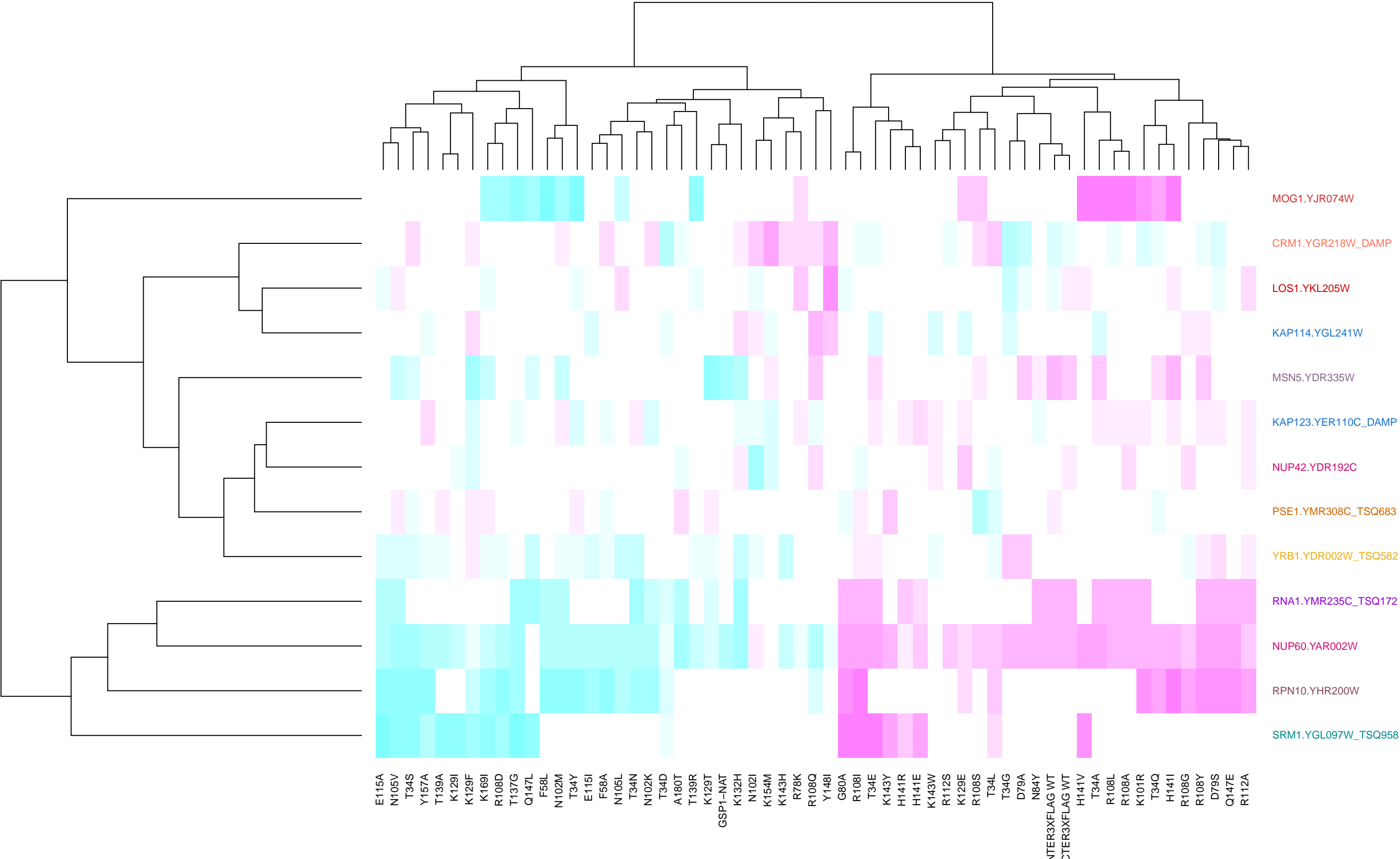
whole_library_13



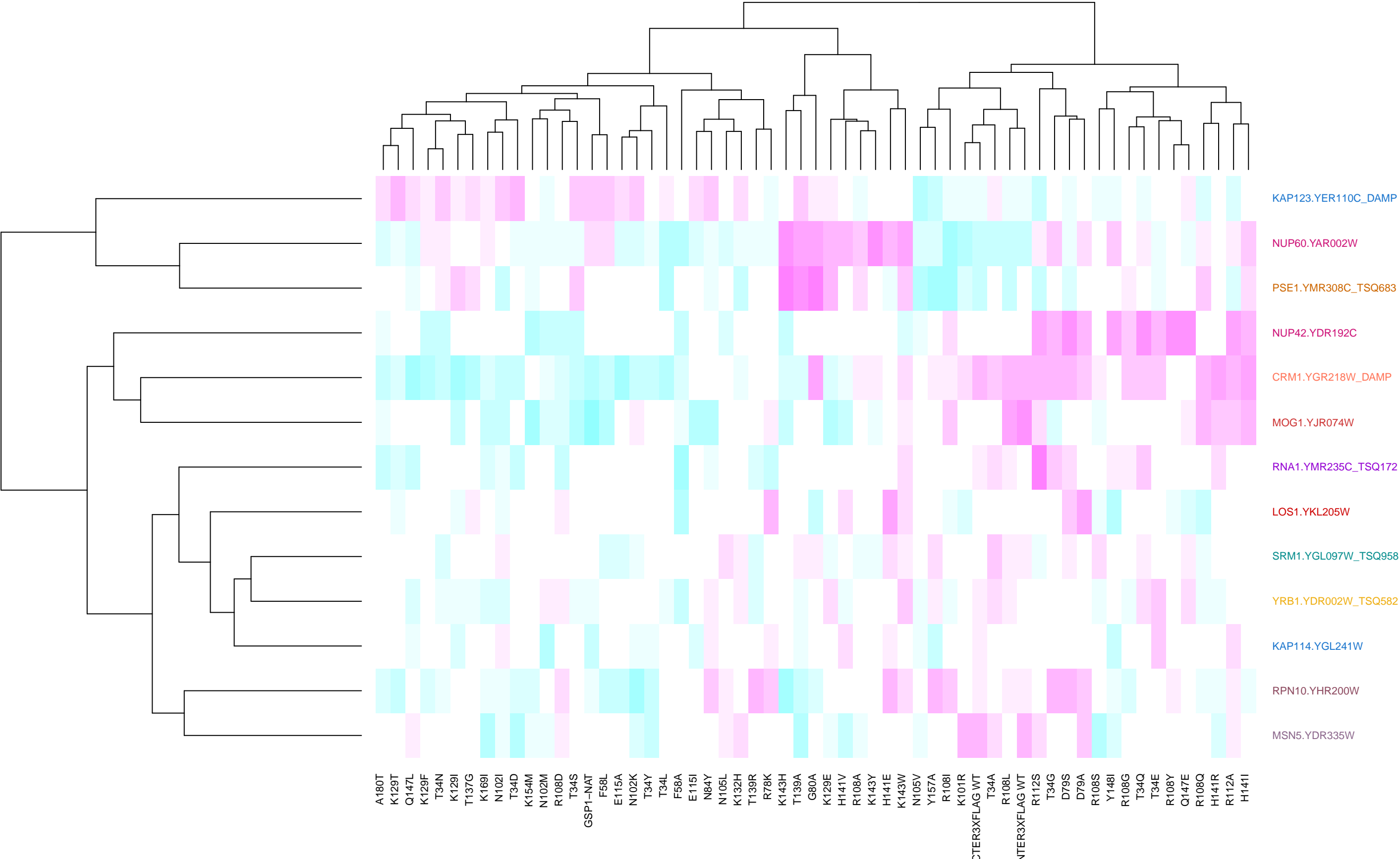
whole_library_17



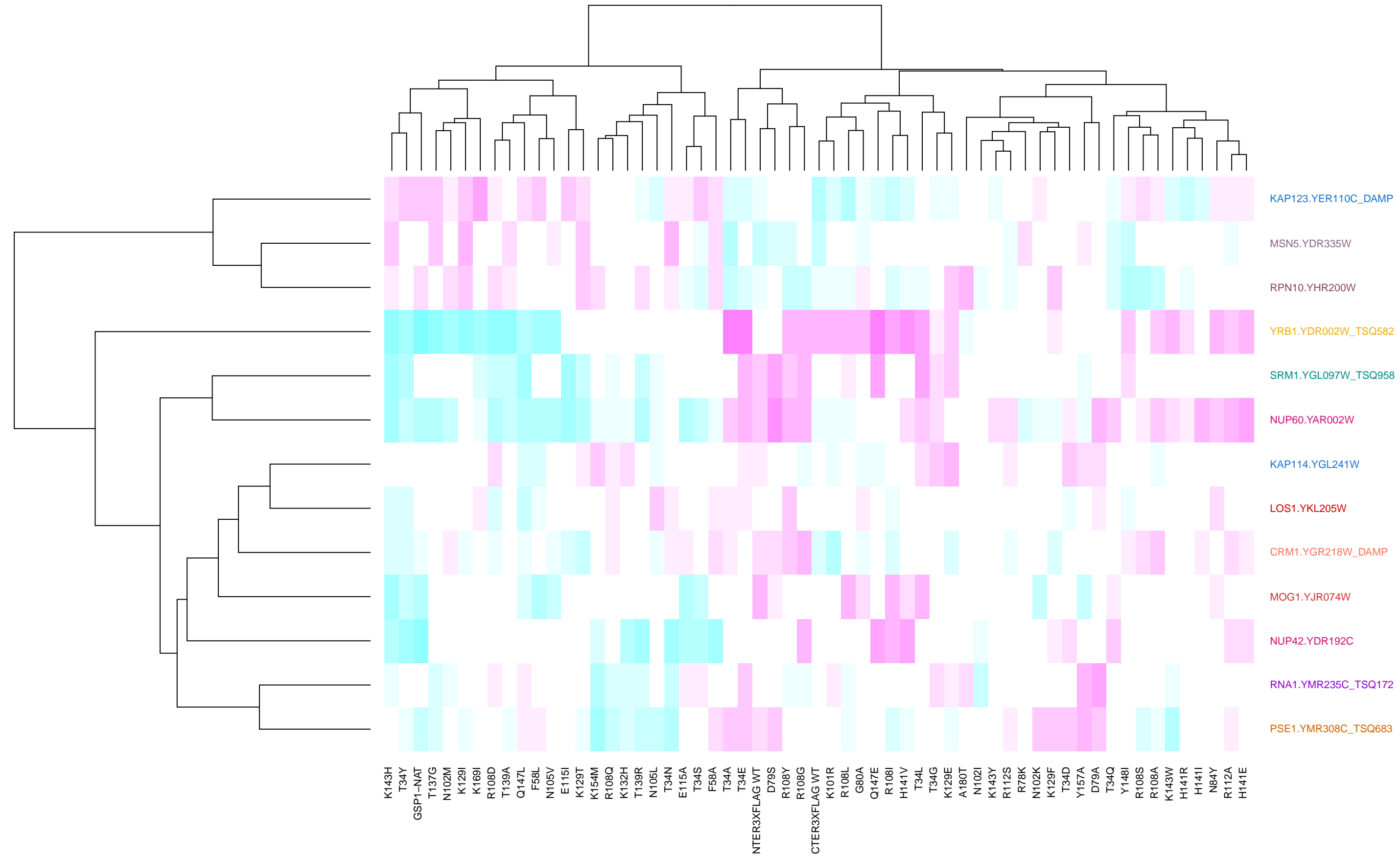
whole_library_20



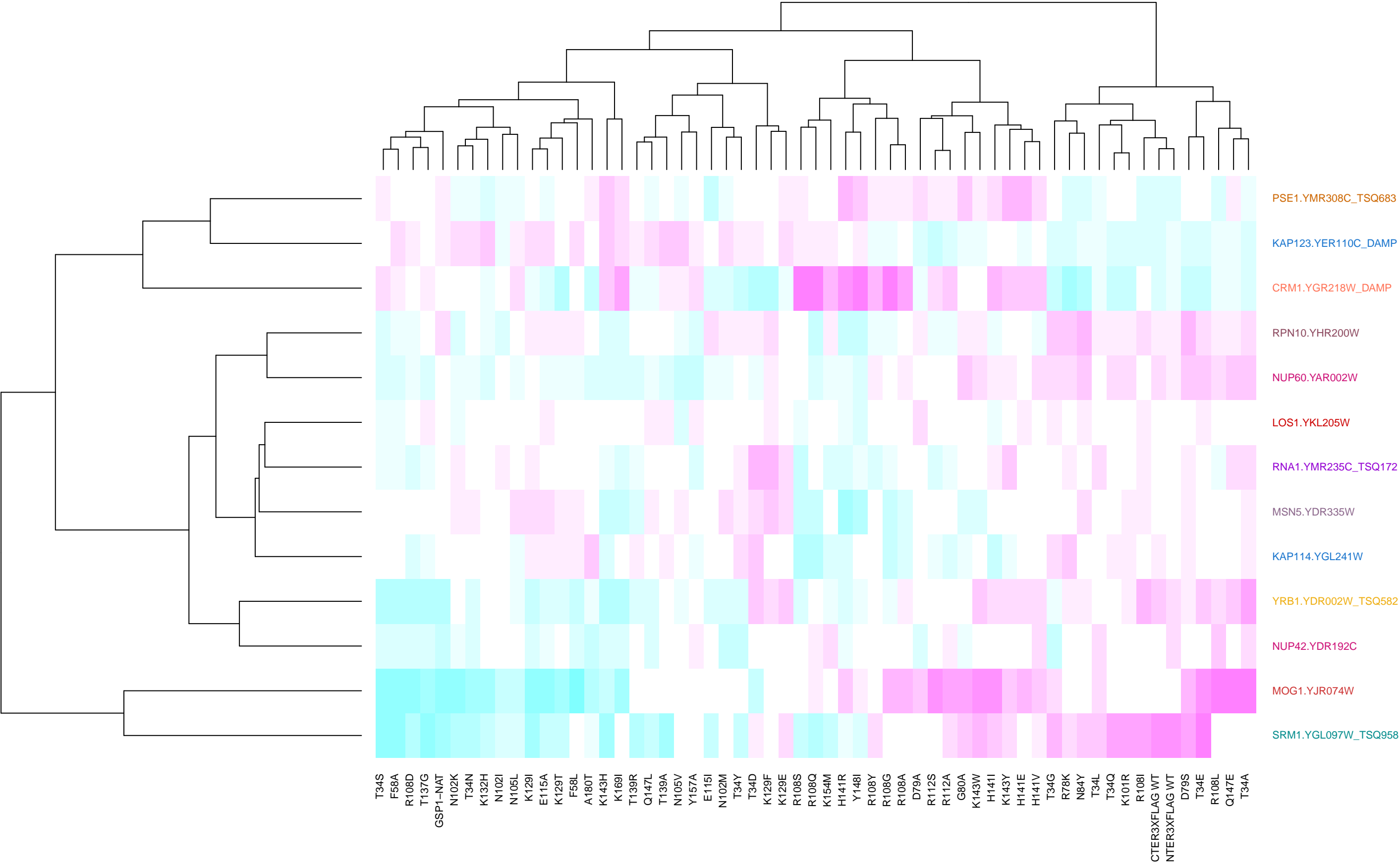
whole_library_4



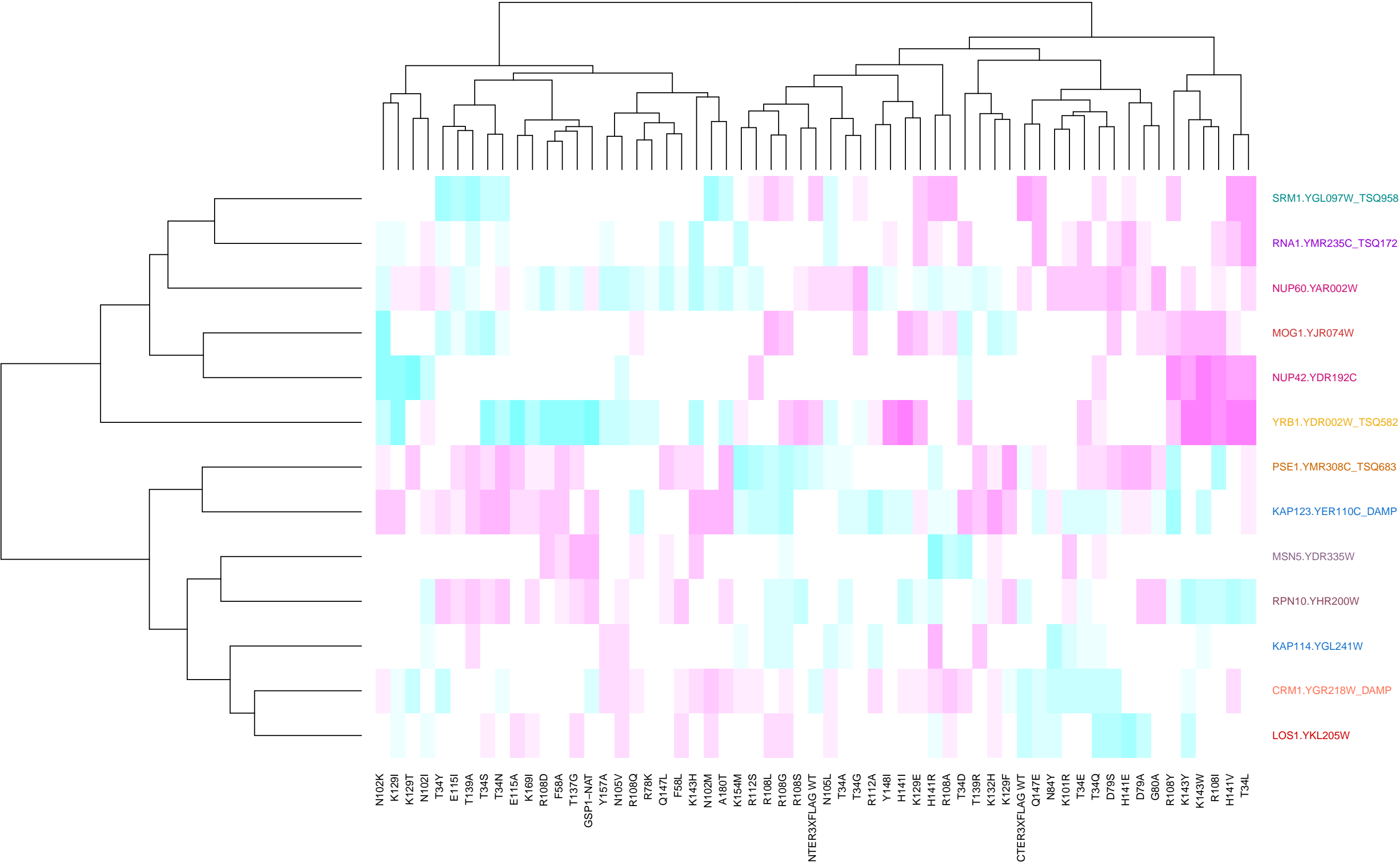
whole_library_6



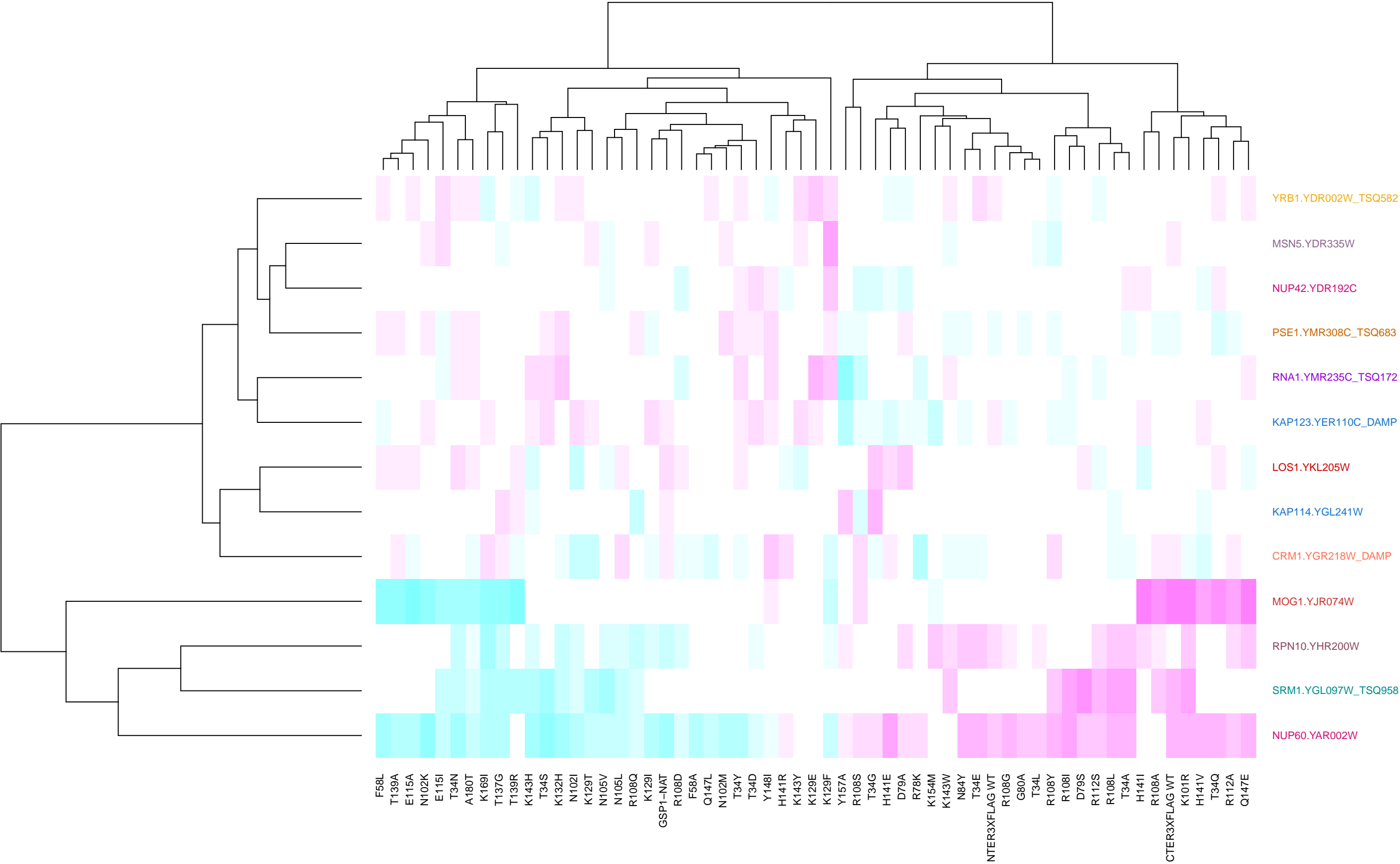
cell cycle



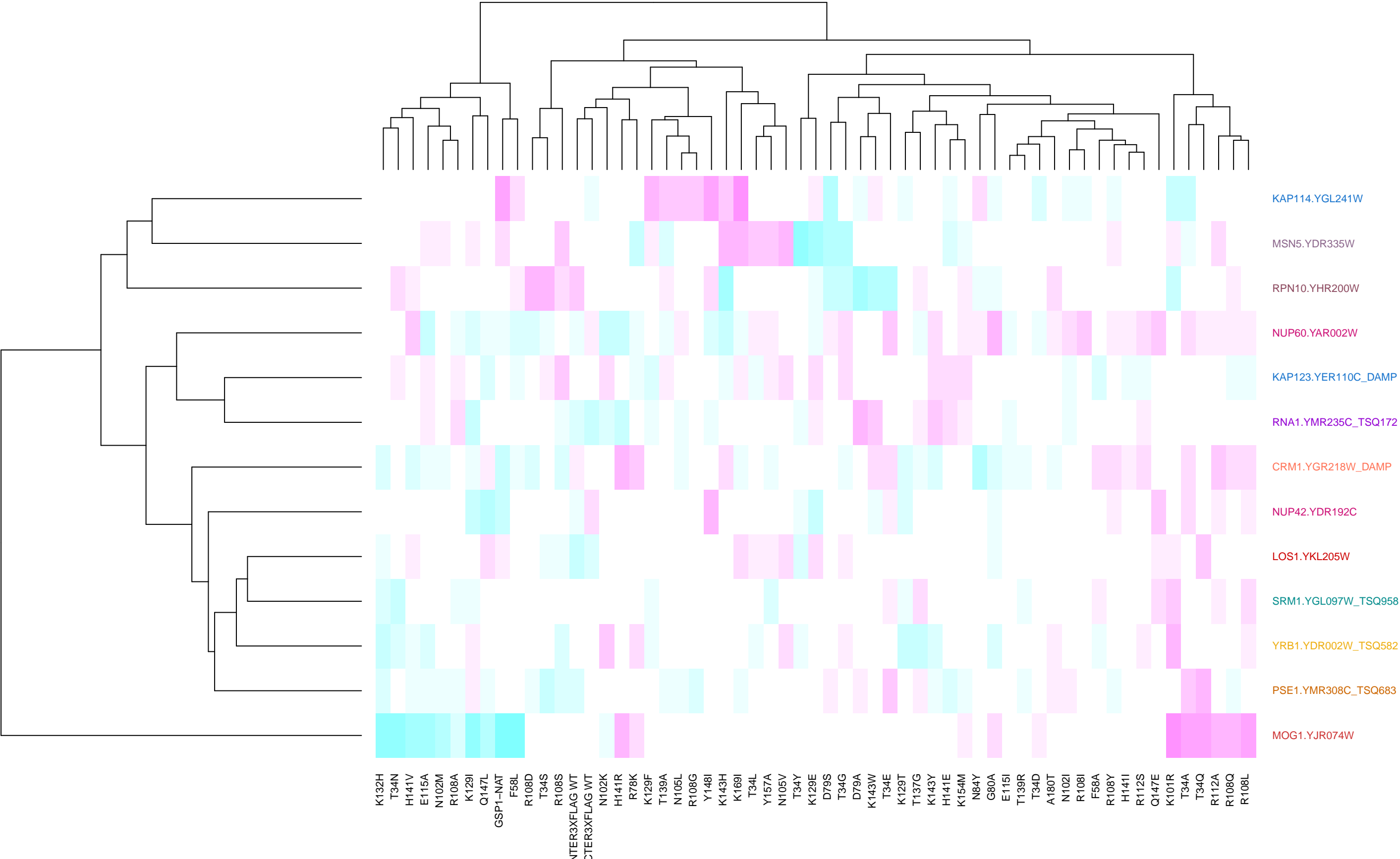
cell cycle_GO_1



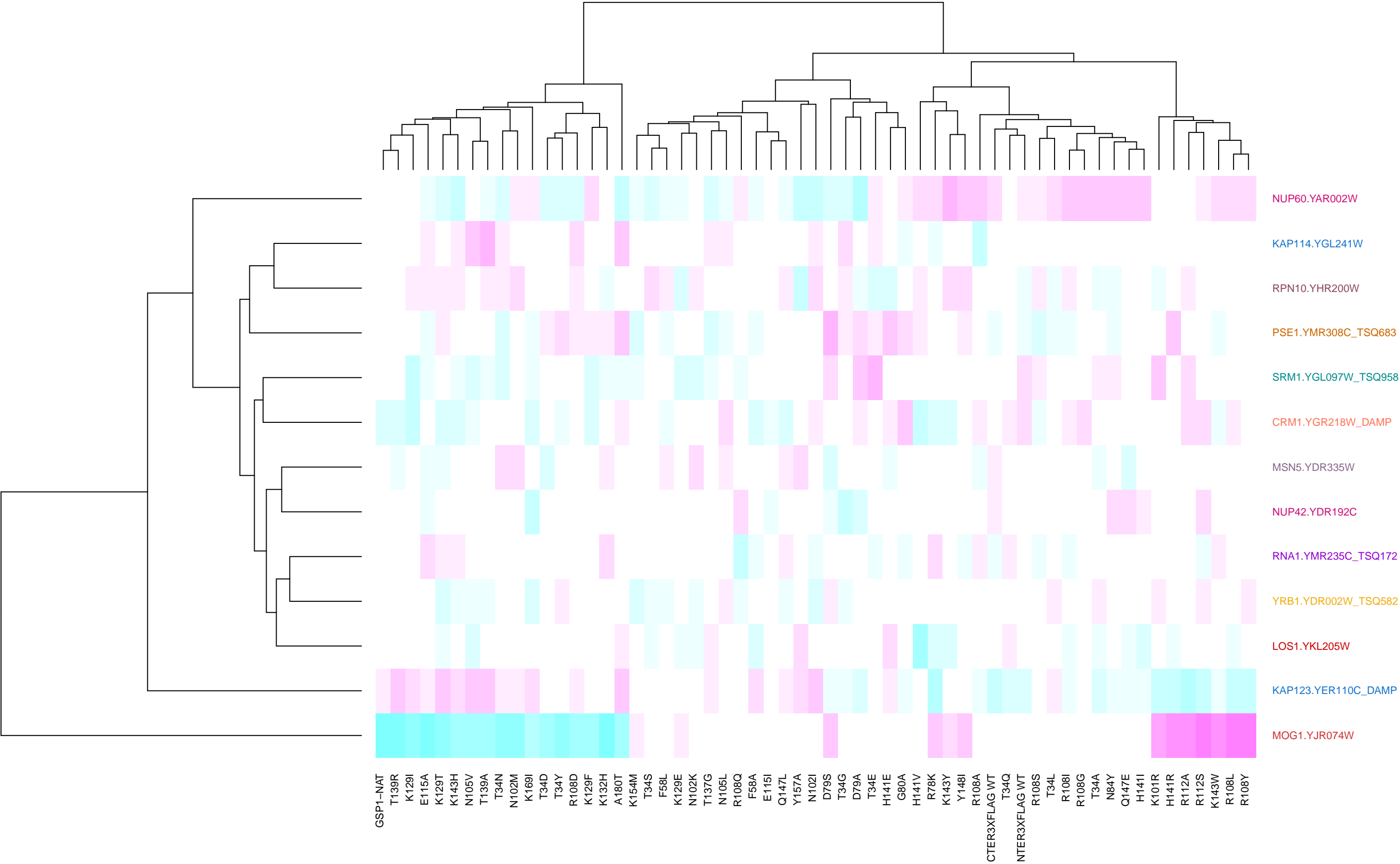
cell cycle_GO_4



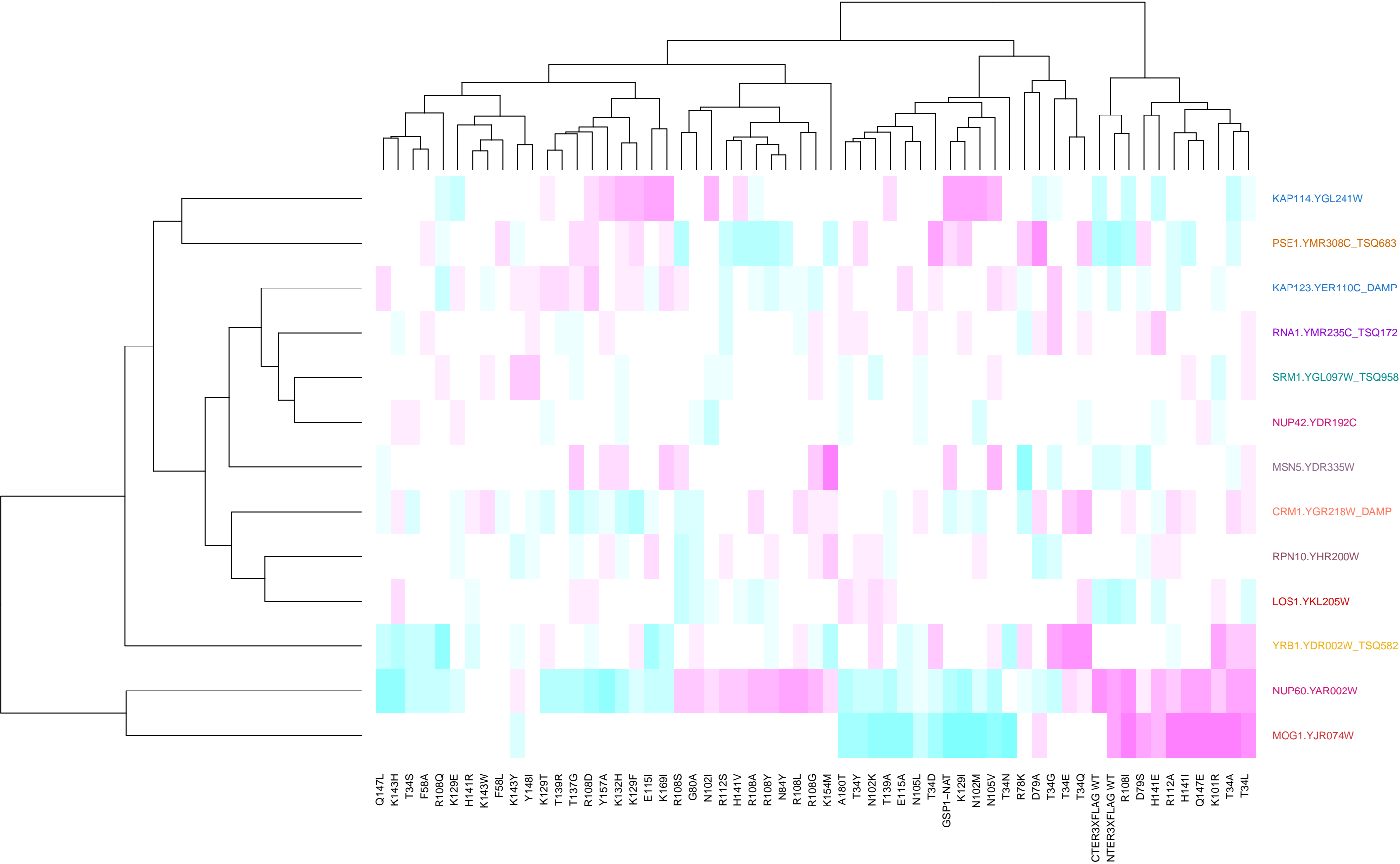
chromatin organization_GO_3



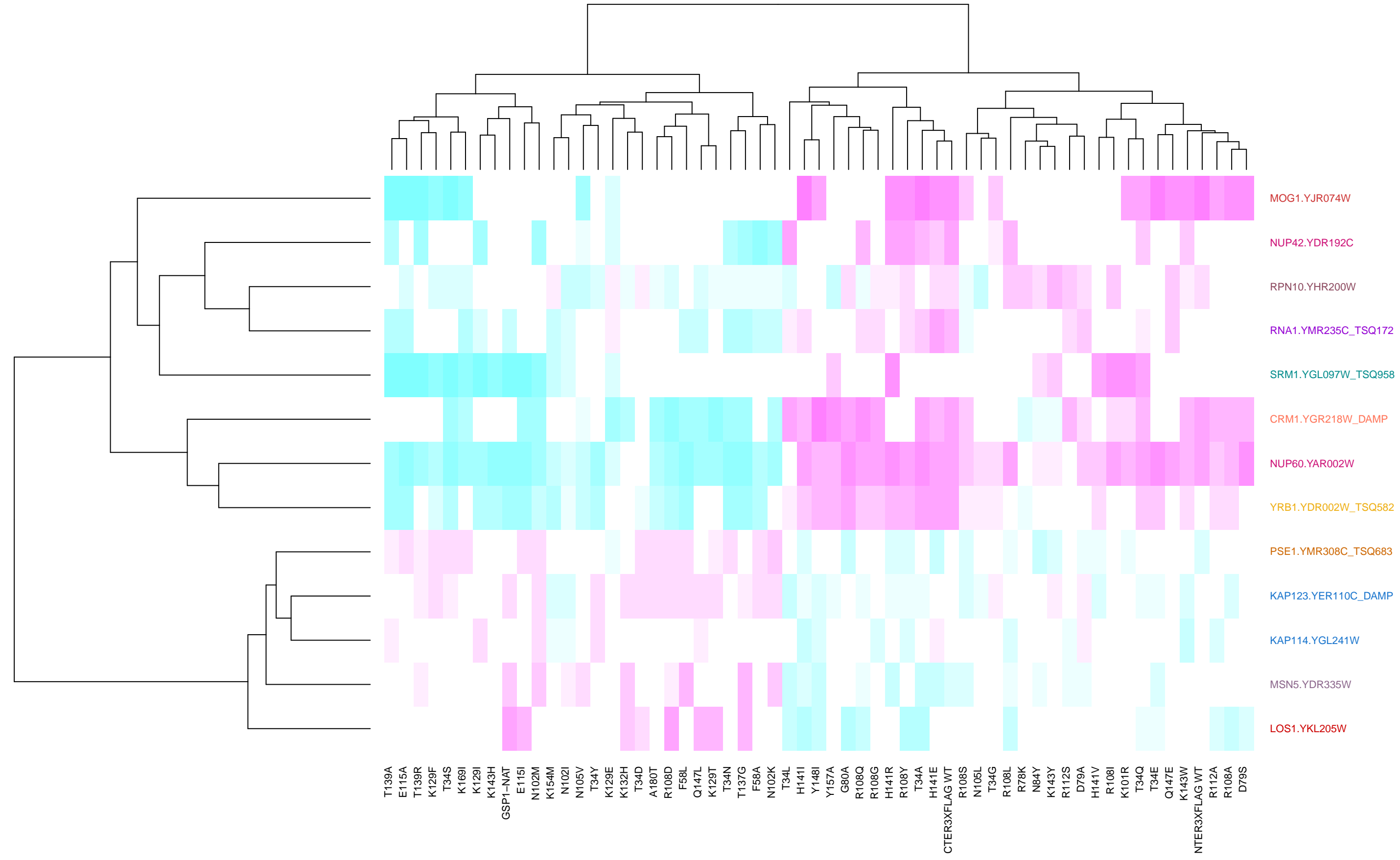
chromatin_GO_3



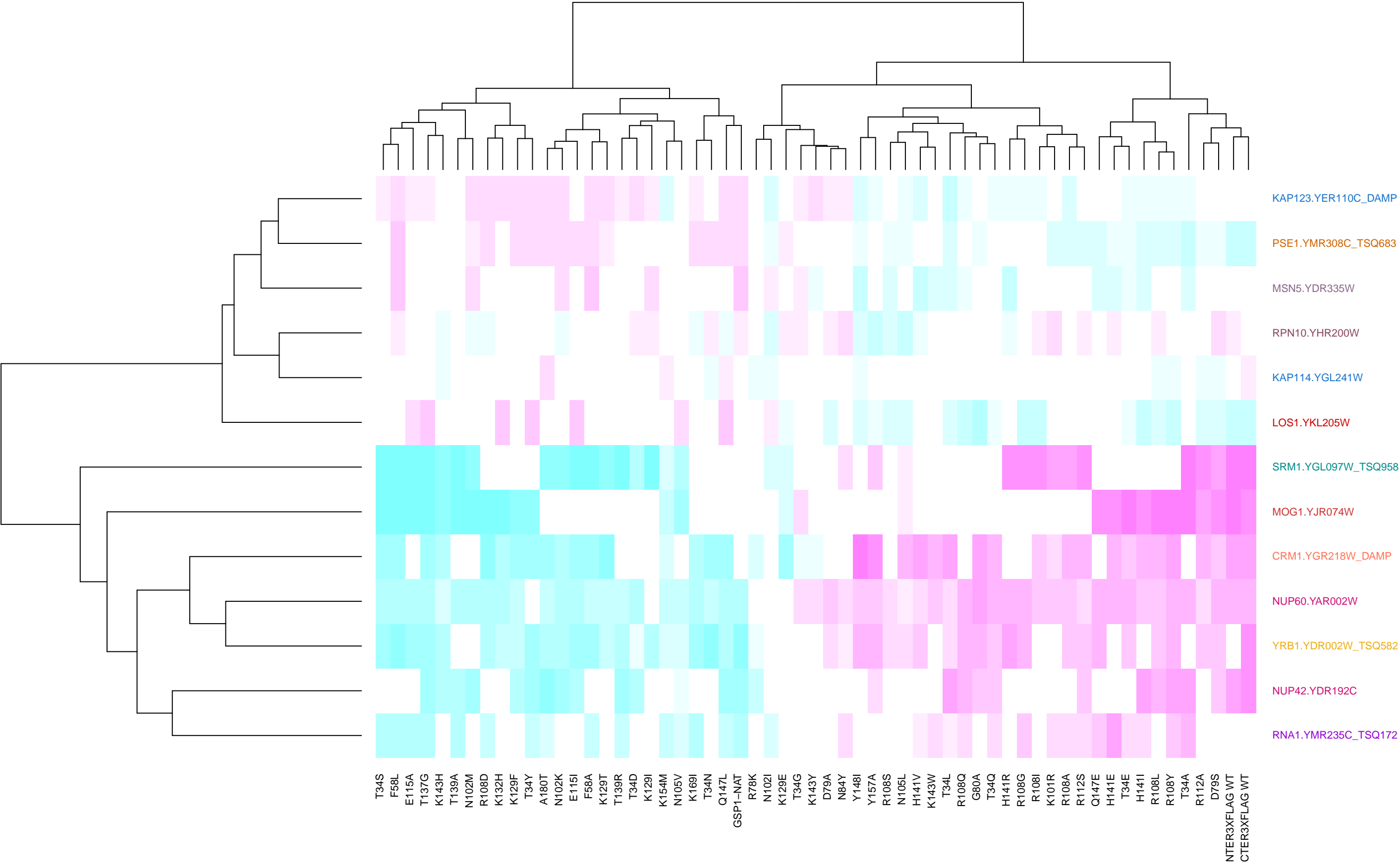
DNA-templated transcription, elongation



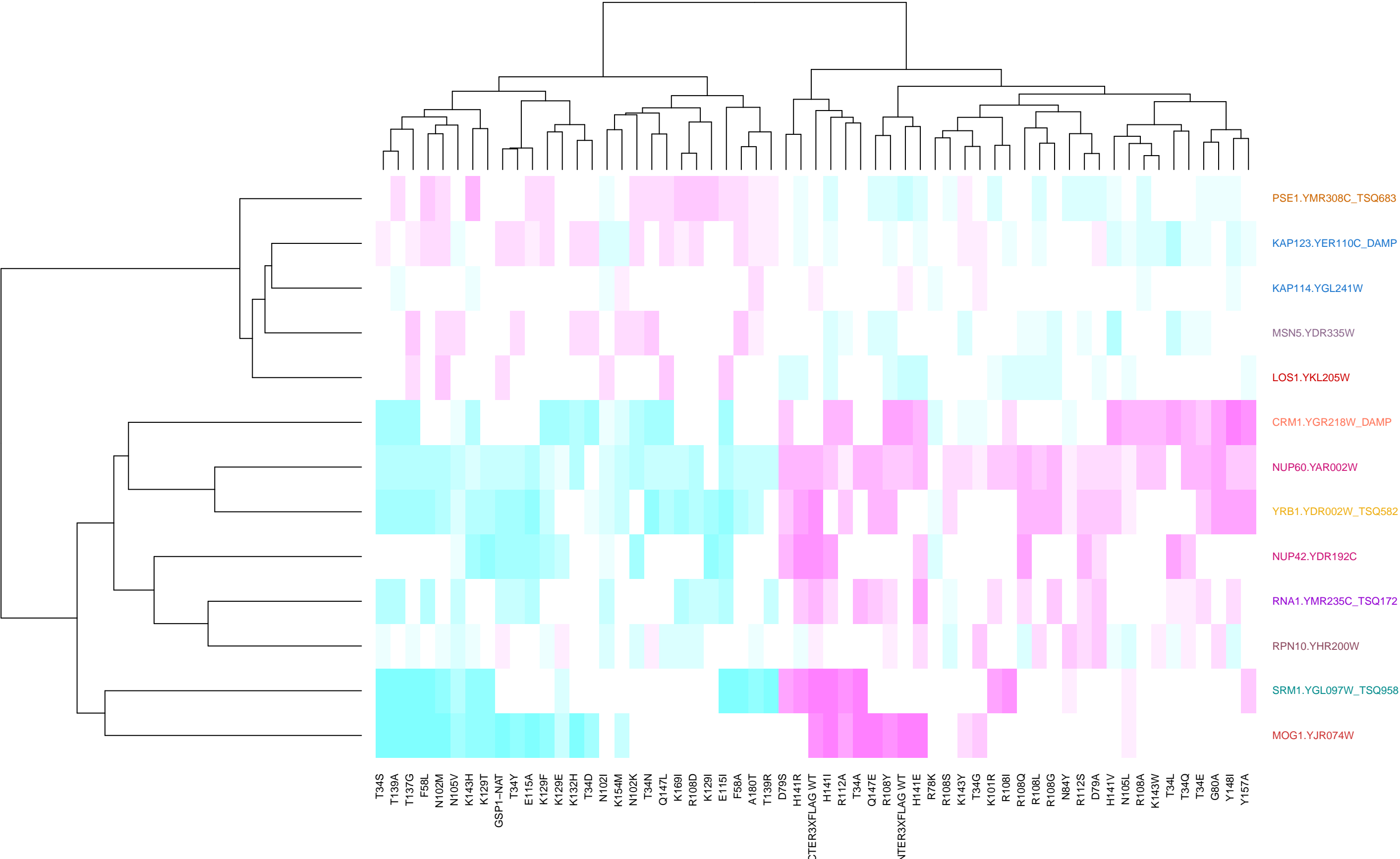
nuclear transport



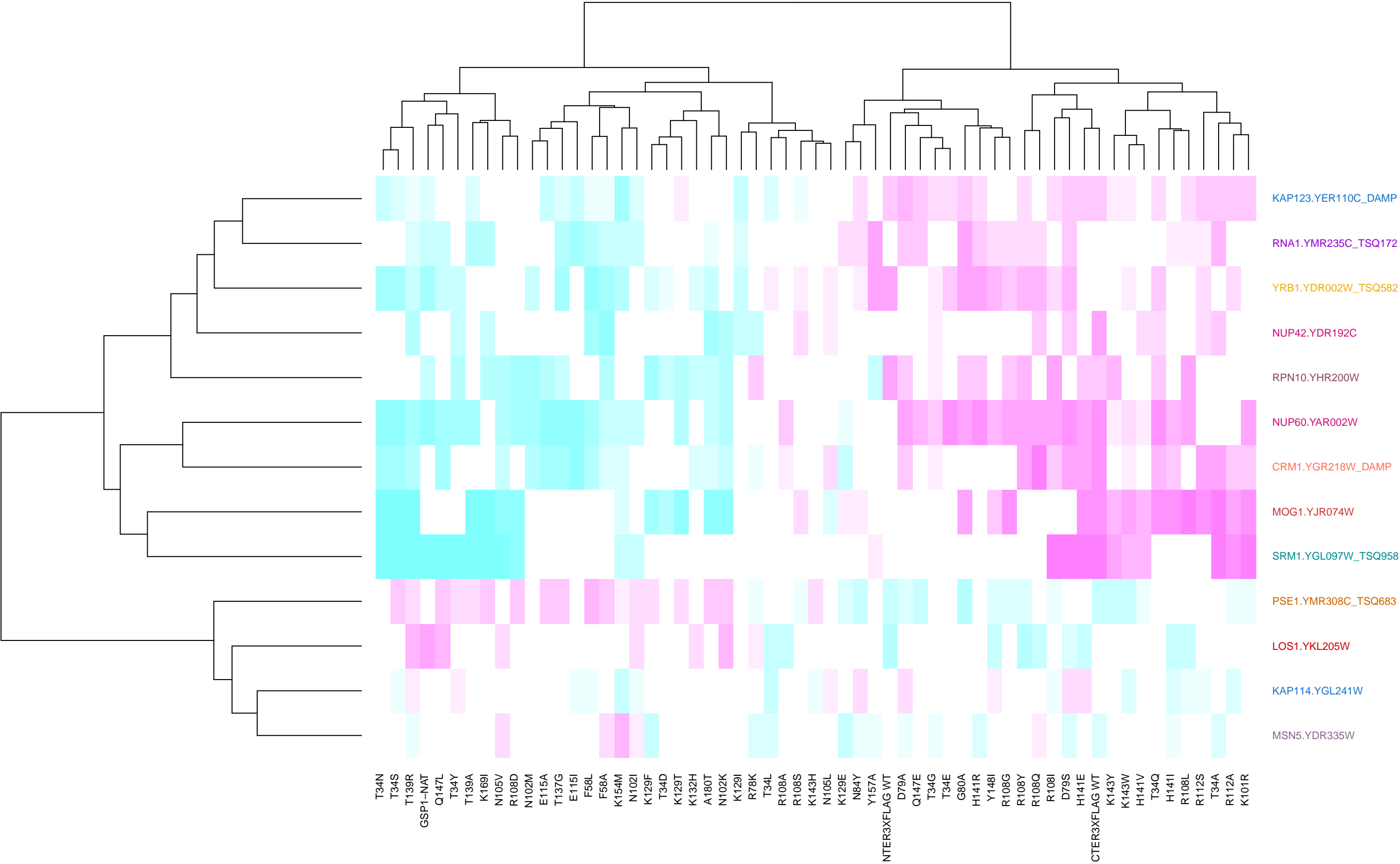
nuclear transport and organization



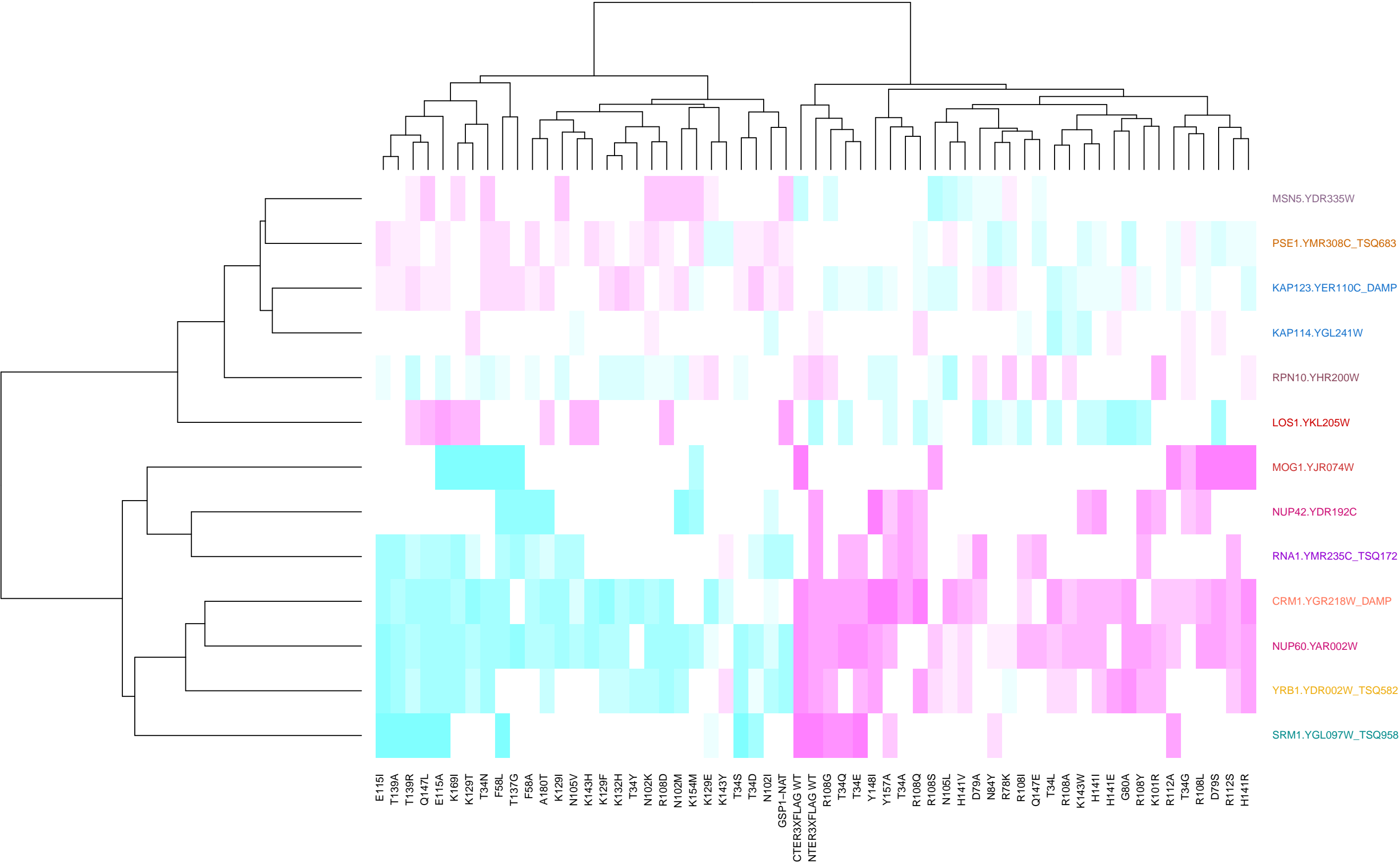
nuclear transport and organization_GO_1



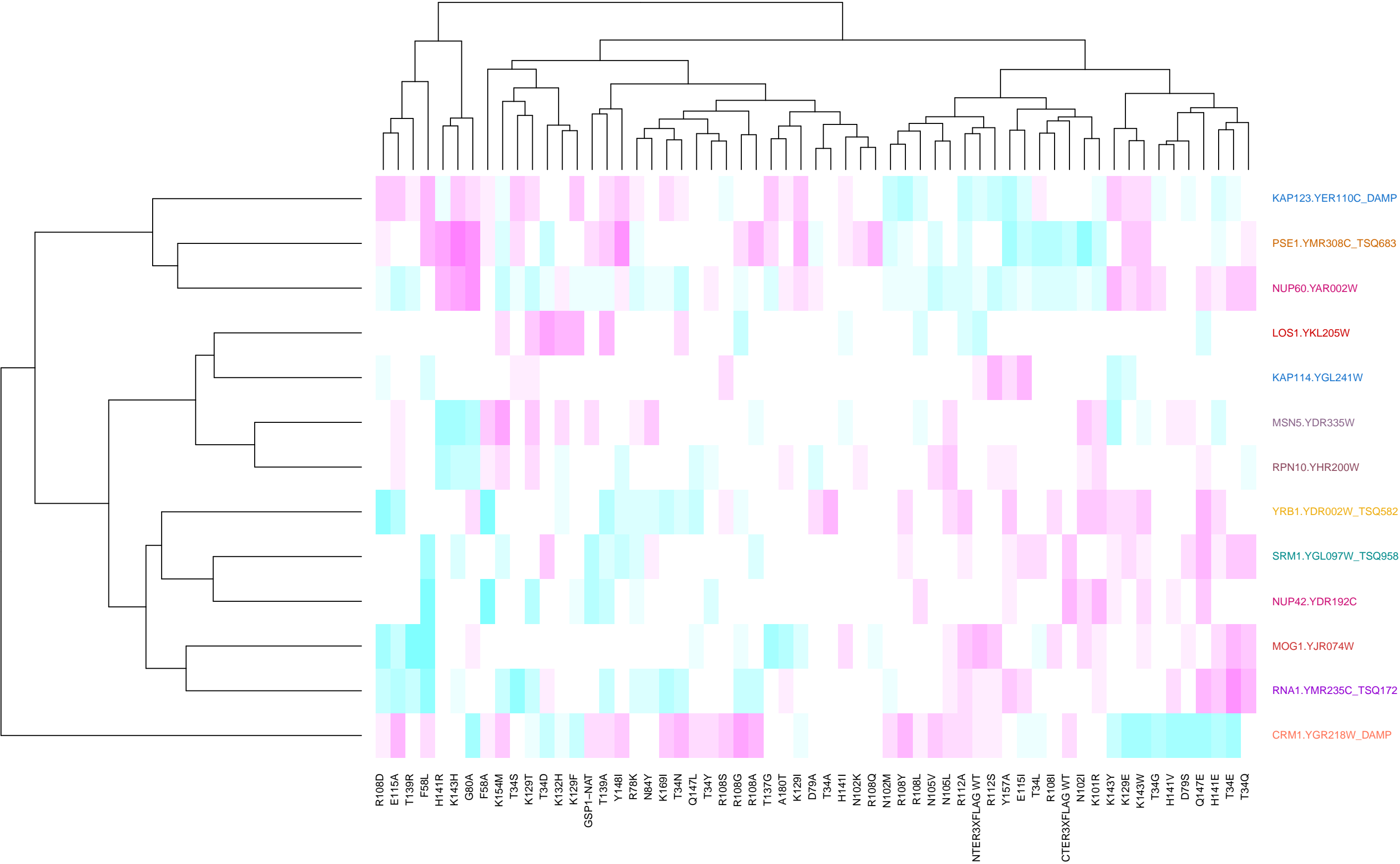
nuclear transport_GO_1



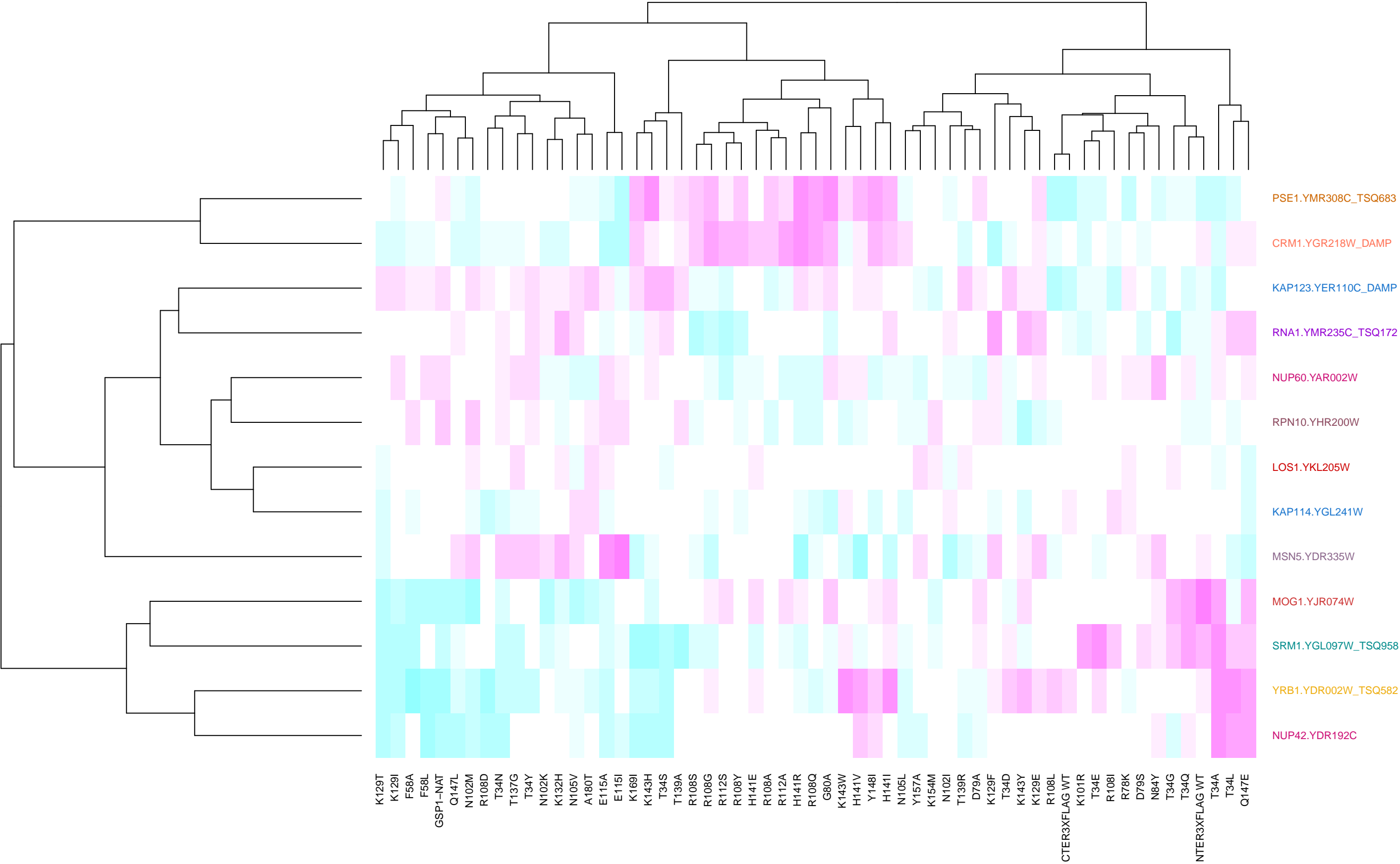
nucleobase-containing compound transport



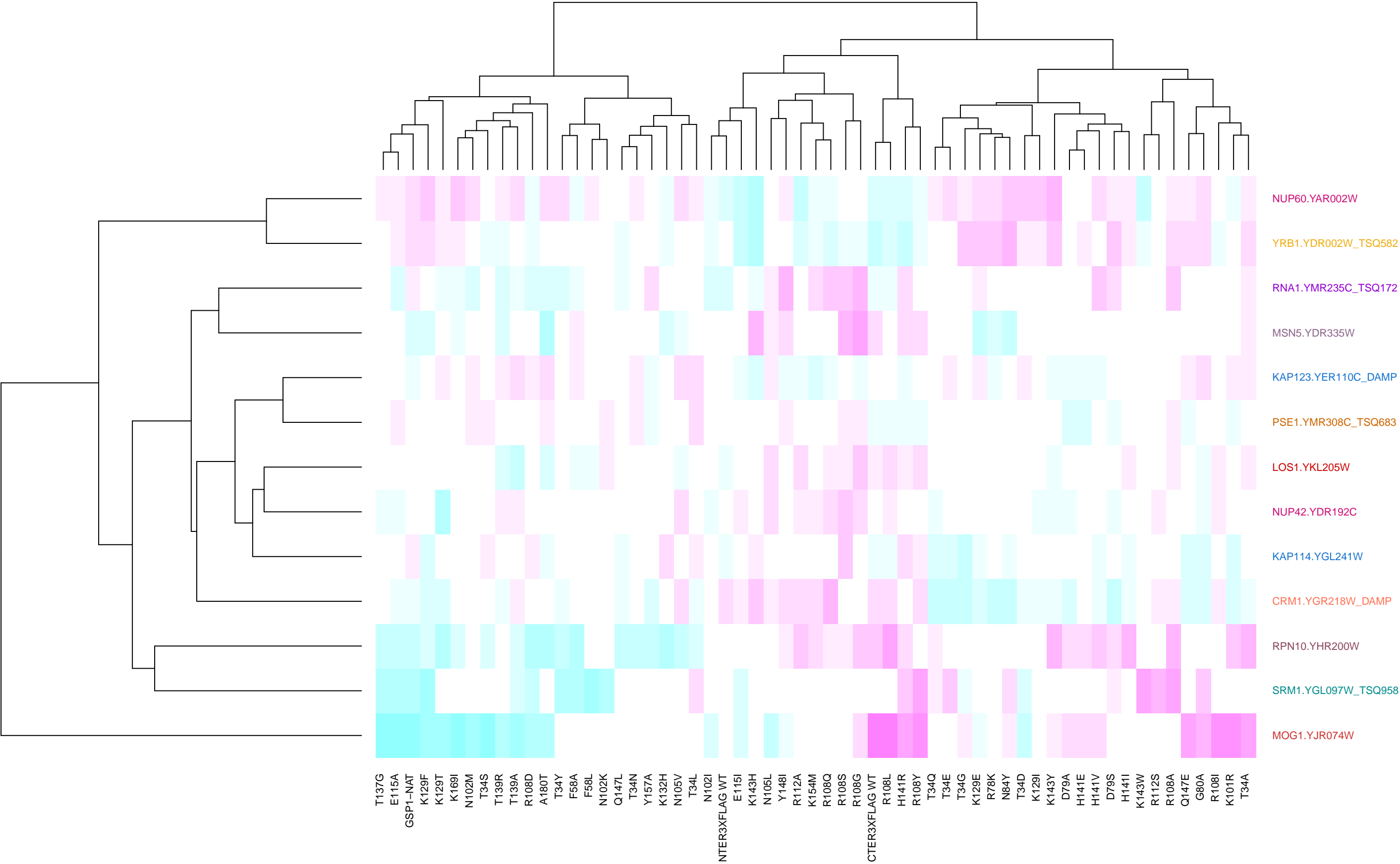
nucleolus



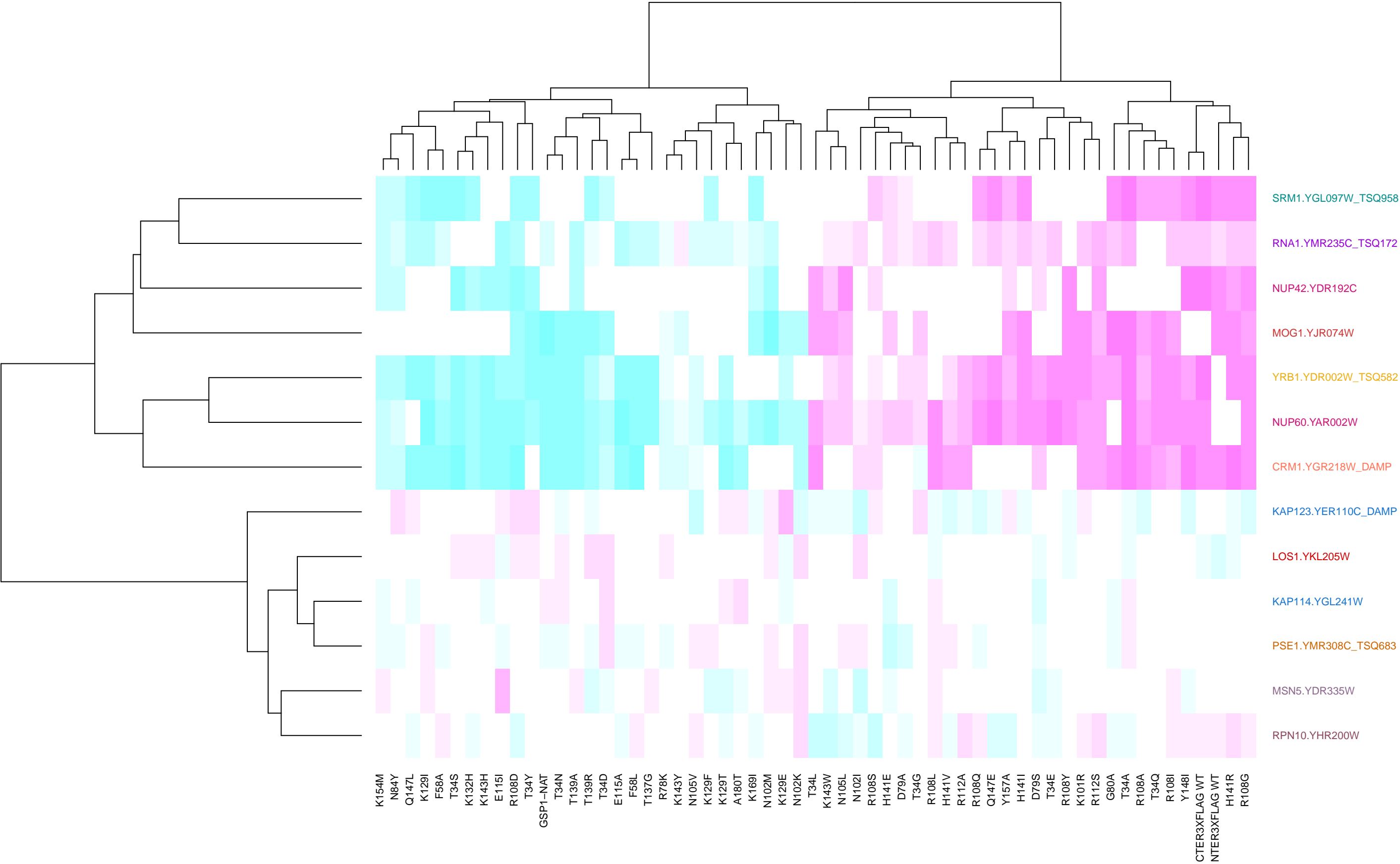
organelle fission



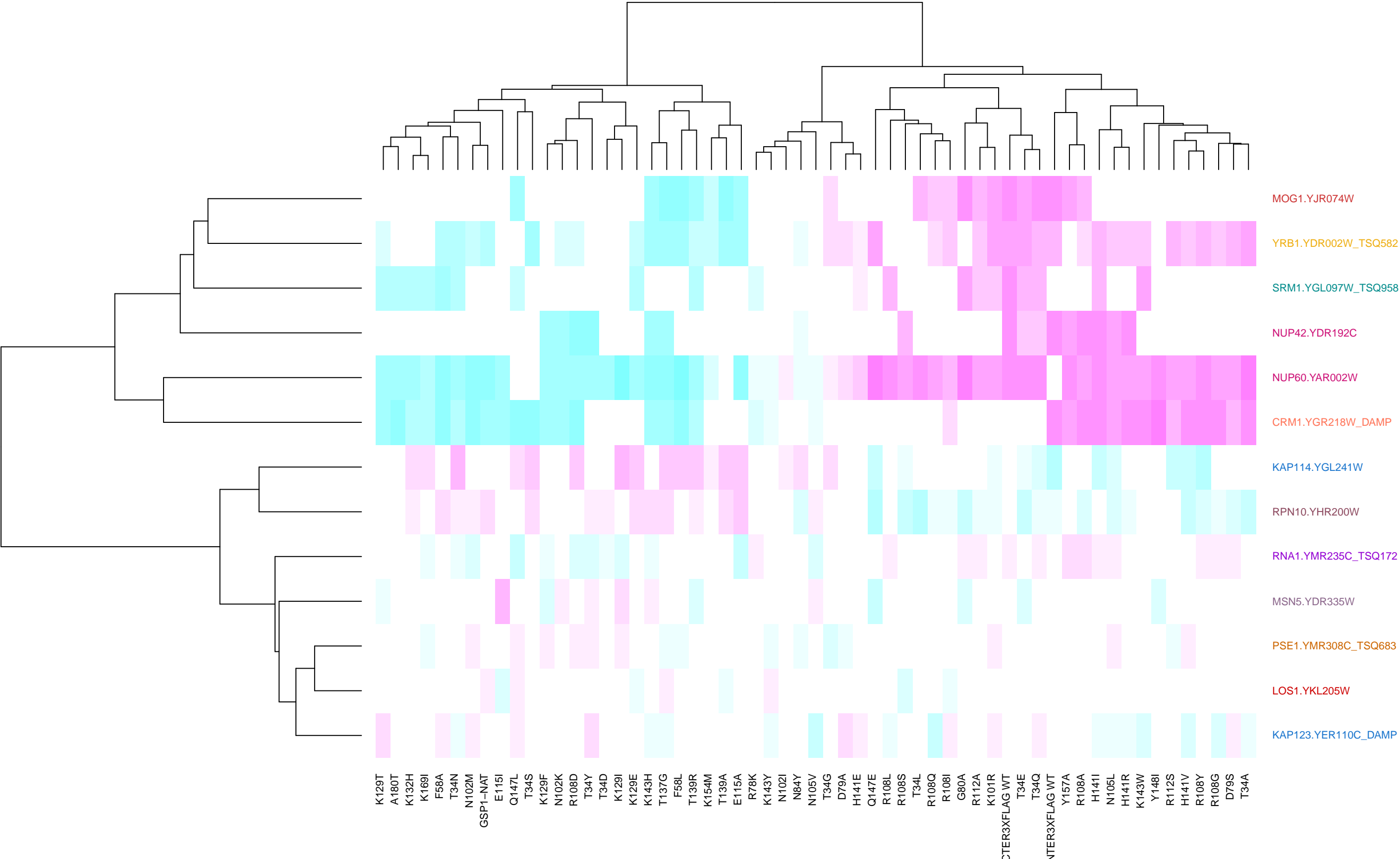
protein modification by small protein conjugation or removal



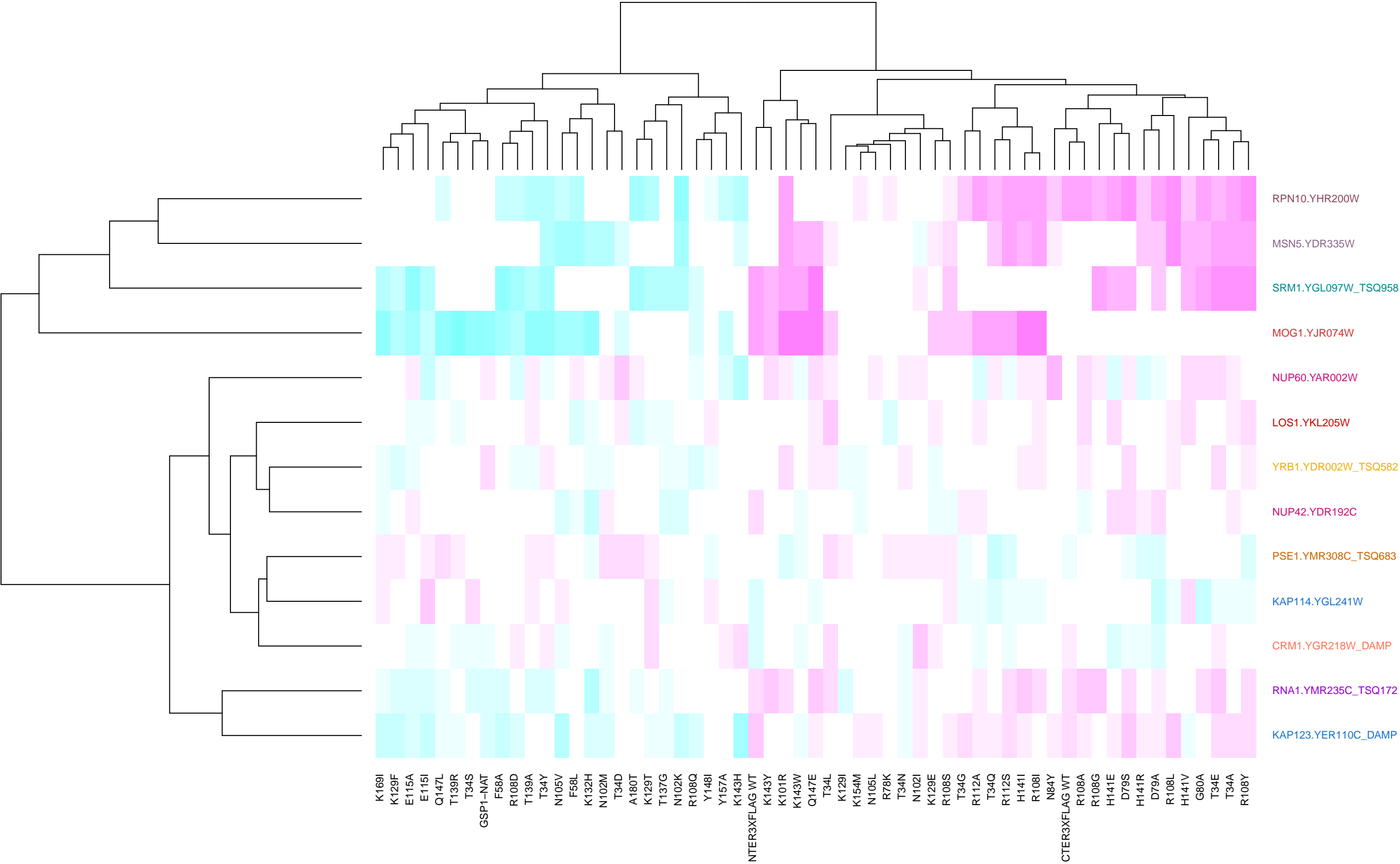
protein targeting



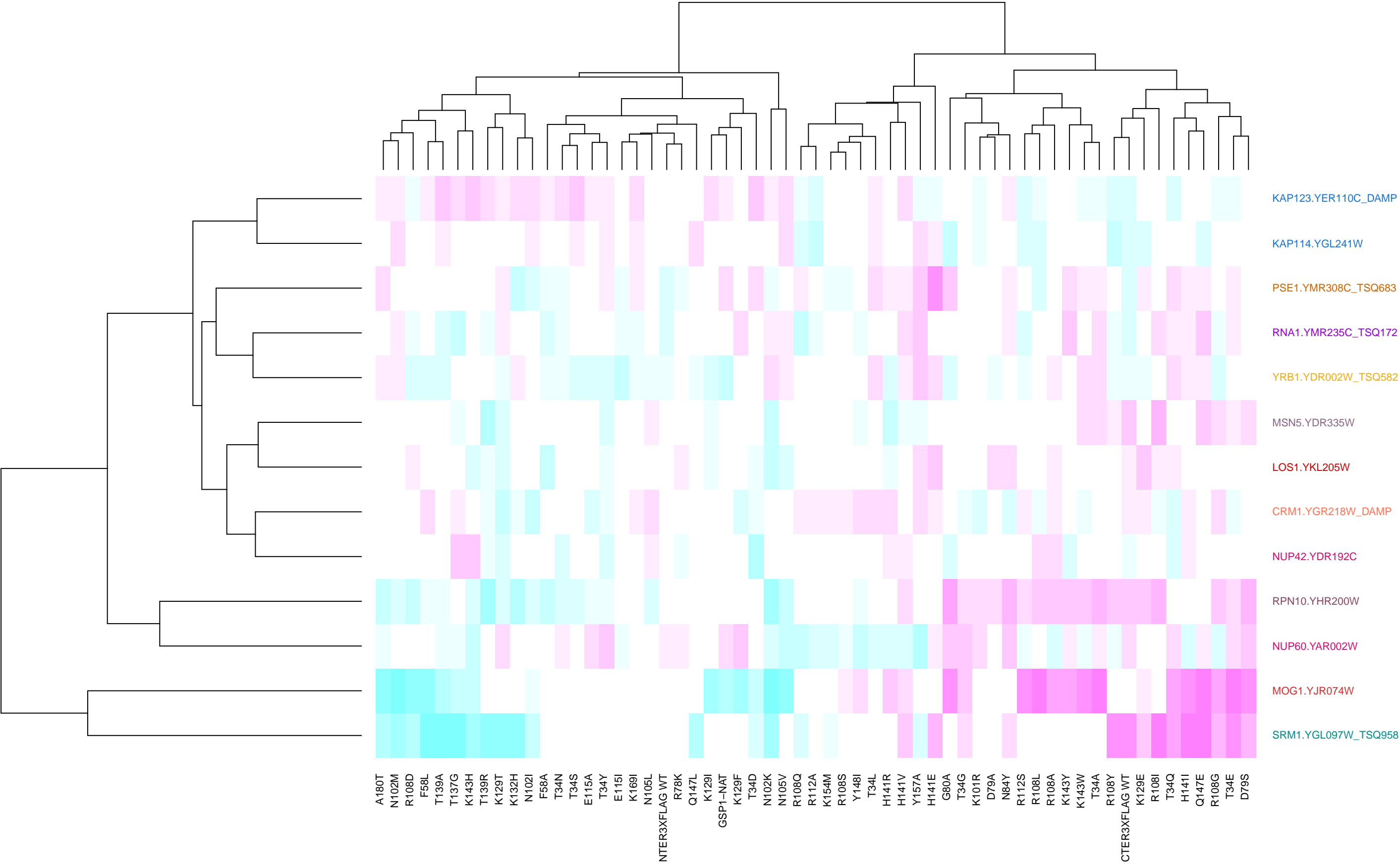
protein targeting_GO_1



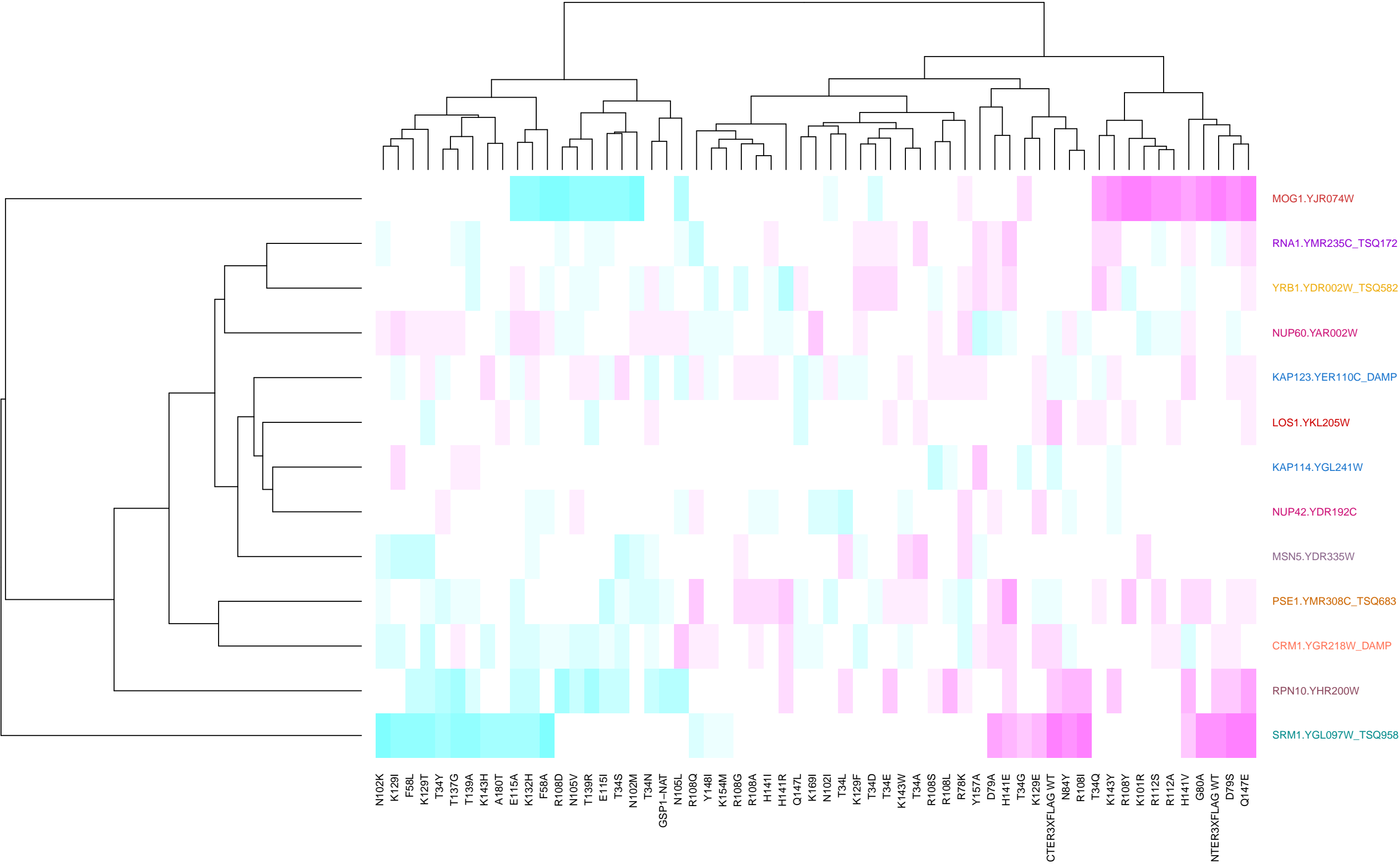
proteolysis involved in cellular protein catabolic process



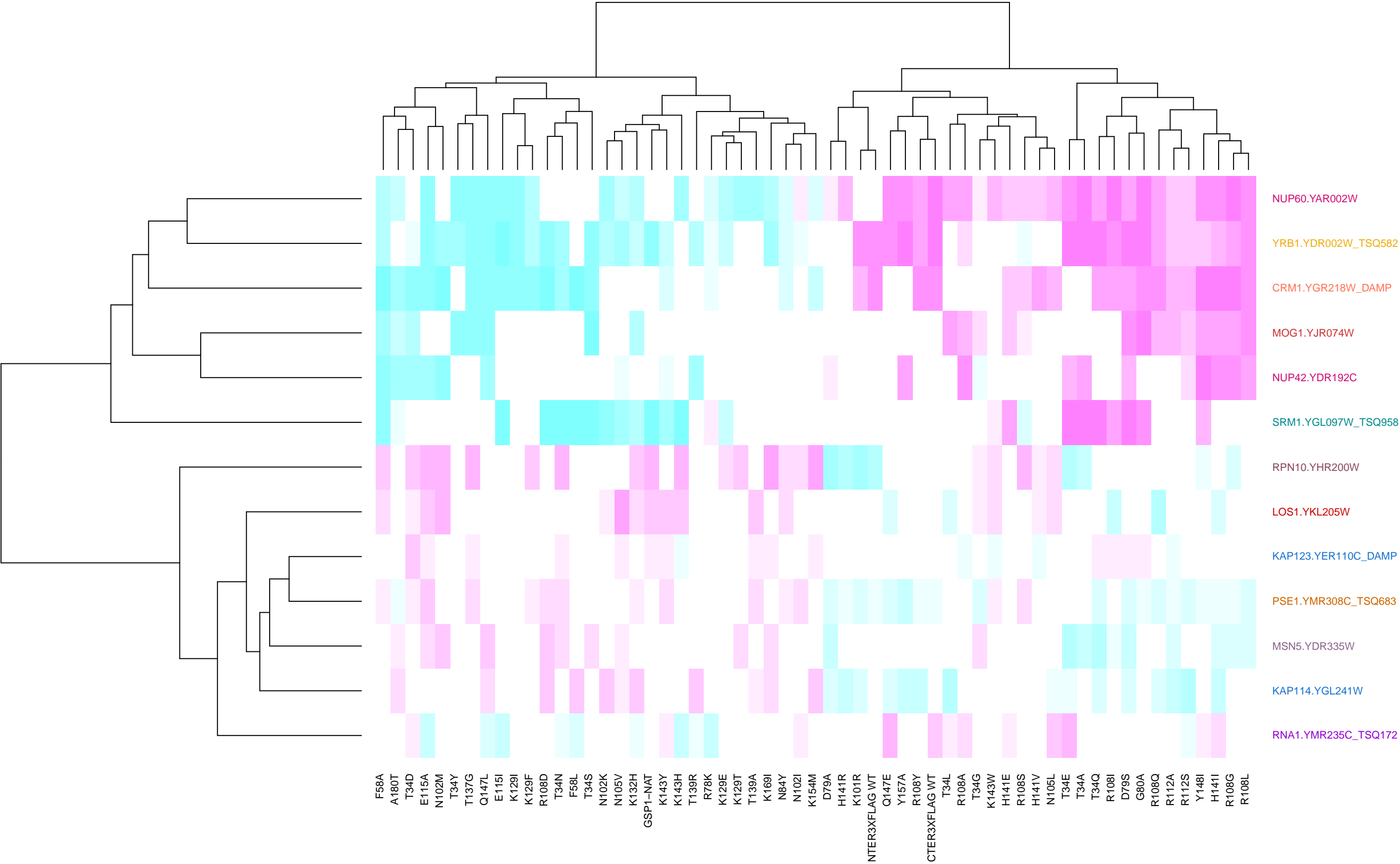
regulation of cell cycle



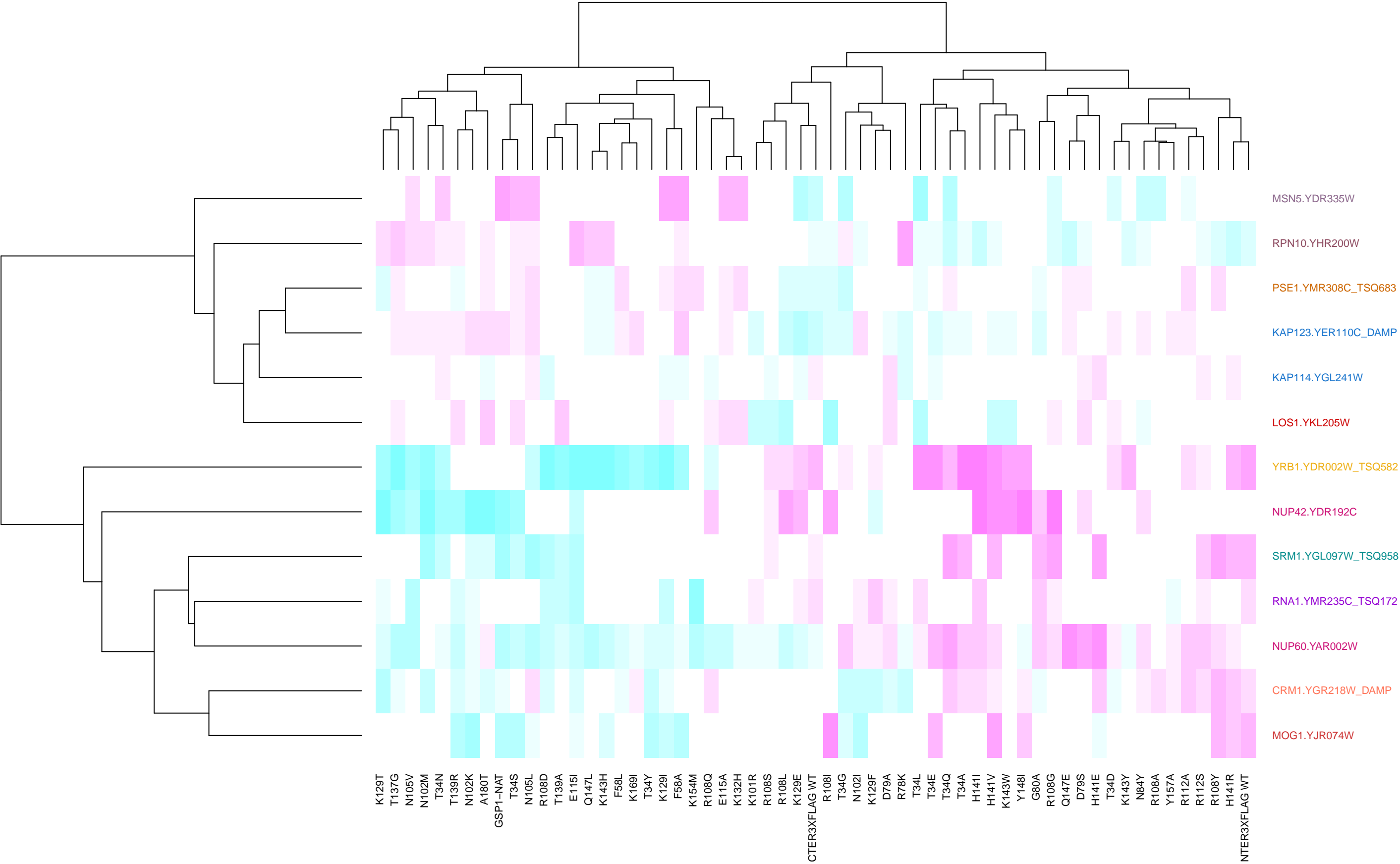
regulation of cell cycle_GO_1



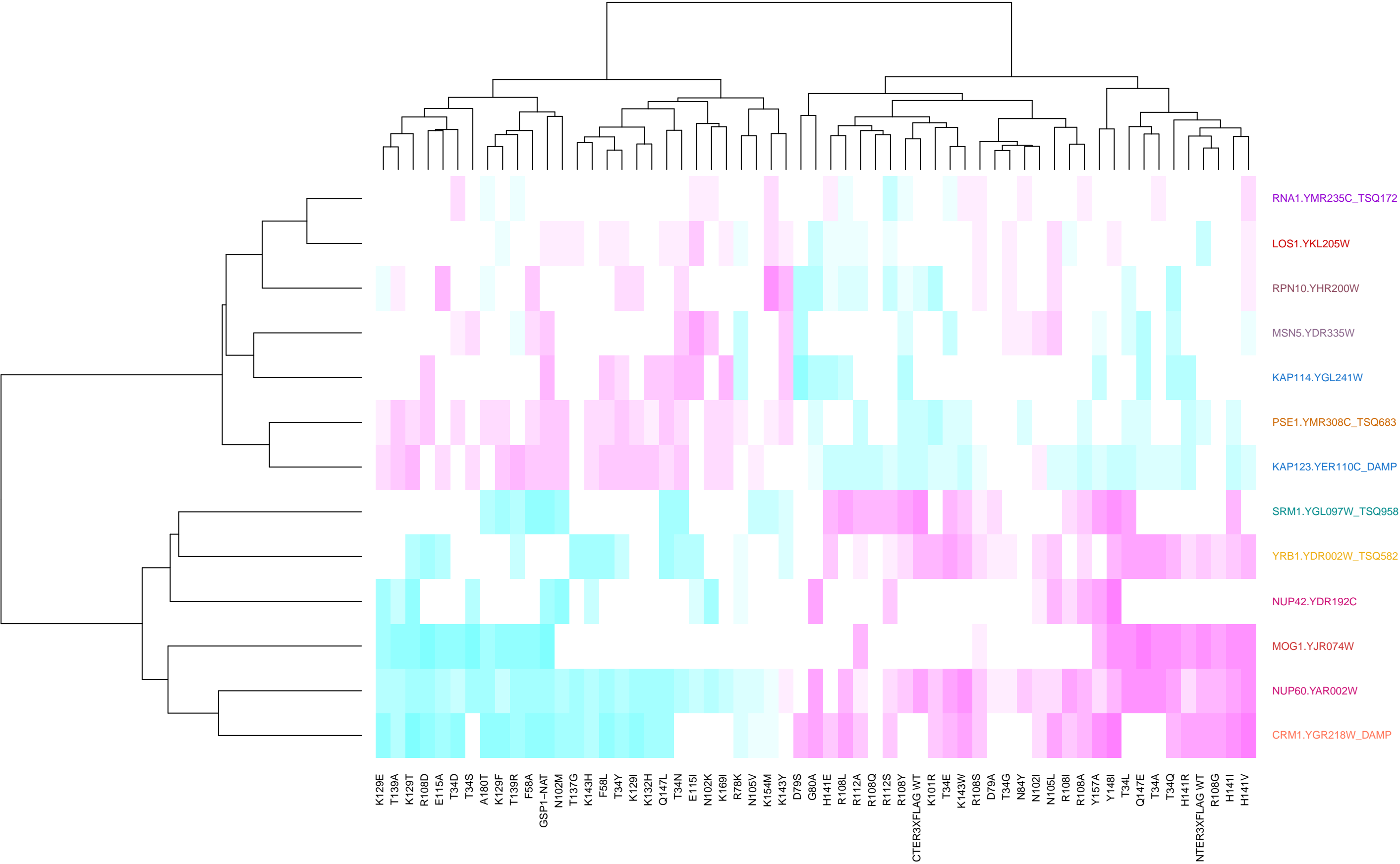
response to heat



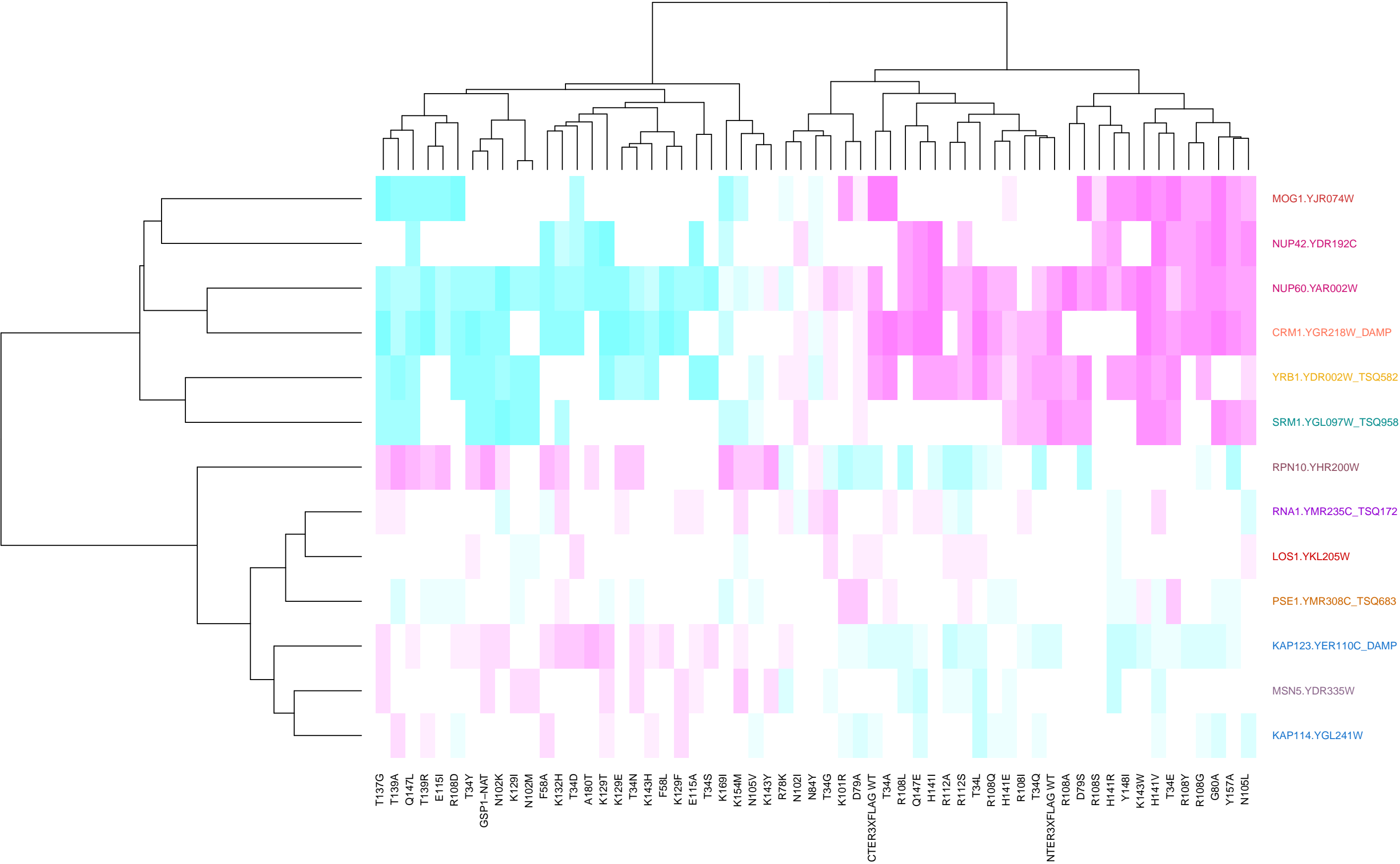
RNA catabolic process



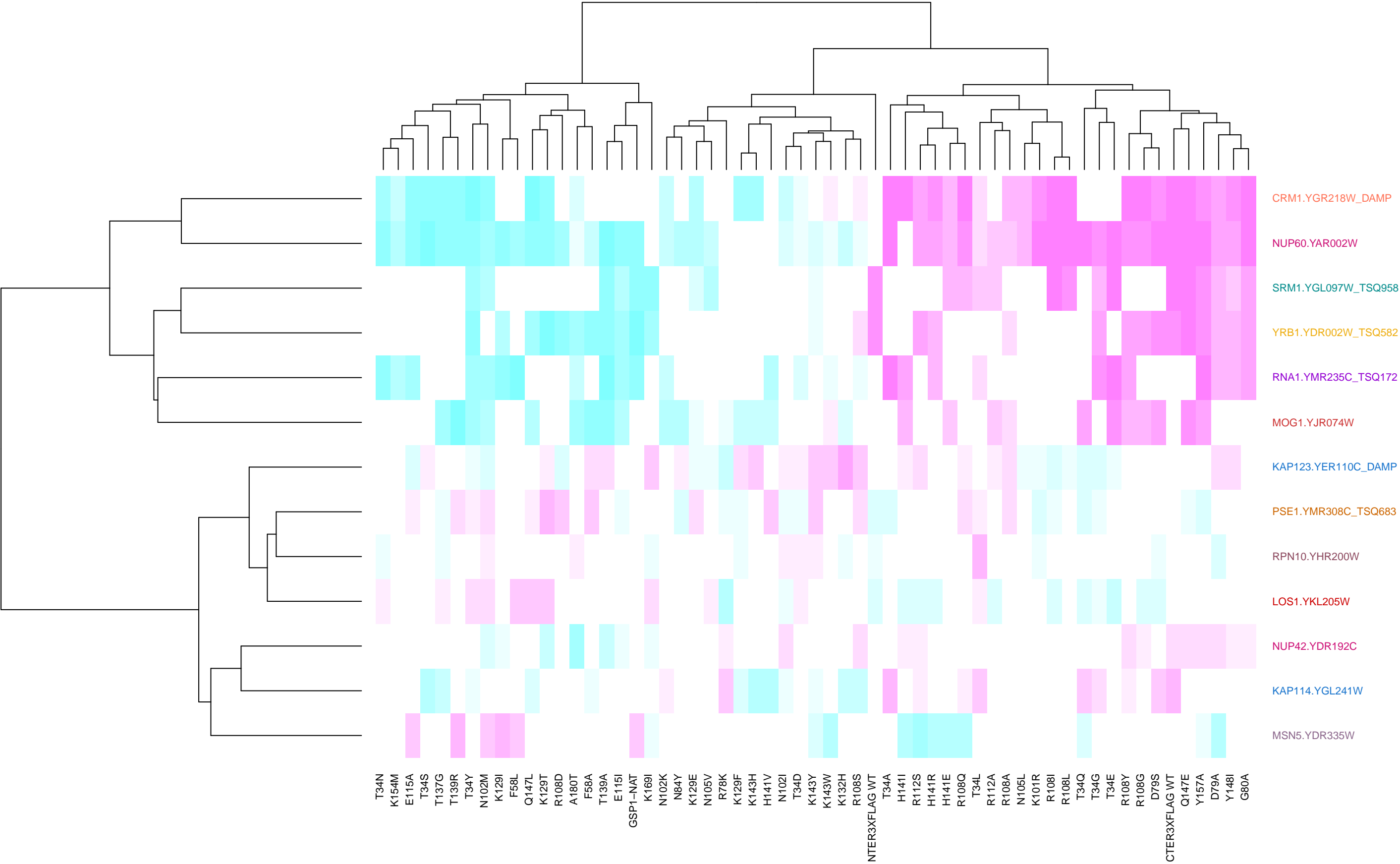
transcription from RNA polymerase II promoter_GO_2



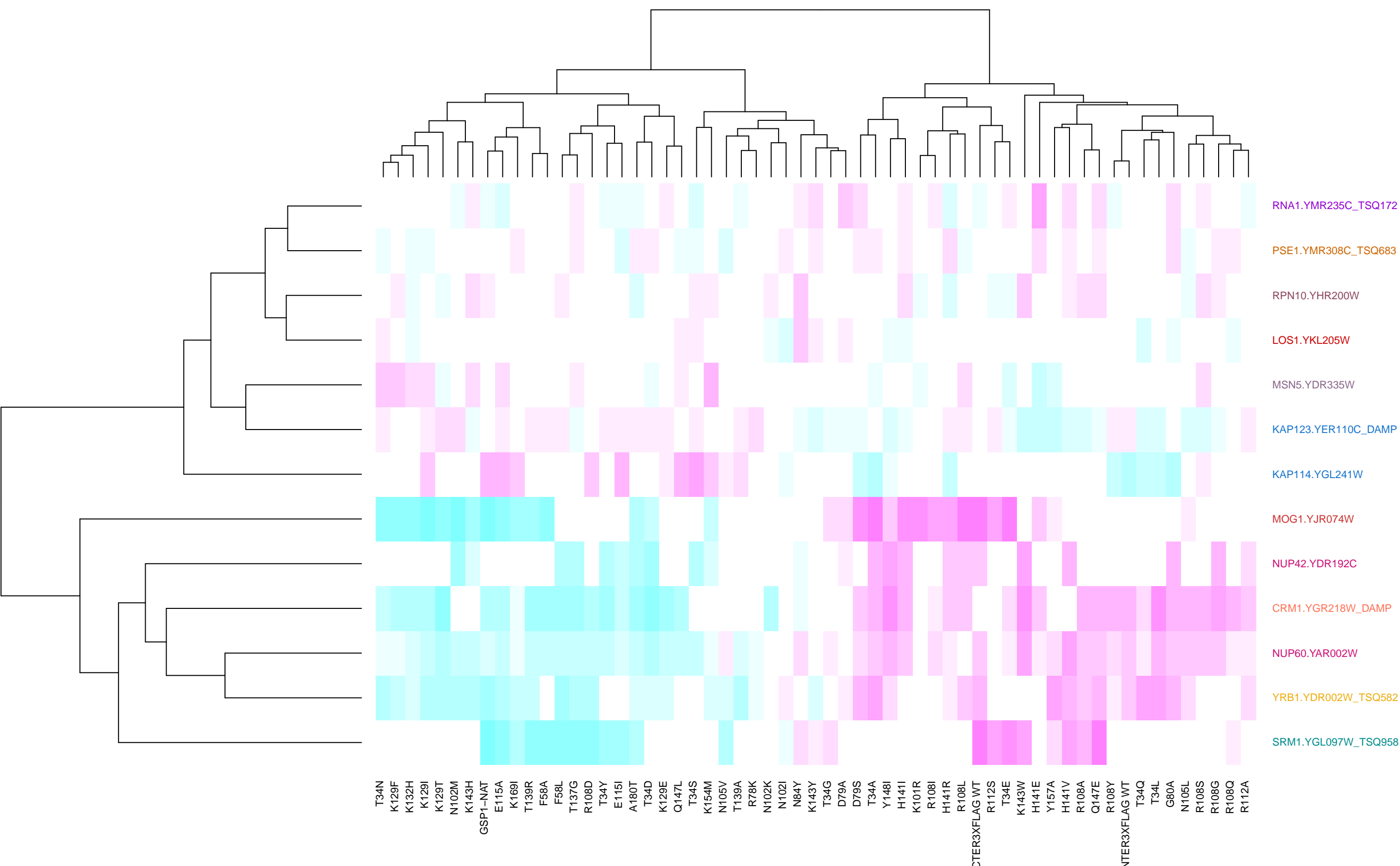
transcription_GO_3



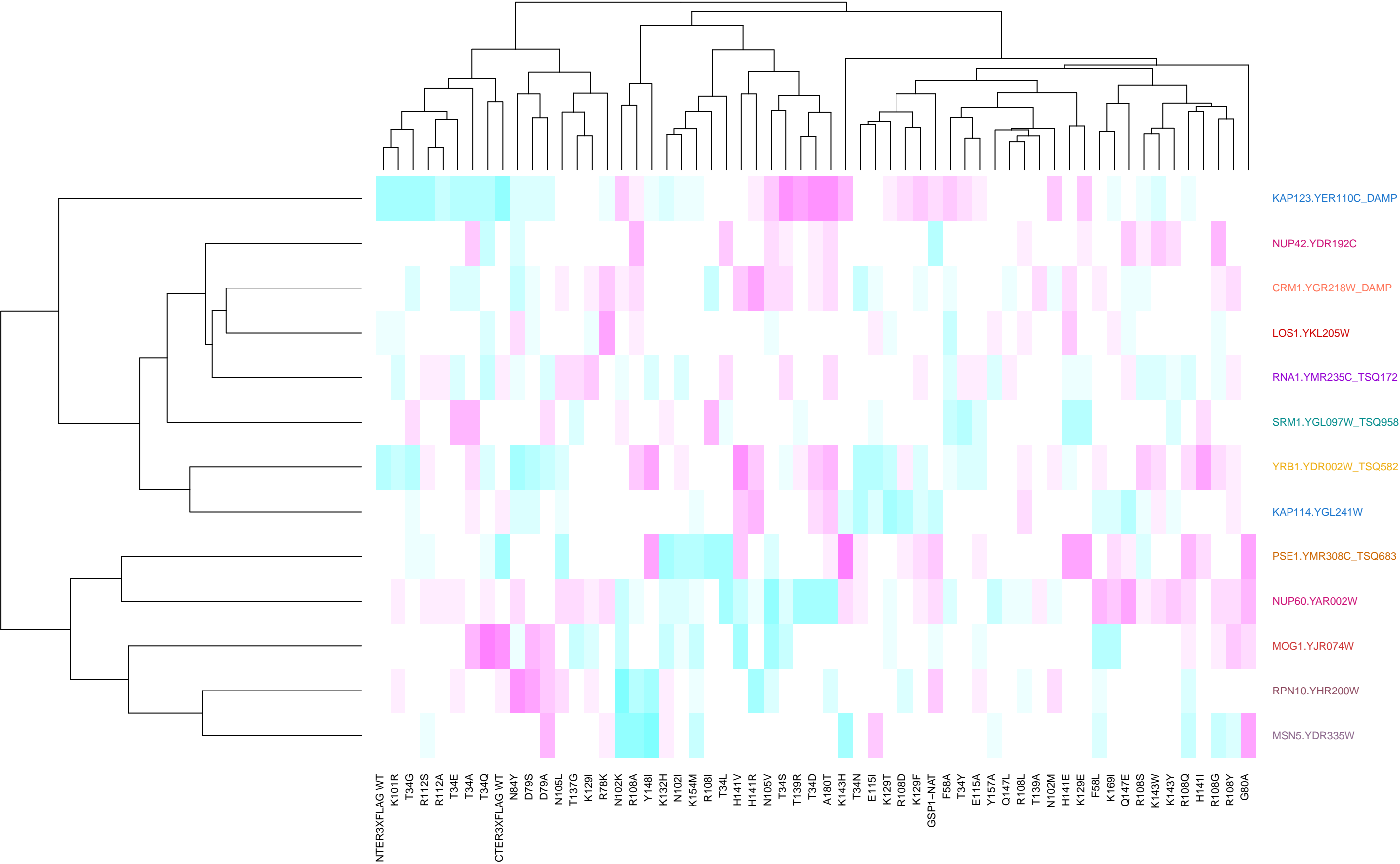
whole_library_22



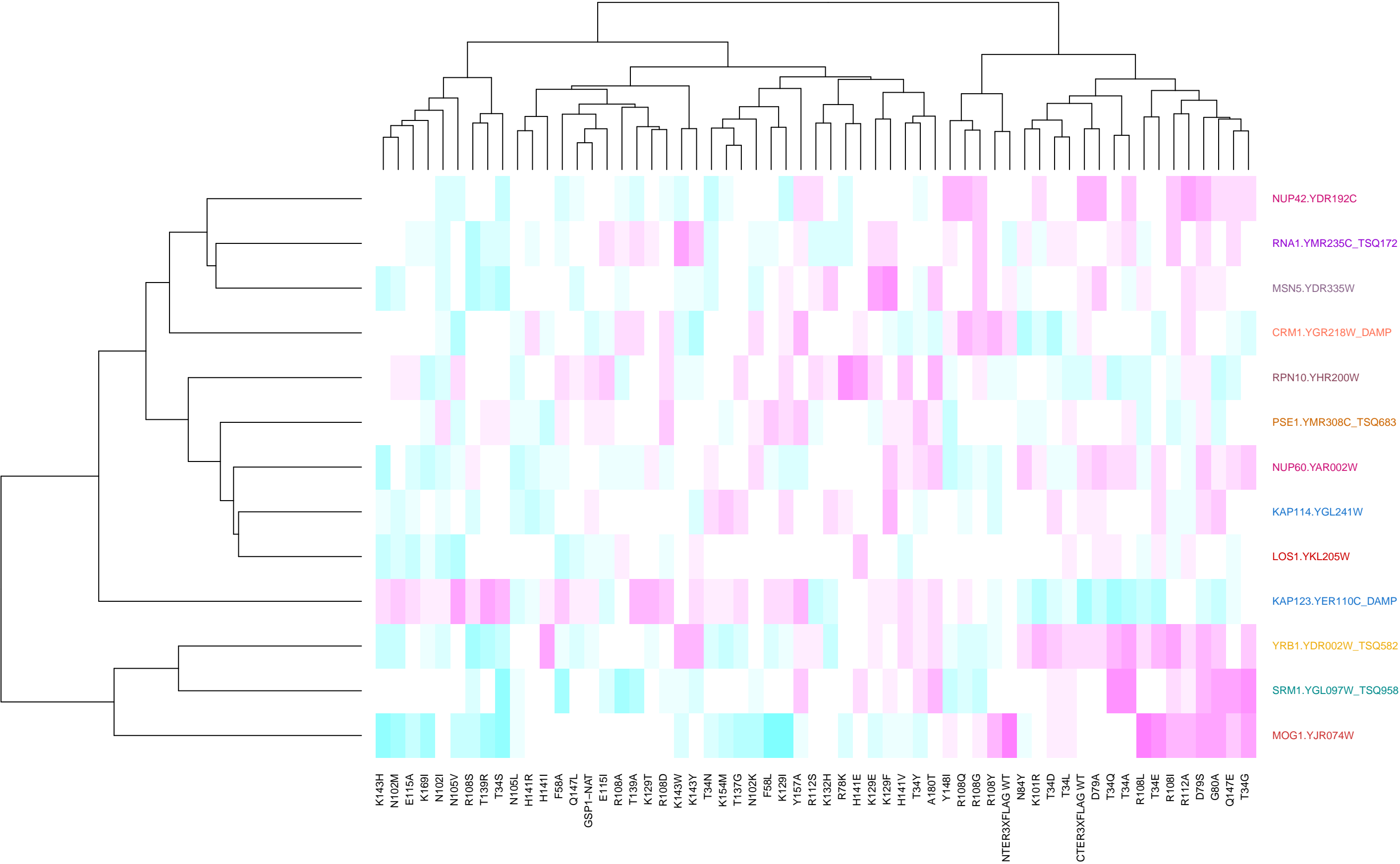
chromatin_GO_1



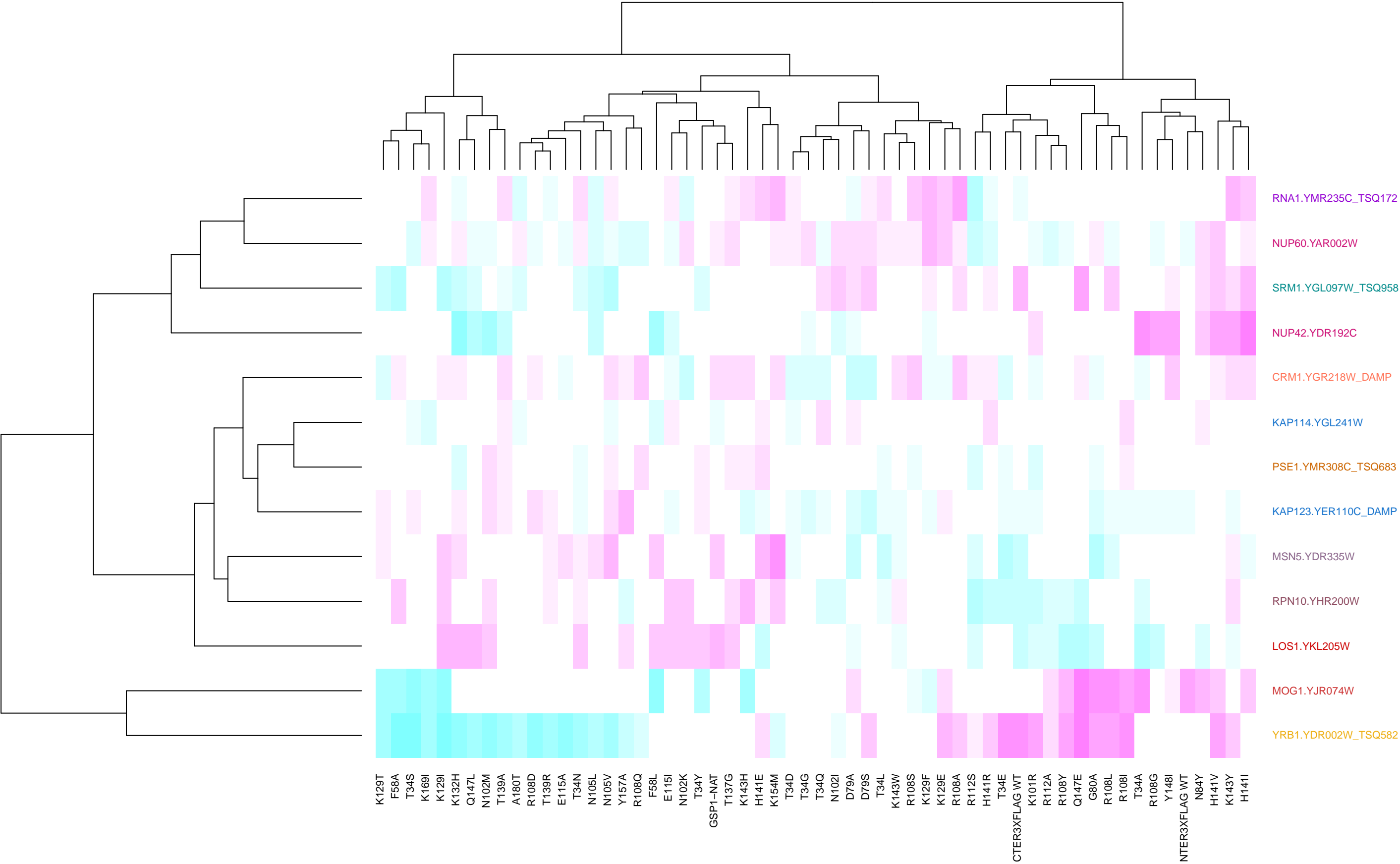
chromosome segregation_GO_1



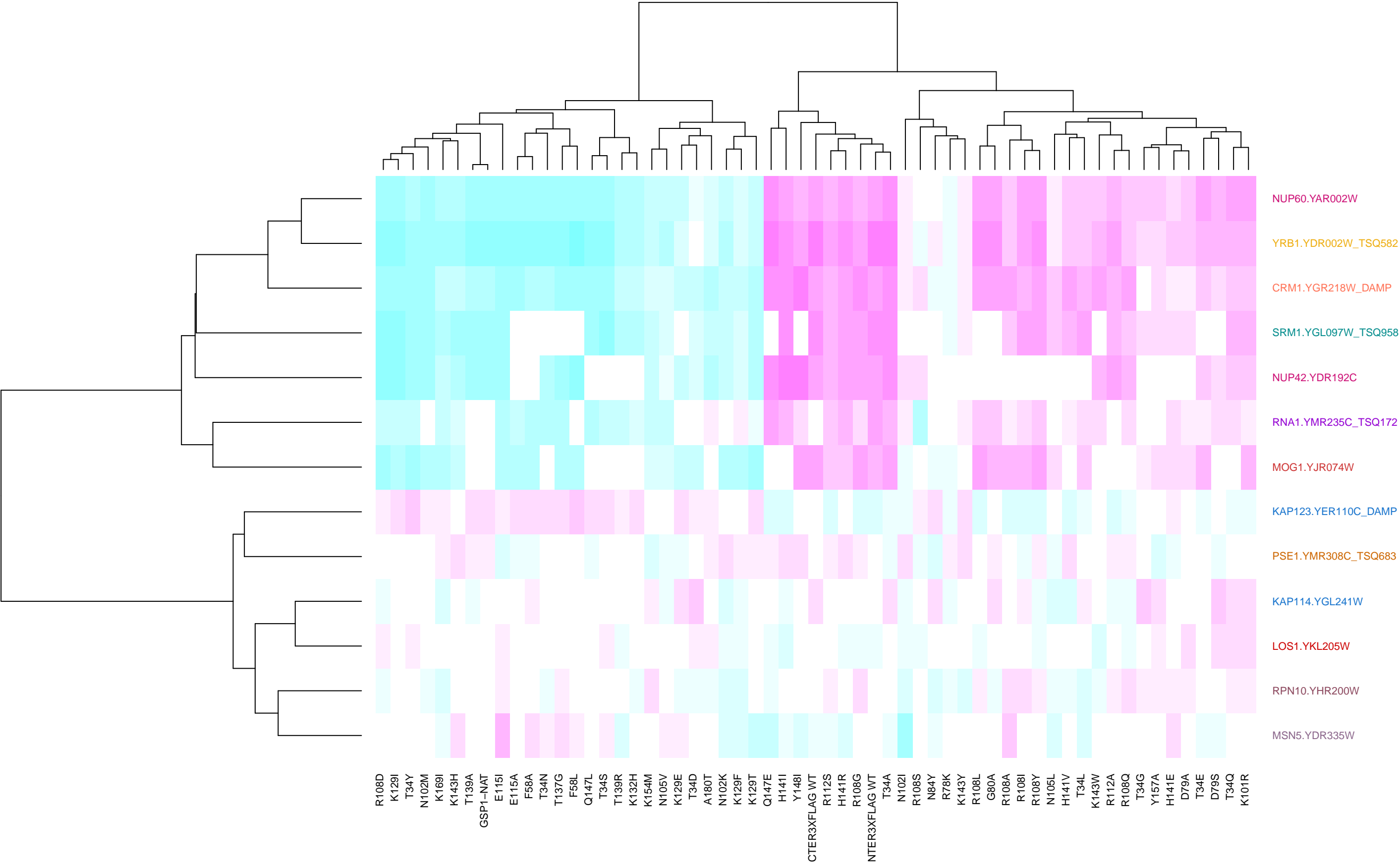
DNA binding



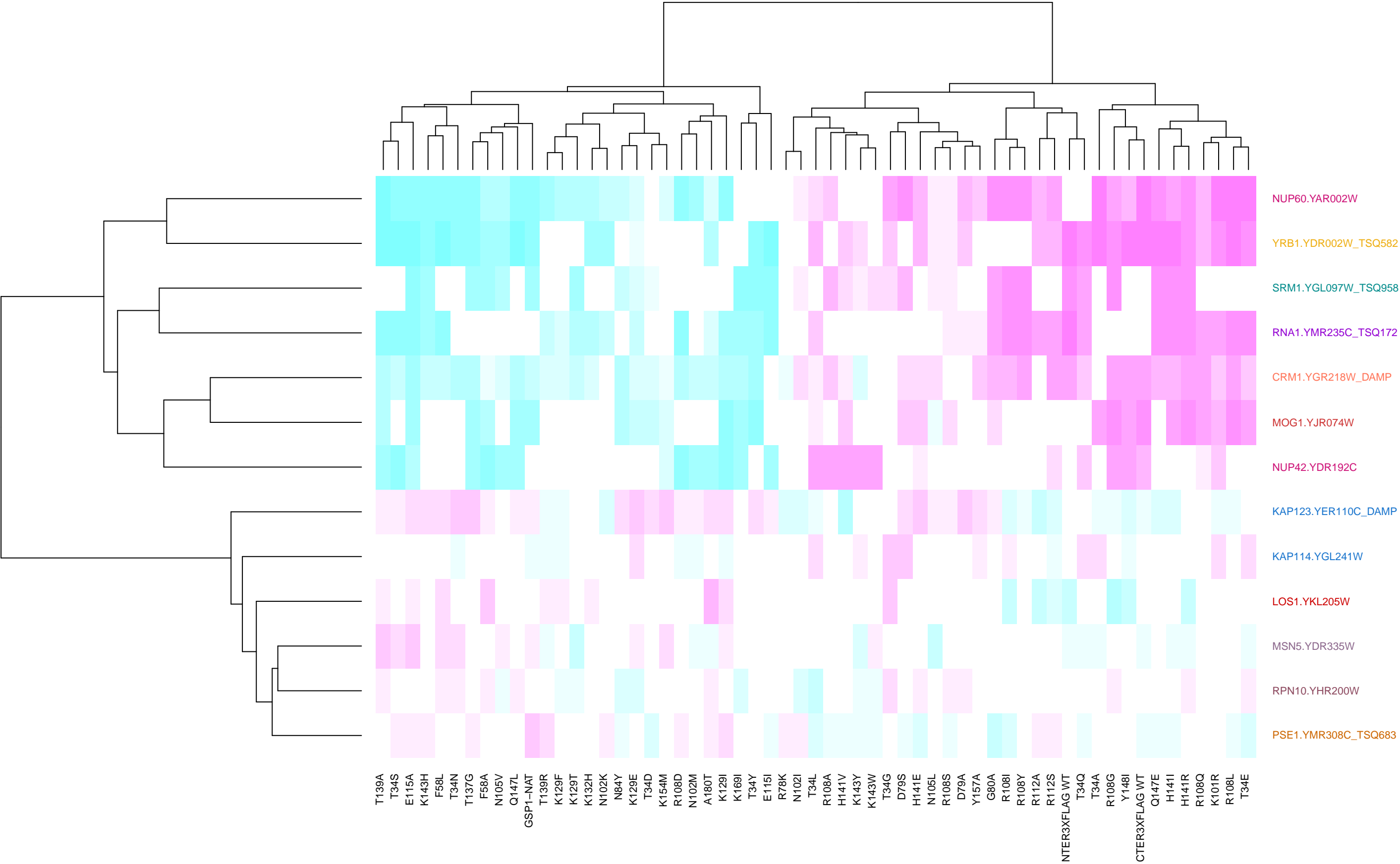
DNA repair_GO_1



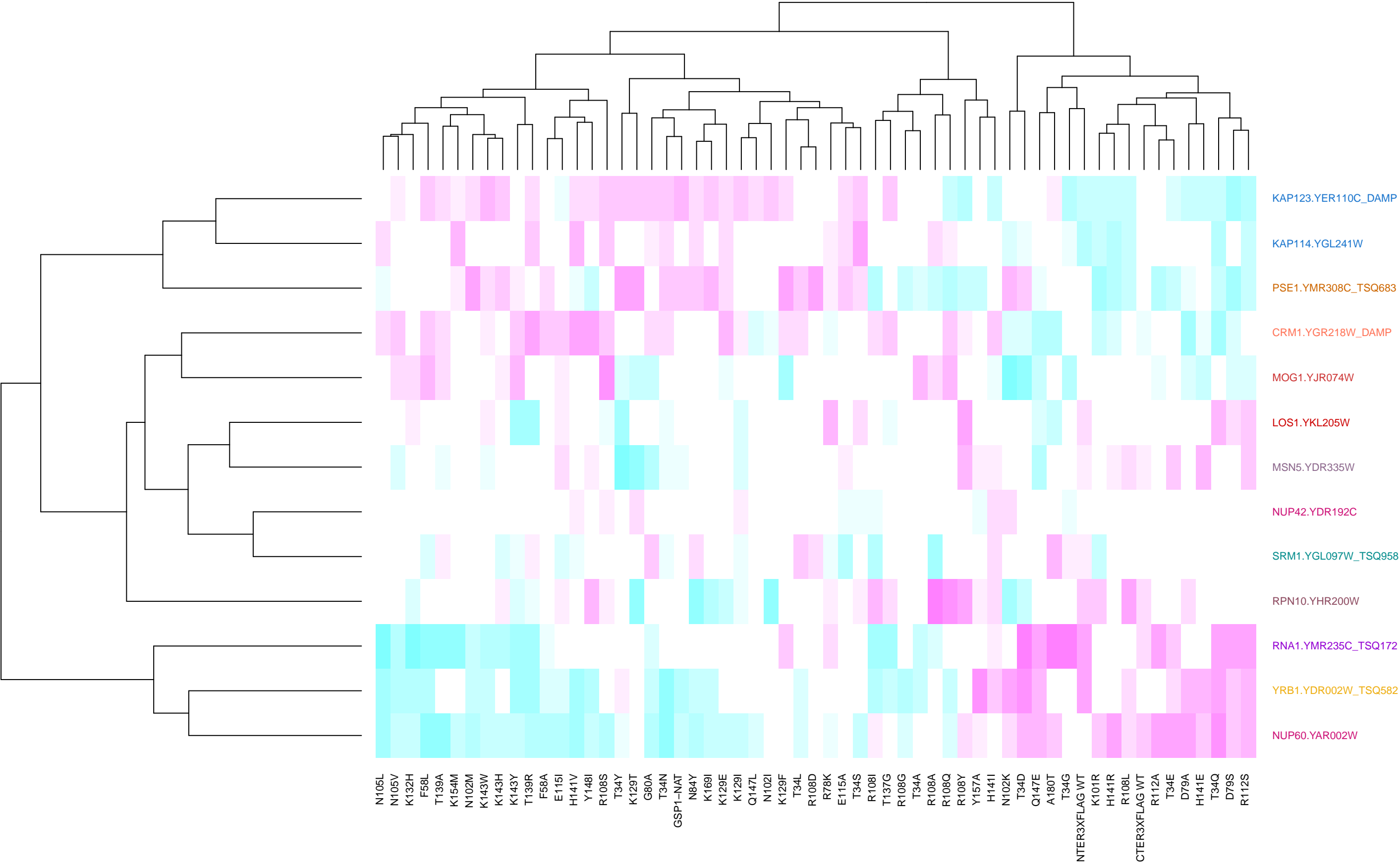
endomembrane system



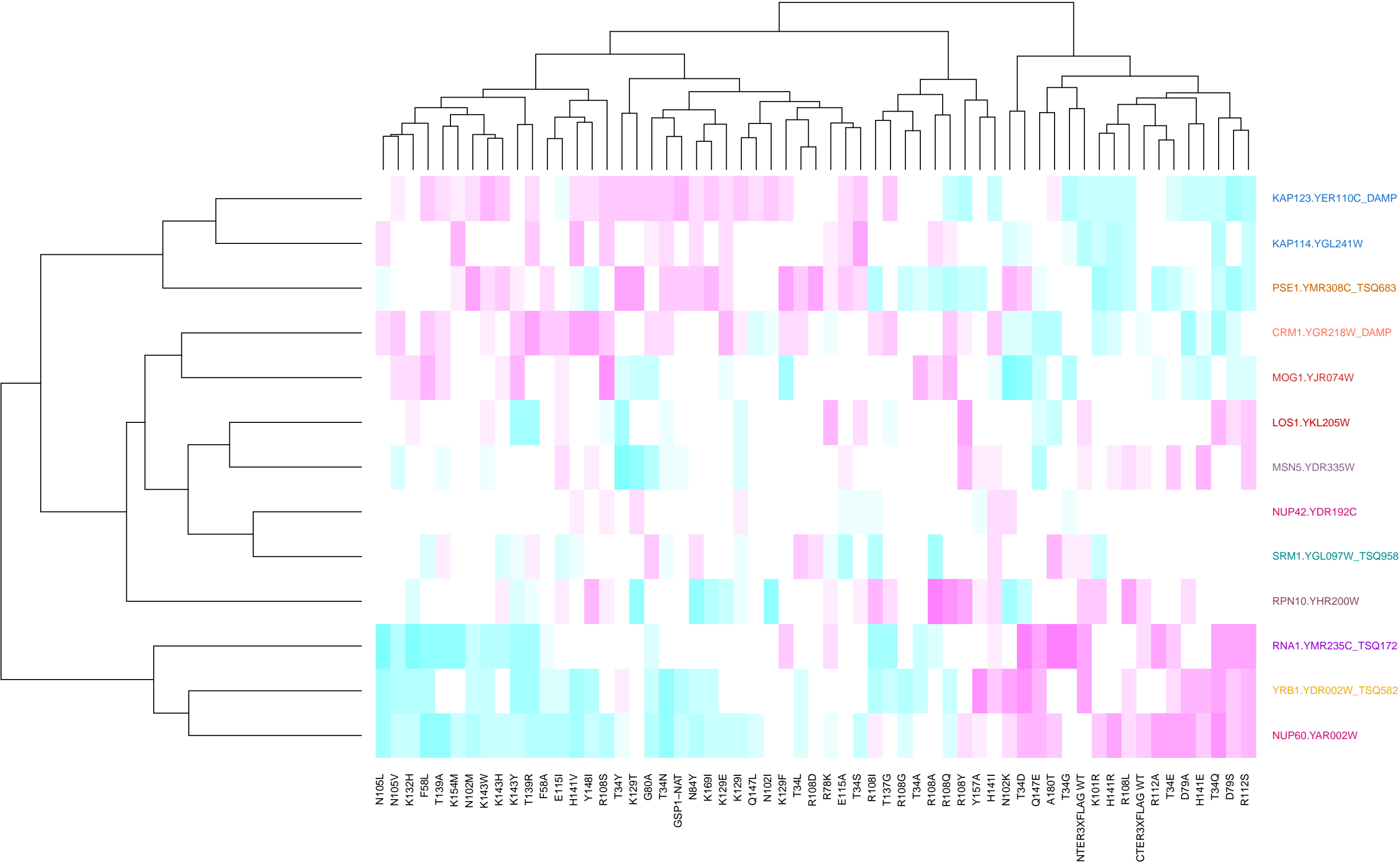
endomembrane system_GO_1



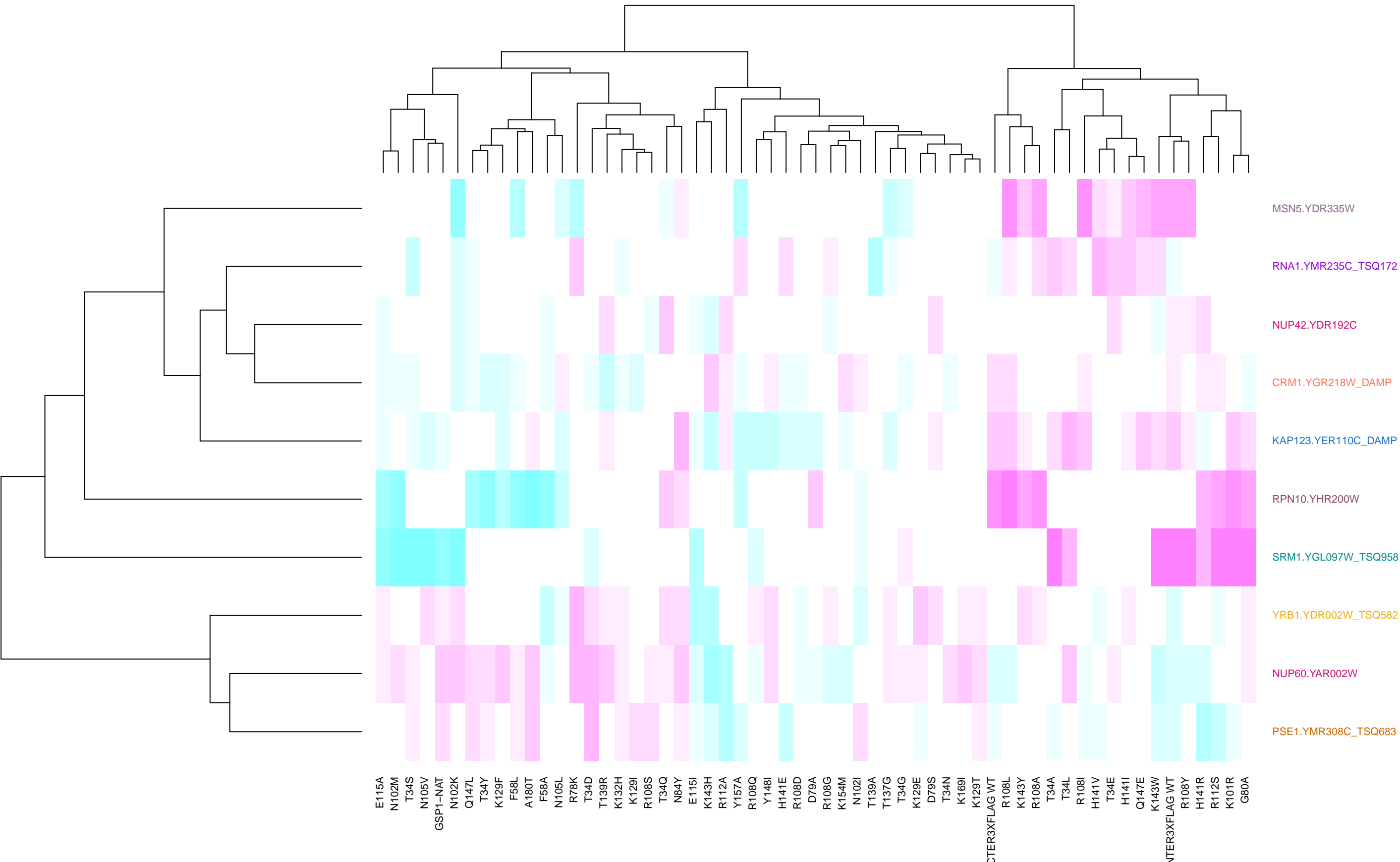
endoplasmic reticulum_GO_2



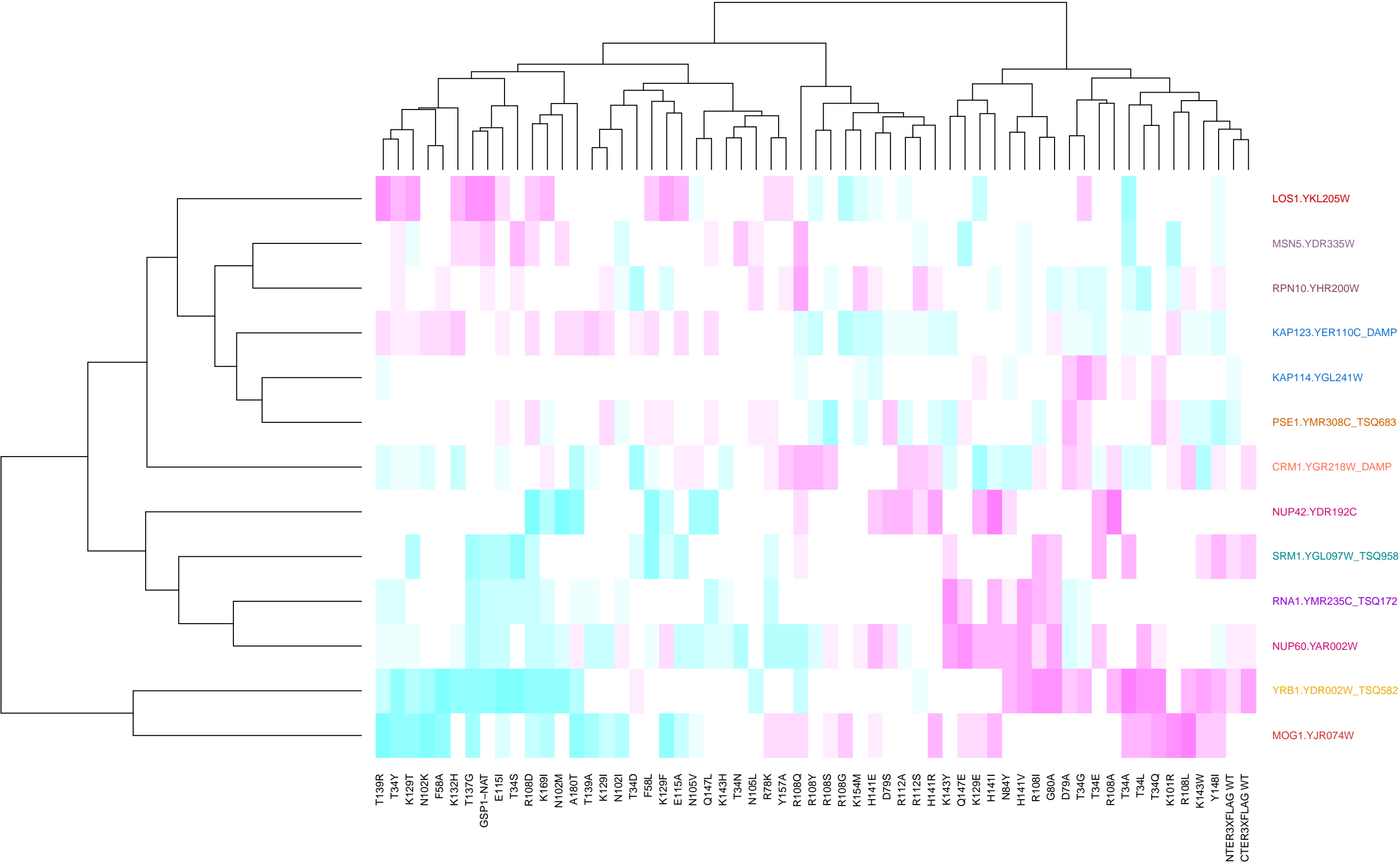
ER_GO_2



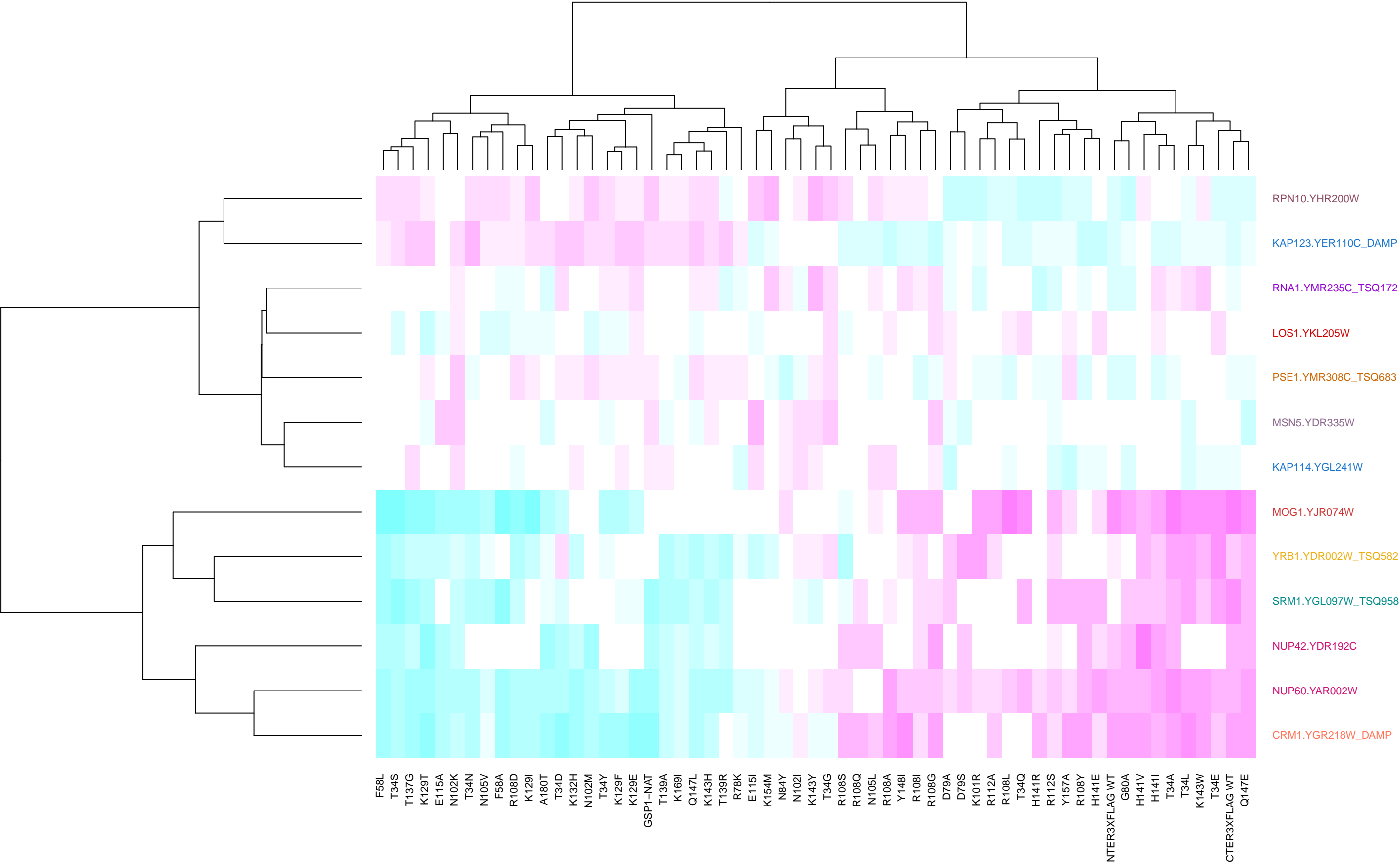
protein modification by small protein conjugation or removal_GO_2



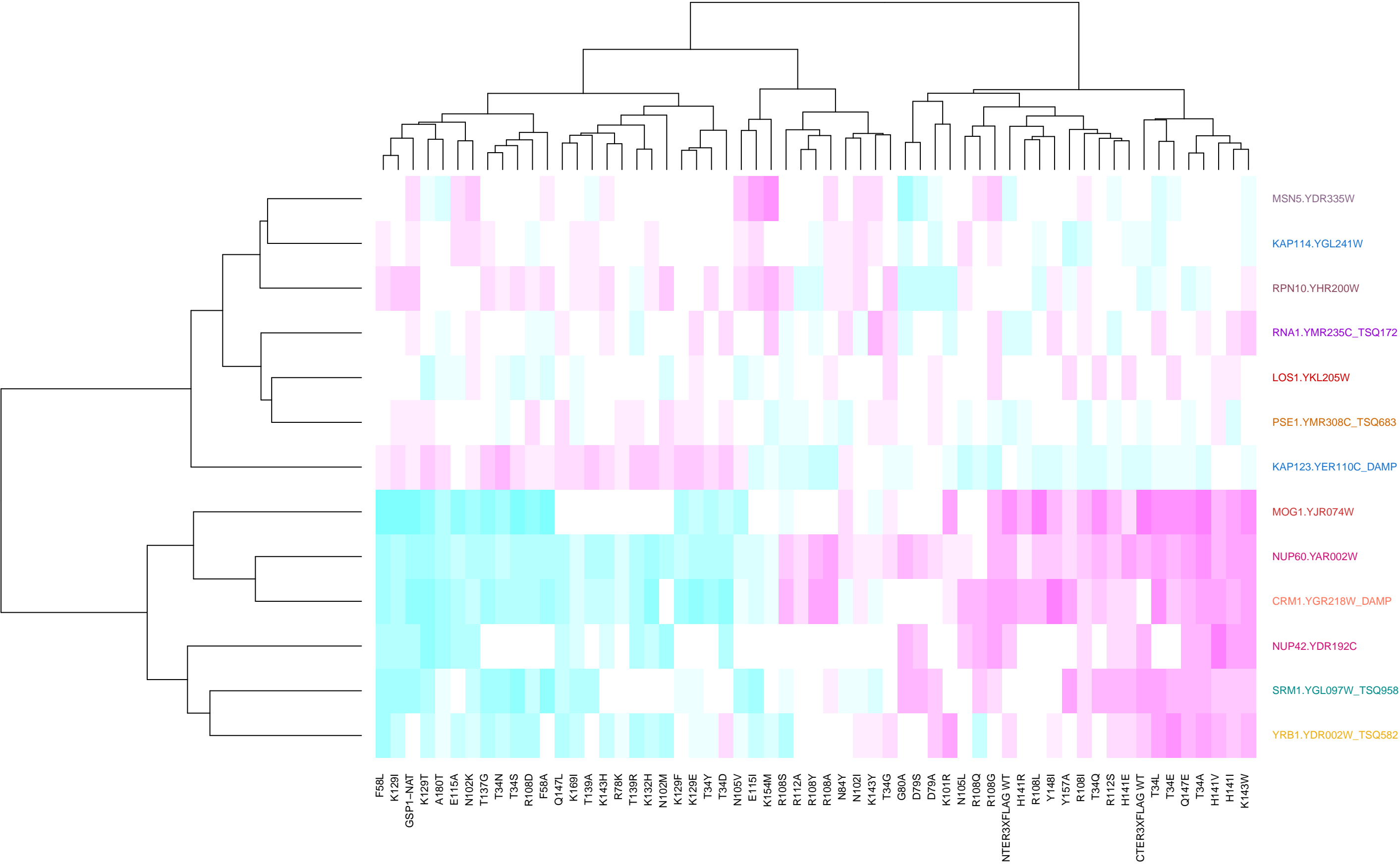
RNA processing



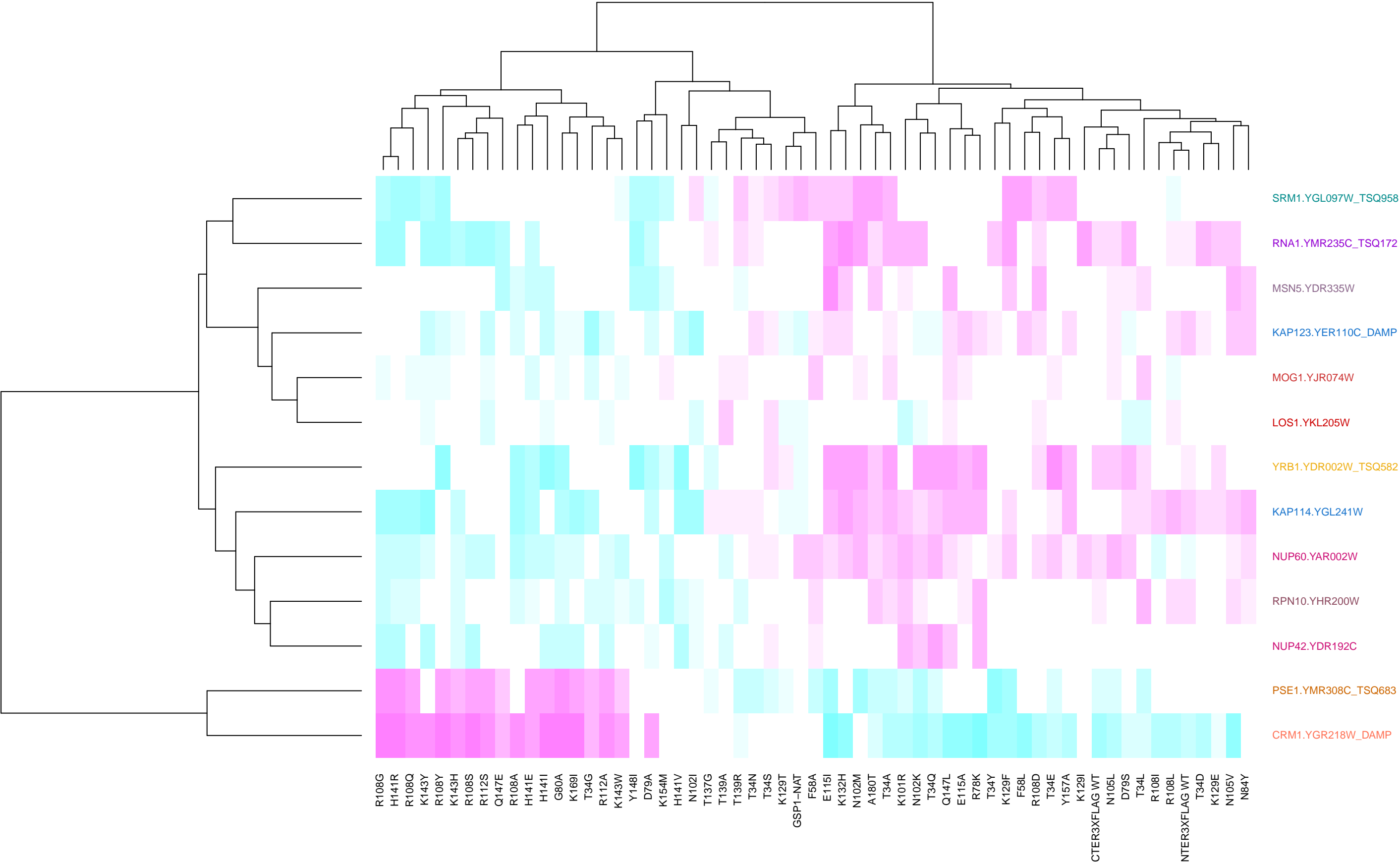
transcription



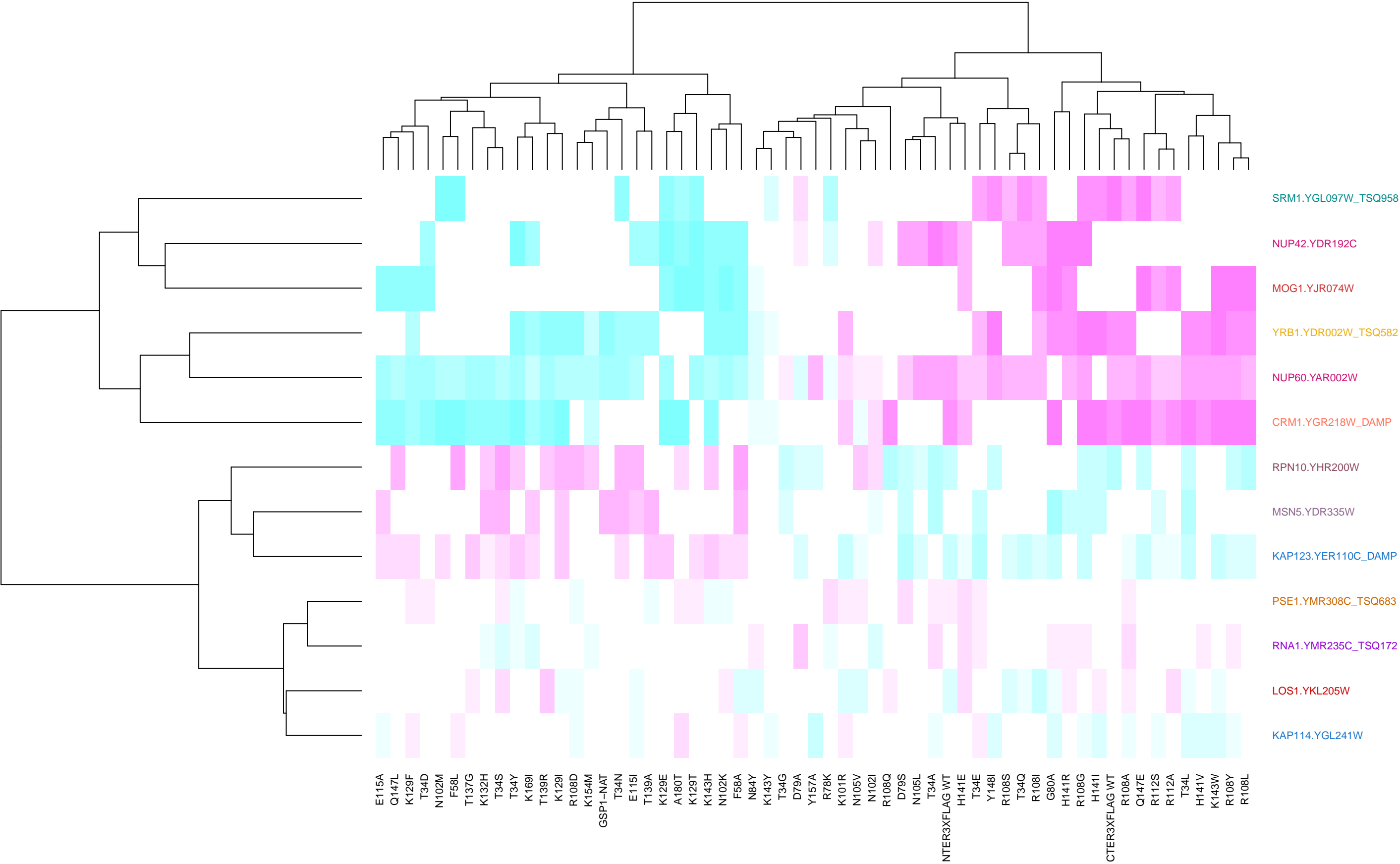
transcription from RNA polymerase II promoter



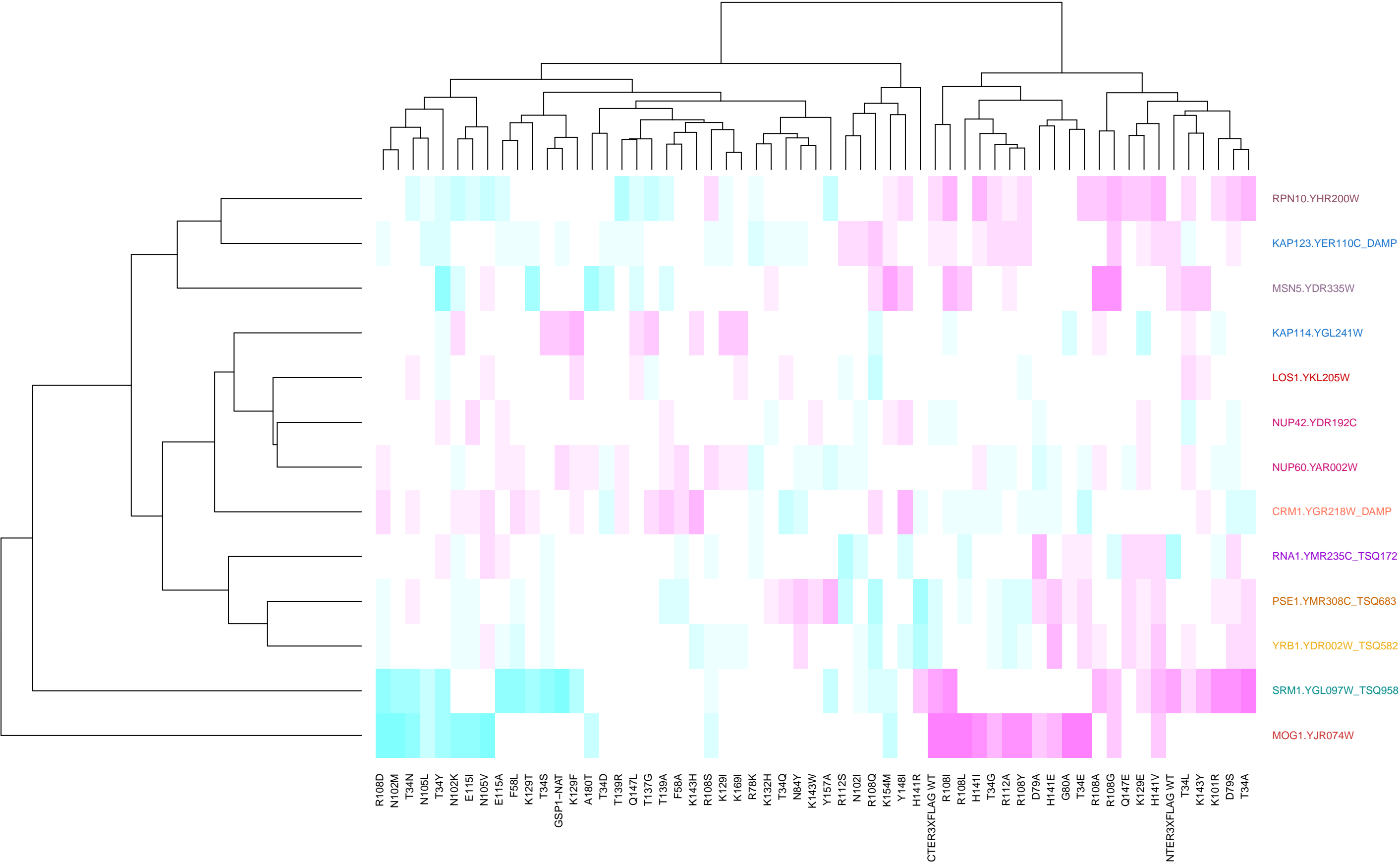
cell cortex_GO_1



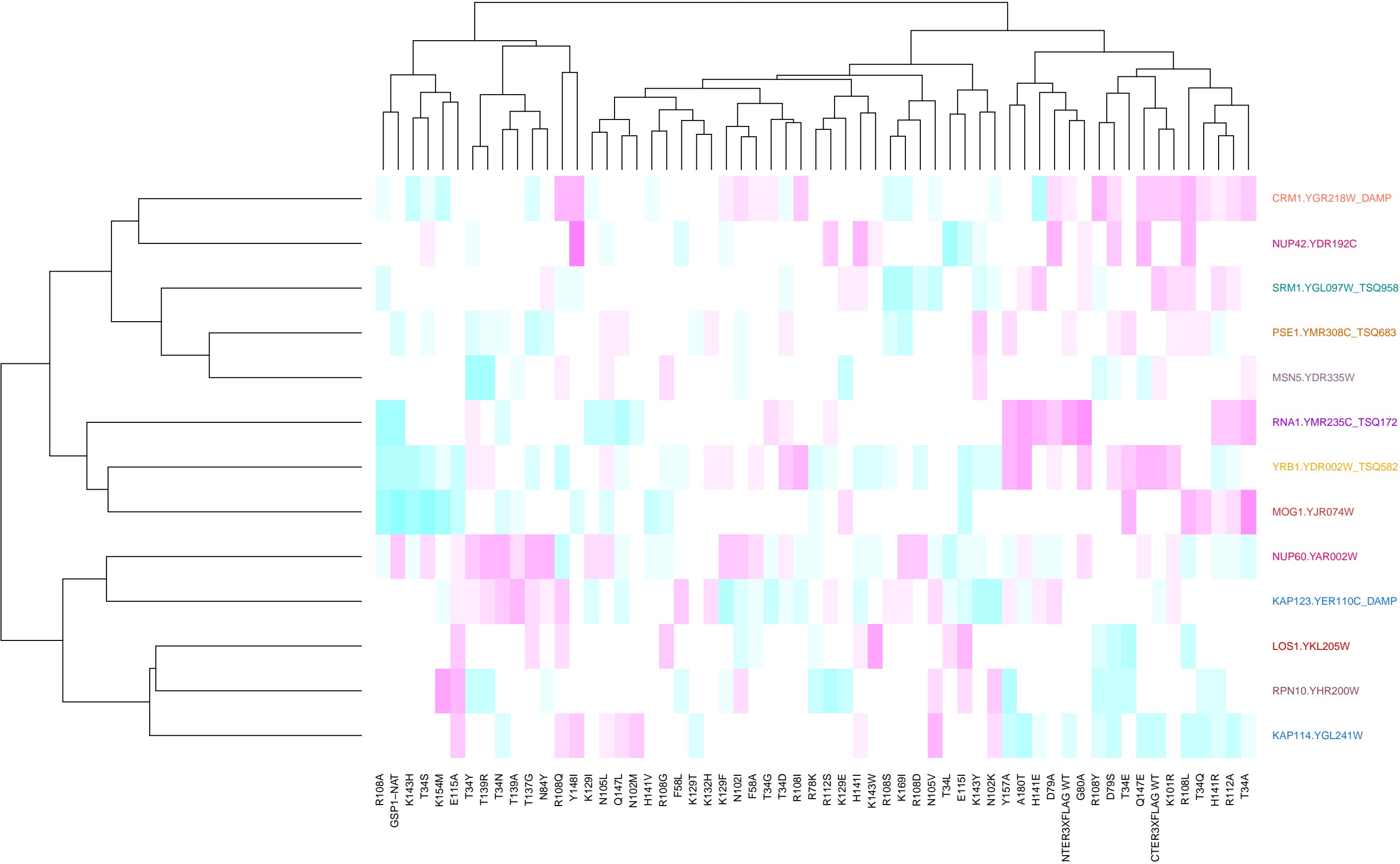
DNA repair_GO_2



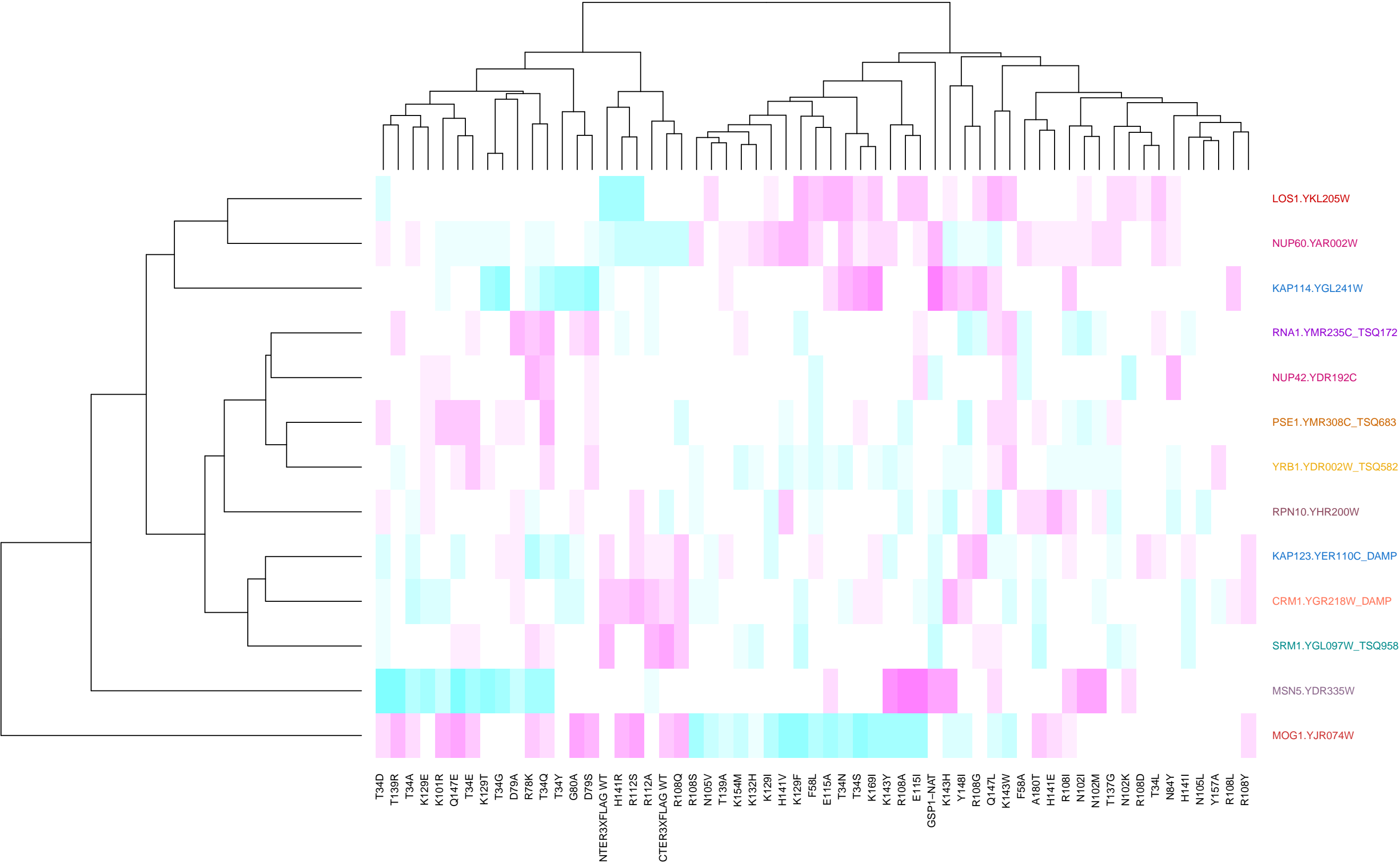
histone modification_GO_1



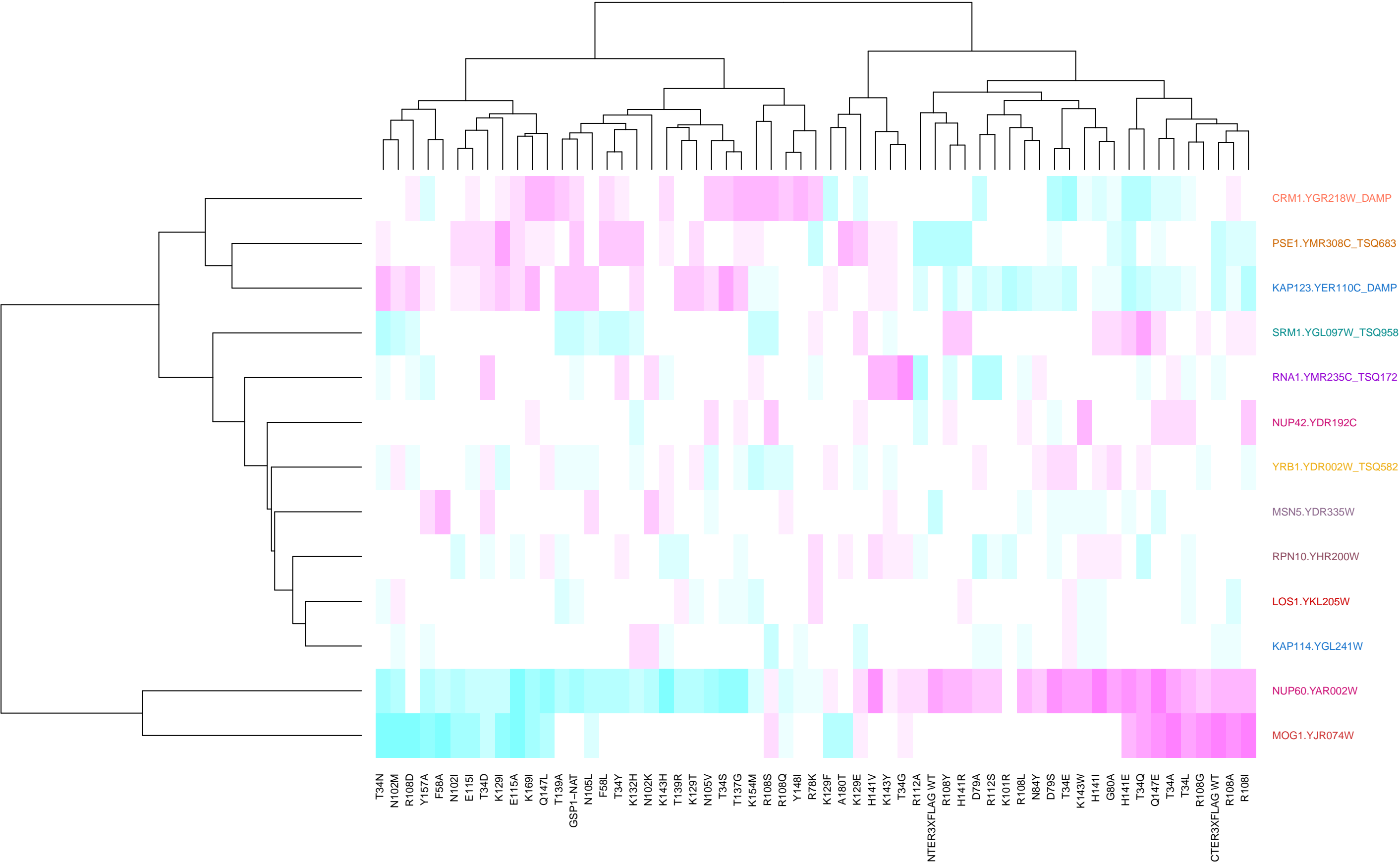
lipids_GO_3



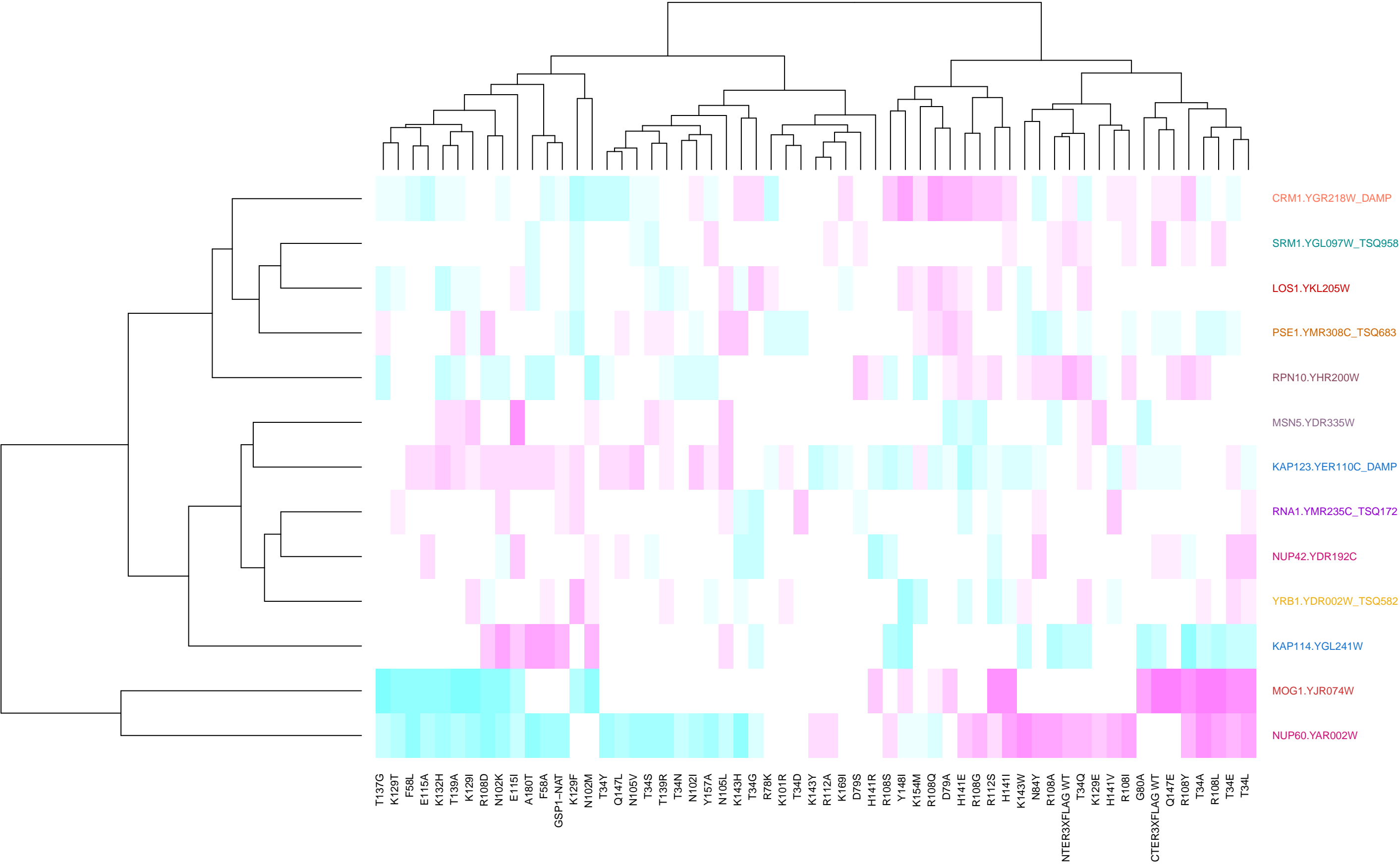
peptidyl-amino acid modification_GO_2



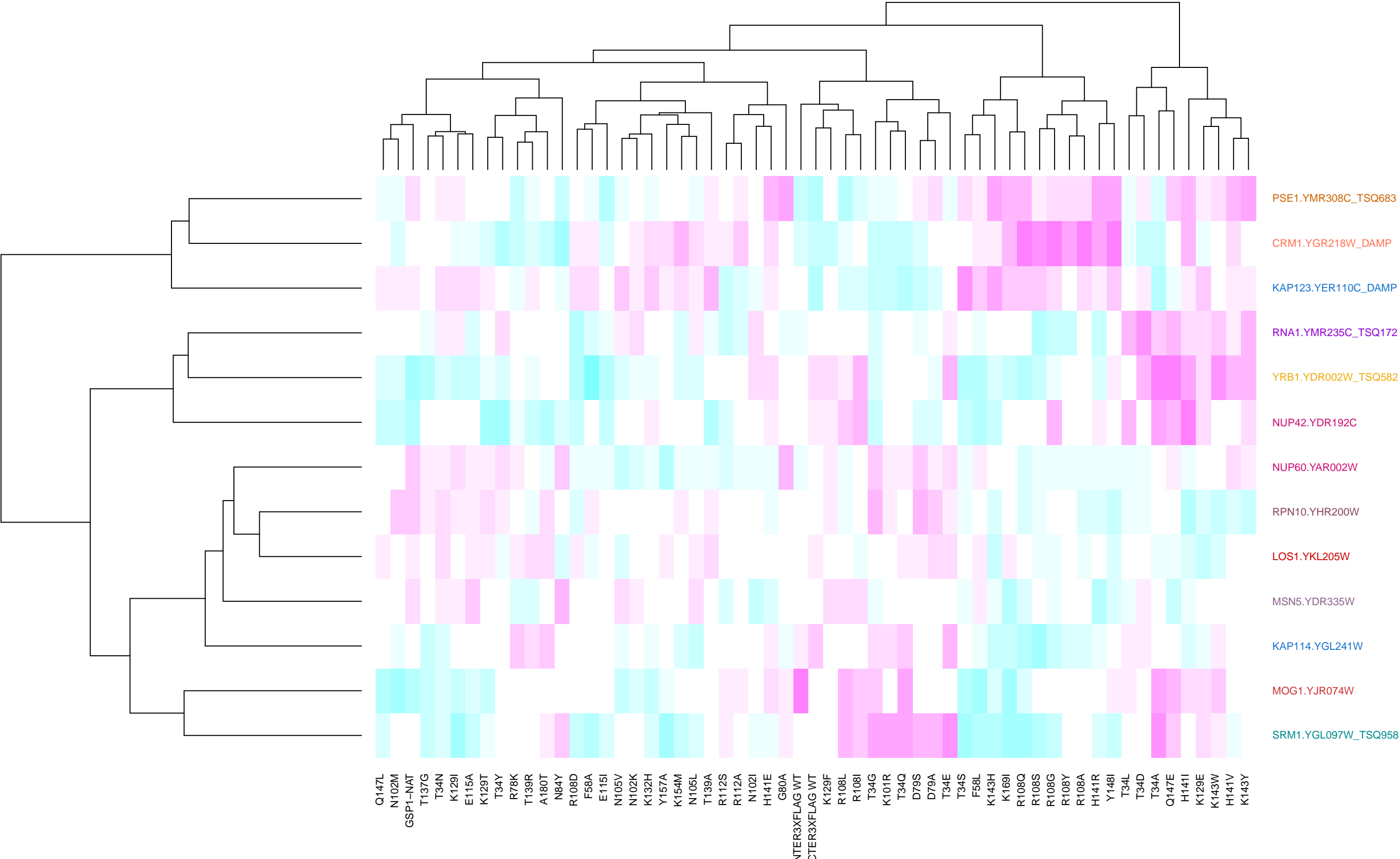
transcription and mRNA processing_GO_5



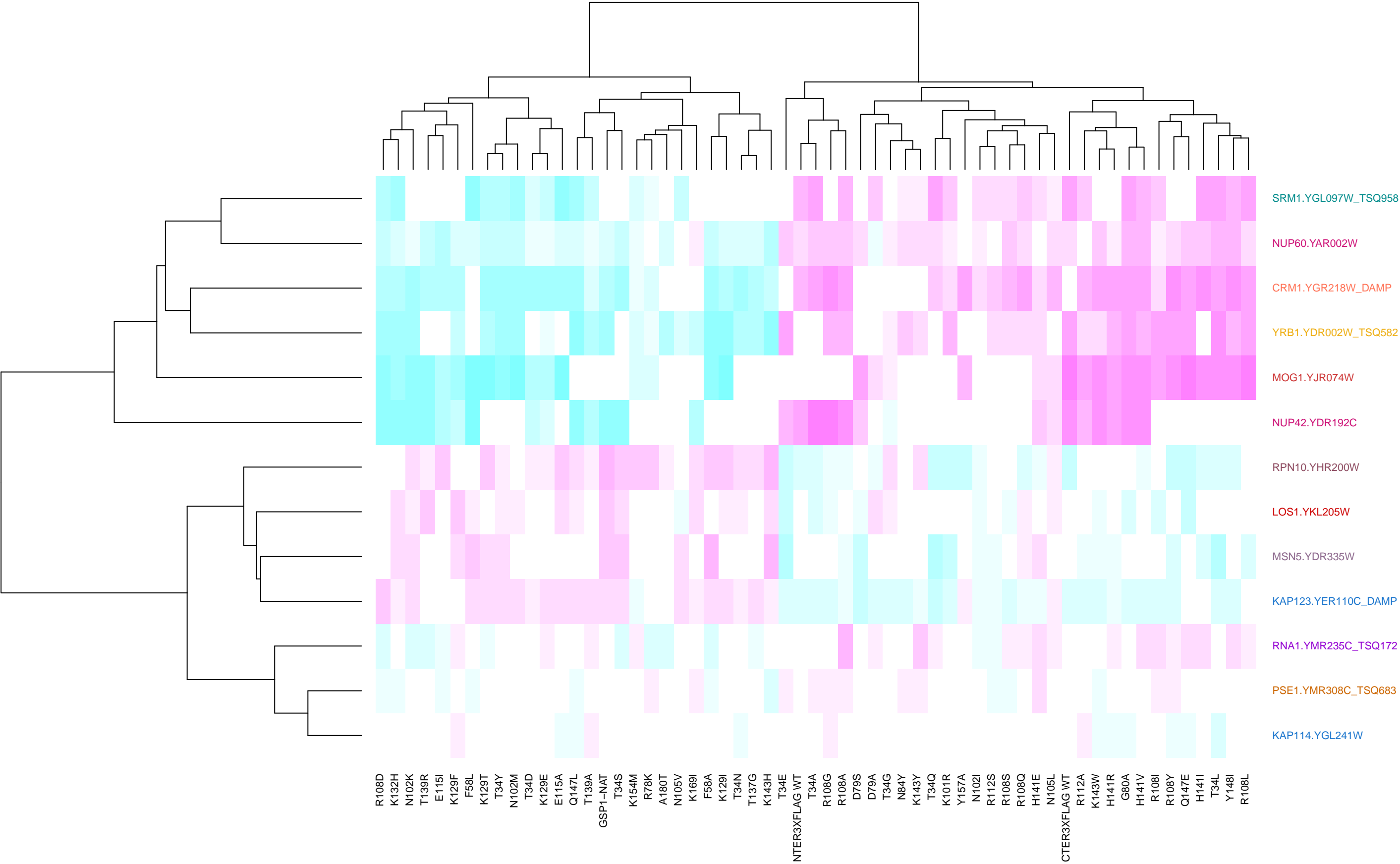
whole_library_24



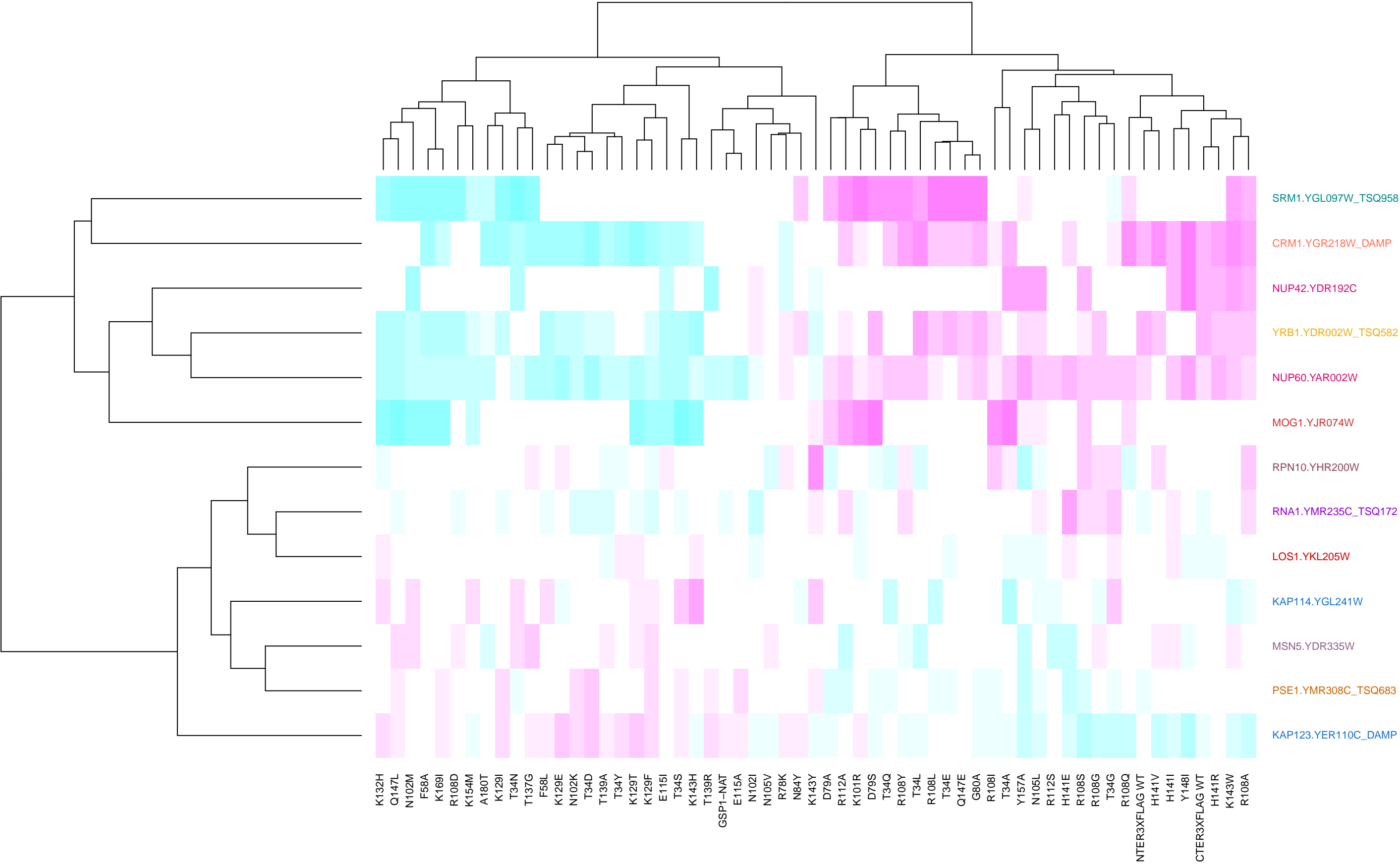
budding



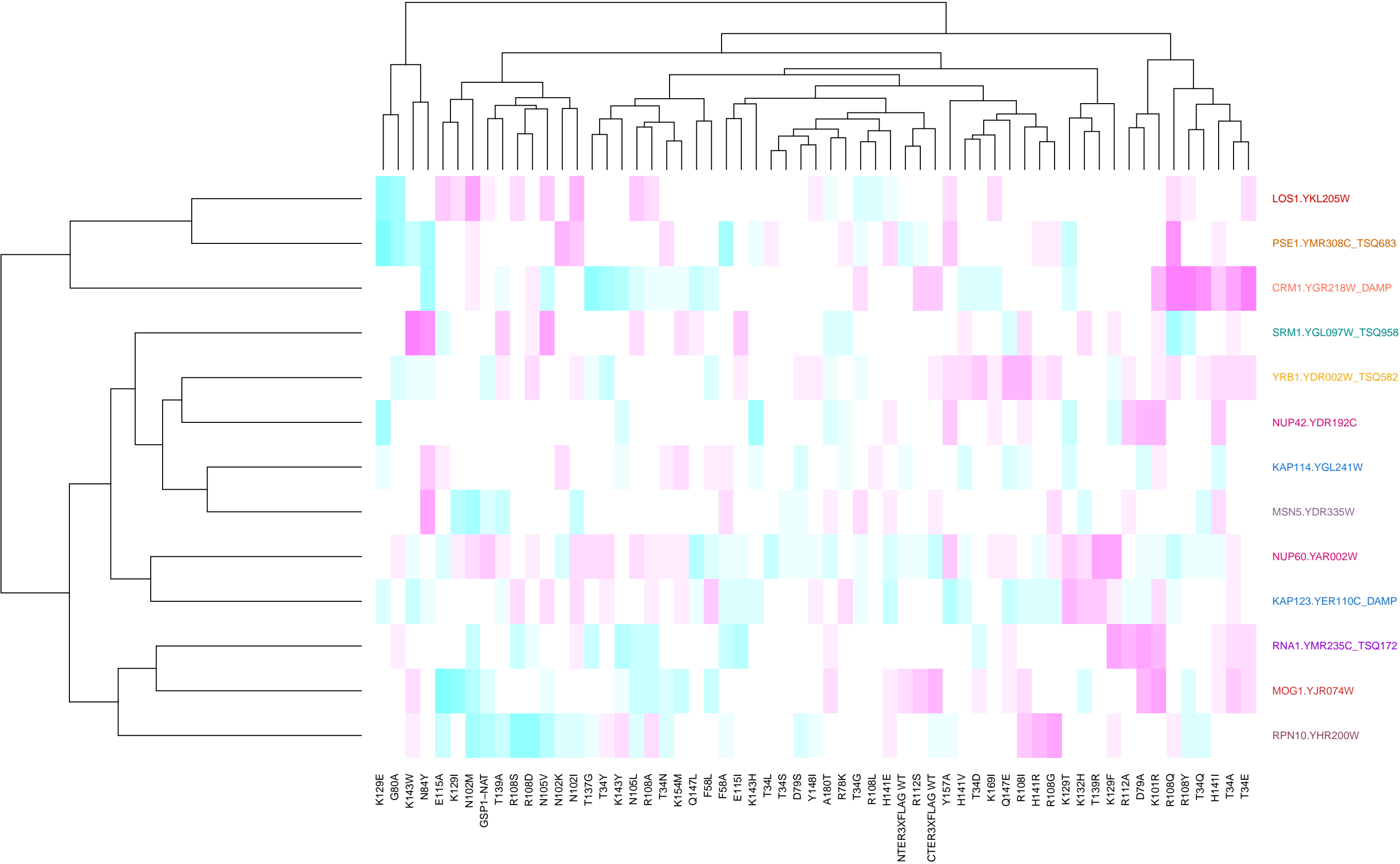
cellular response to DNA damage stimulus



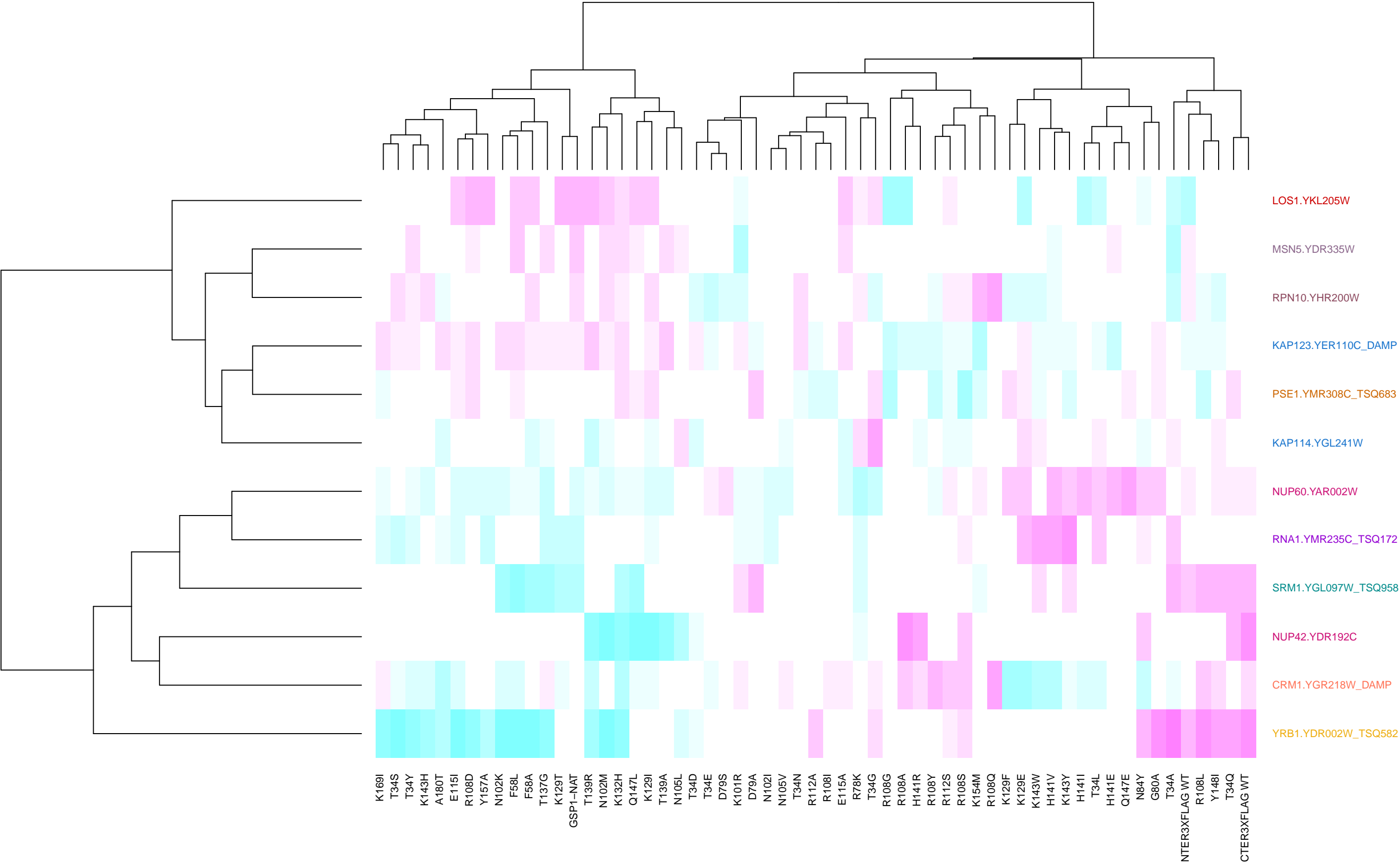
chromatin organization_GO_2



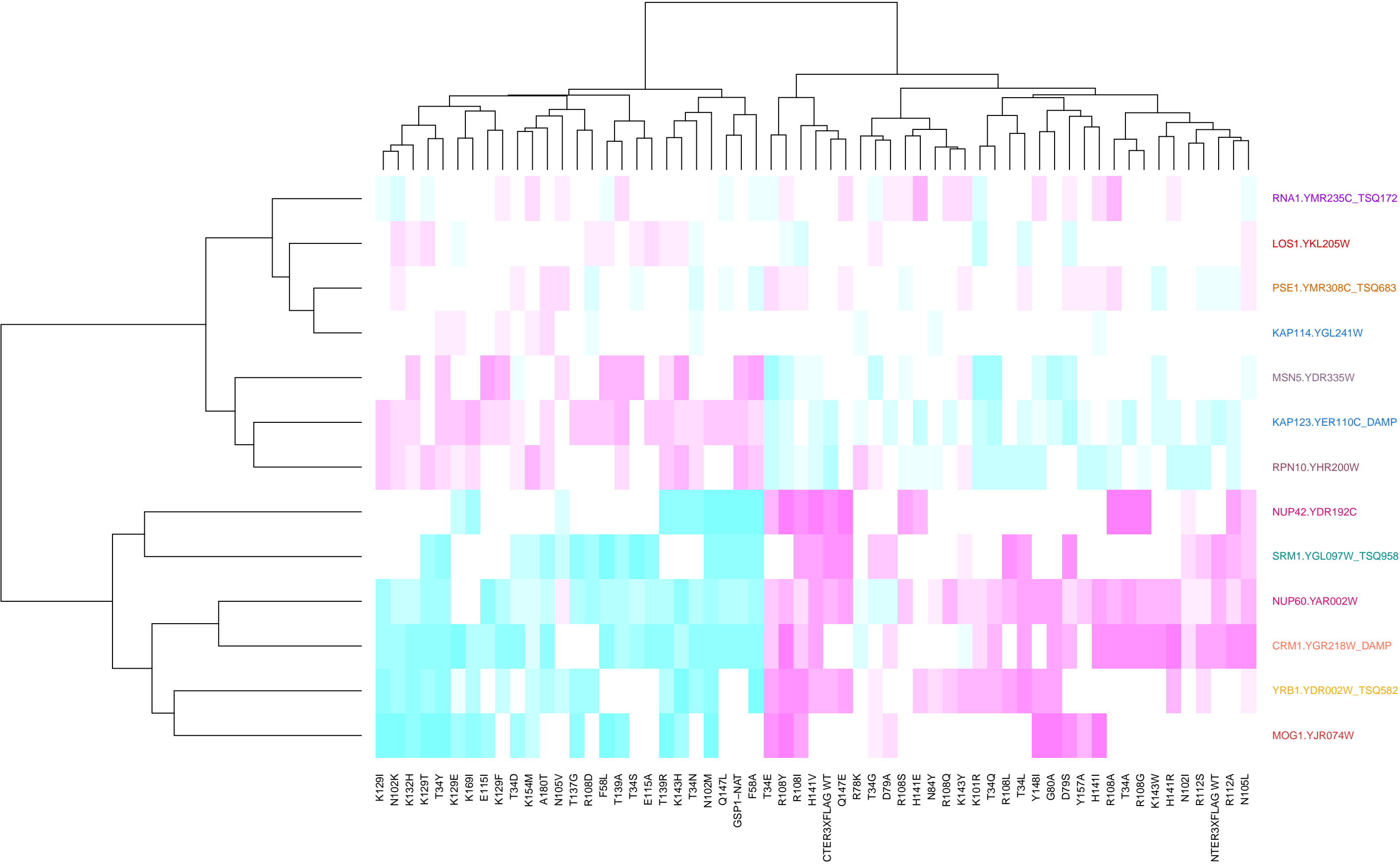
lipid binding



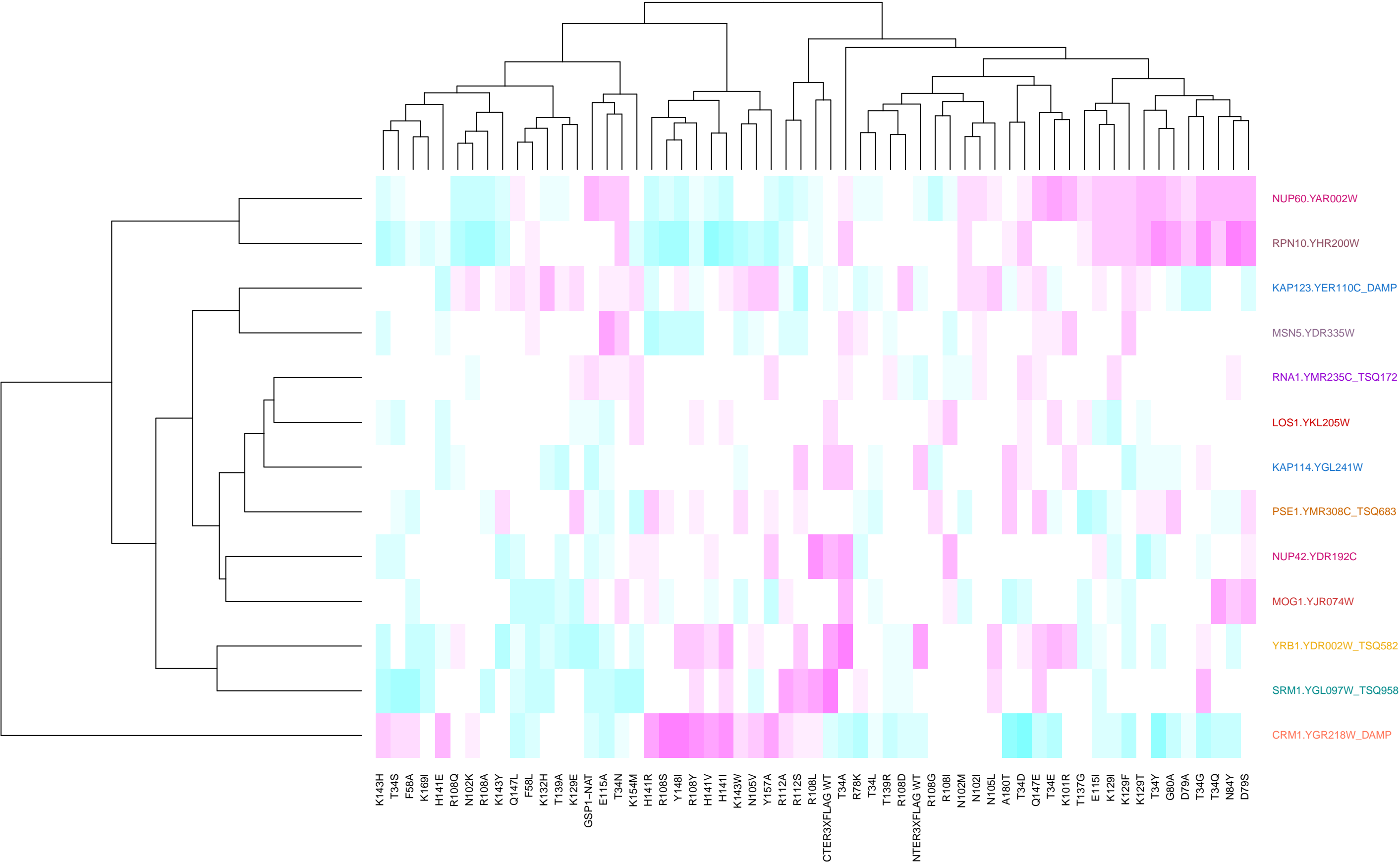
mRNA processing



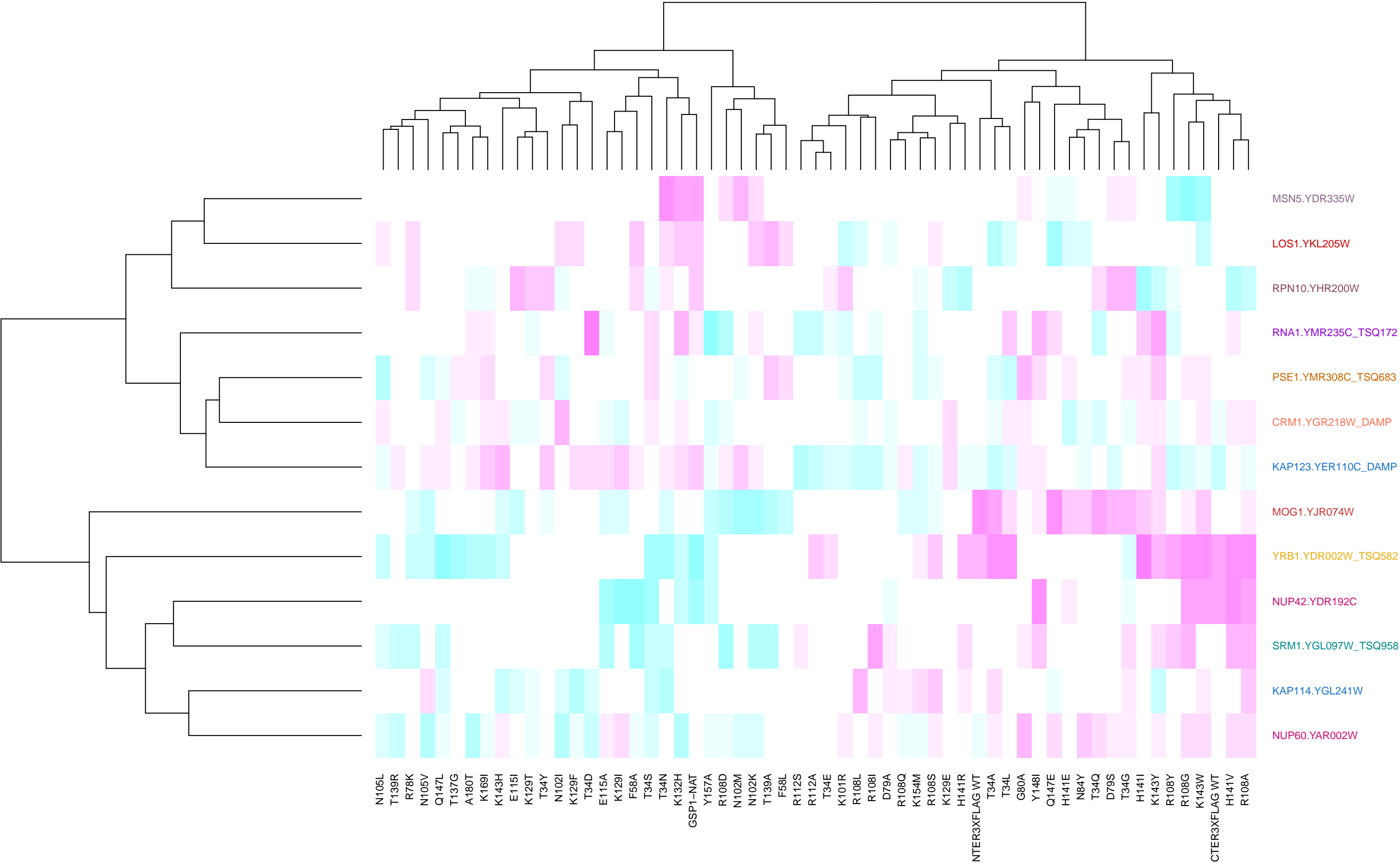
cellular response to DNA damage stimulus_GO_1



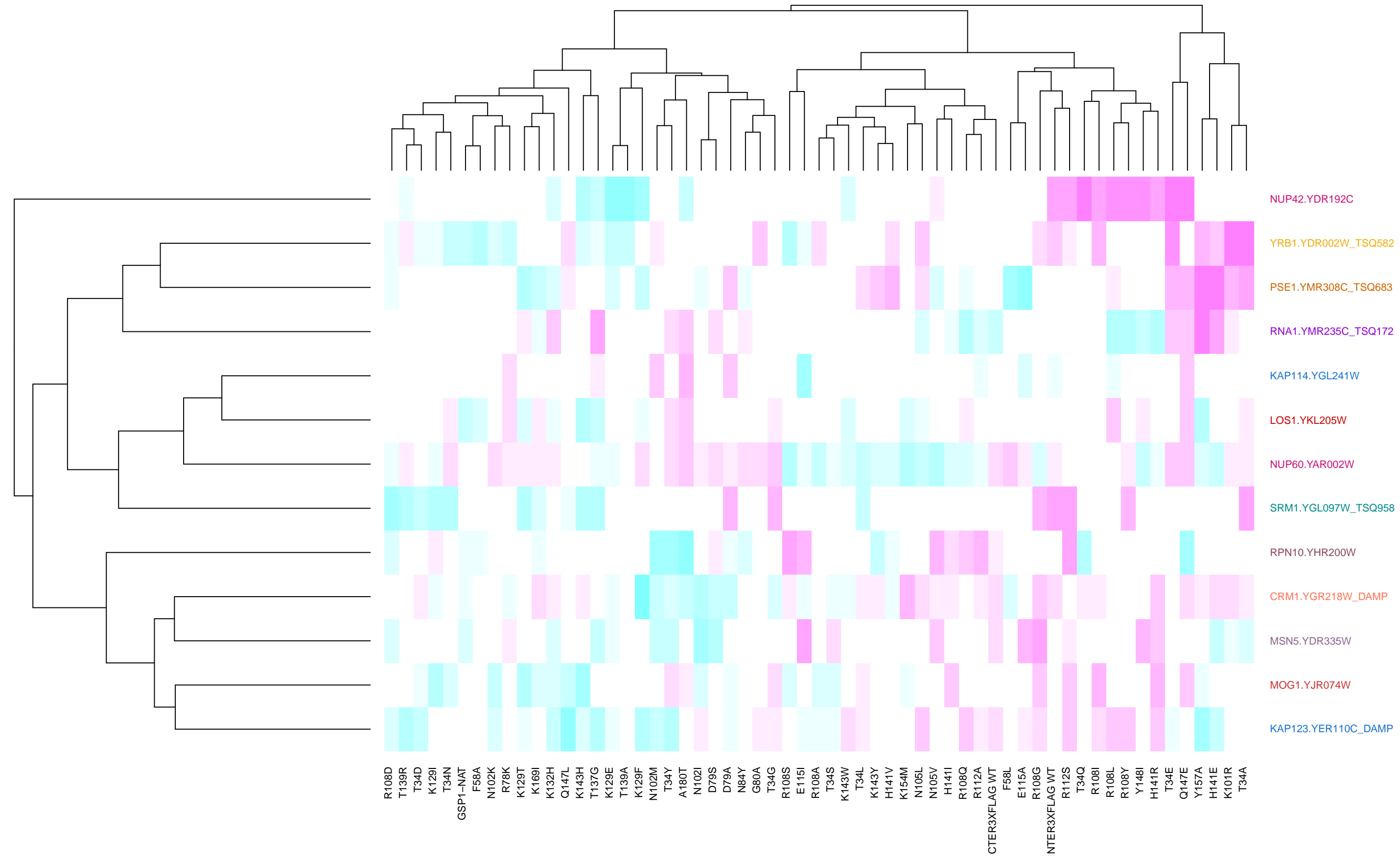
cytoskeleton and microtubules_GO_2



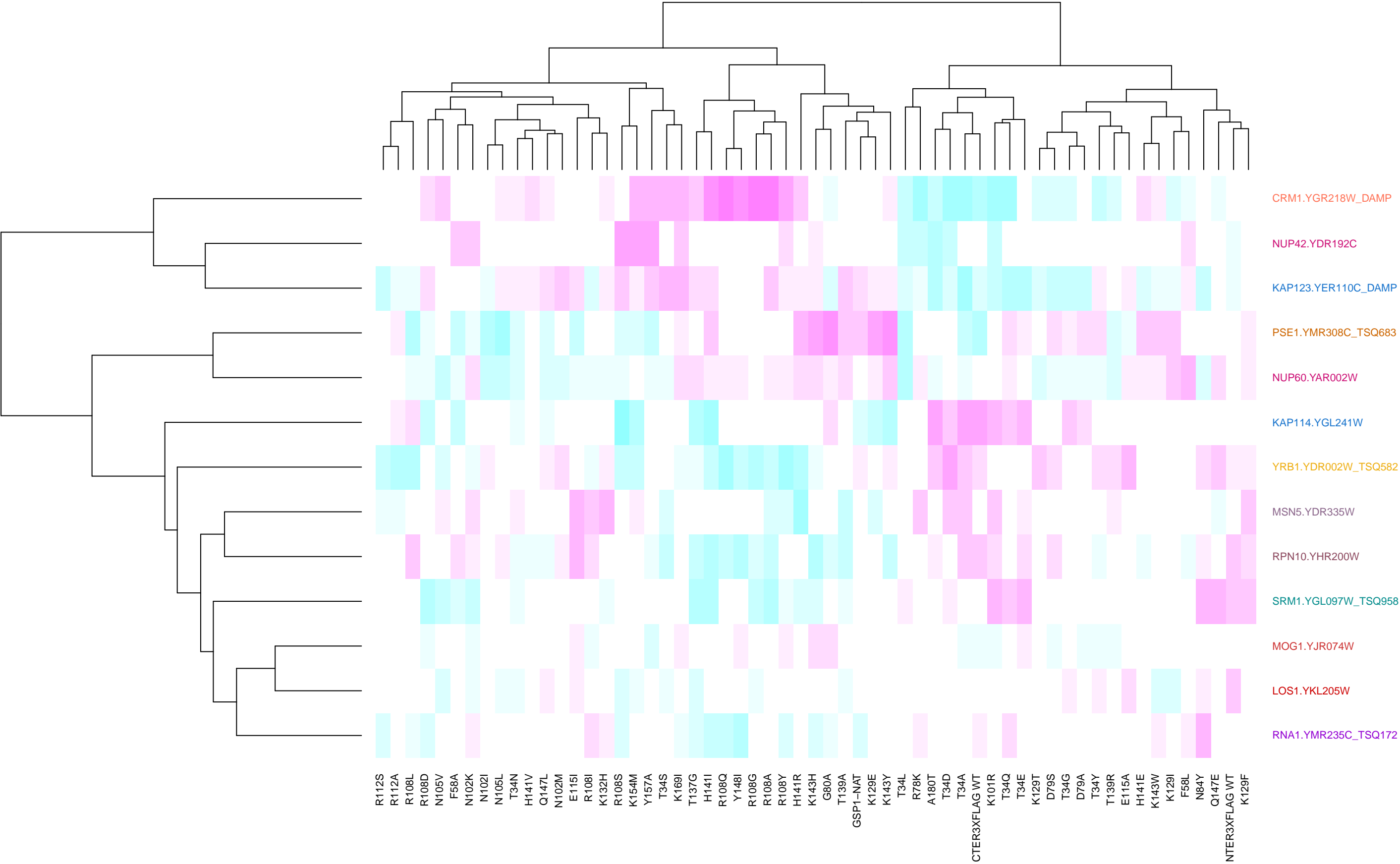
organelle fission_GO_3



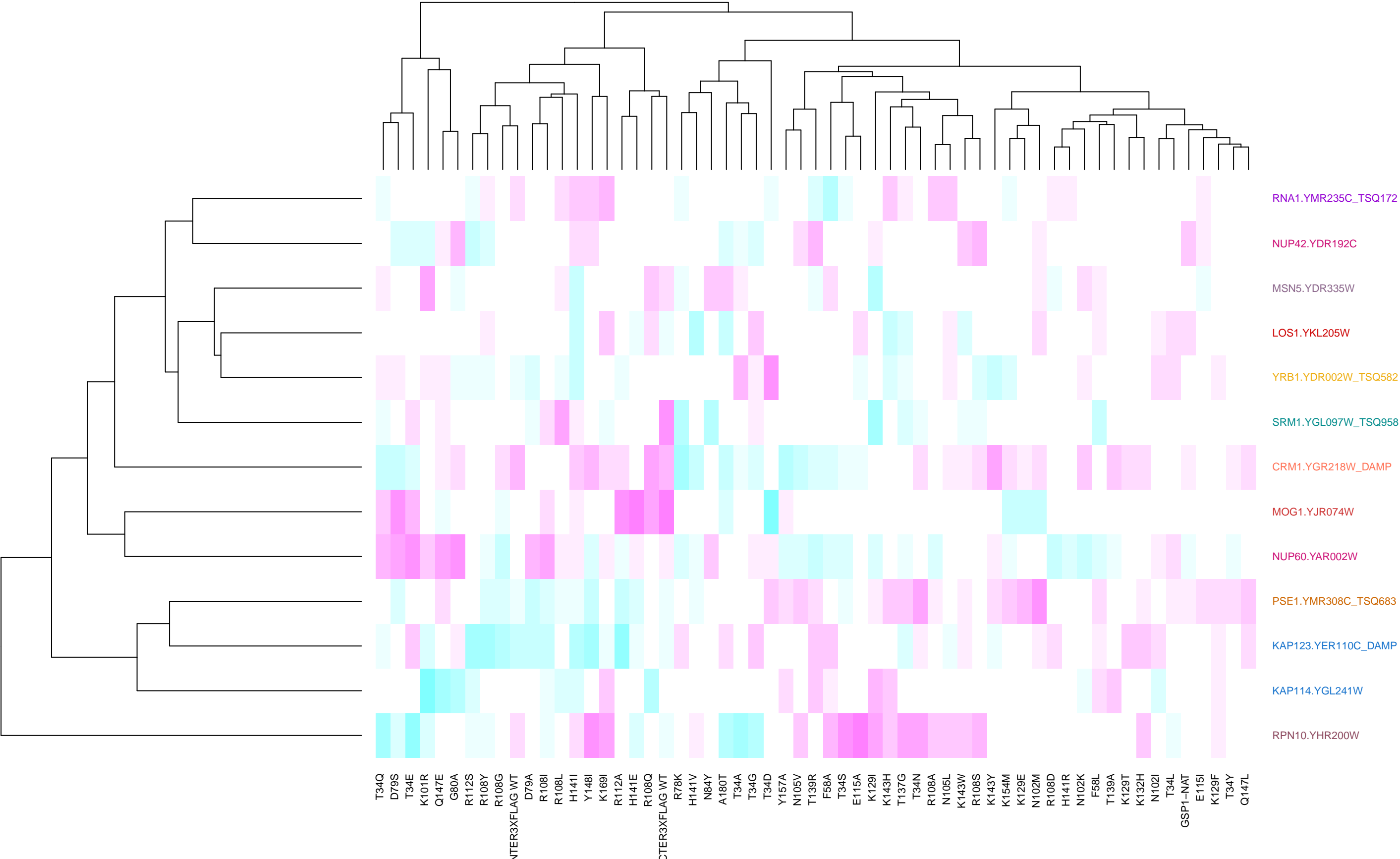
whole_library_19



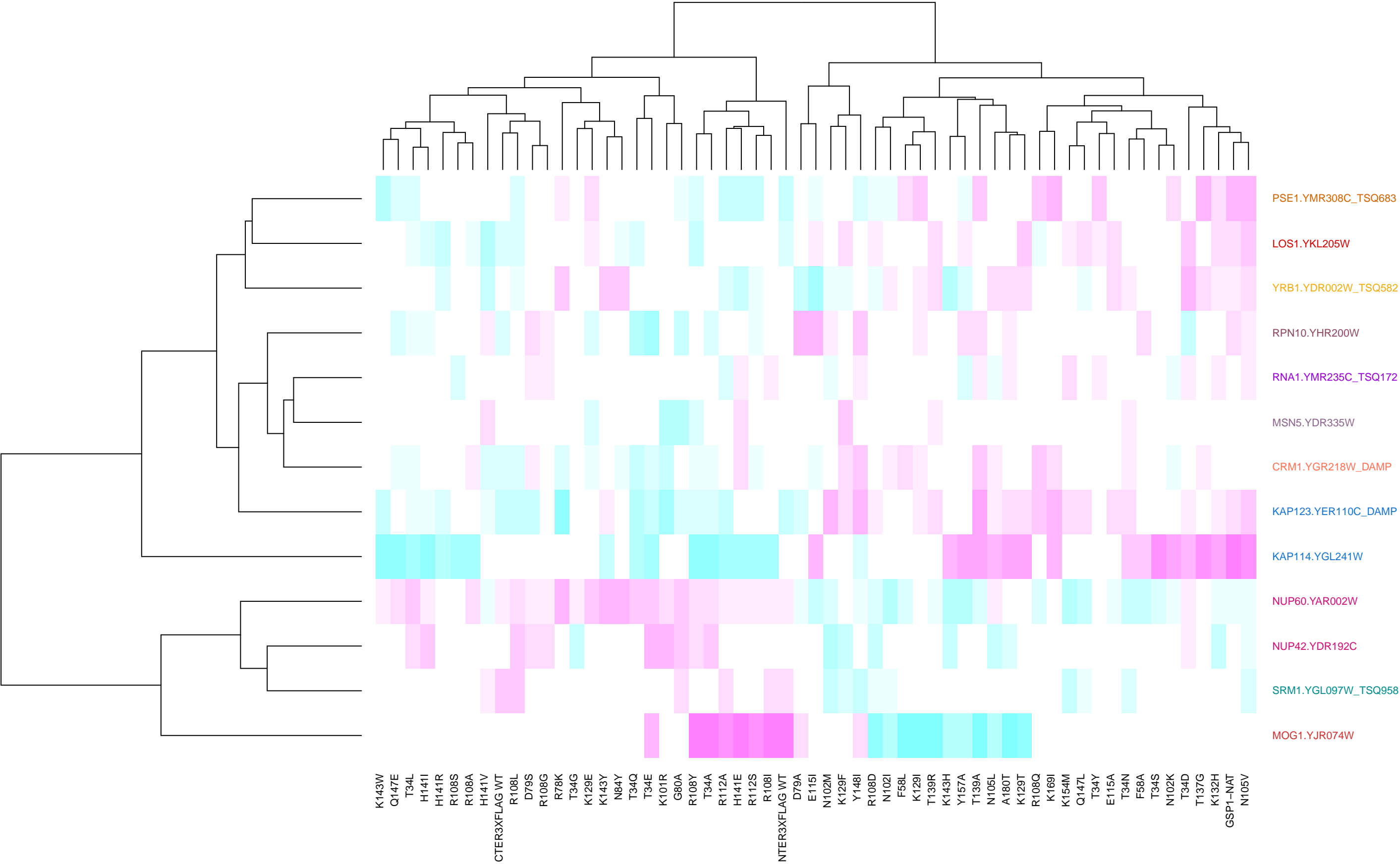
cell cycle_GO_3



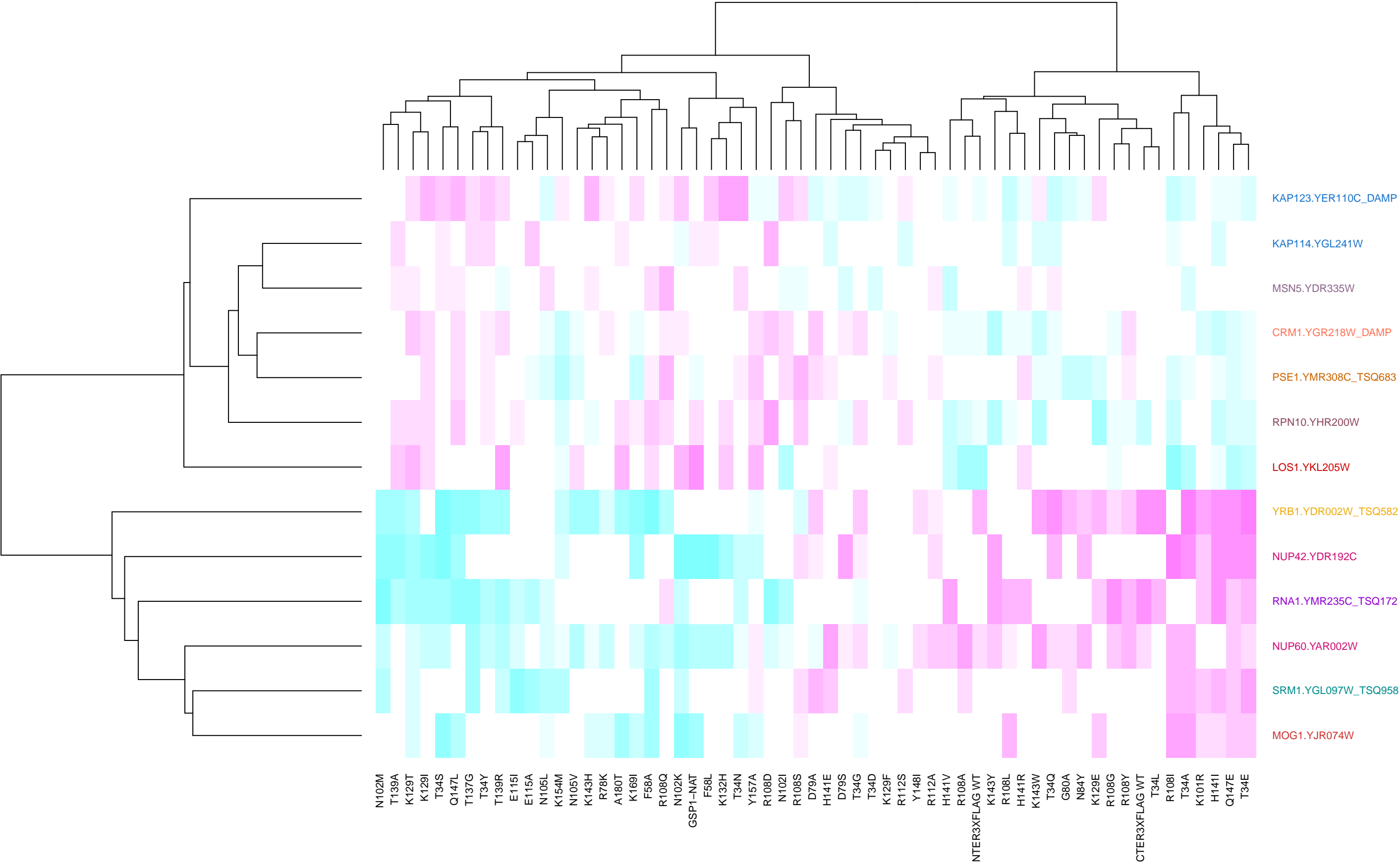
transcription and mRNA processing_GO_6



telomere organization



transcription and mRNA processing_GO_3



proteolysis involved in cellular protein catabolic process_GO_1

