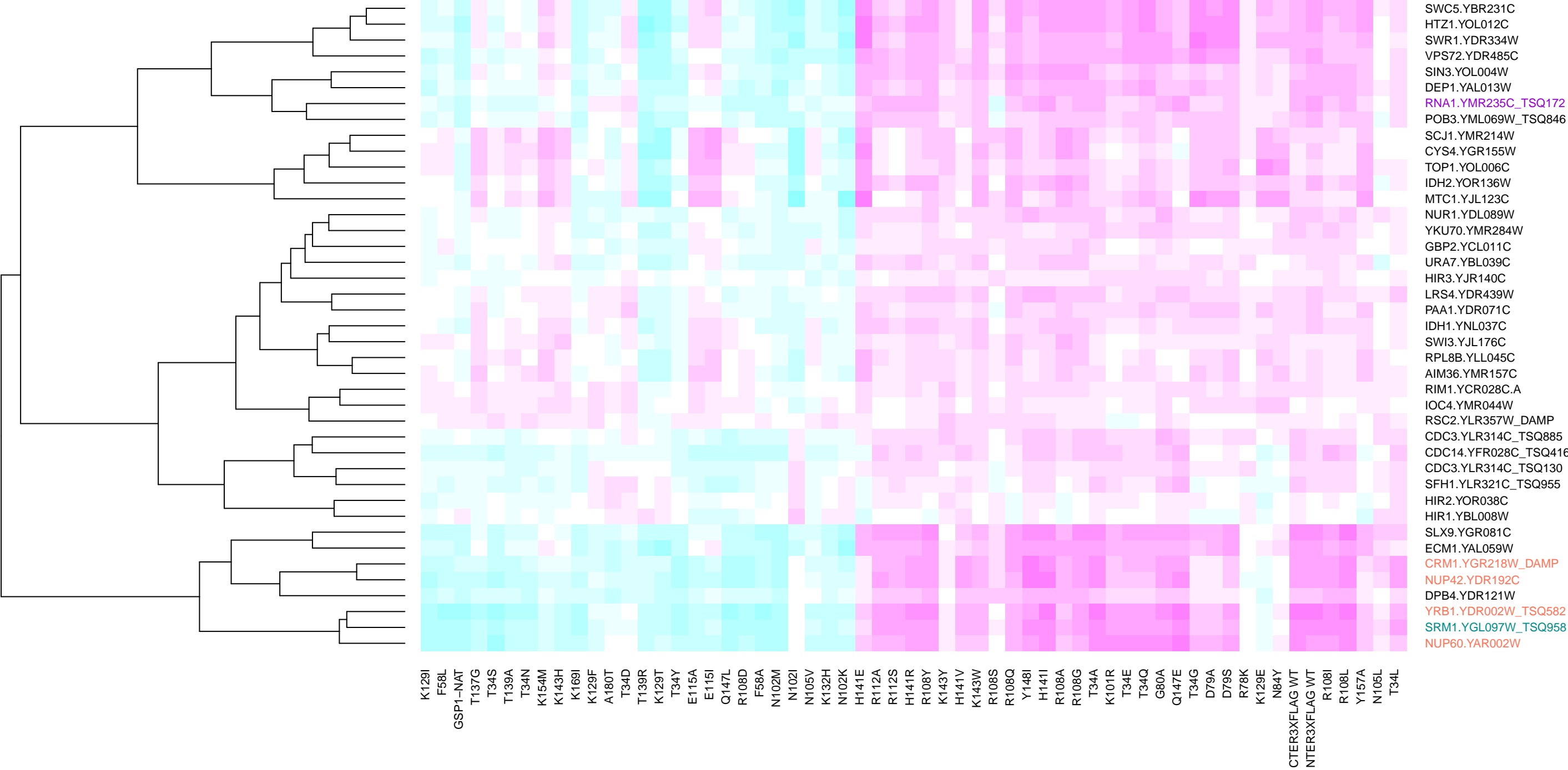
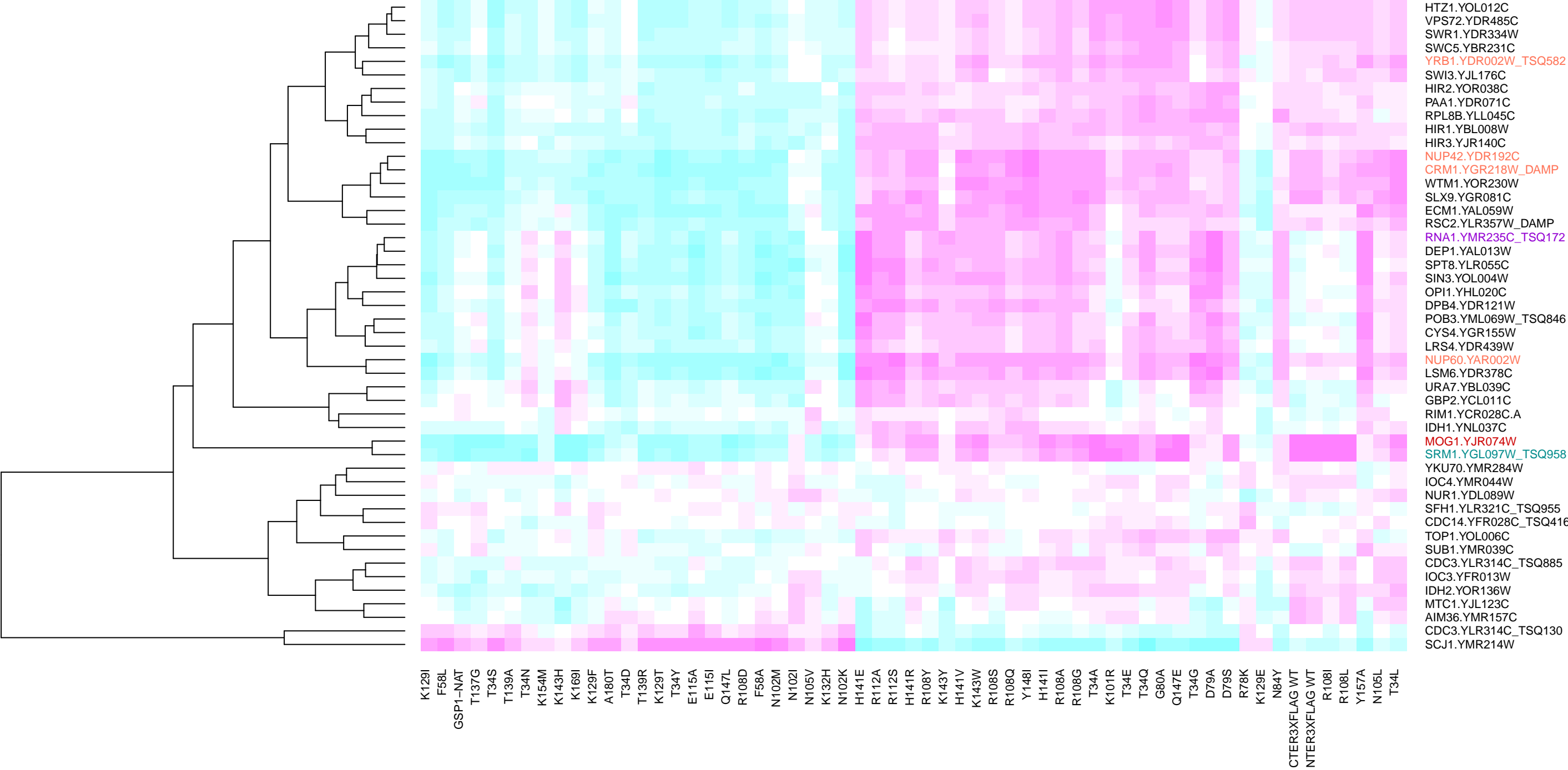


endomembrane system



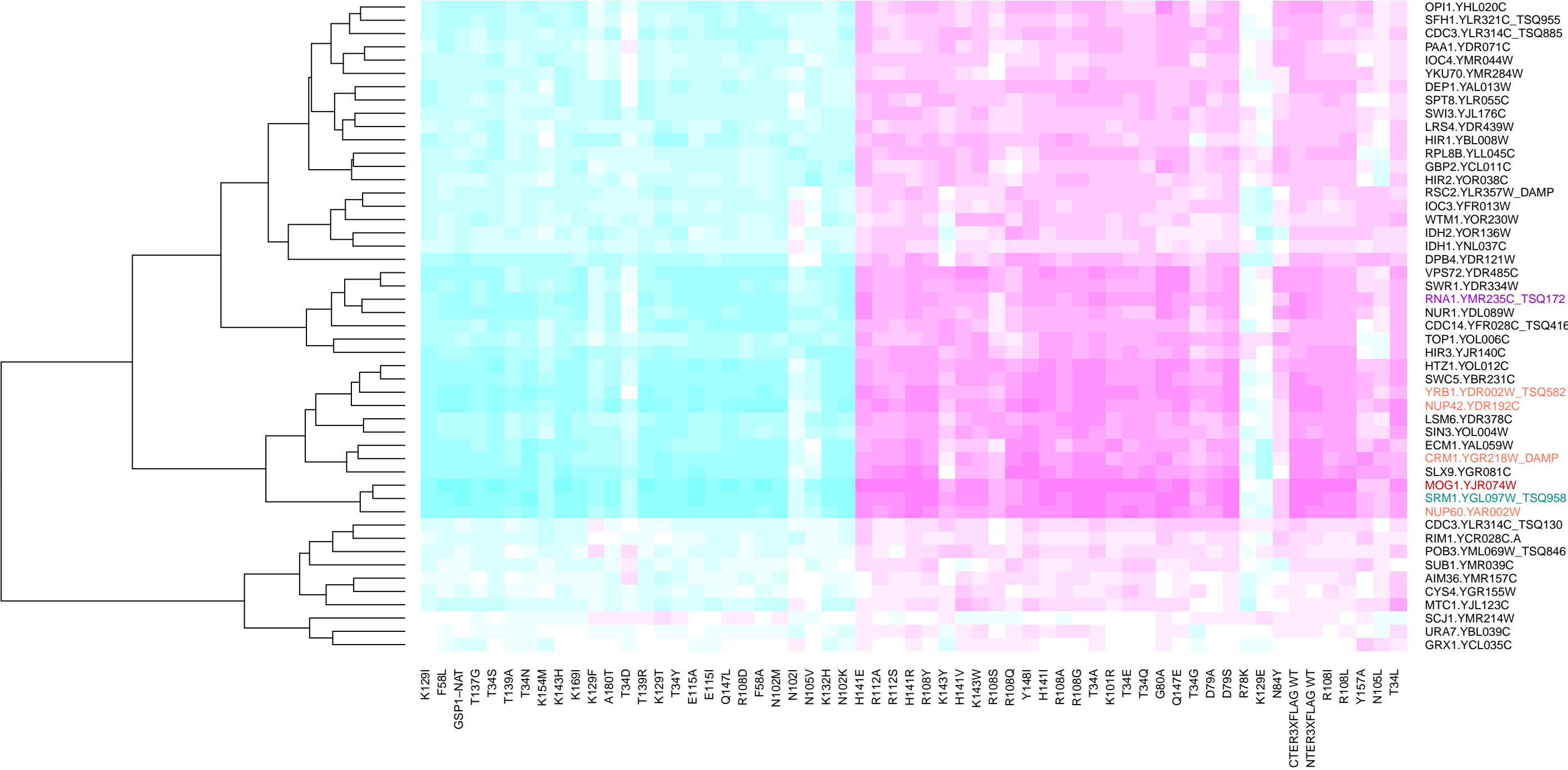
chromatin organization_GO_15_1_mut



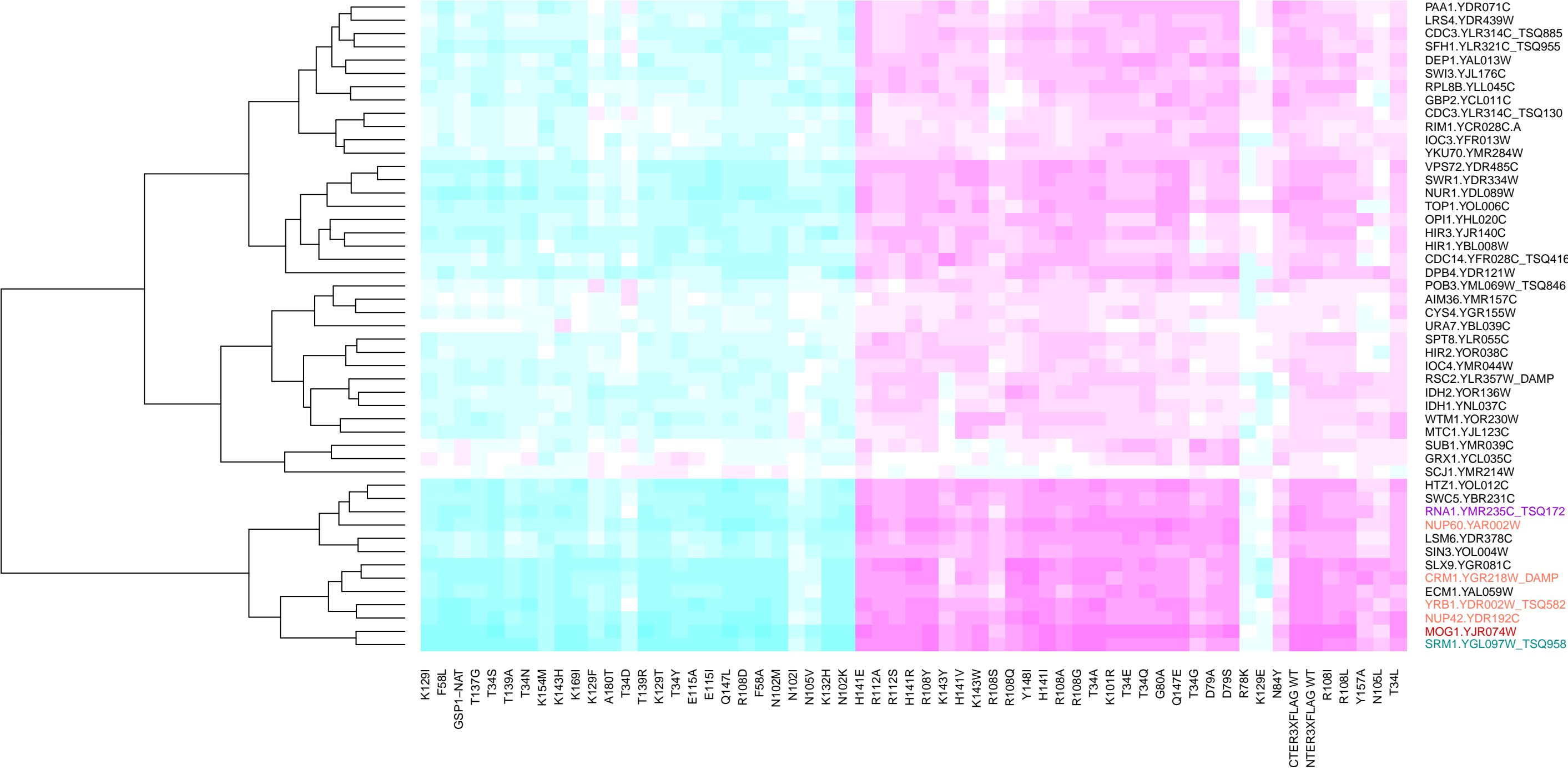
lipids_GO_15_2_all



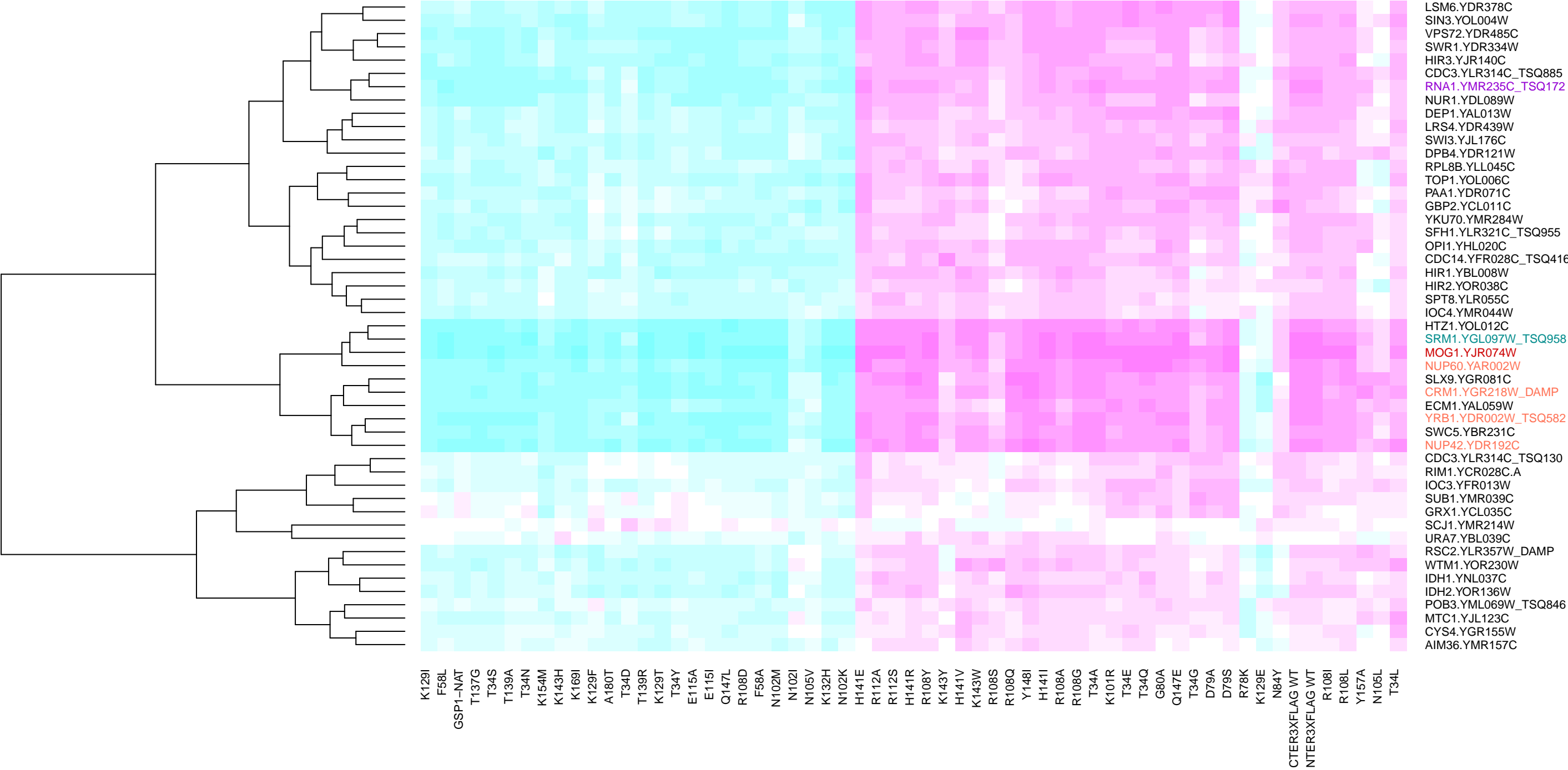
nuclear transport



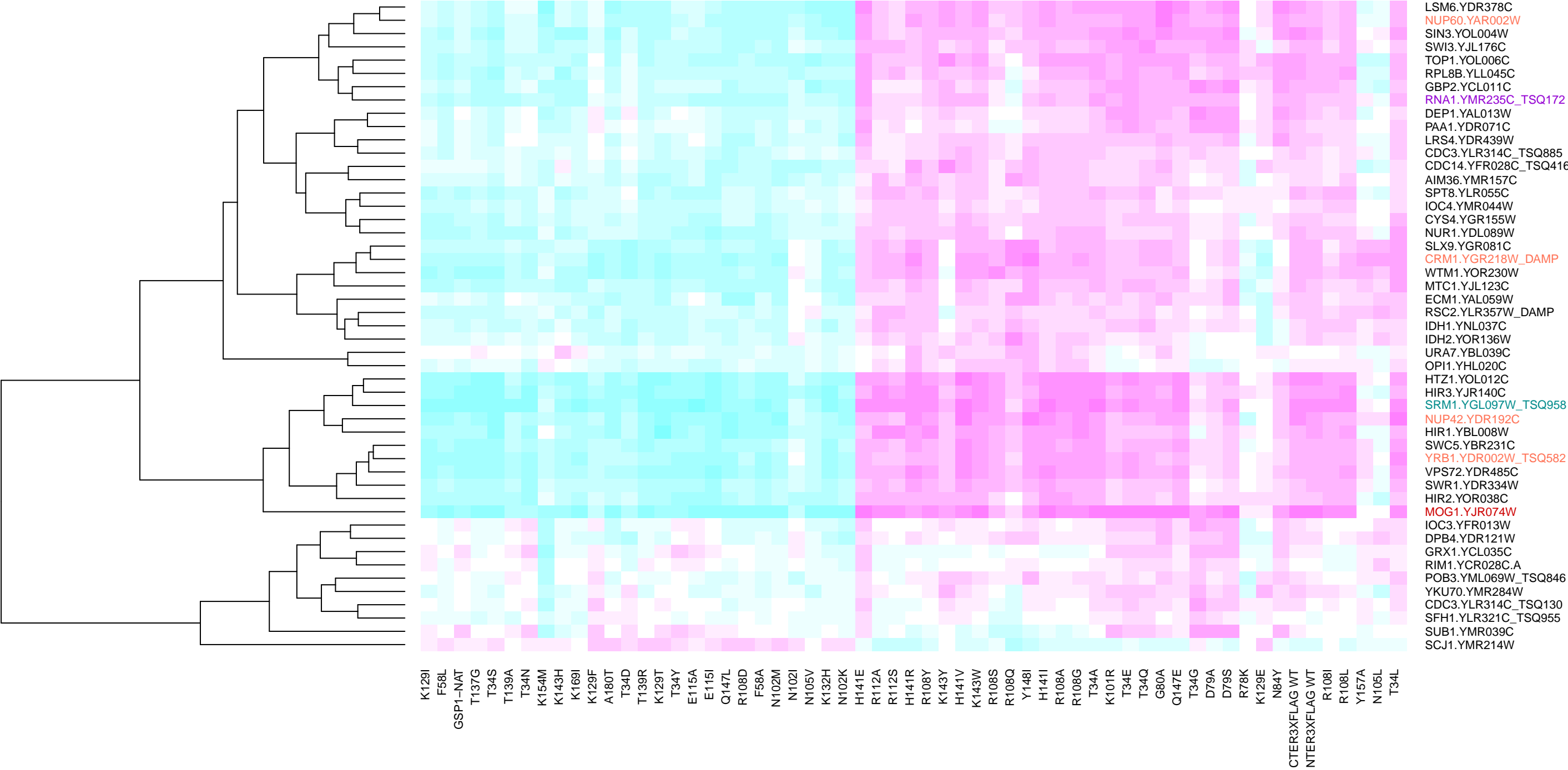
nuclear transport and organization



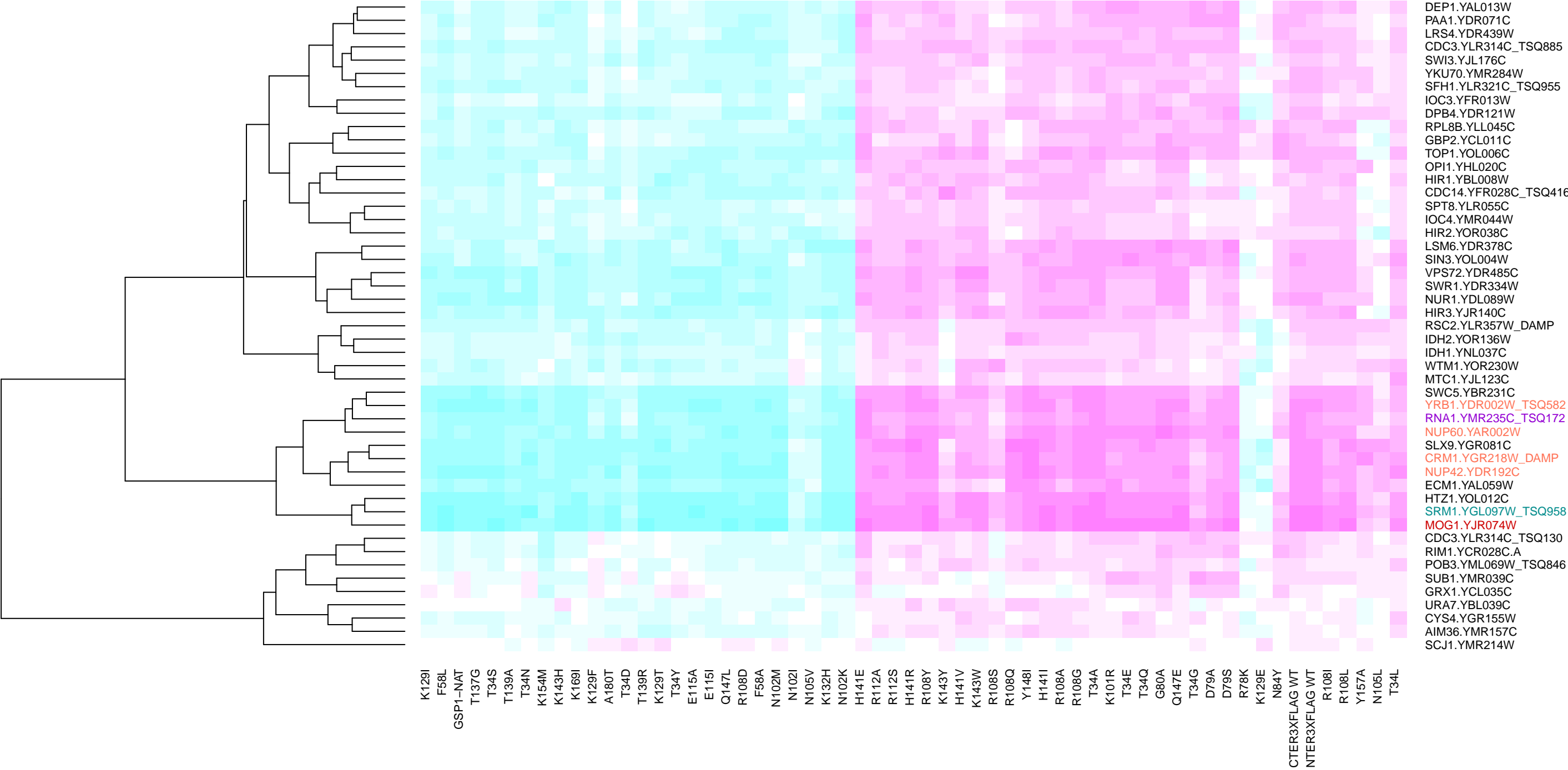
nuclear transport and organization_GO_15_1_mut



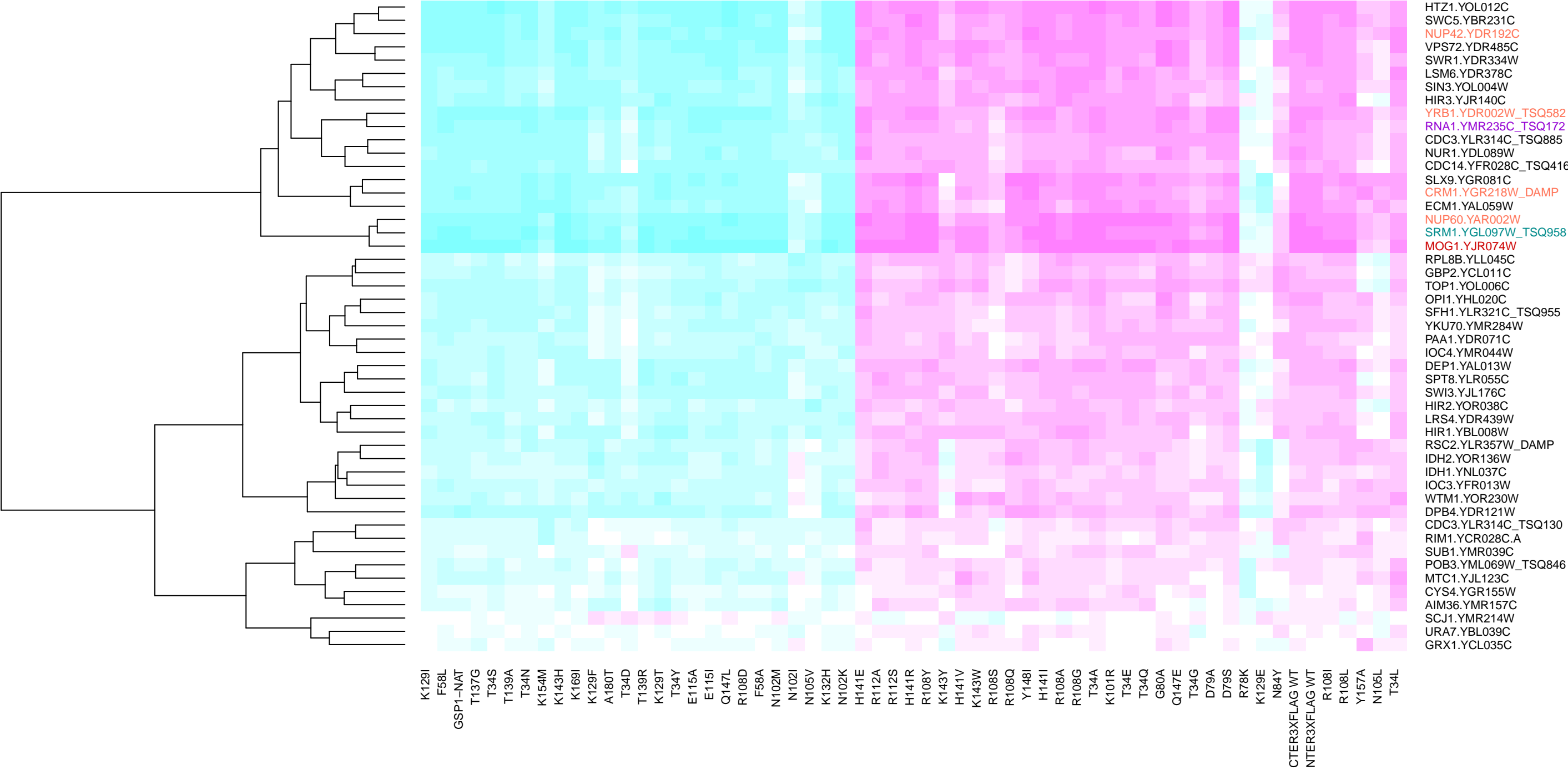
nuclear transport and organization_GO_15_2_all



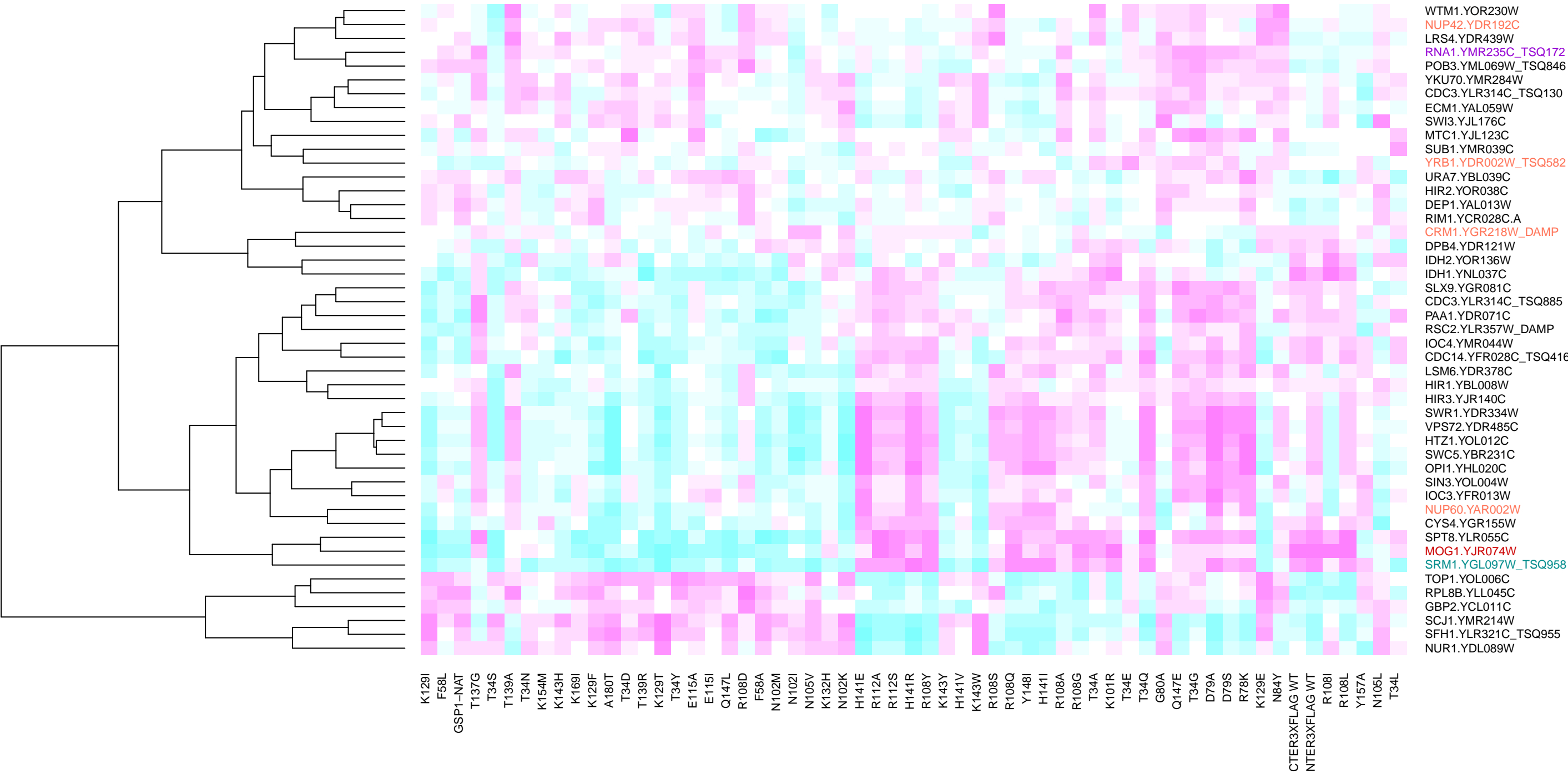
nuclear transport and organization_GO_30_1_all



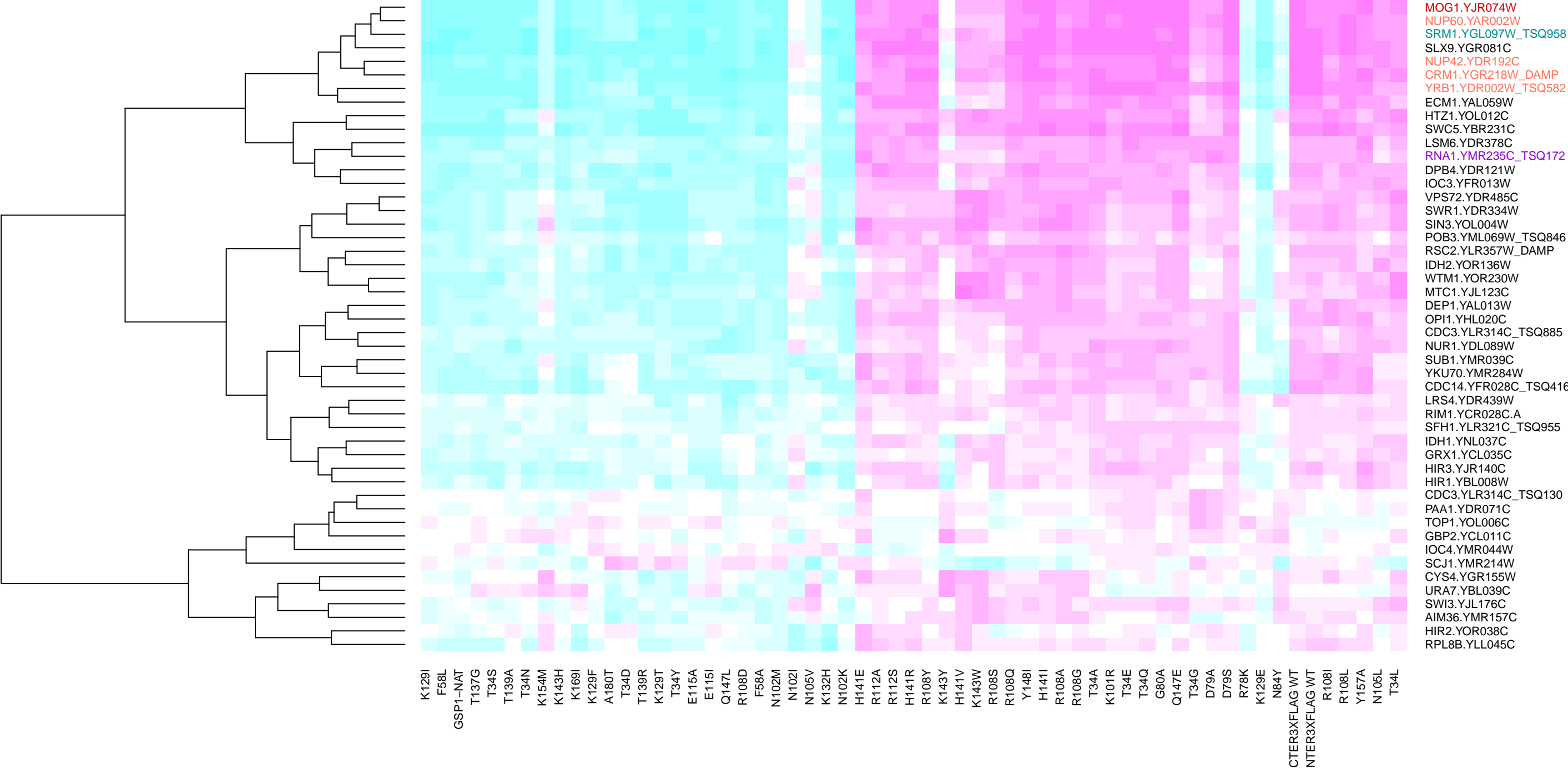
nuclear transport_GO_15_1_mut



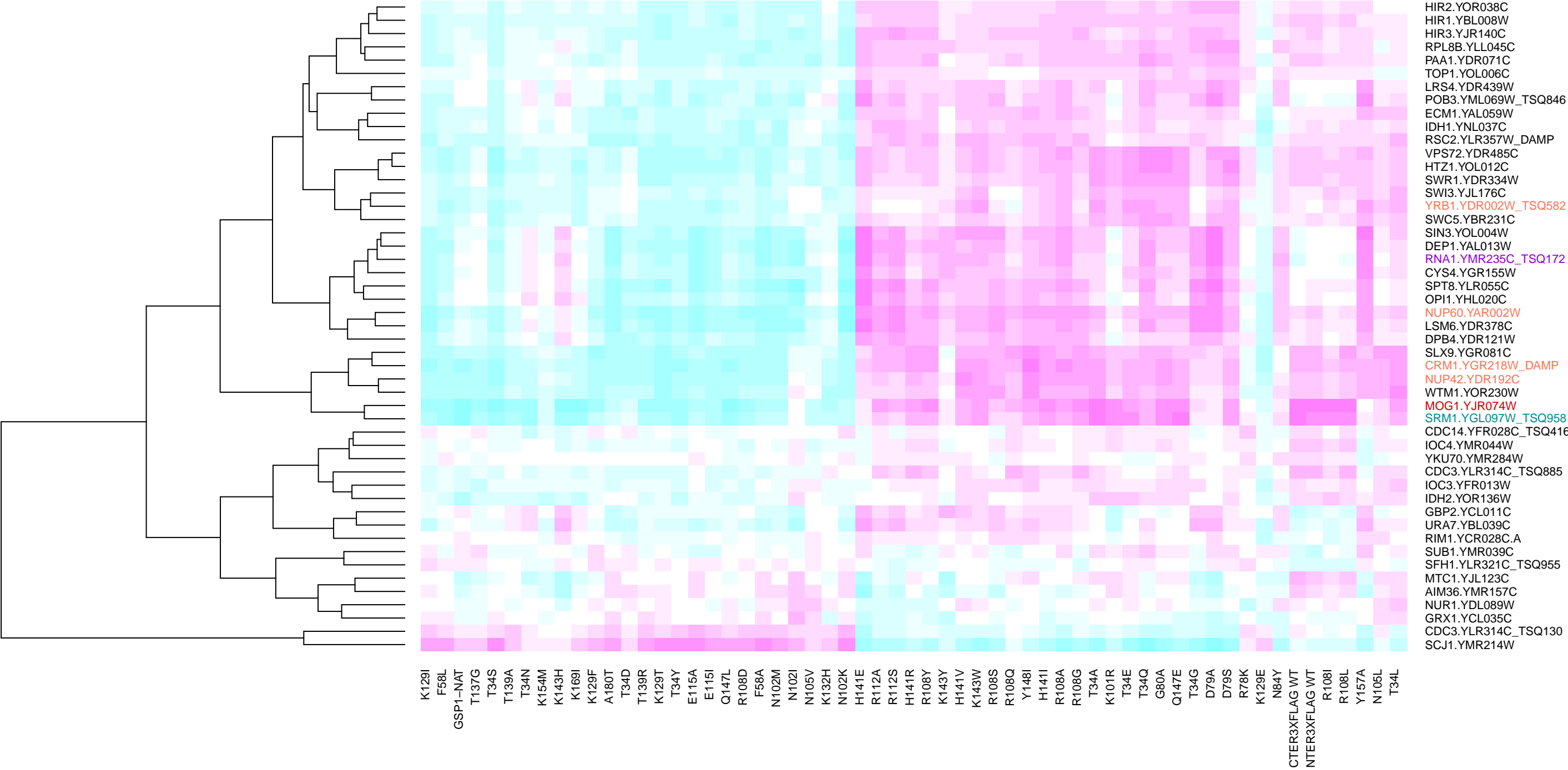
peptidyl-amino acid modification_GO_15_2_mut



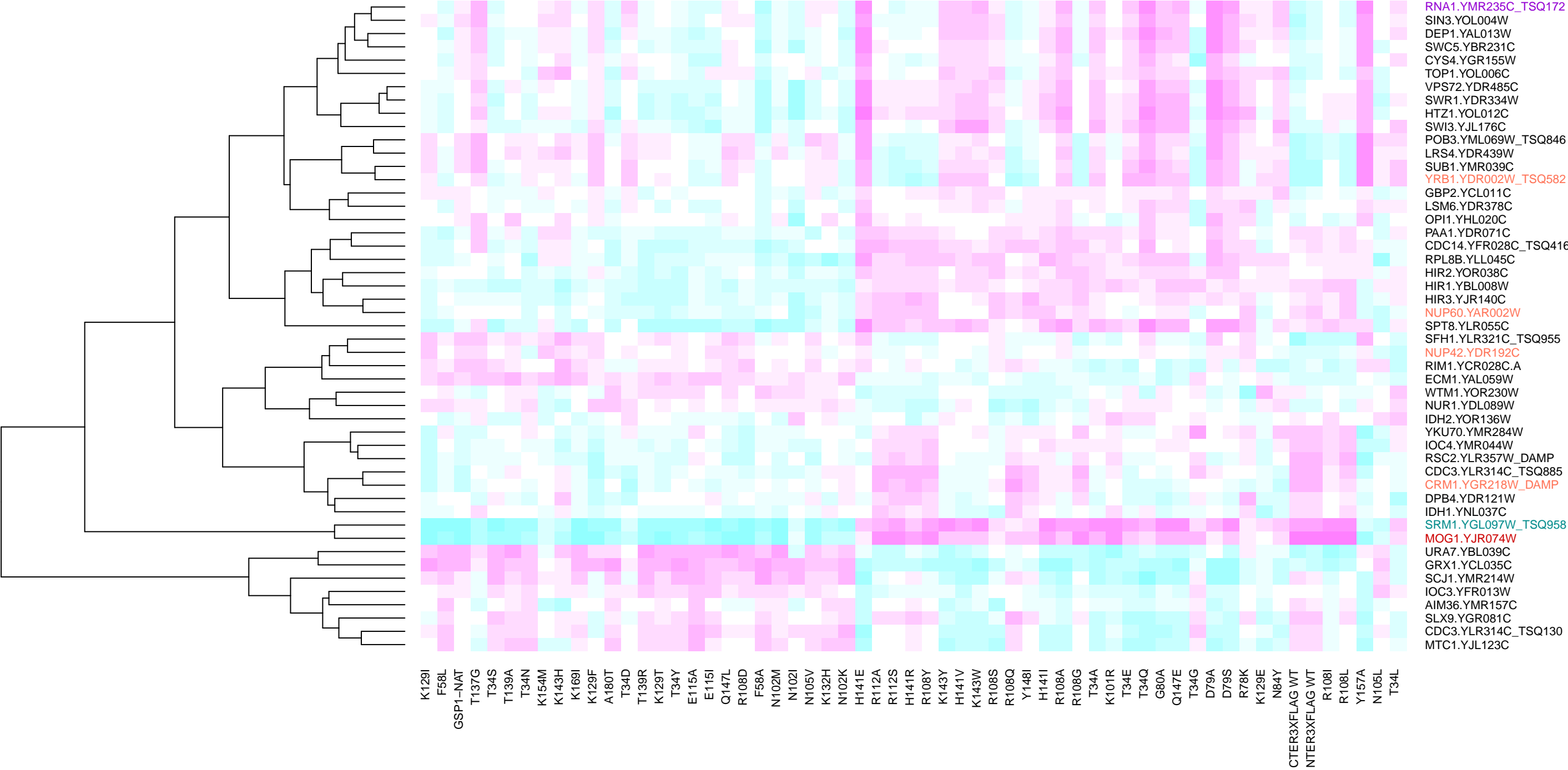
protein targeting_GO_15_3_all



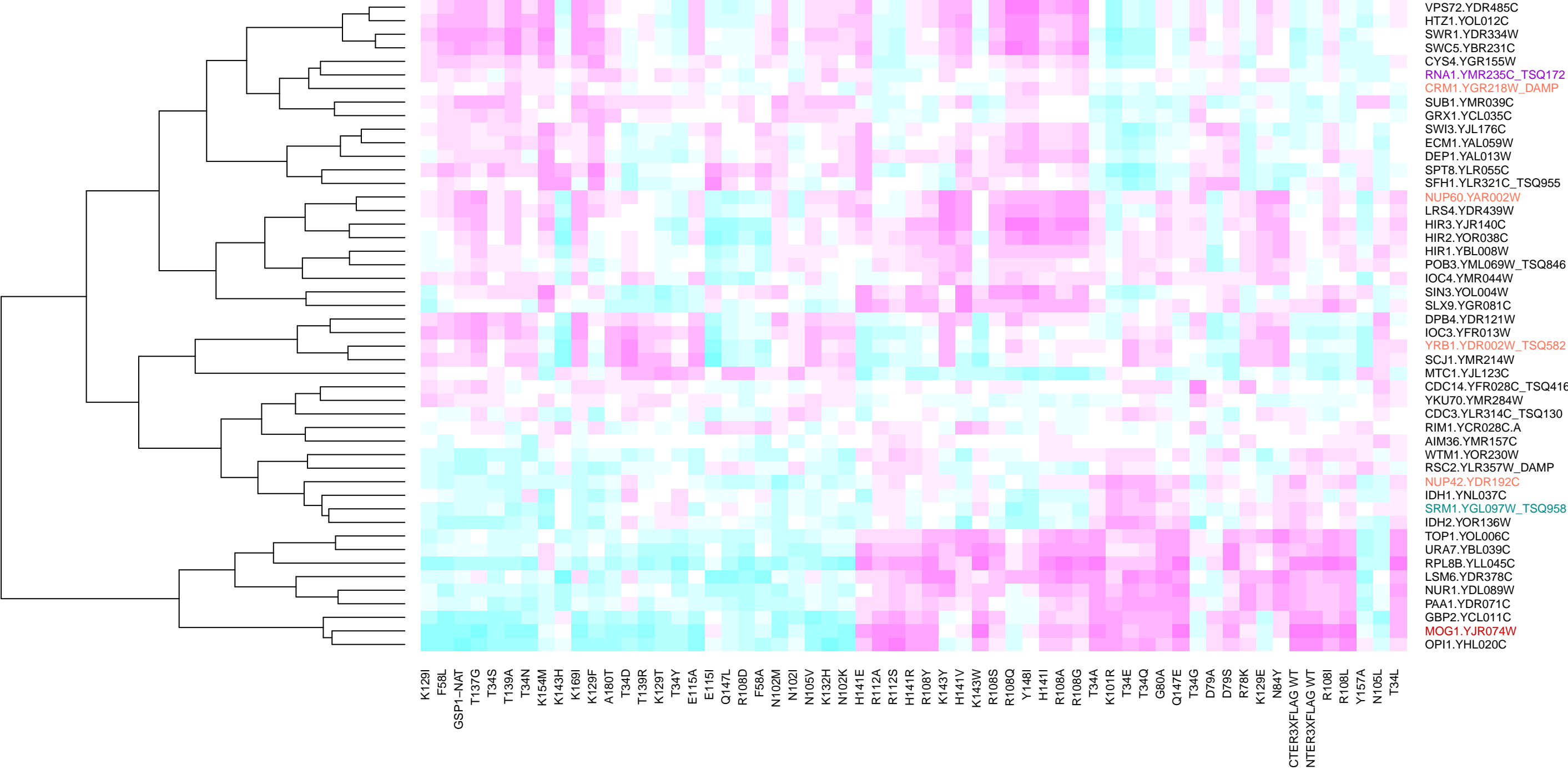
chromatin organization_GO_30_1_mut



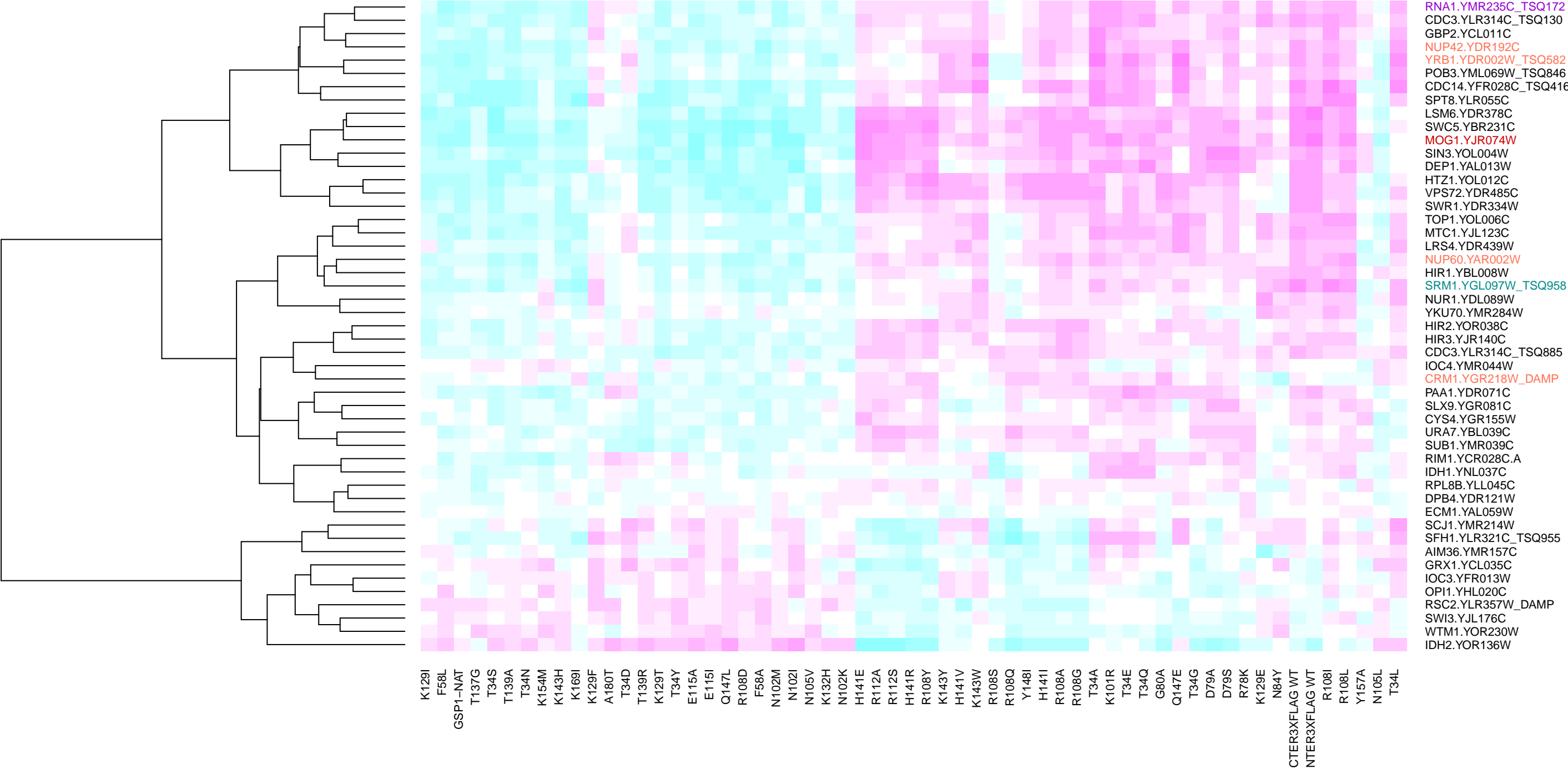
histone modification_GO_15_1_mut



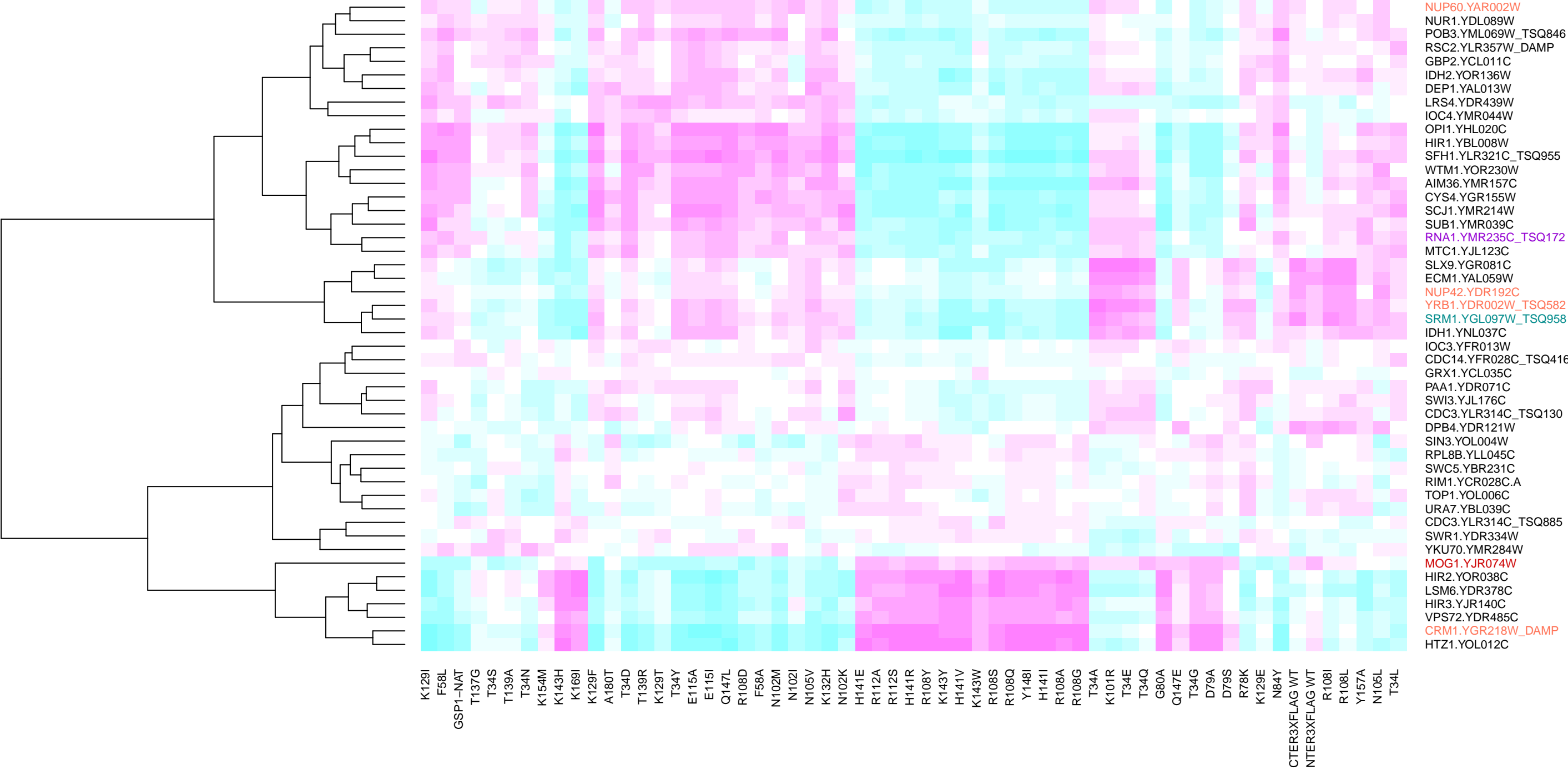
telomere organization



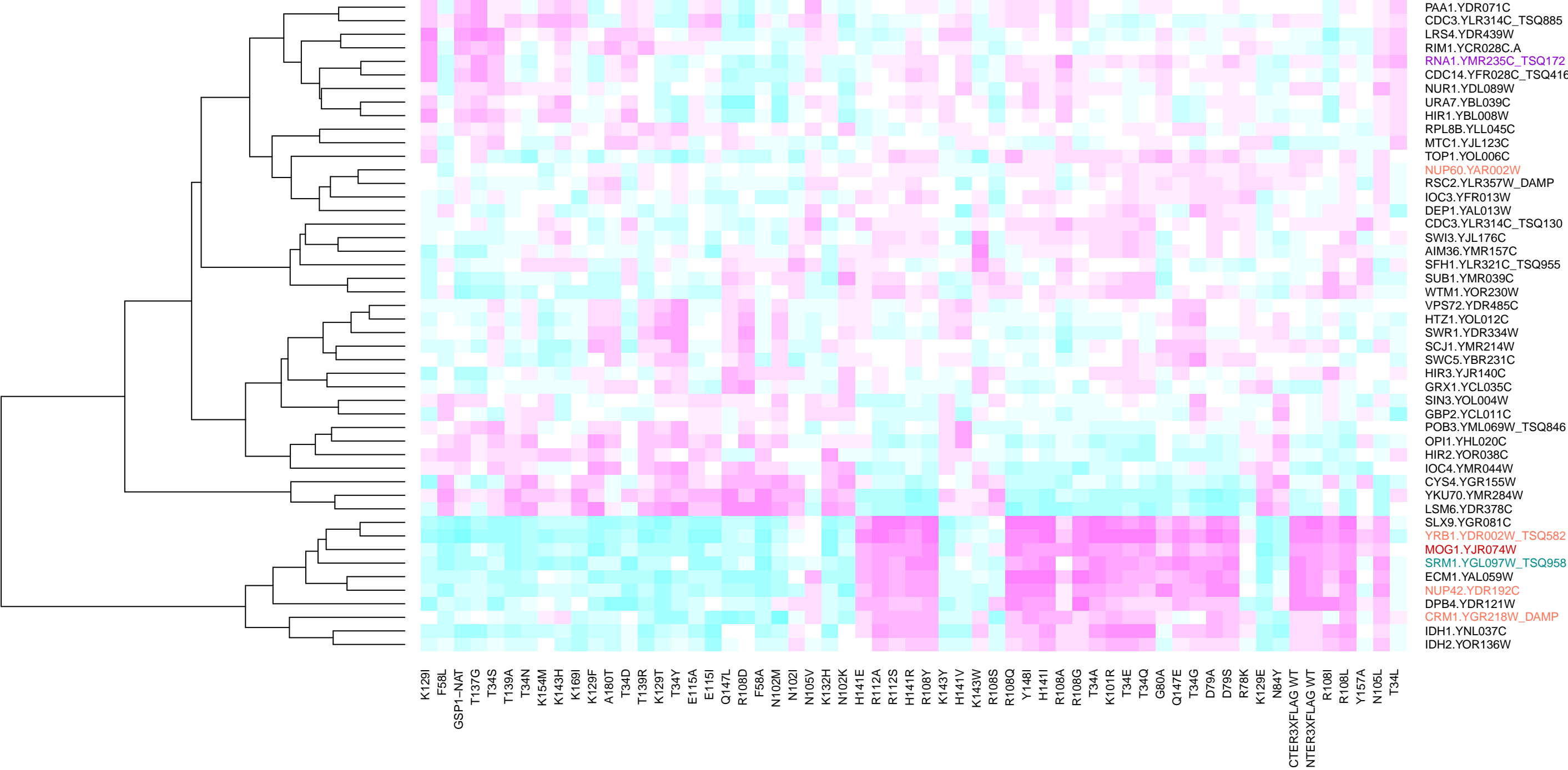
budding_GO_30_1_mut



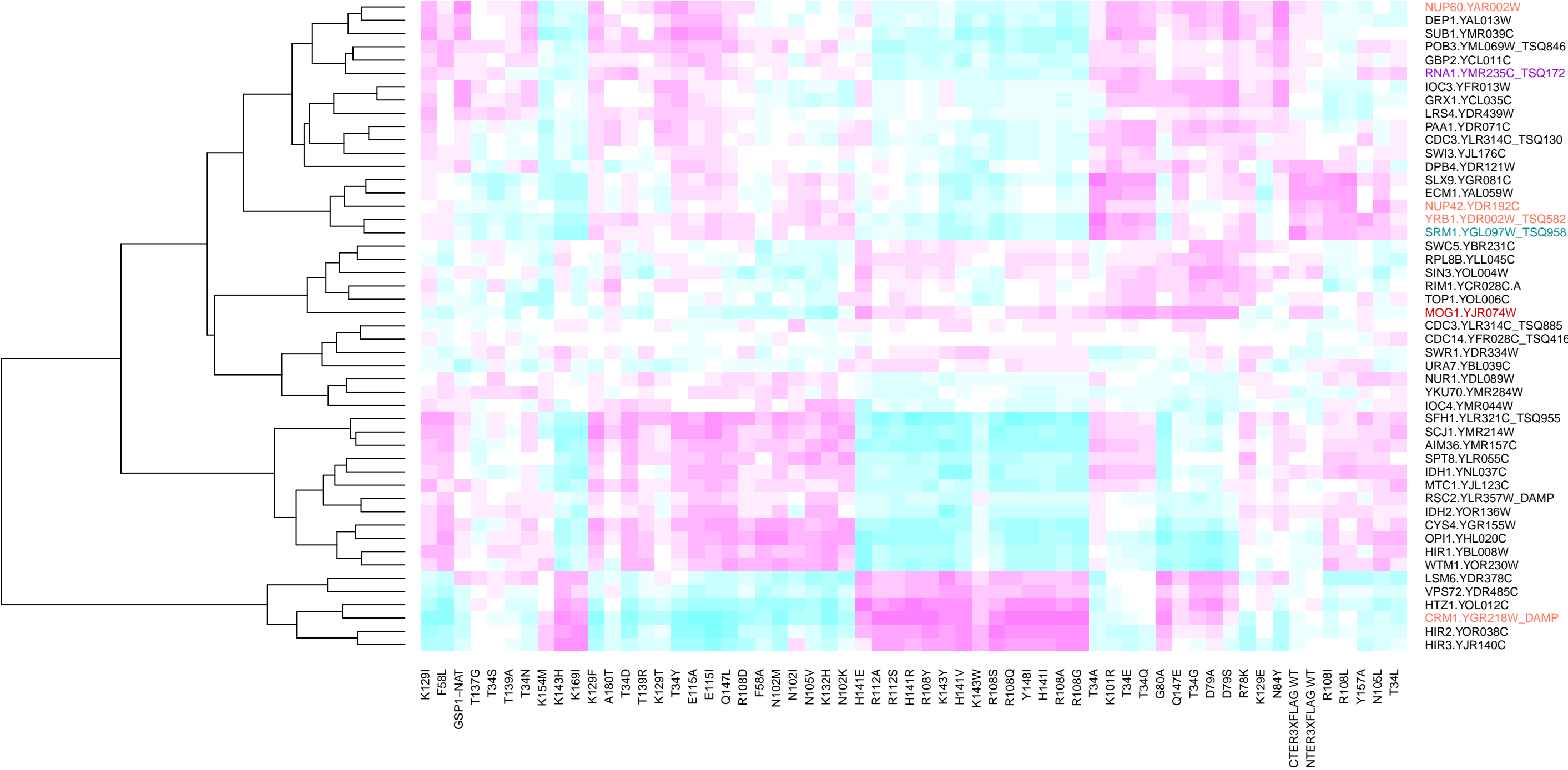
cytoskeleton organization_GO_15_1_mut



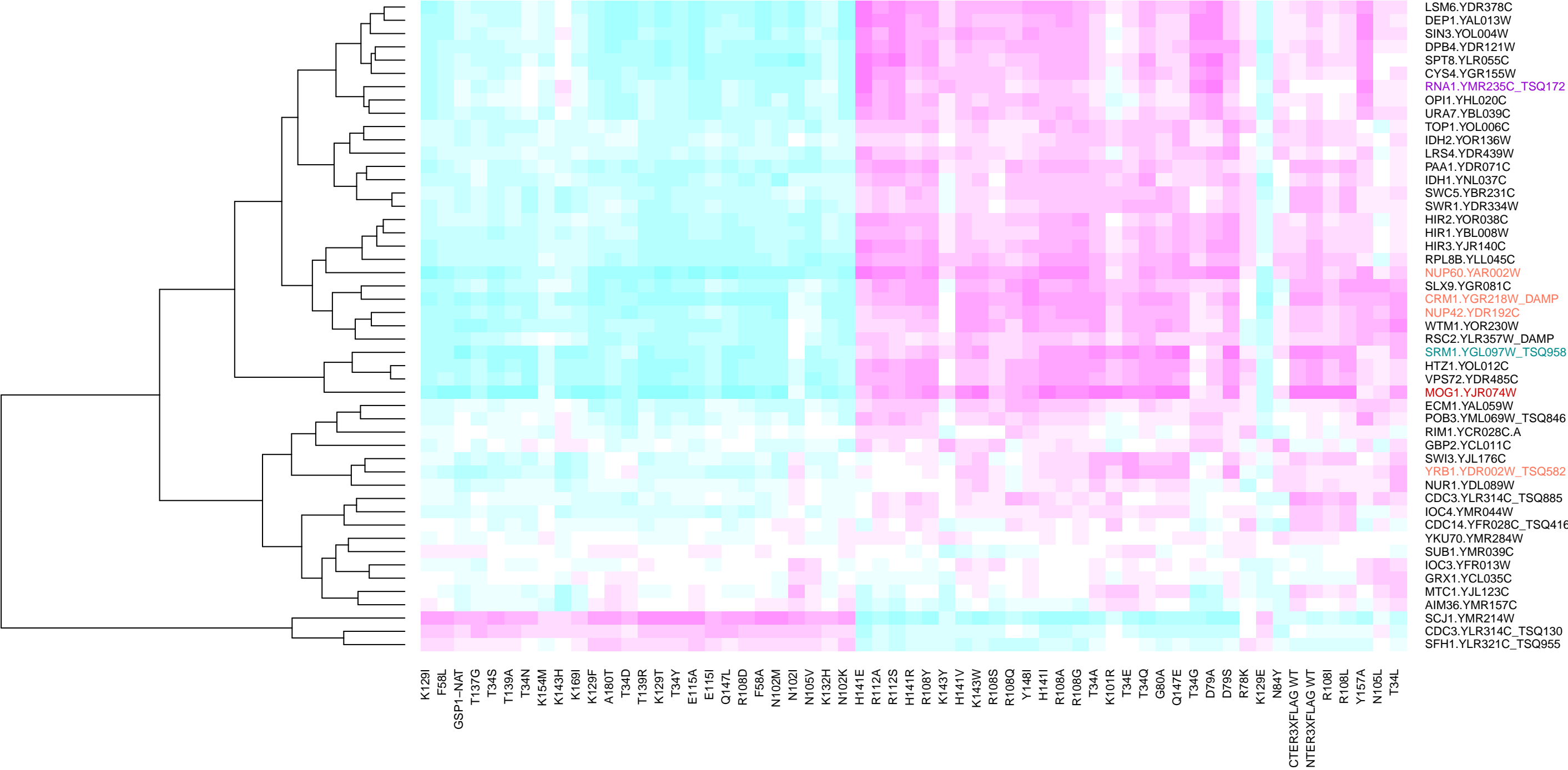
cytoskeleton organization_GO_15_3_all



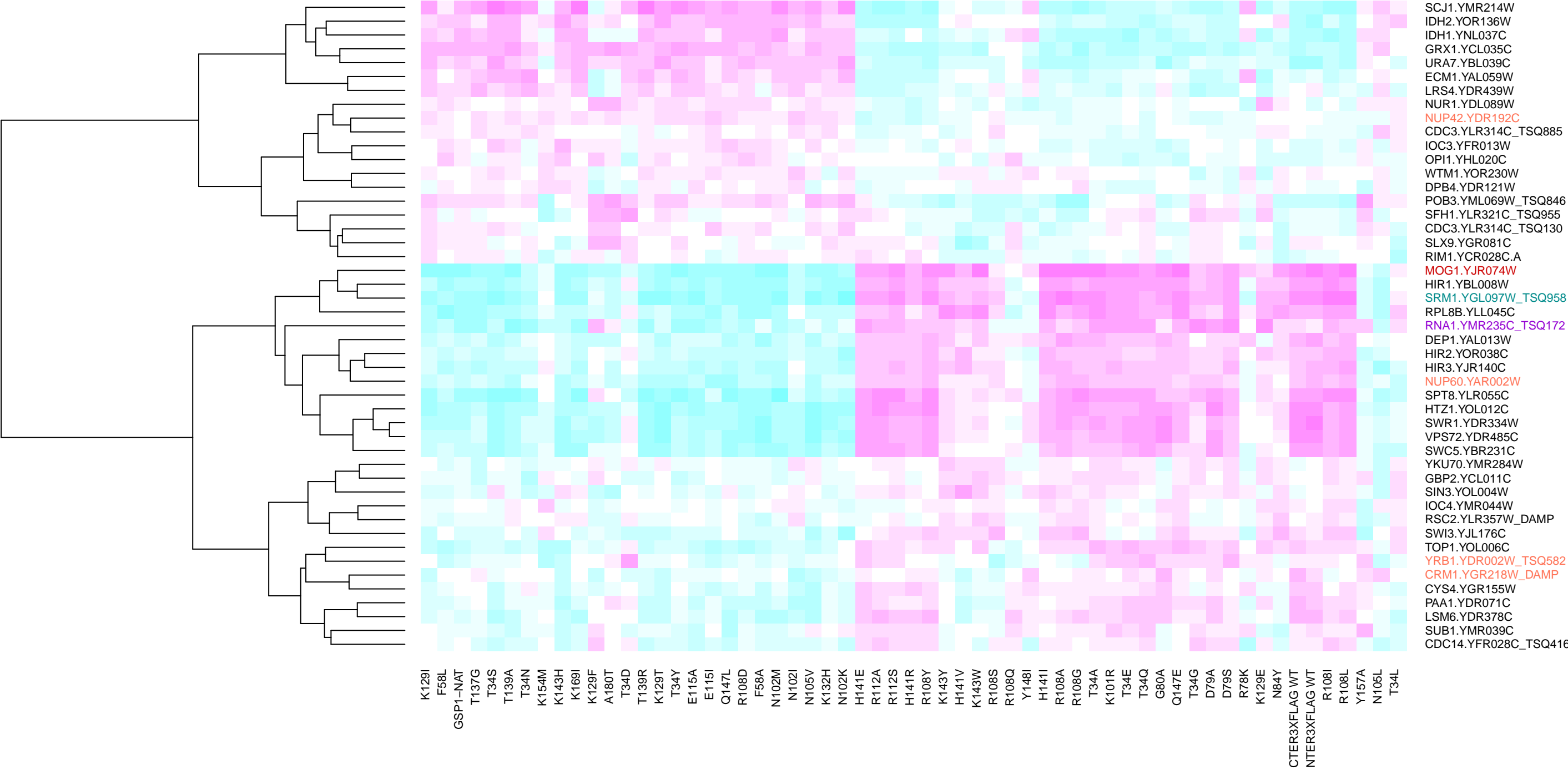
cytoskeleton organization_GO_30_1_mut



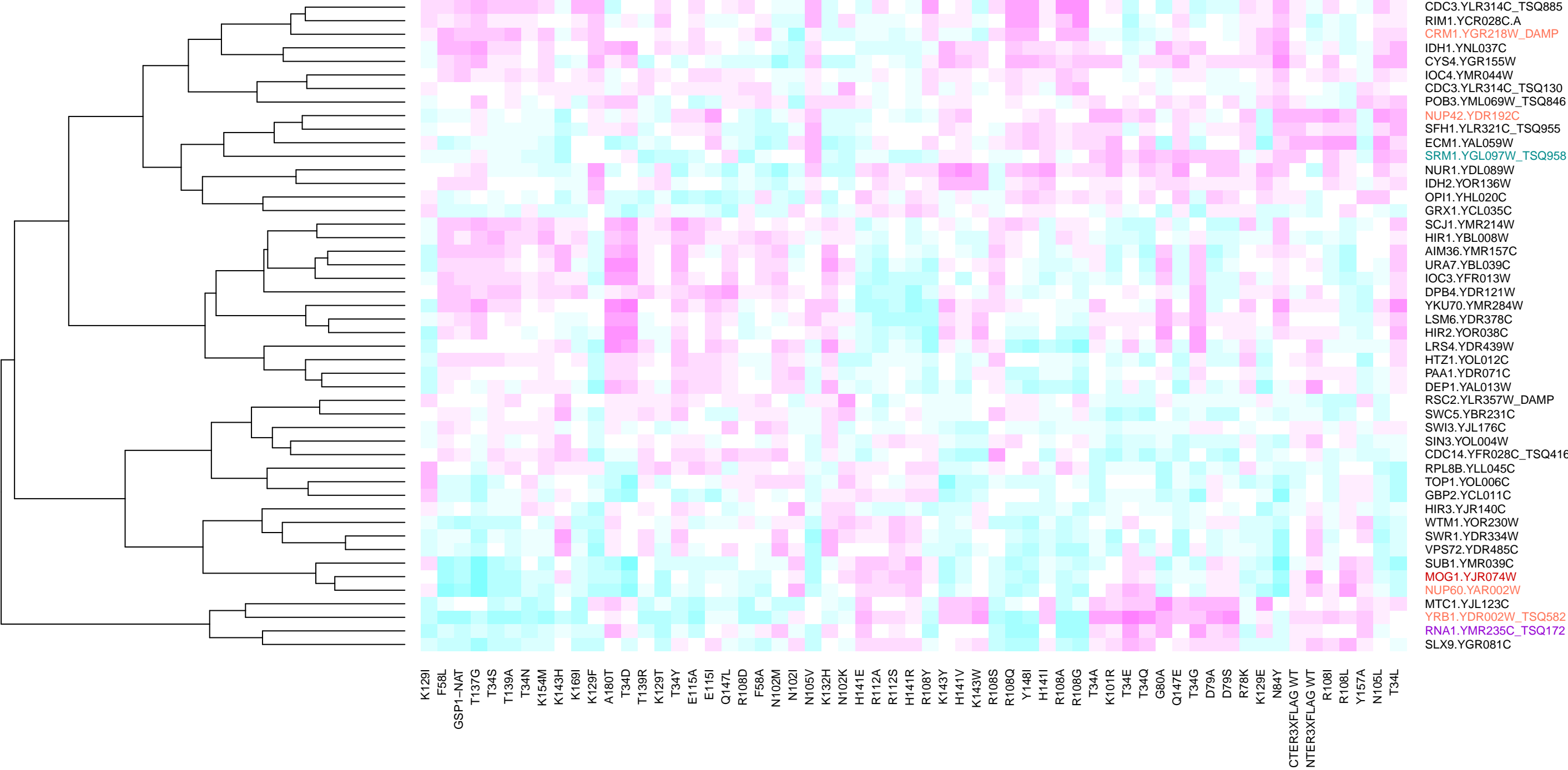
chromatin_GO_30_3_mut



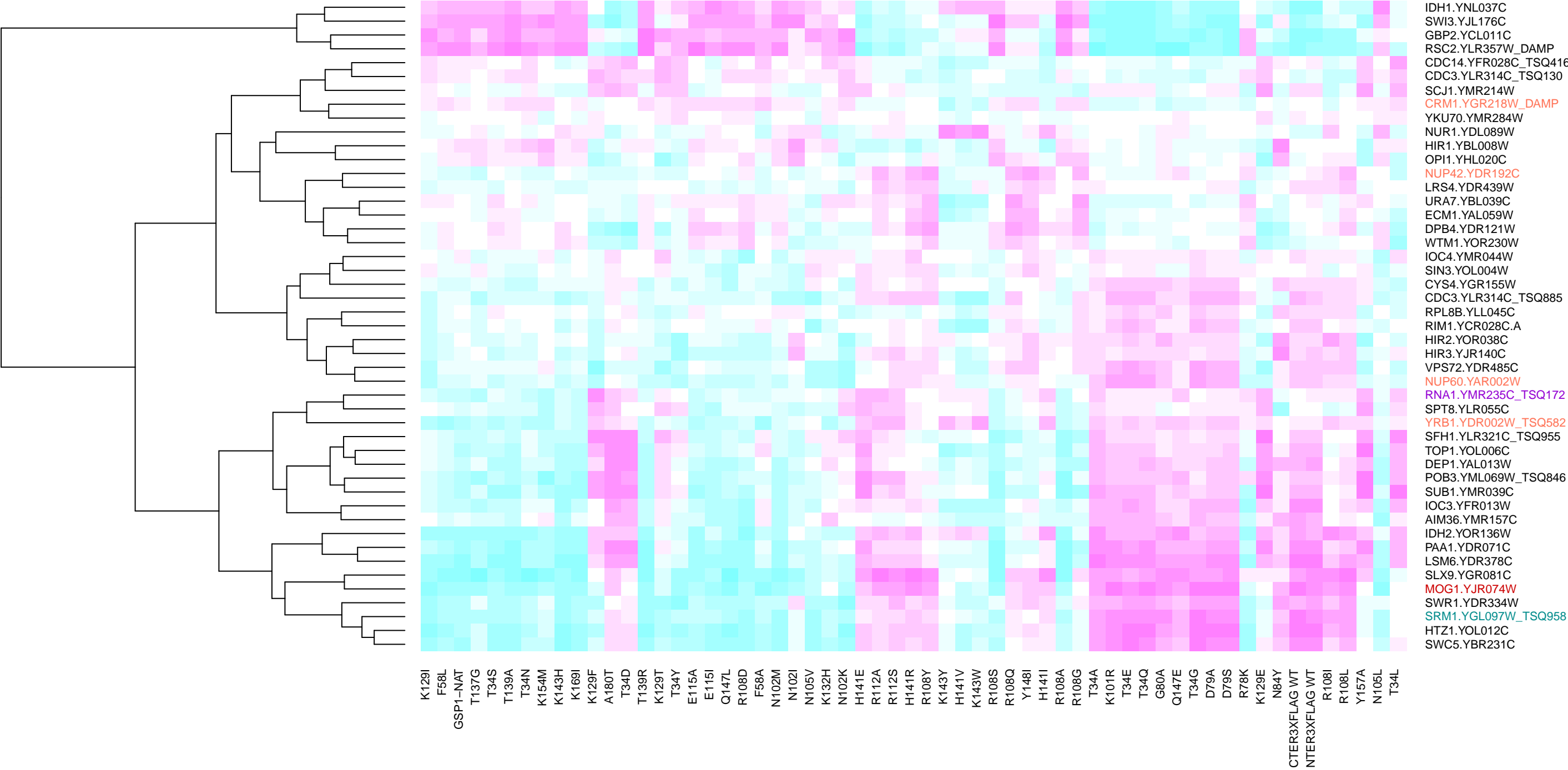
regulation of cell cycle_GO_15_2_mut



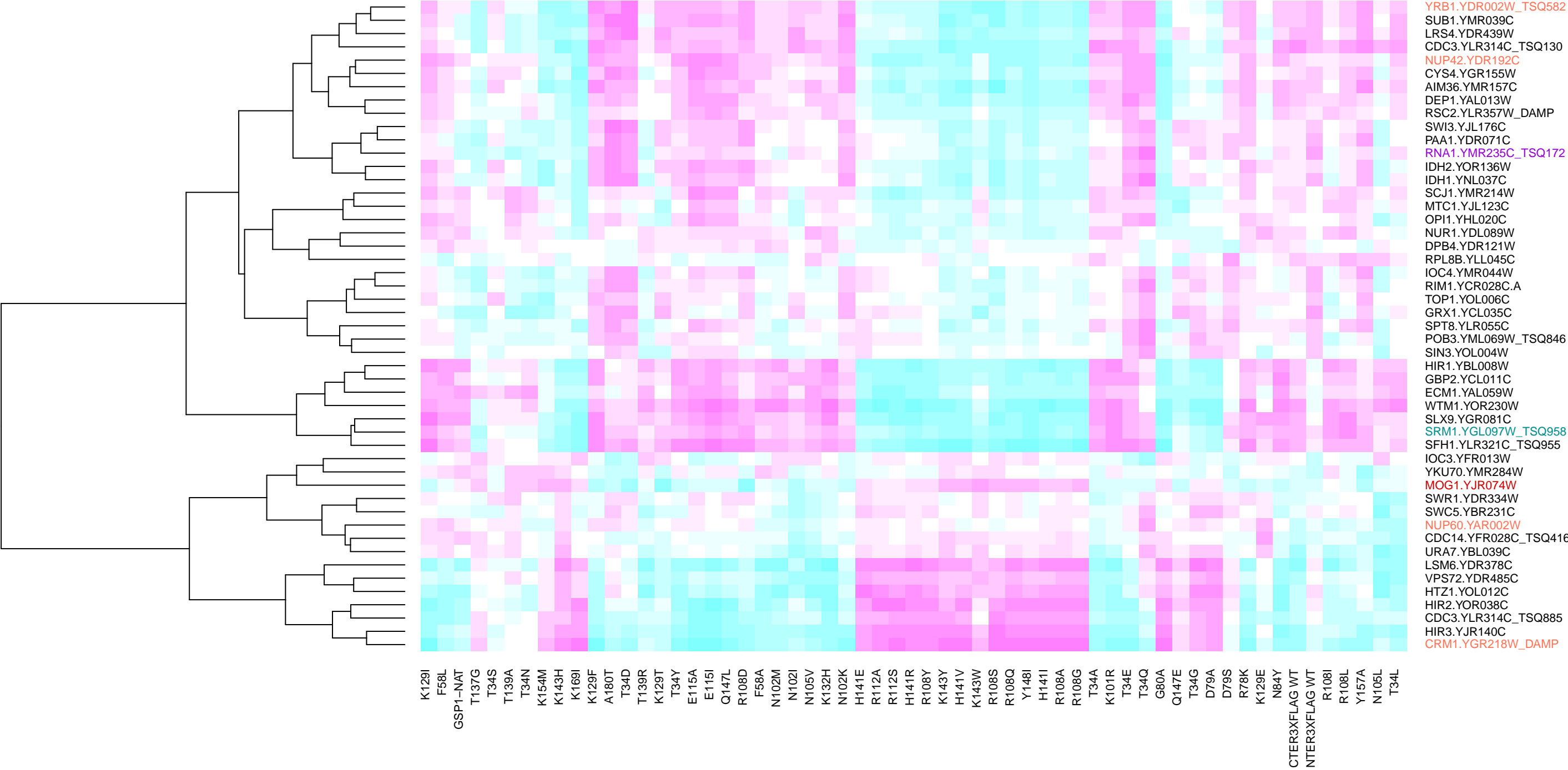
sig_Gsp1_Gl_15_5_all



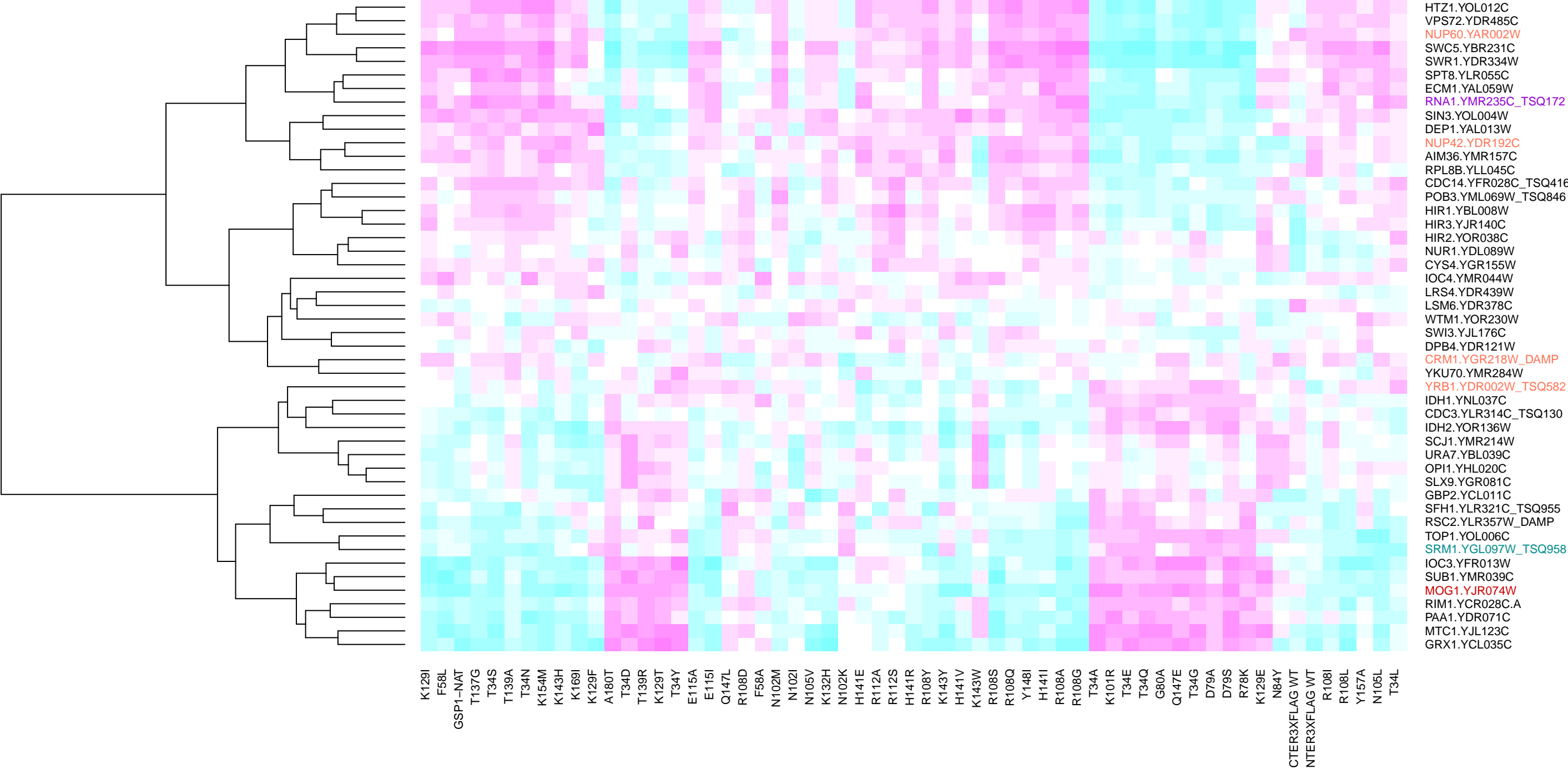
sig_Gsp1_Gl_15_6_mut



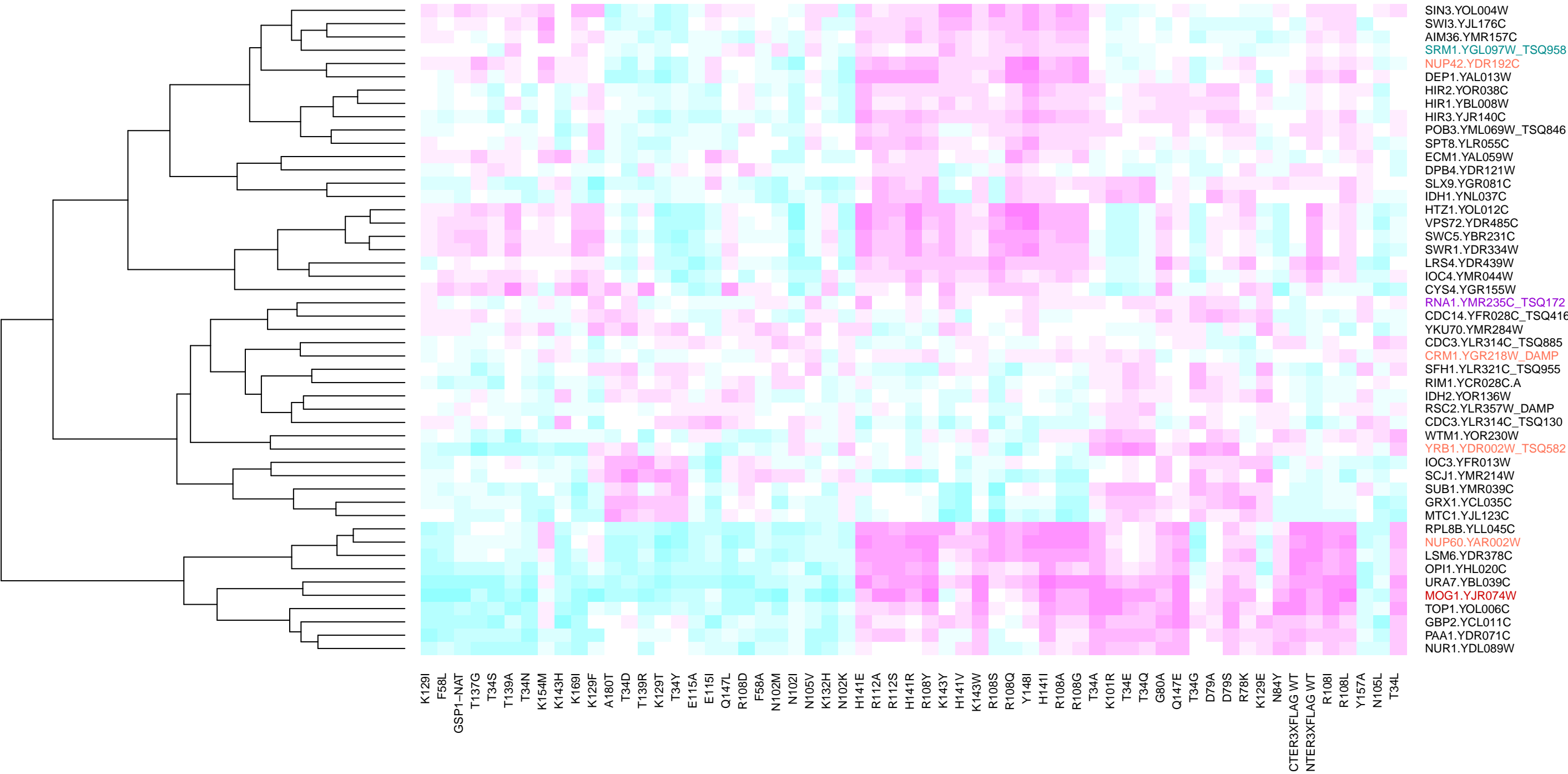
sig_Gsp1_Gl_15_3_mut



chromatin_GO_15_3_mut



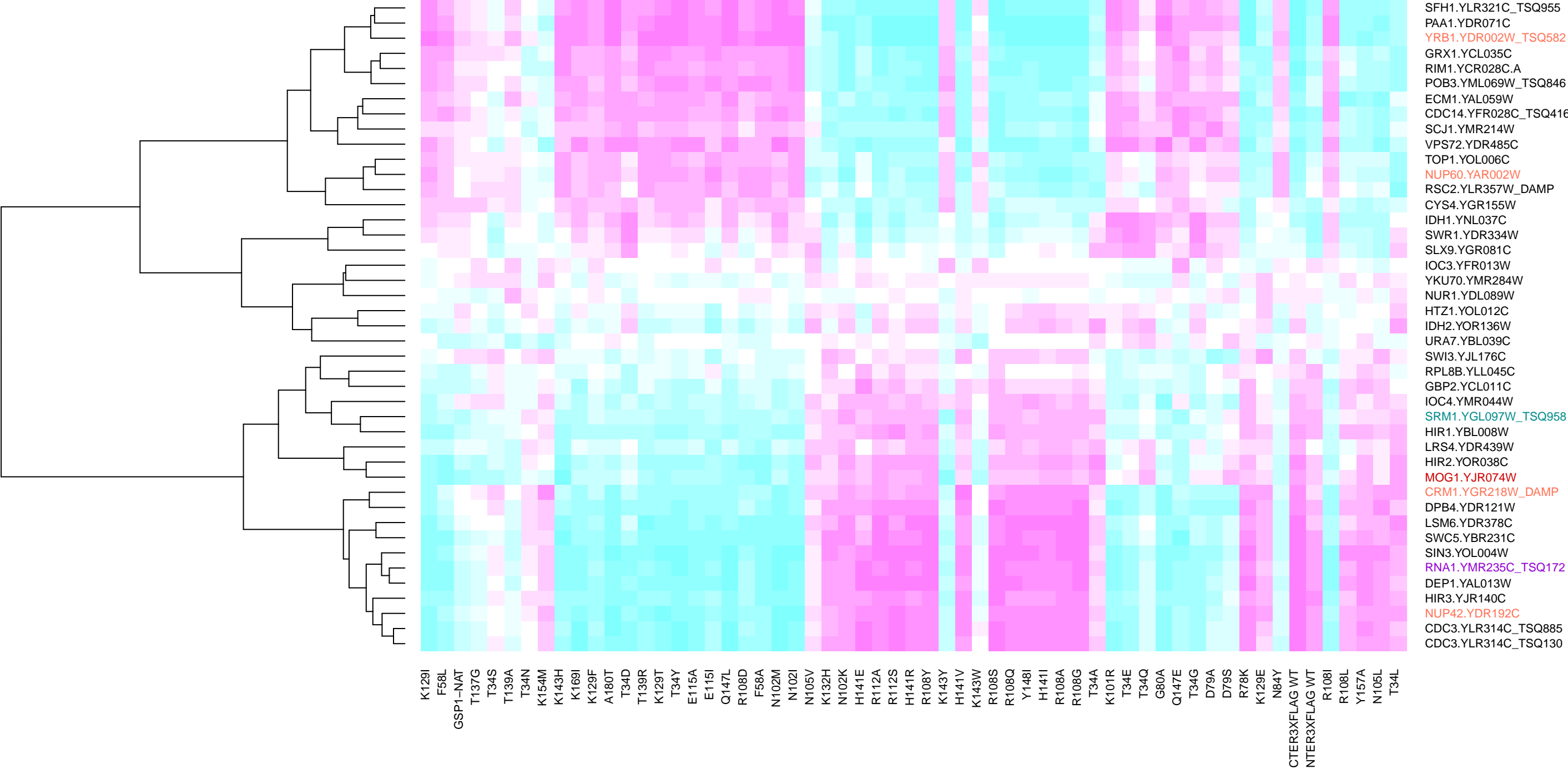
DNA-templated transcription, elongation_GO_15_1_all



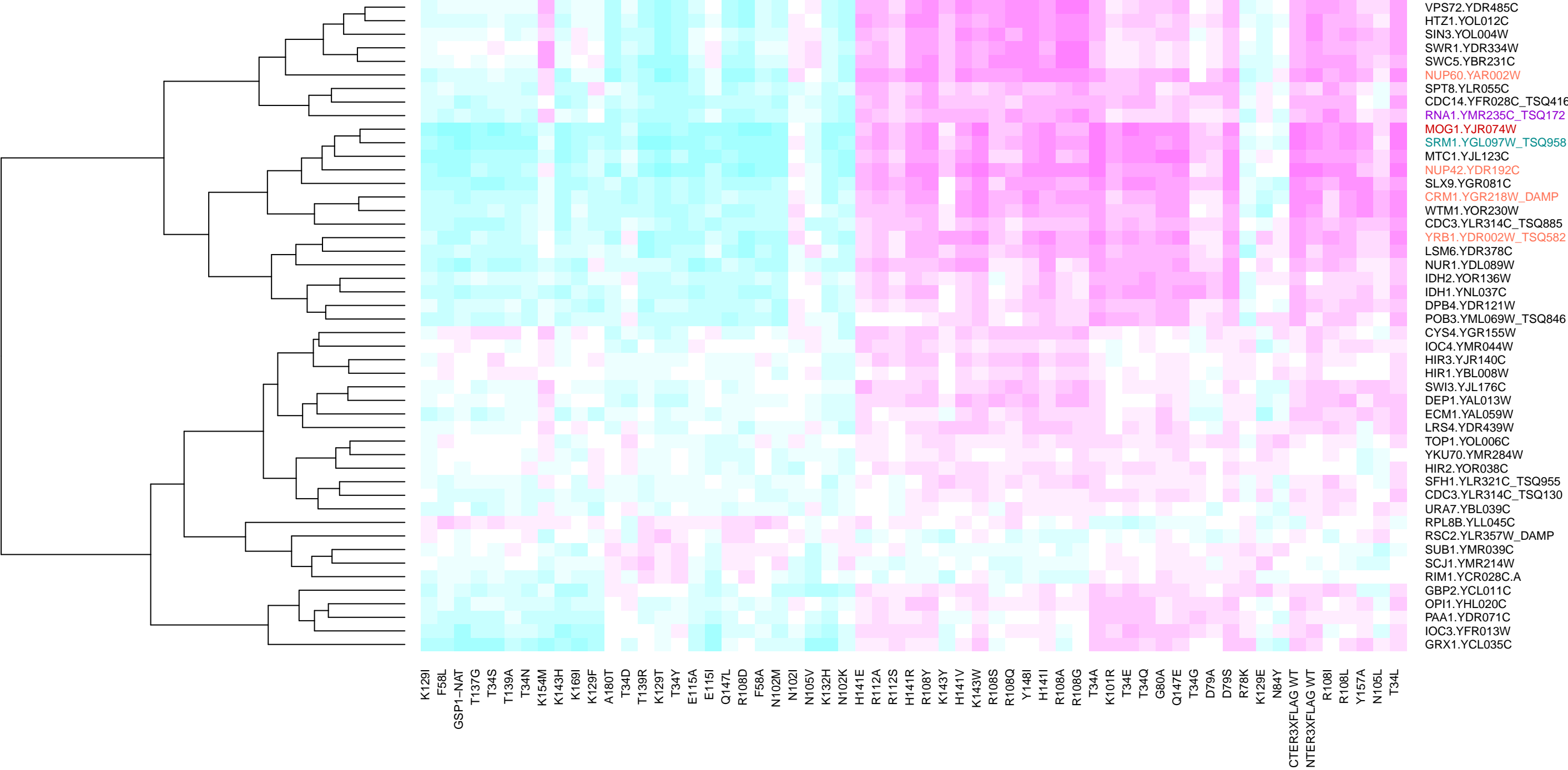
transcription and mRNA processing_GO_15_3_all



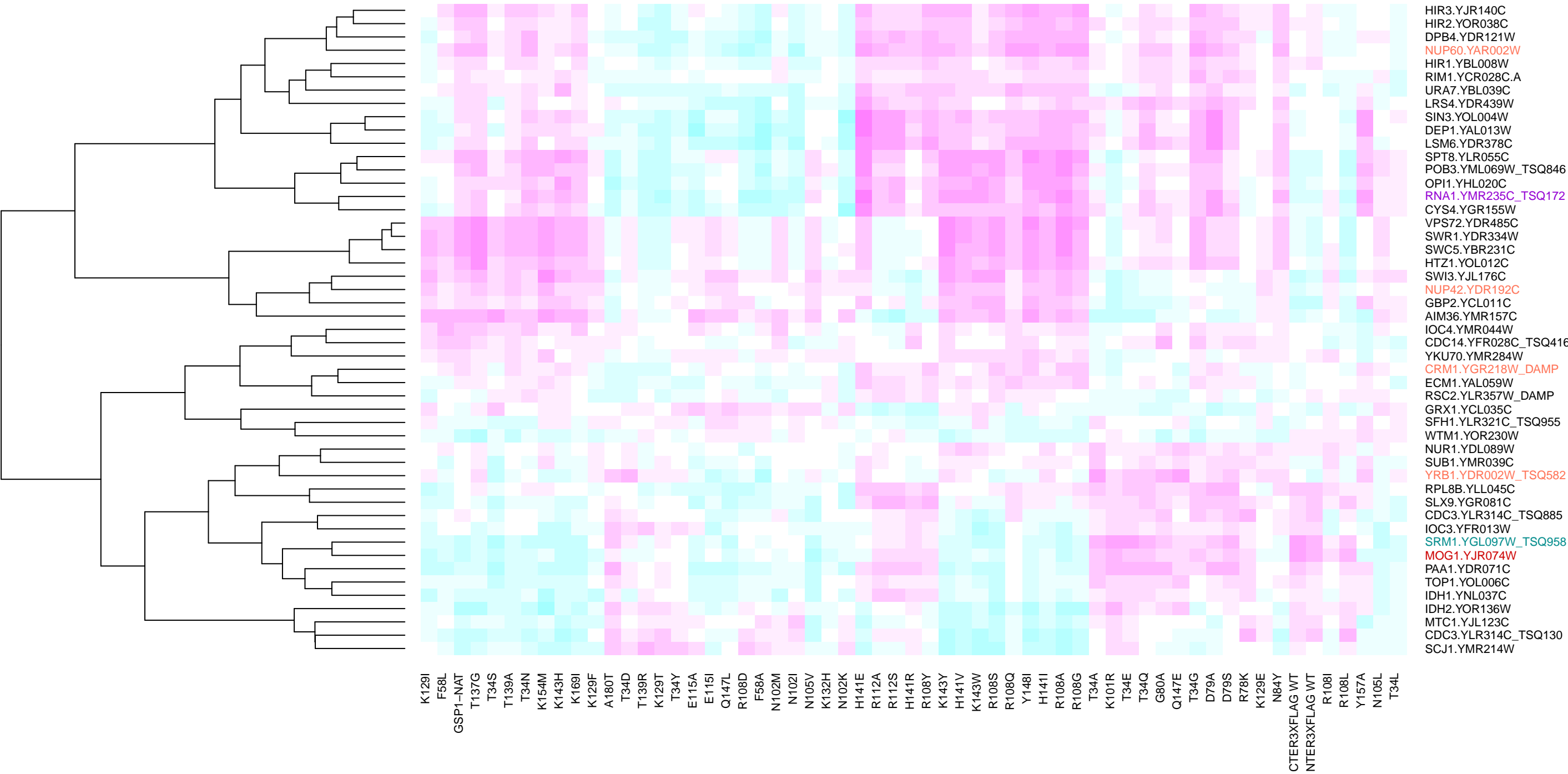
RNA modification_GO_15_1_mut



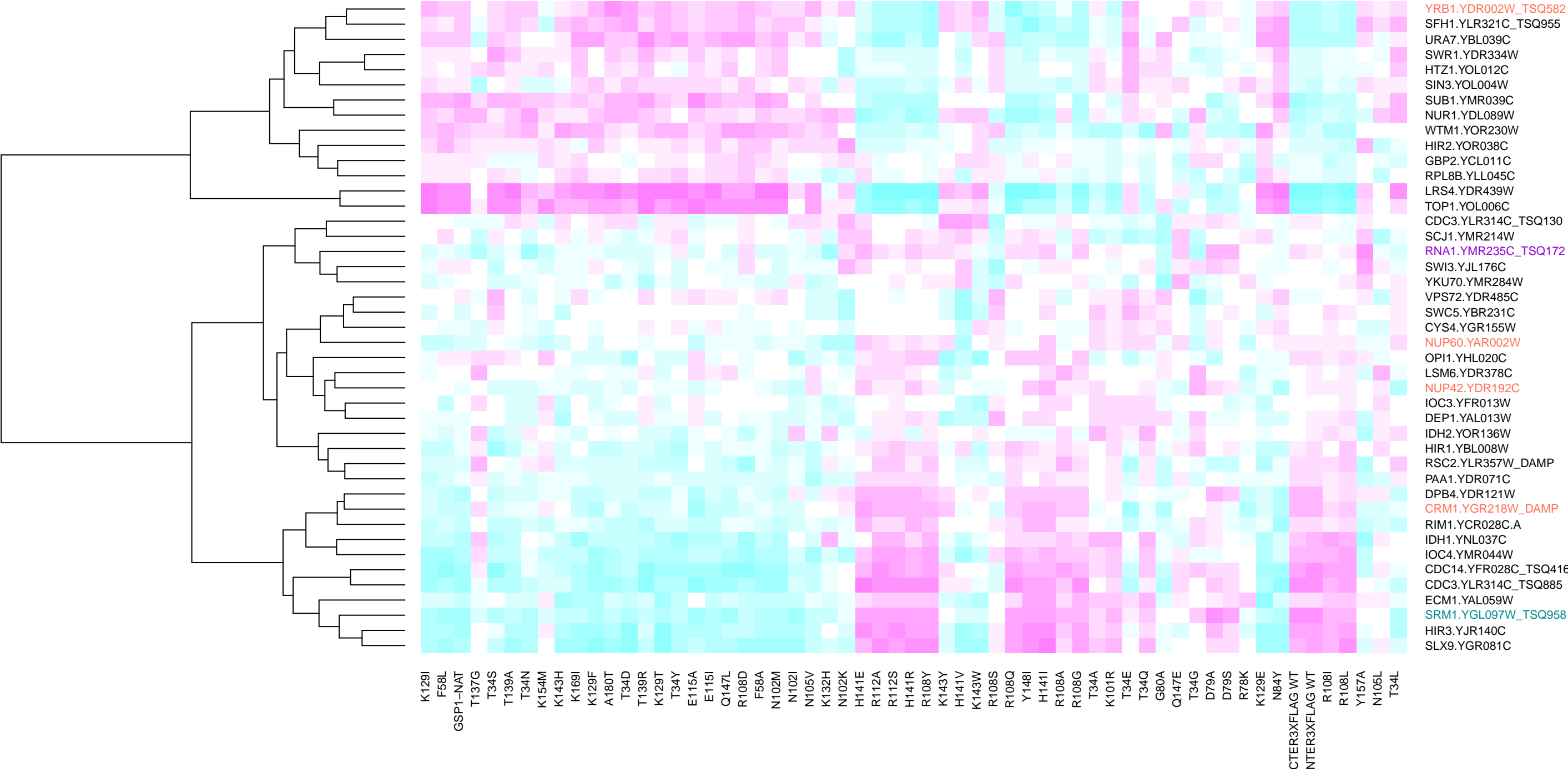
transcription and mRNA processing_GO_30_1_all



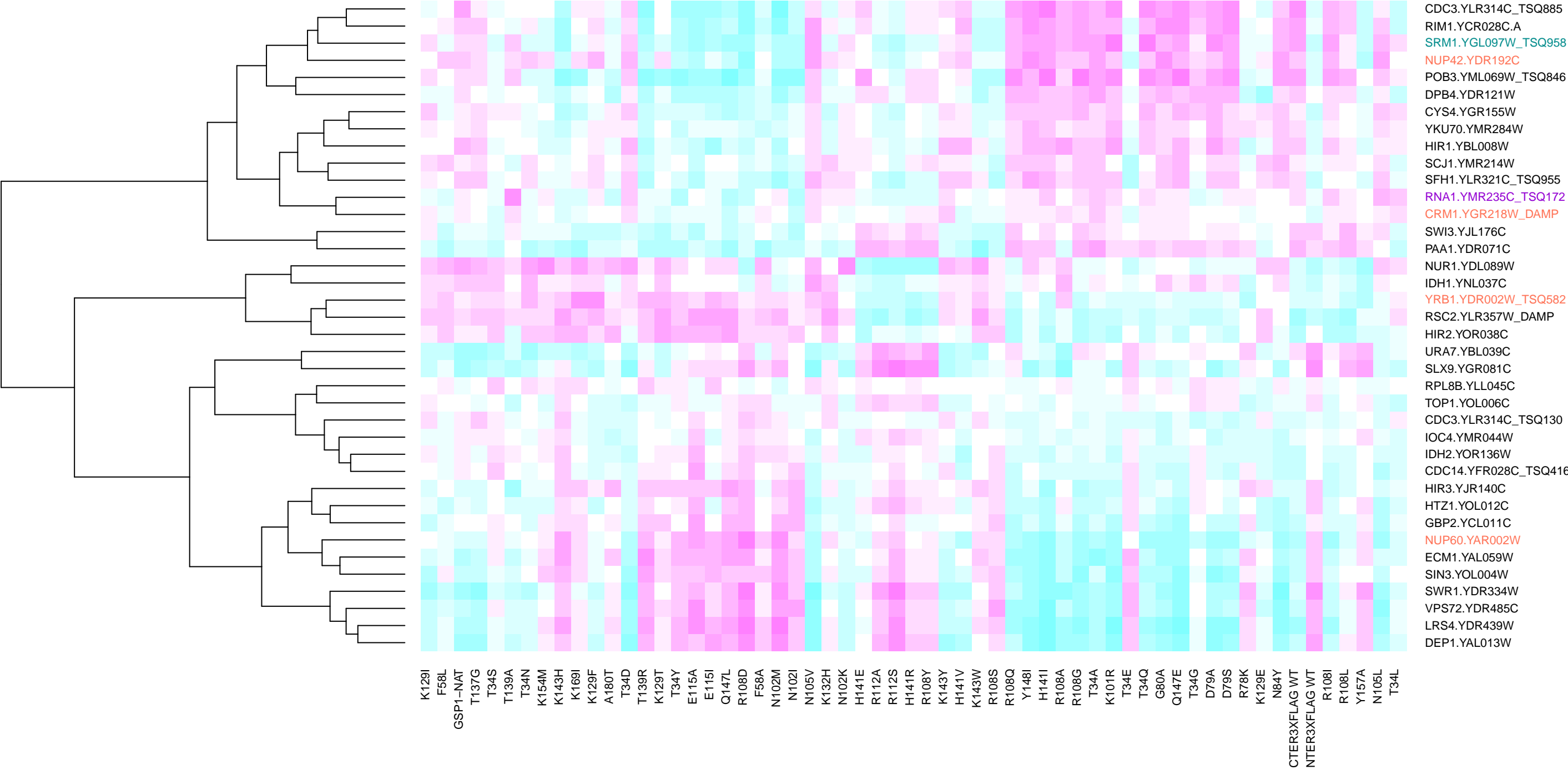
chromosome



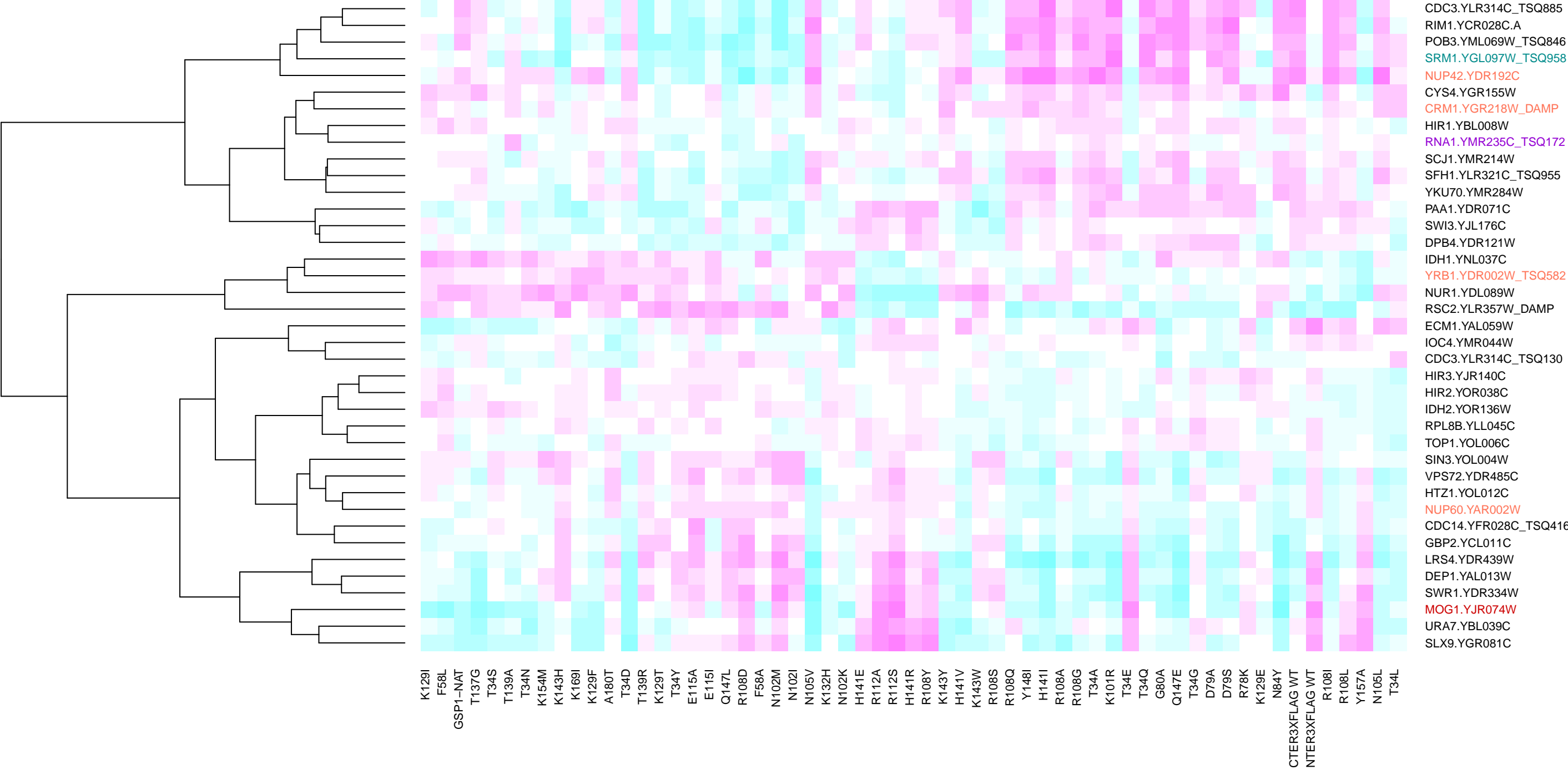
whole_library_15_24_mut



cytoplasmic translation



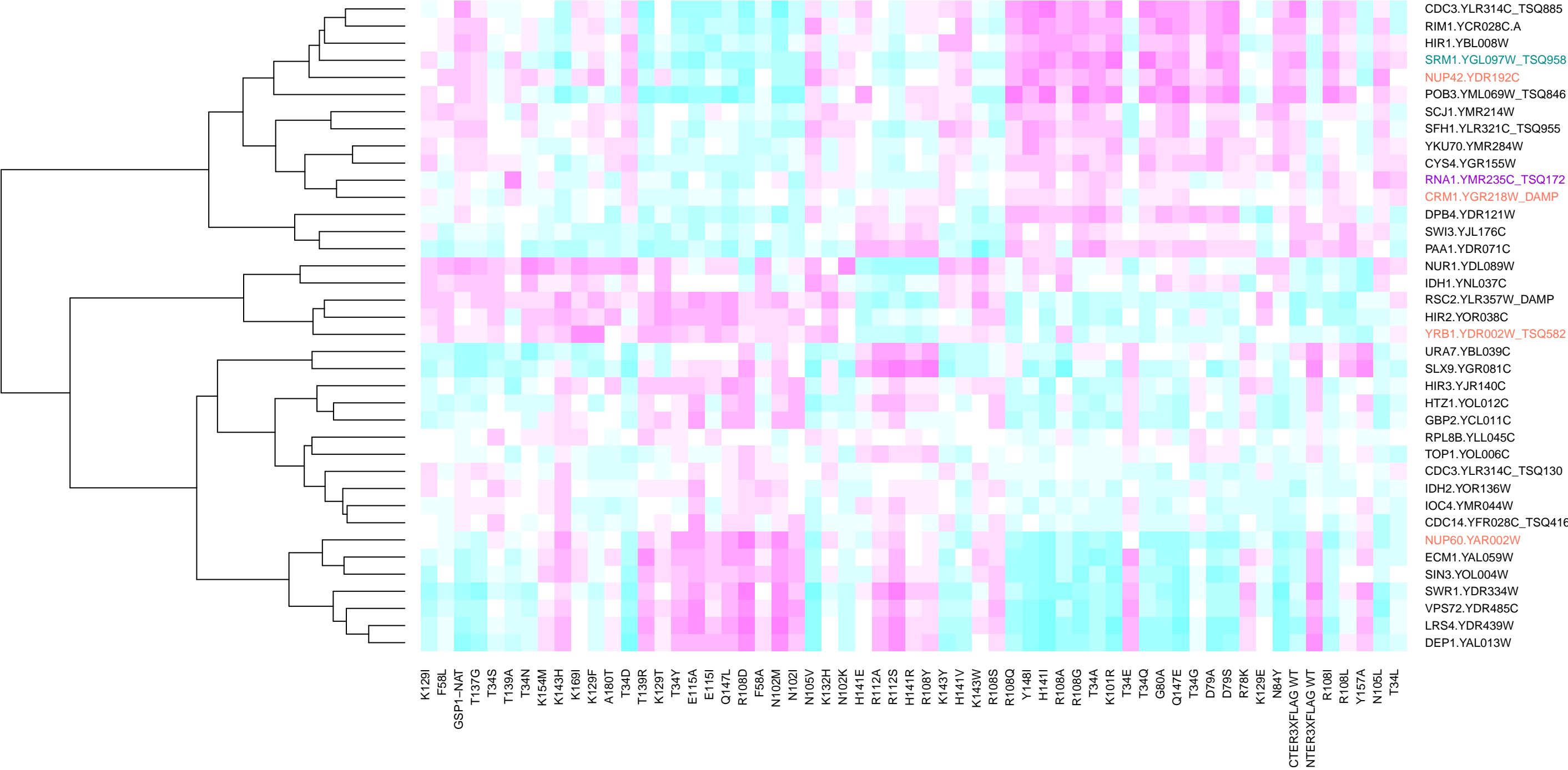
ribosome



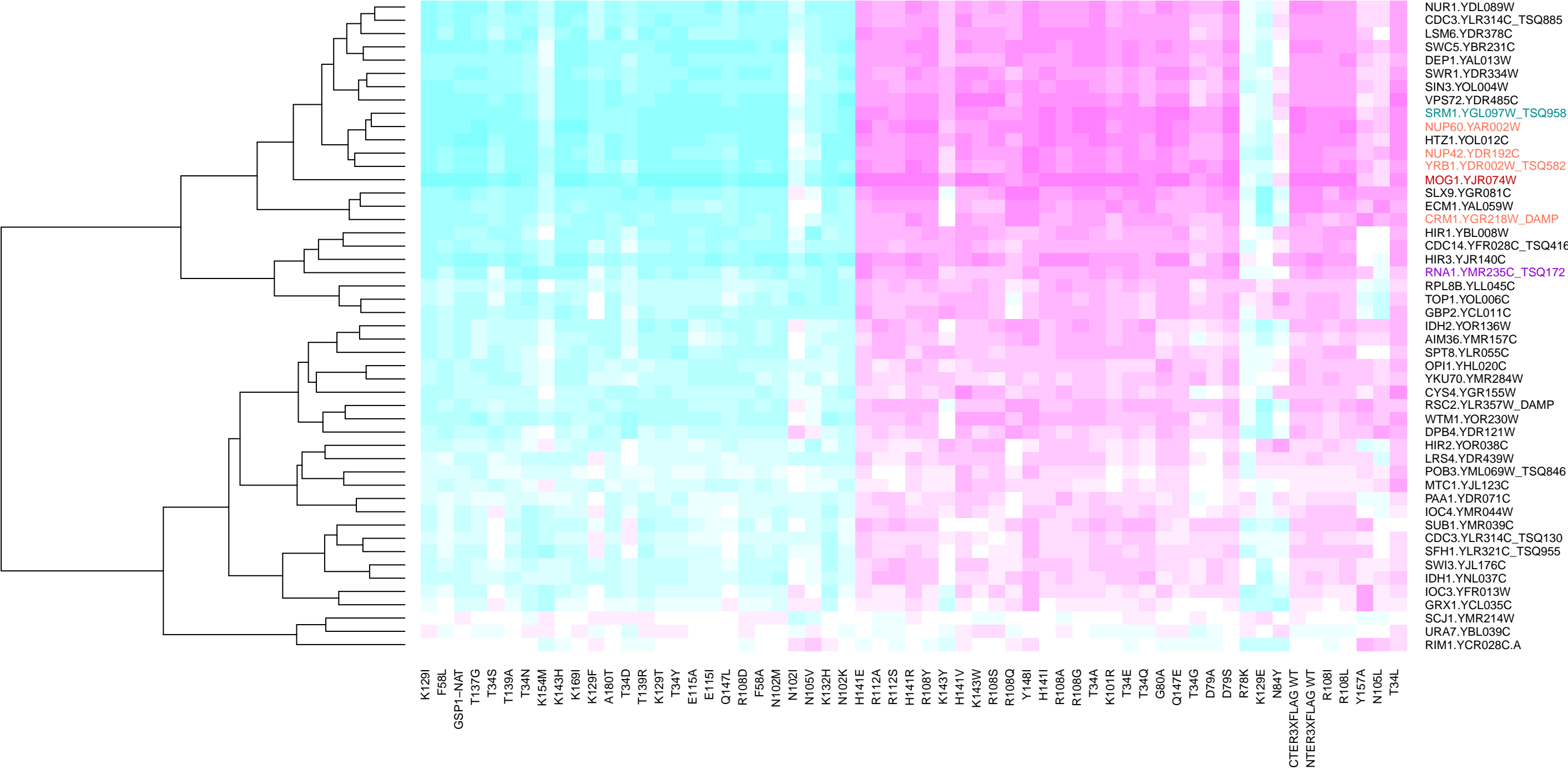
ribosome_GO_15_1_all



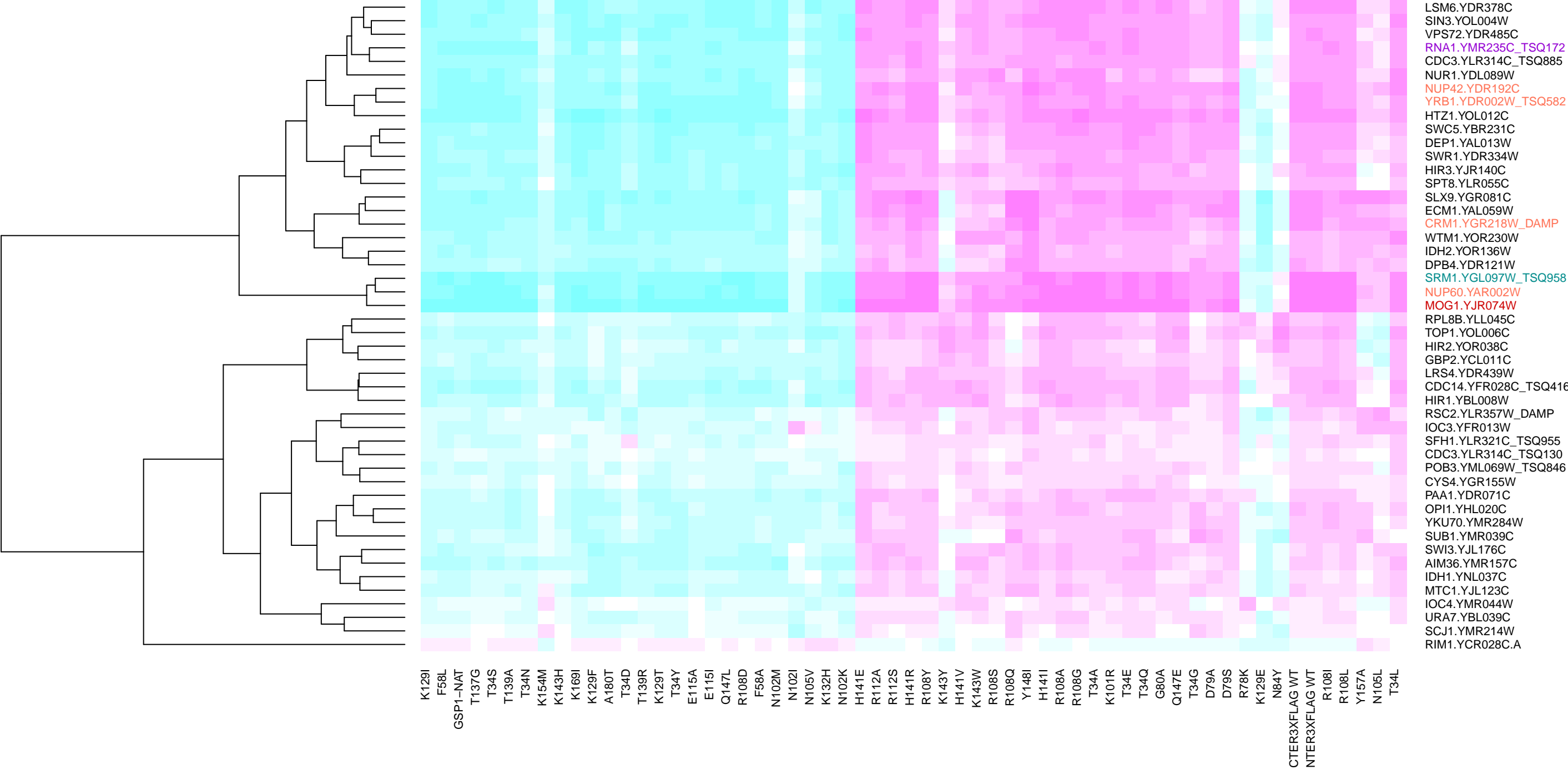
structural constituent of ribosome



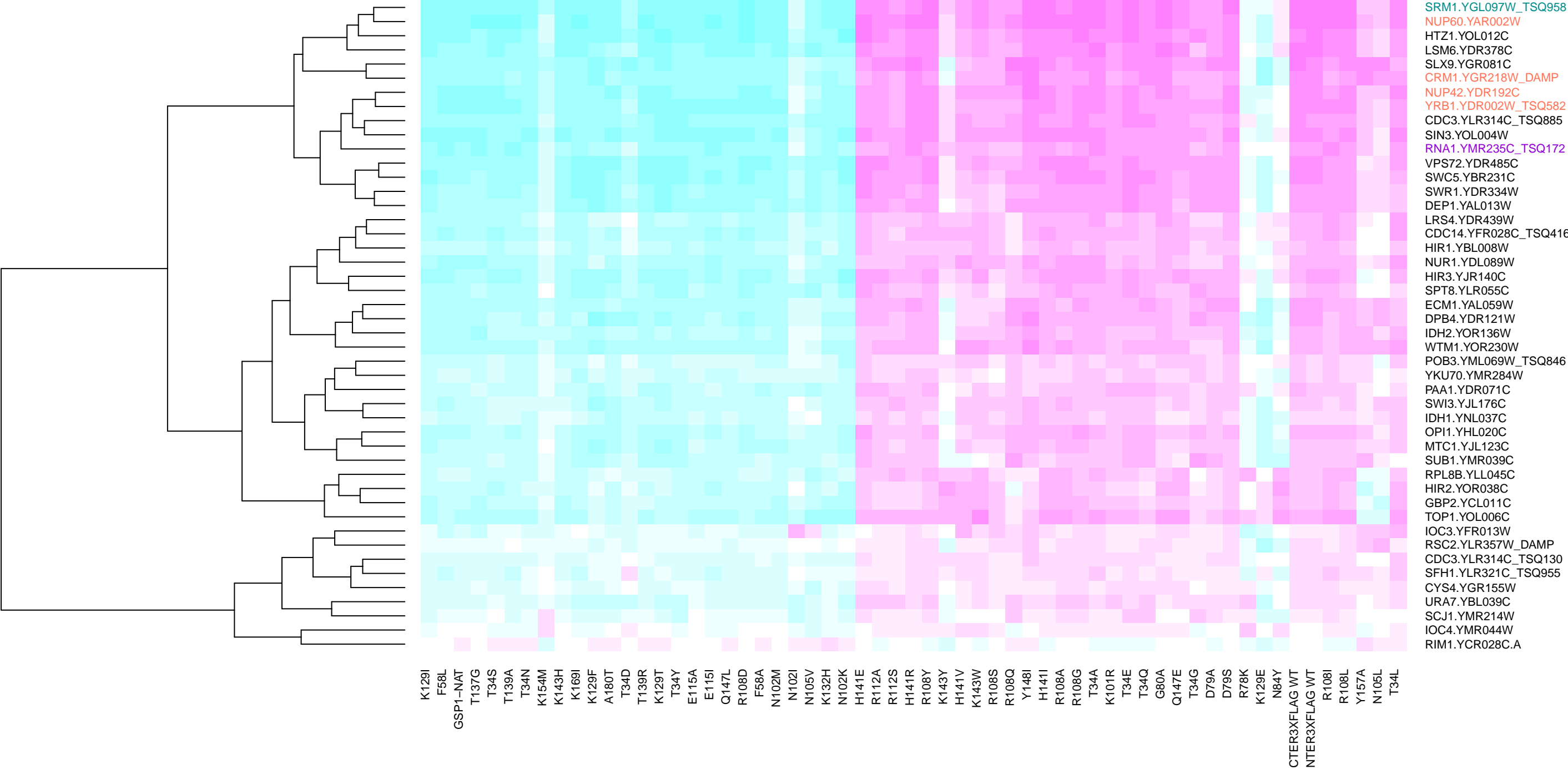
sig_Gsp1_Gl_15_4_mut



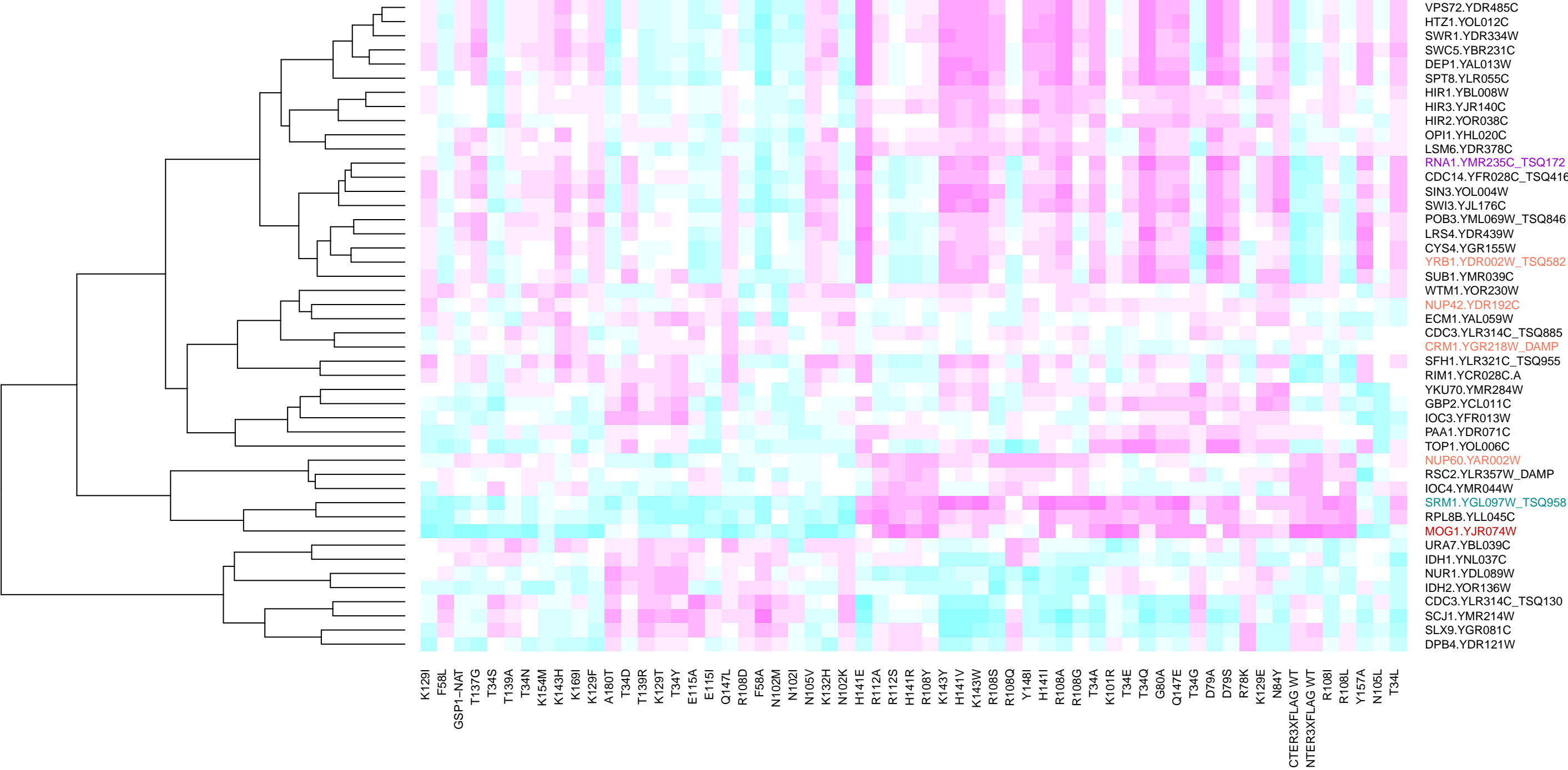
whole_library_15_14_mut



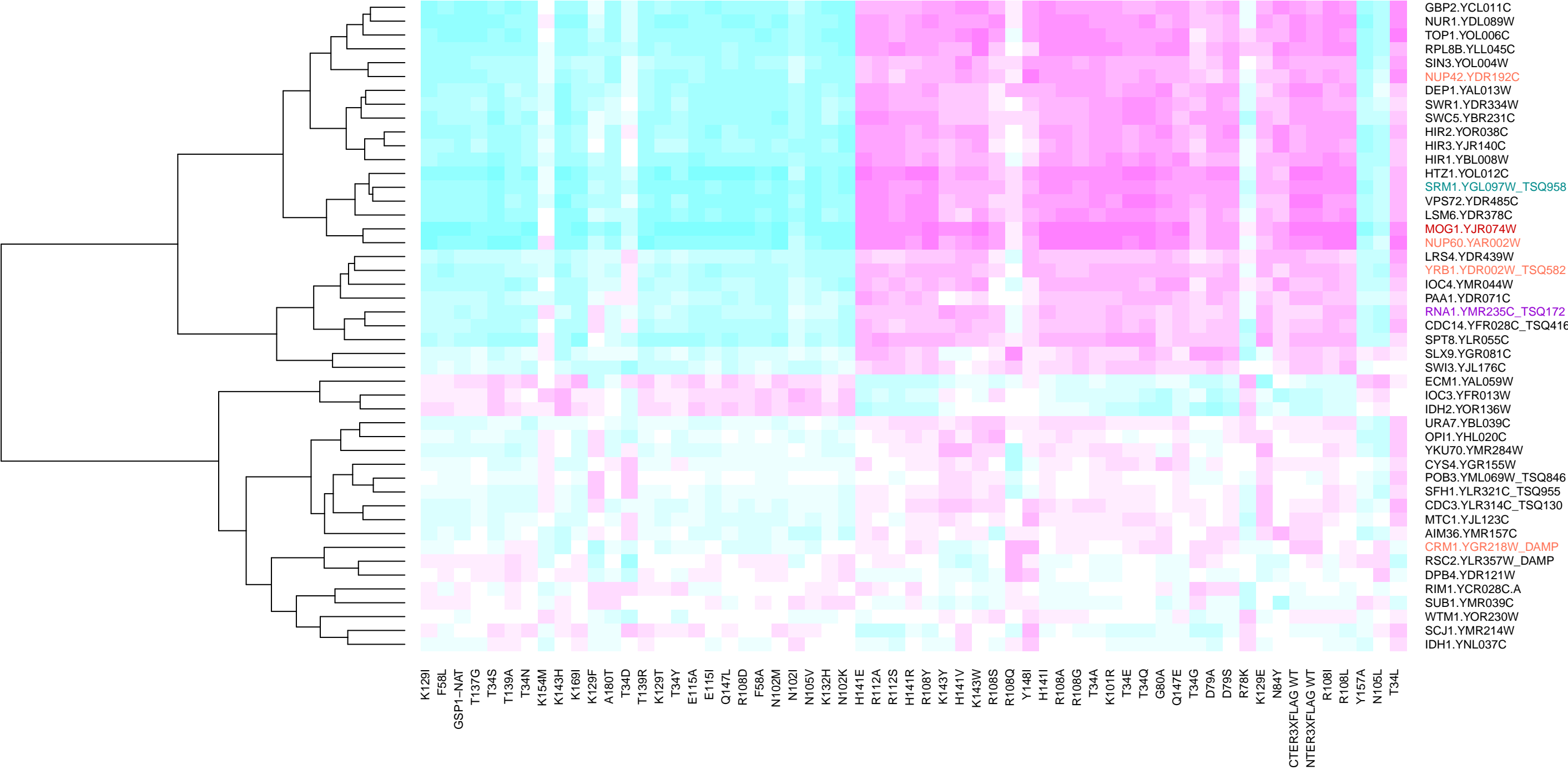
whole_library_30_12_mut



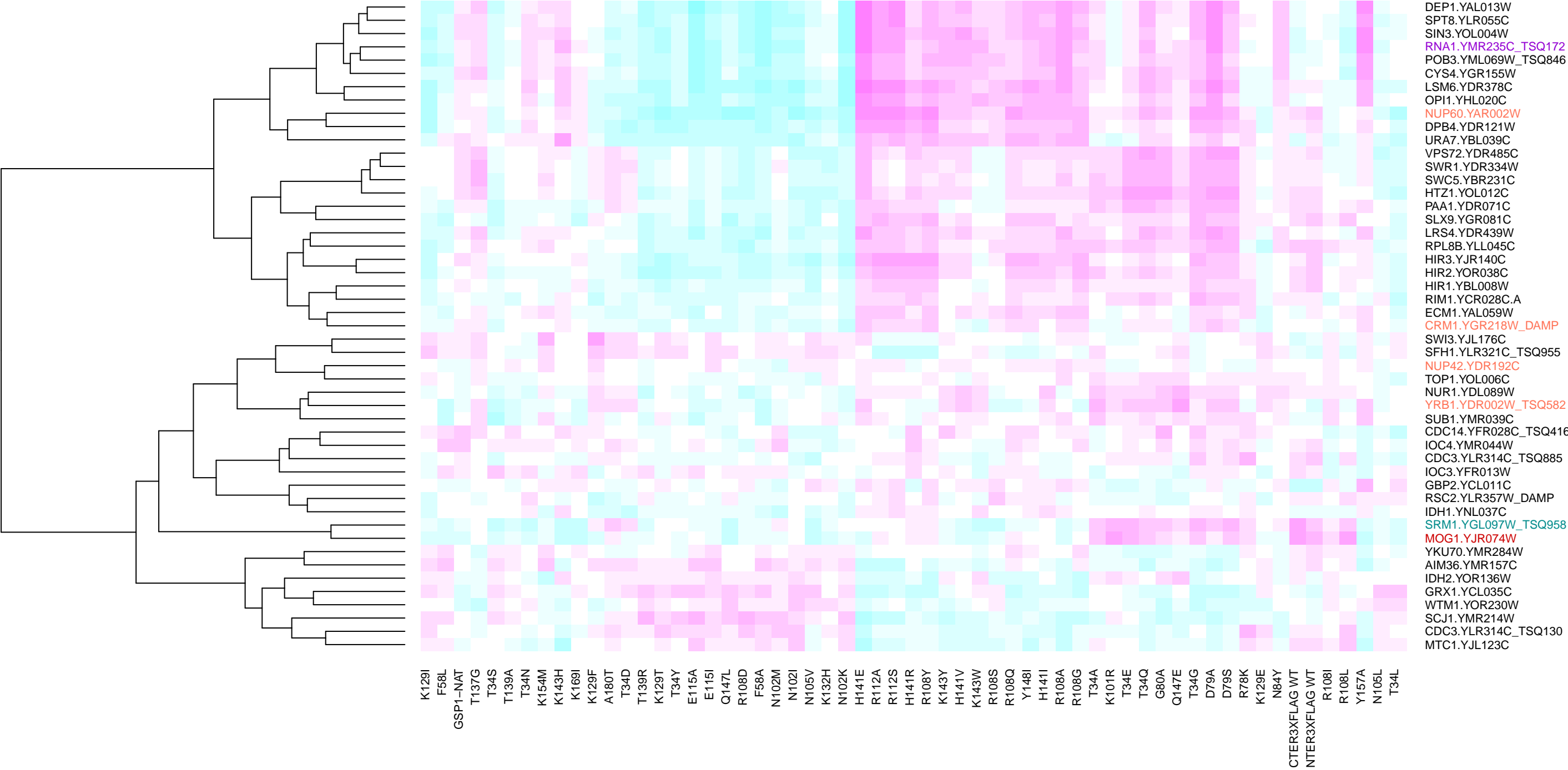
histone modification_GO_15_1_all



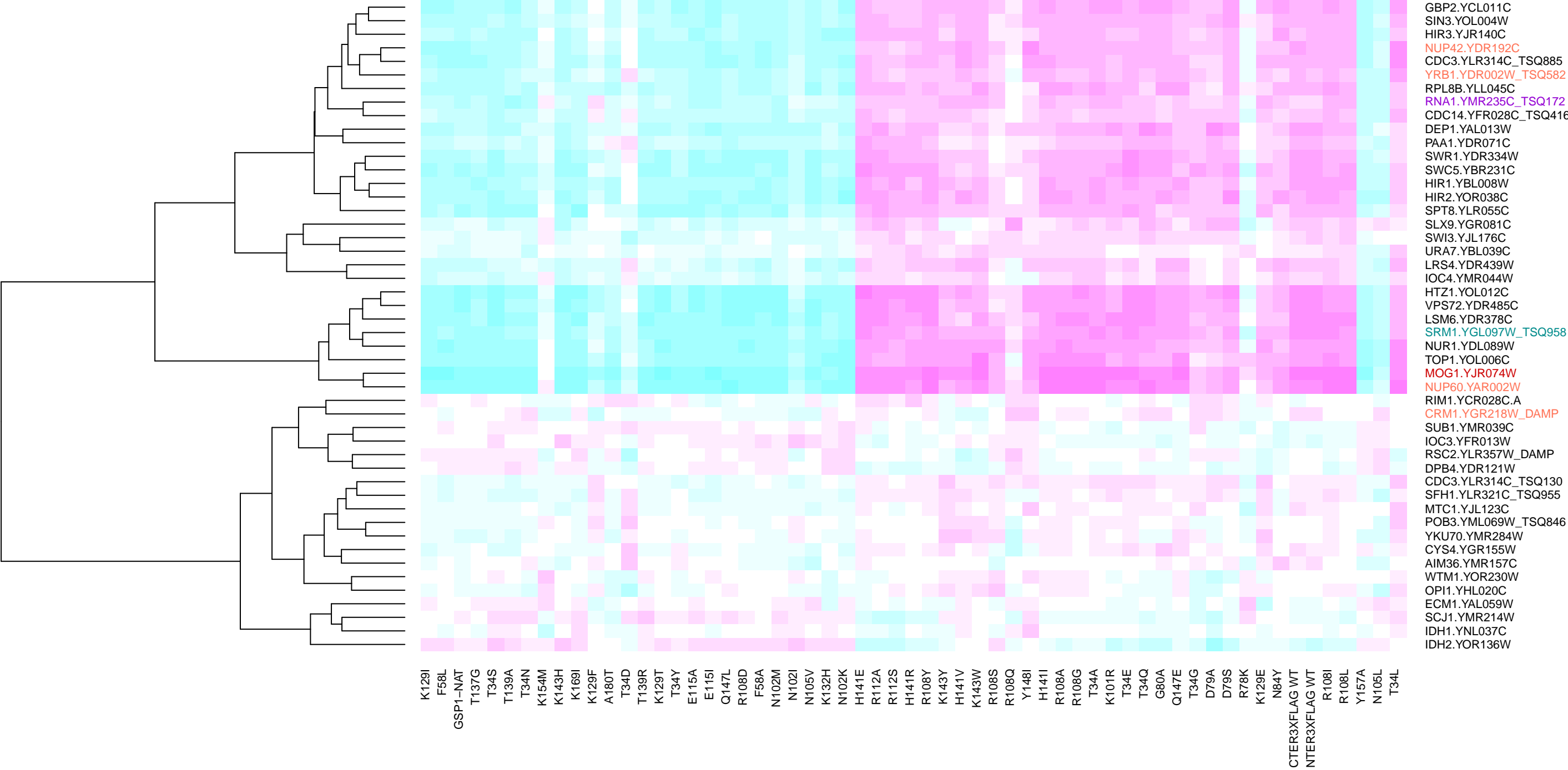
cell cycle_GO_15_9_mut



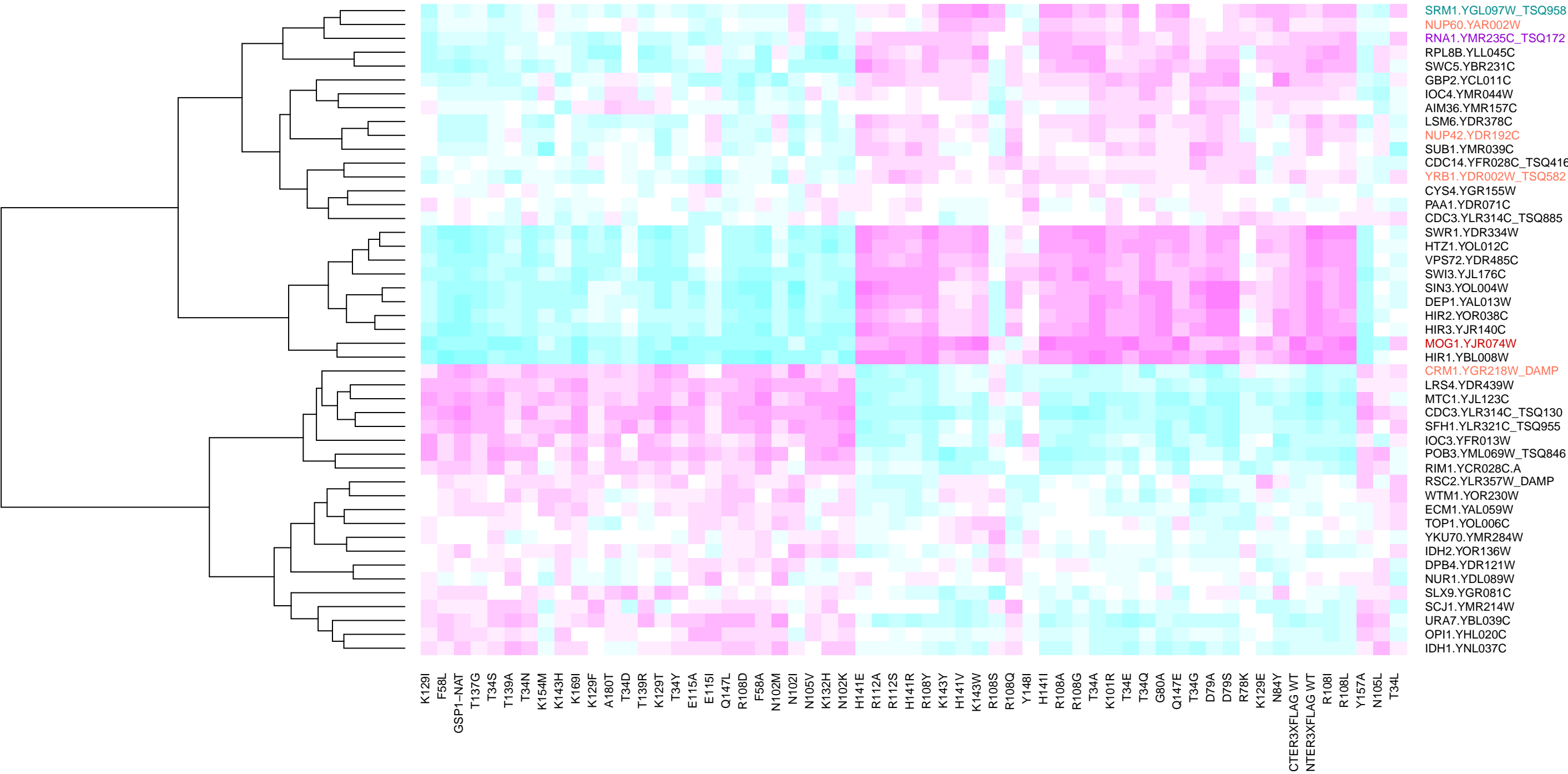
chromosome_GO_30_1_mut



cell cycle_GO_30_3_mut



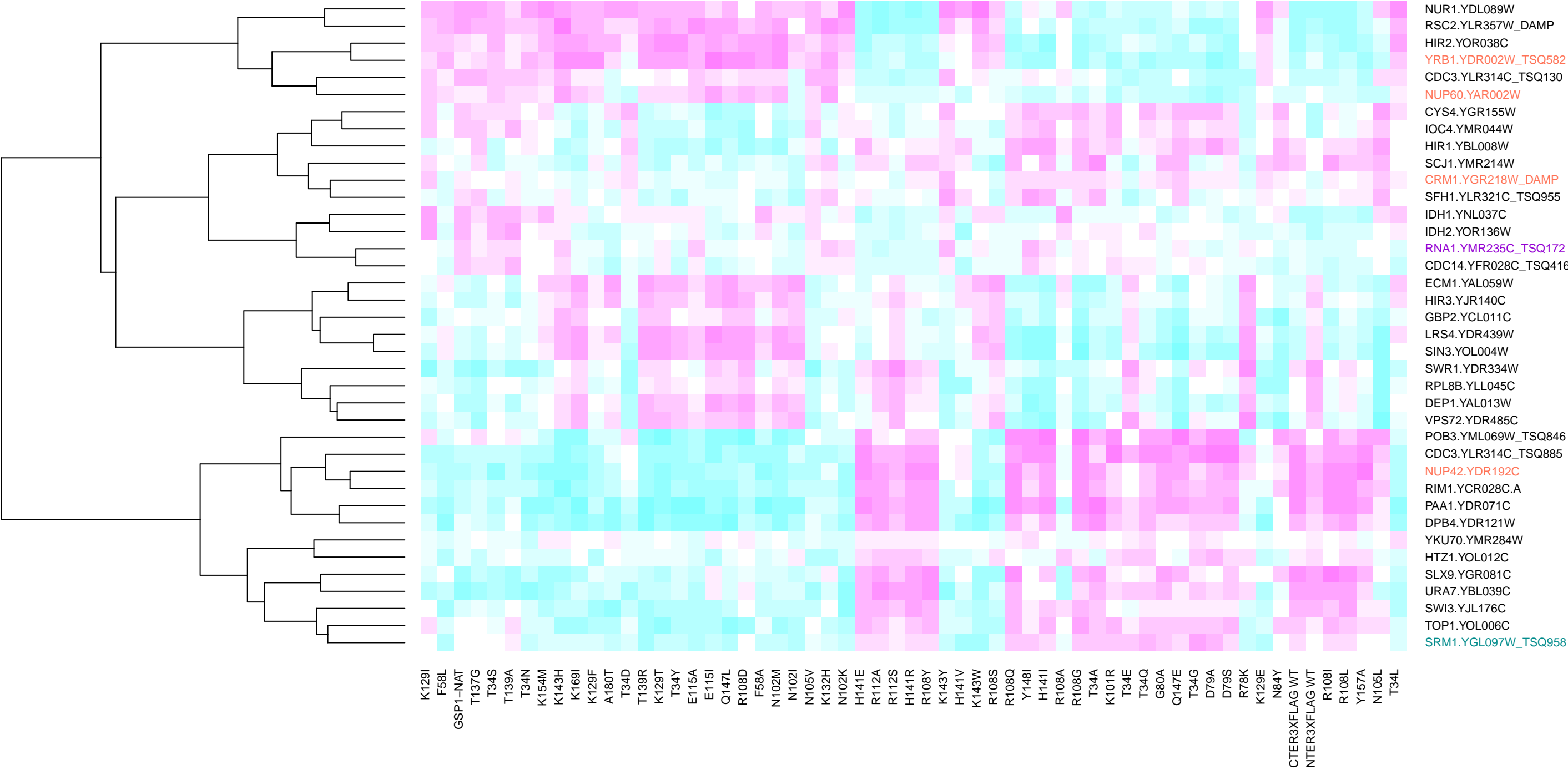
proteolysis involved in cellular protein catabolic process_GO_15_2_mut



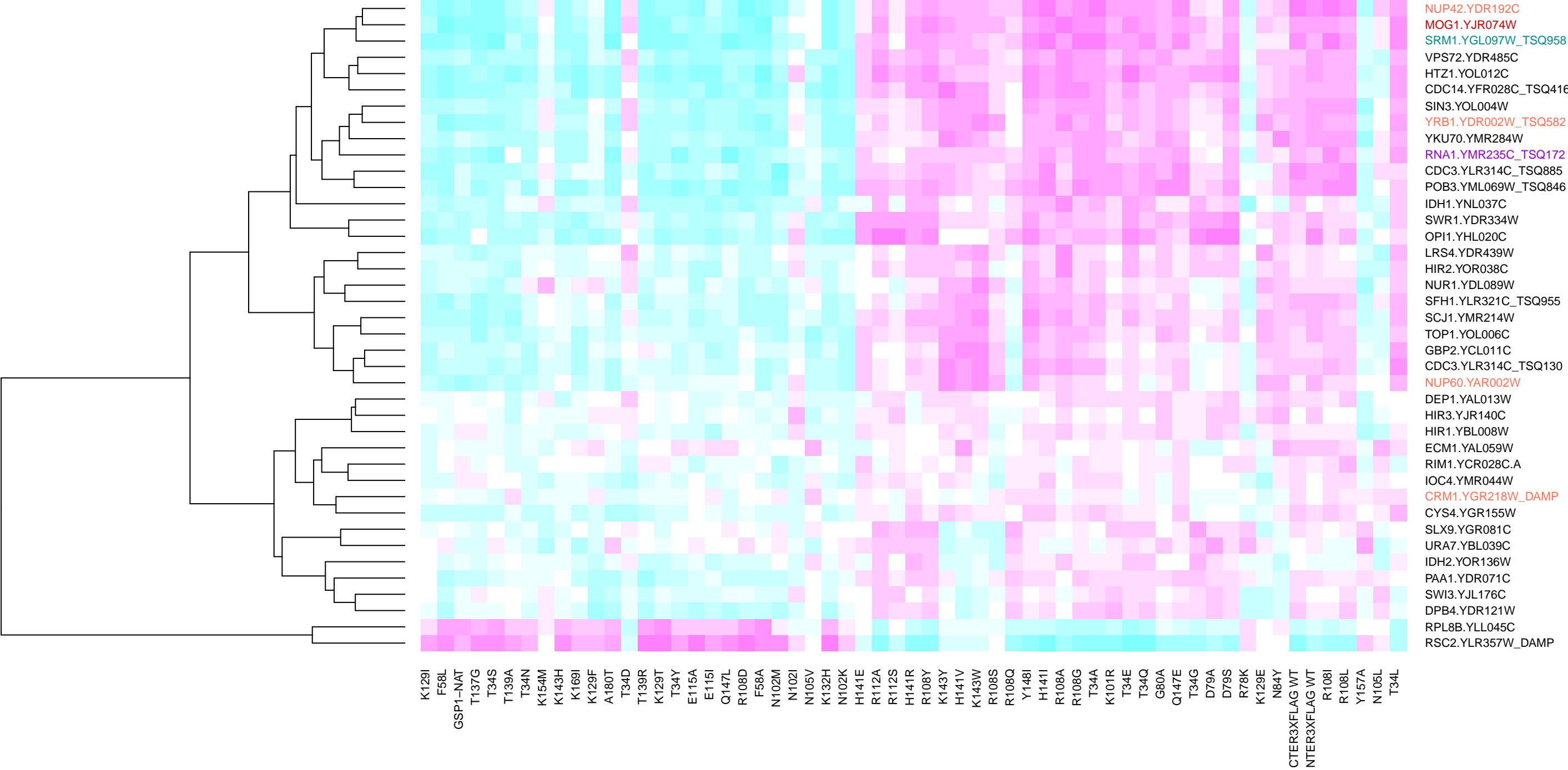
sig_Gsp1_Gl_15_1_all



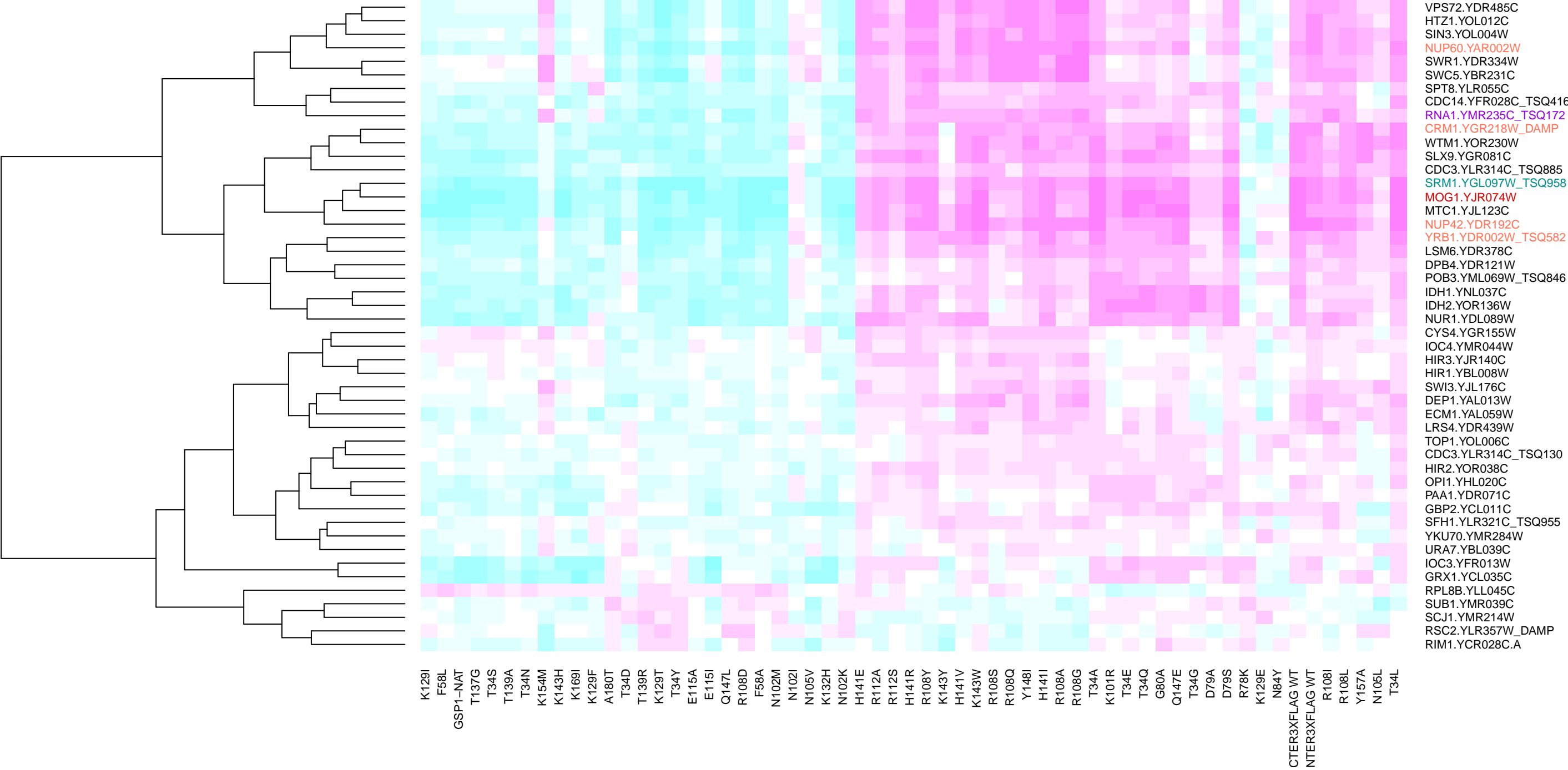
cytoplasmic translation_GO_15_1_all



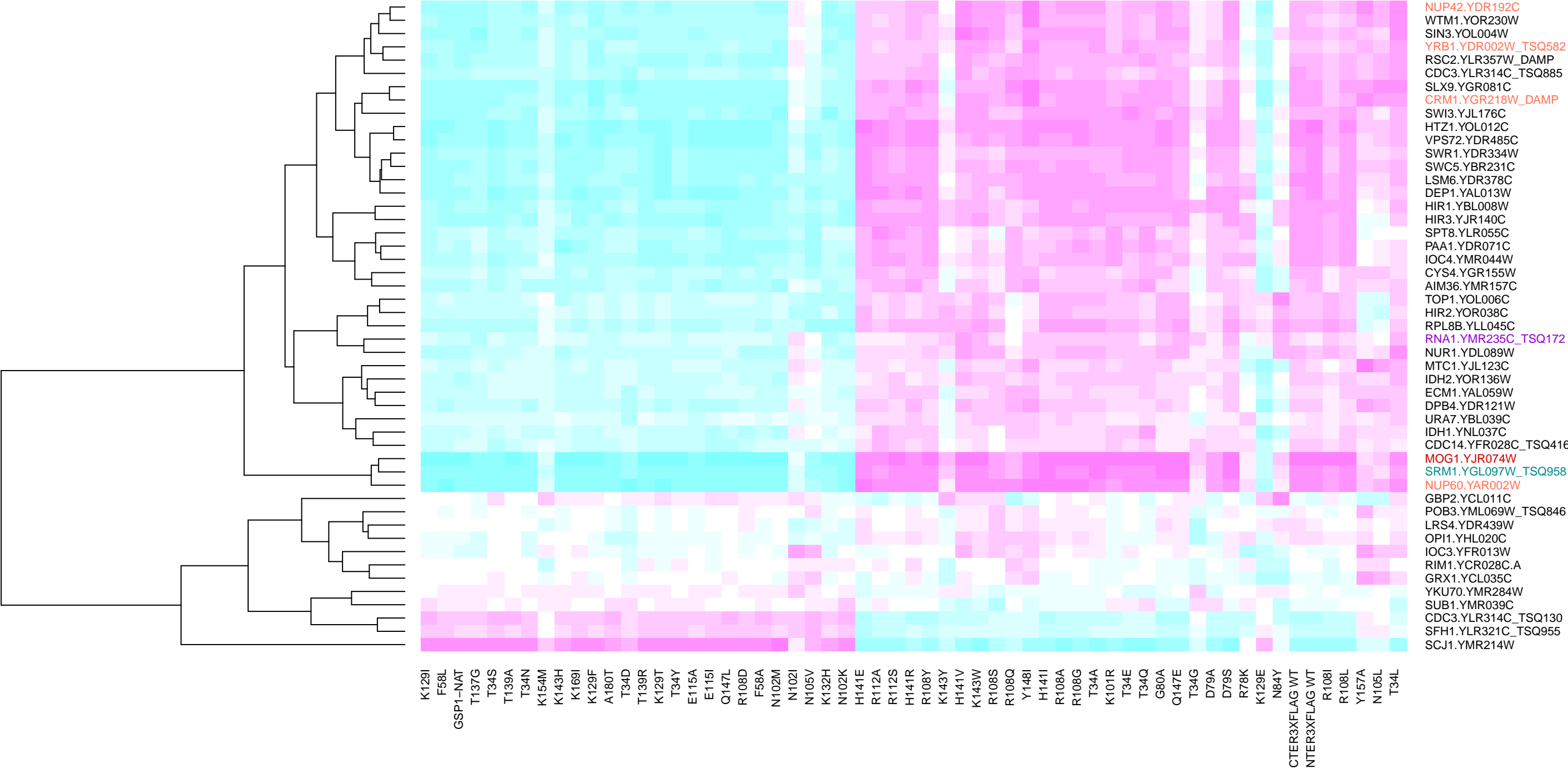
ribosomes and translation_GO_15_2_mut



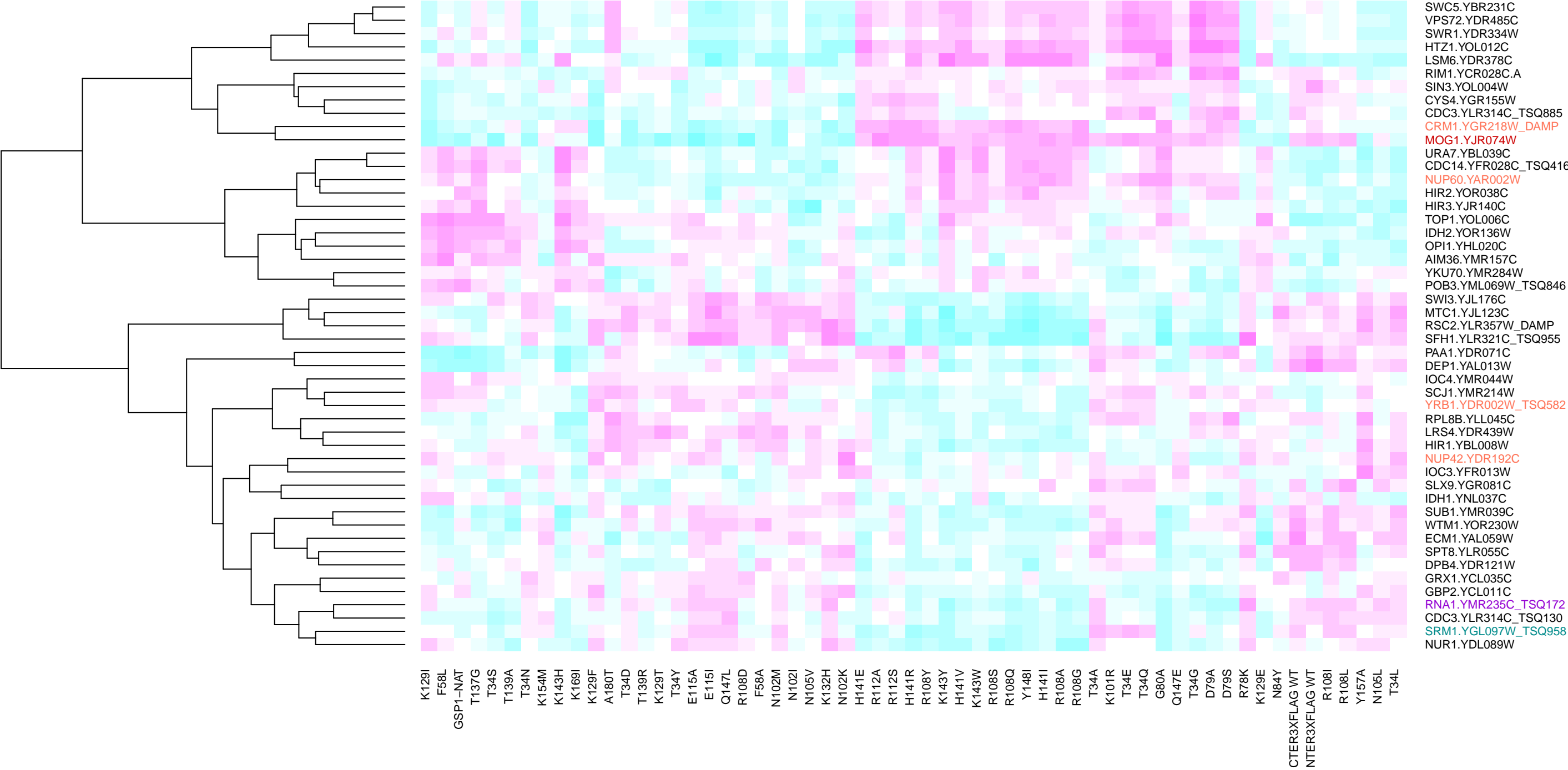
transcription and mRNA processing_GO_15_1_all



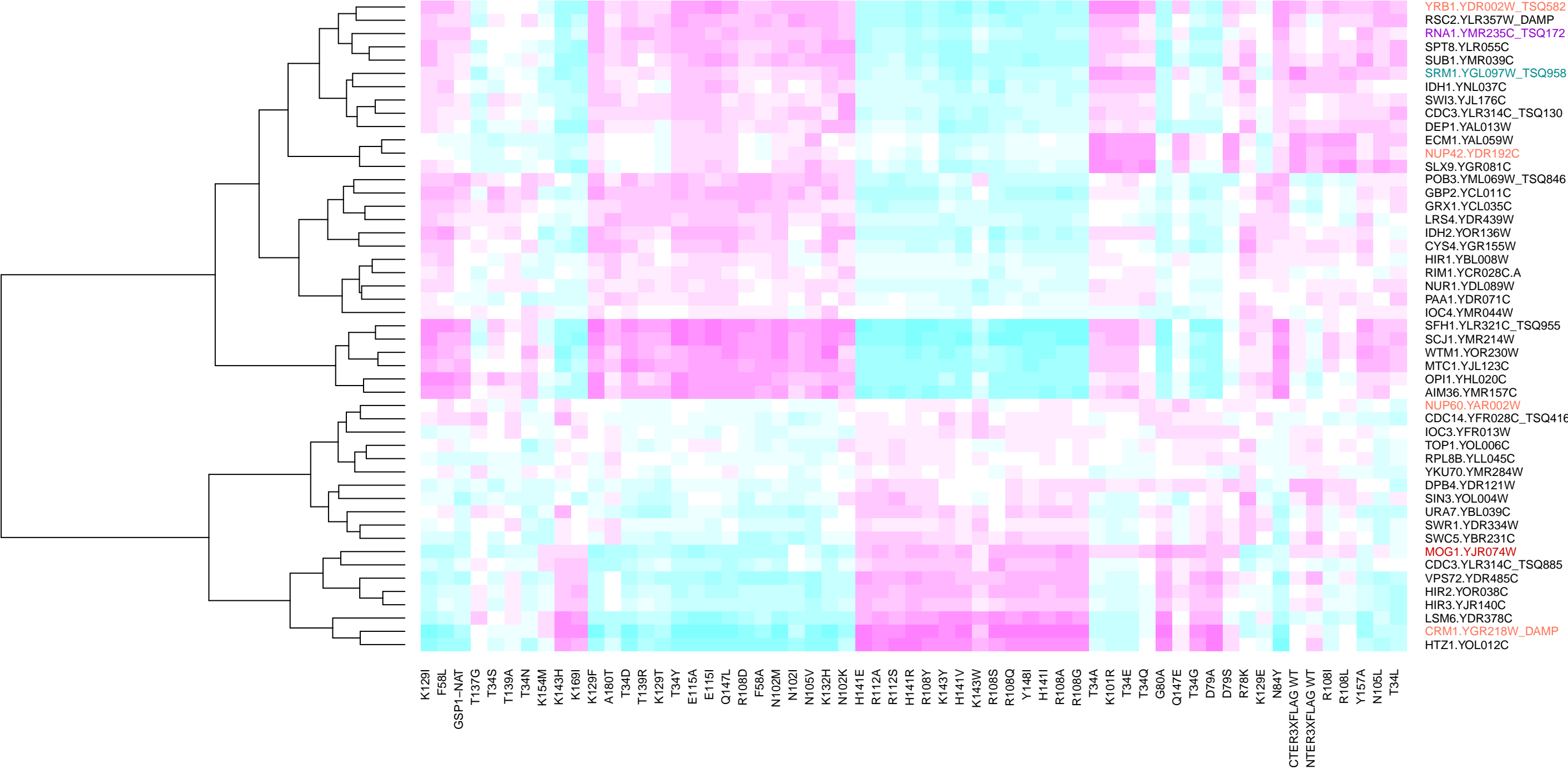
chromatin_GO_15_7_mut



chromosome segregation_GO_15_1_all



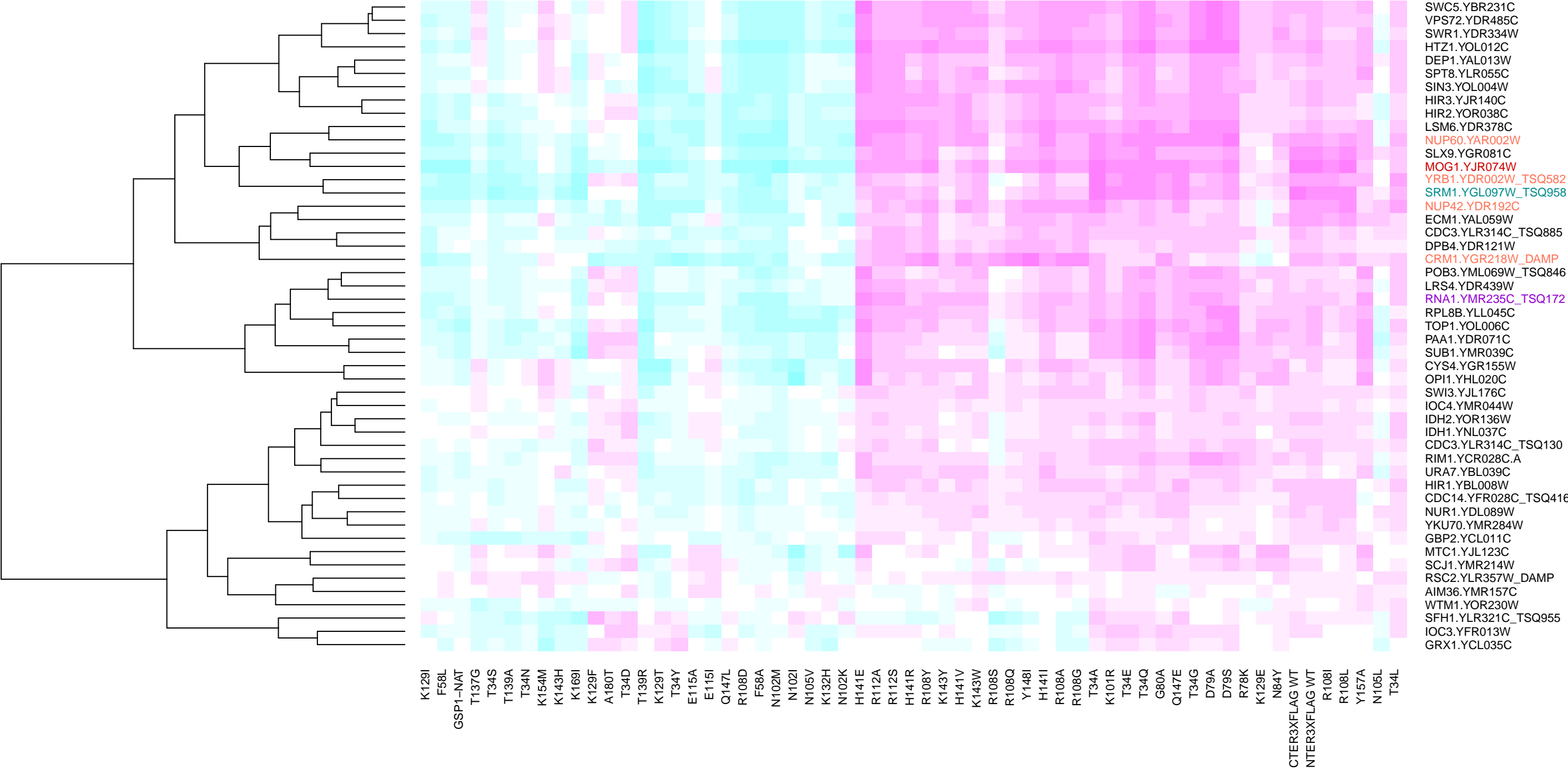
cytoskeleton and microtubules_GO_15_2_mut



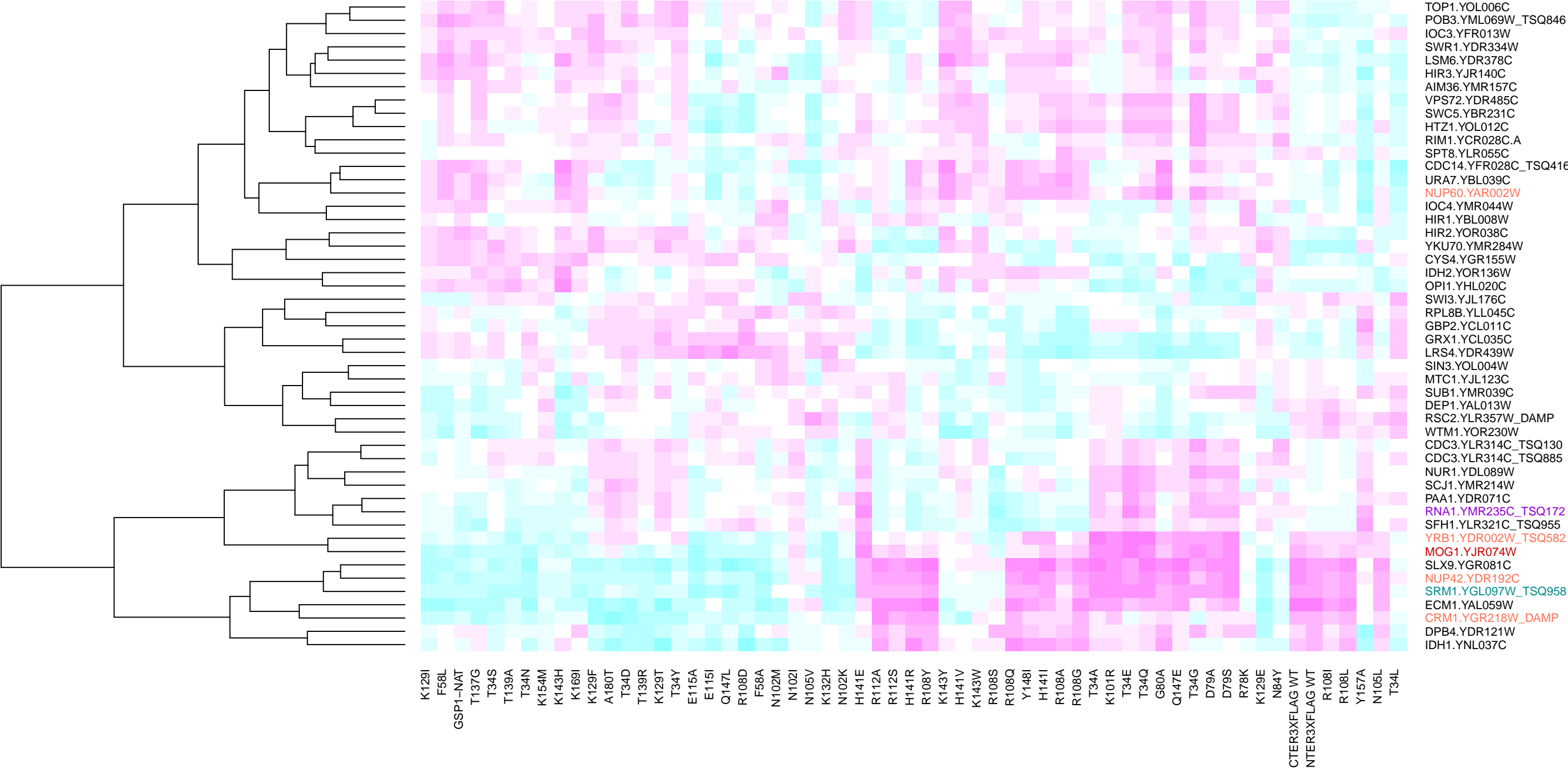
cytoskeleton and microtubules_GO_15_3_all



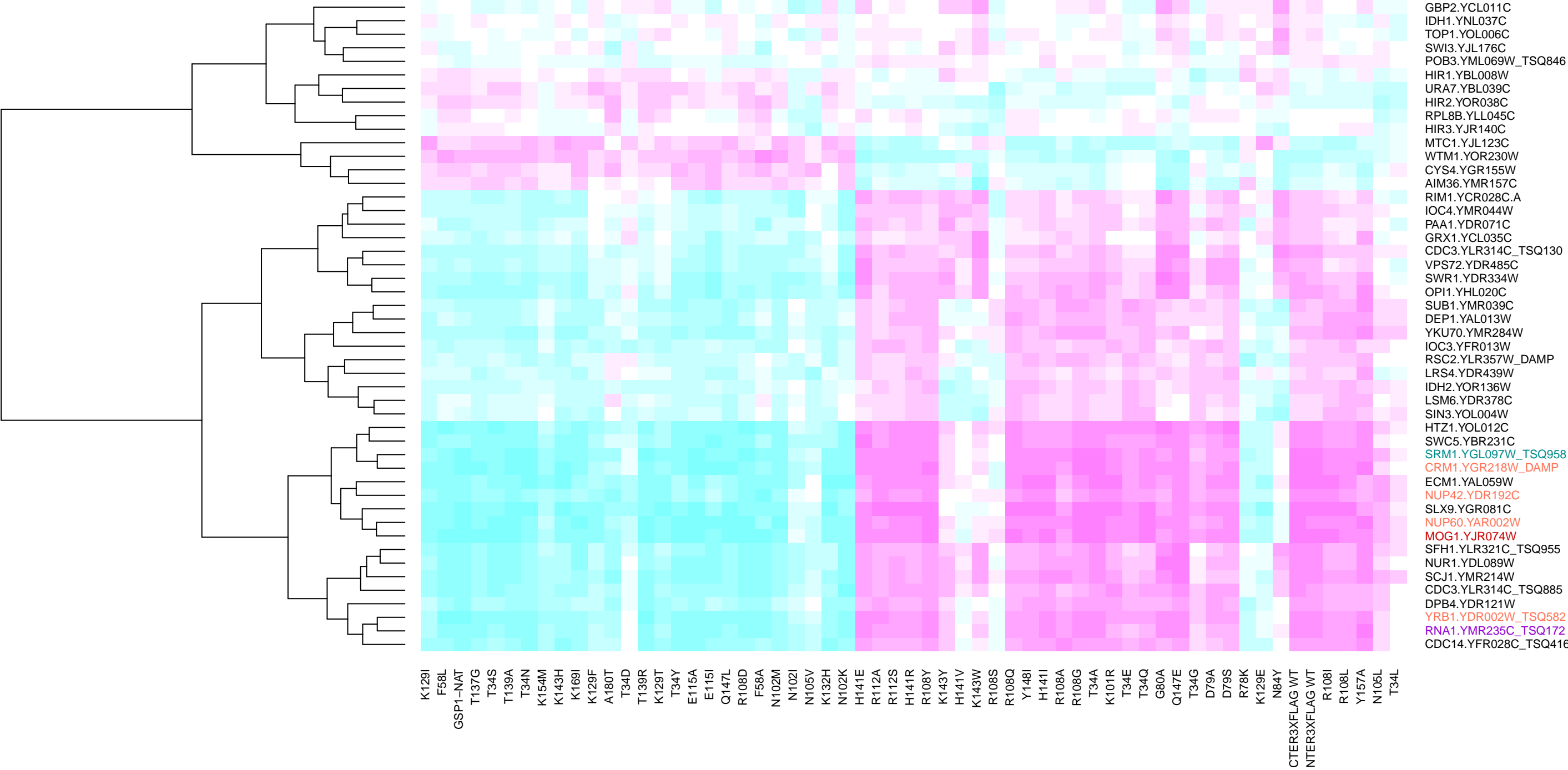
sig_Gsp1_GI



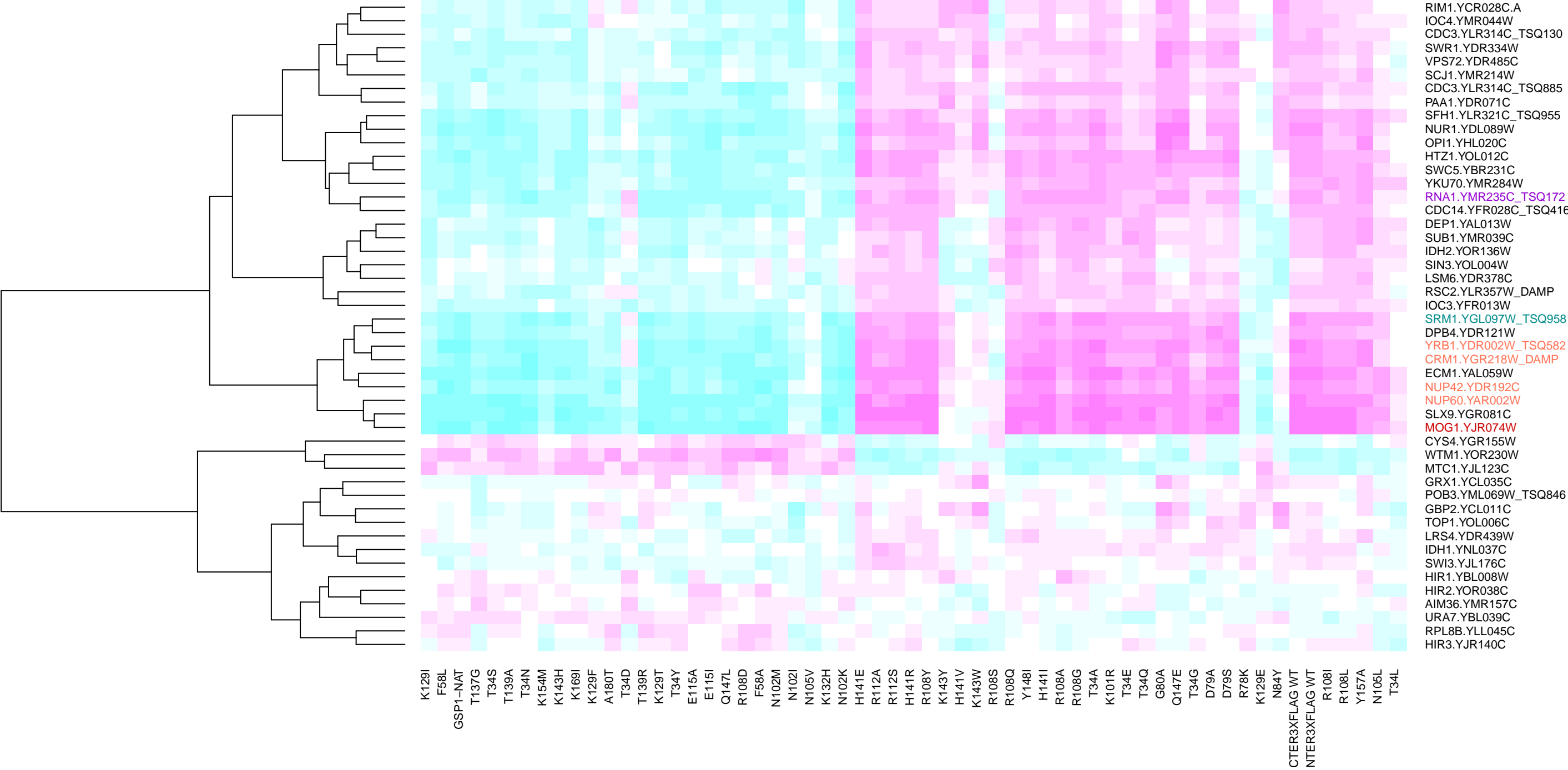
cytoskeleton and microtubules_GO_30_3_all



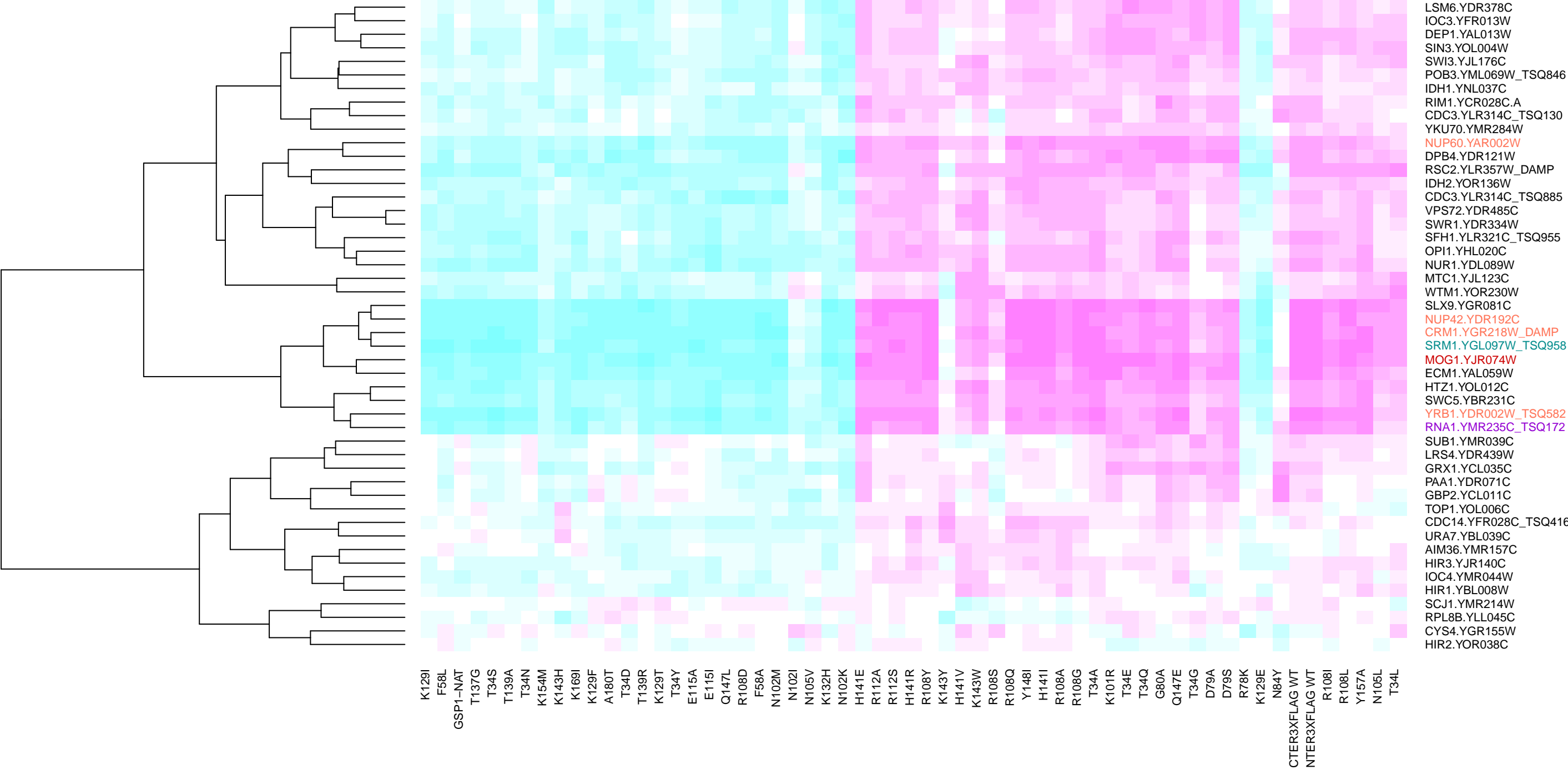
nuclear transport and organization_GO_15_3_all



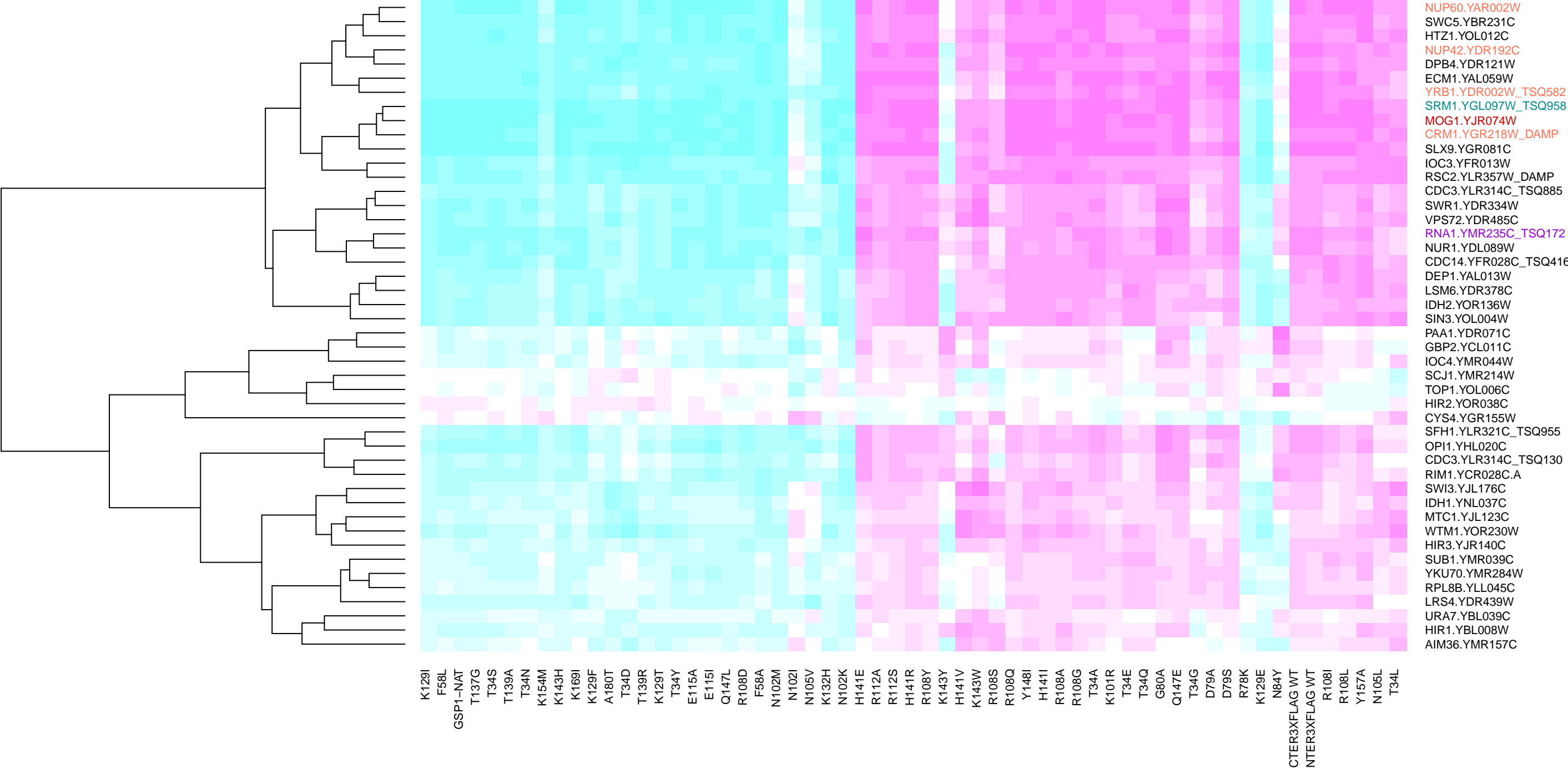
nuclear transport_GO_15_1_all



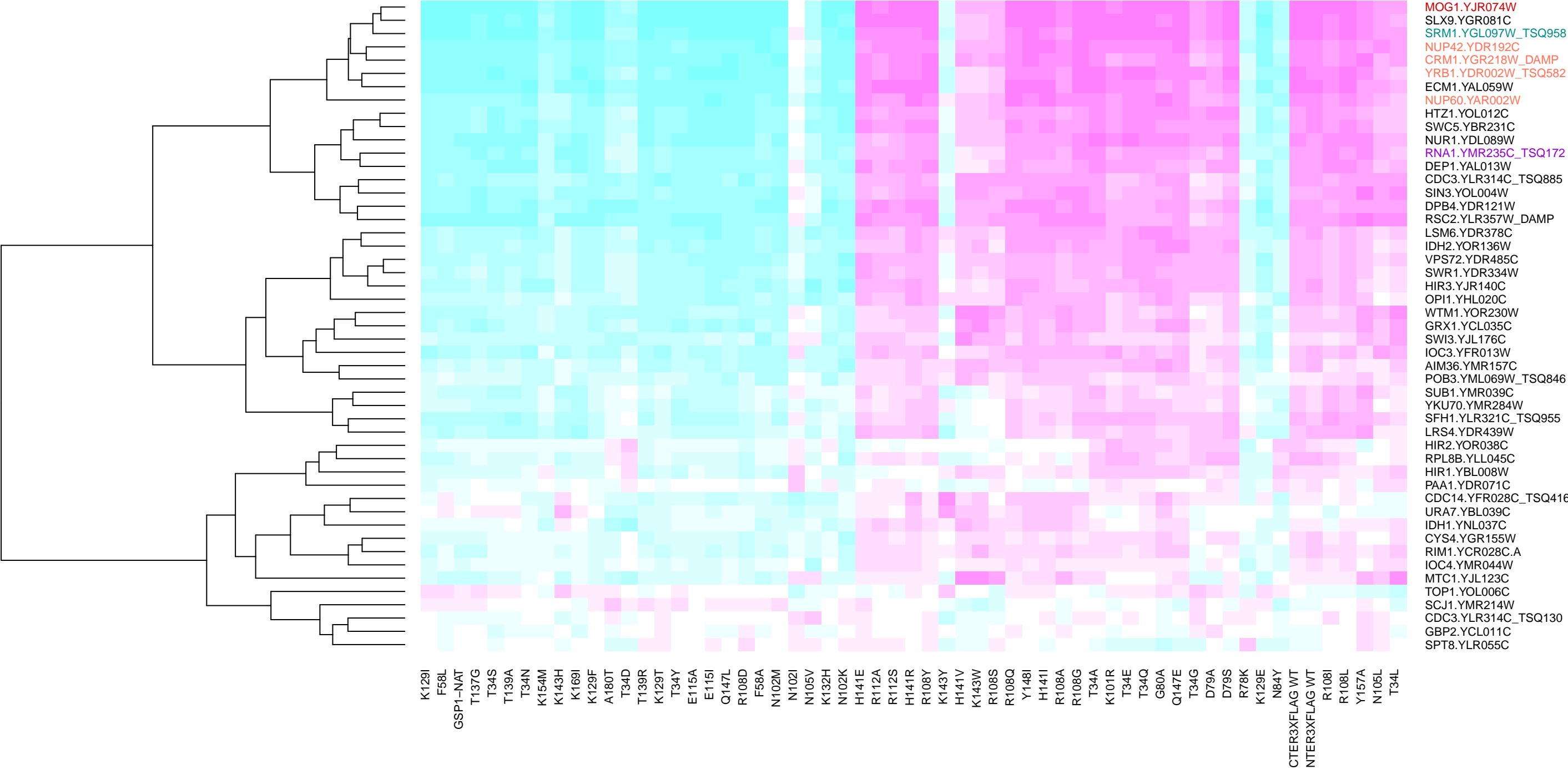
nucleus organization



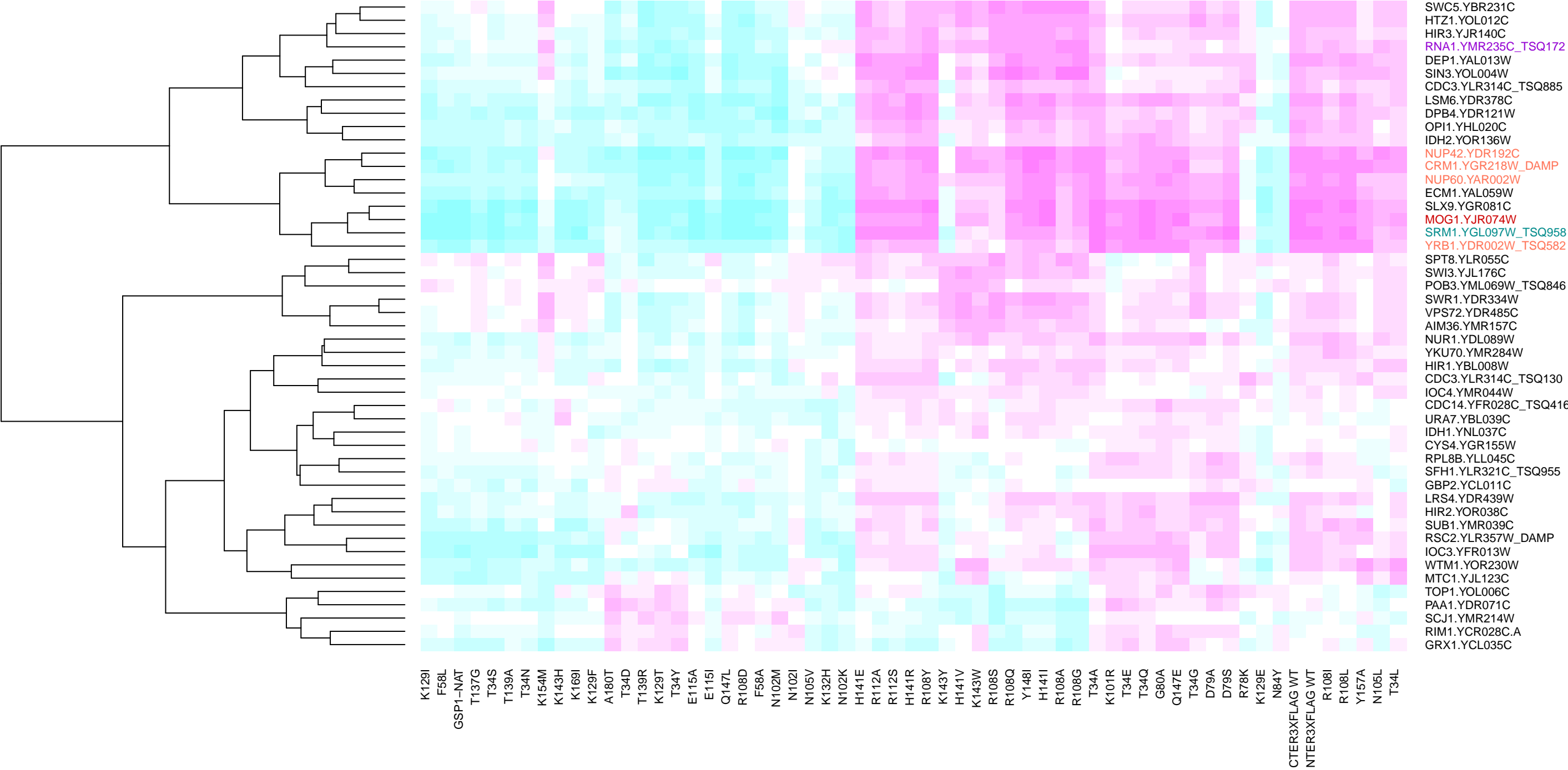
nucleus organization_GO_15_1_all



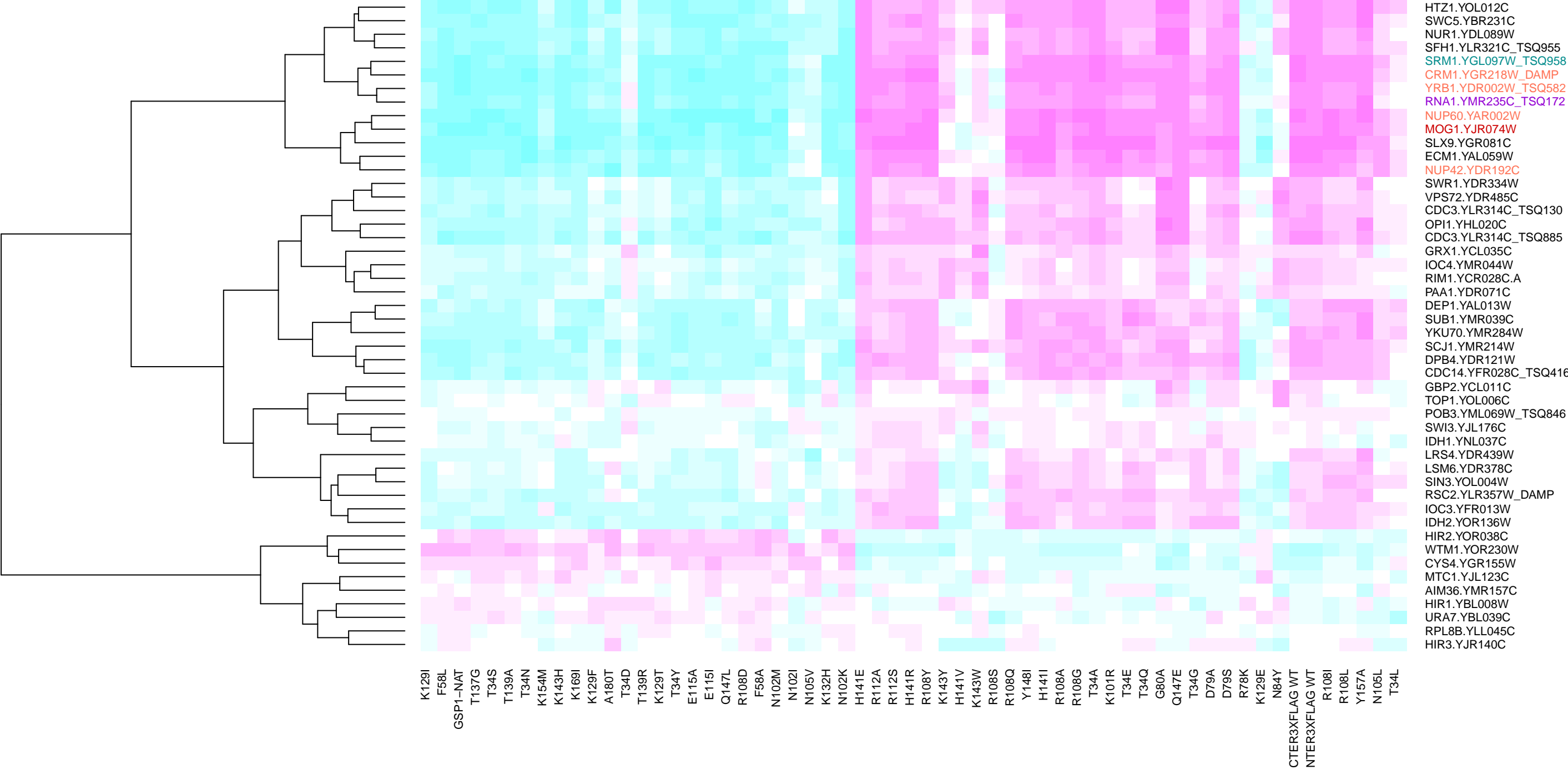
sig_Gsp1_Gl_15_9_all



sig_Gsp1_Gl_30_1_all



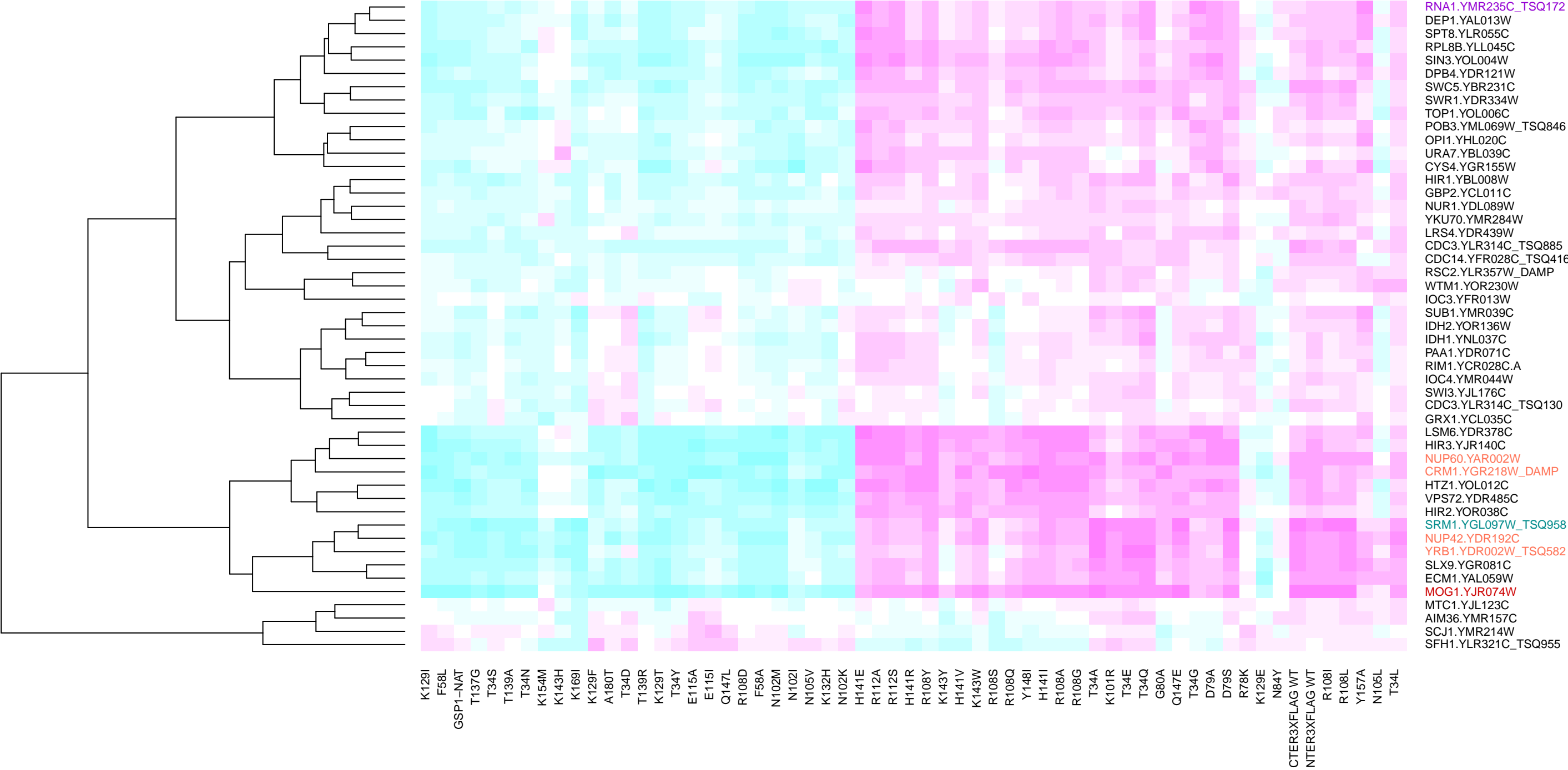
whole_library_15_36_all



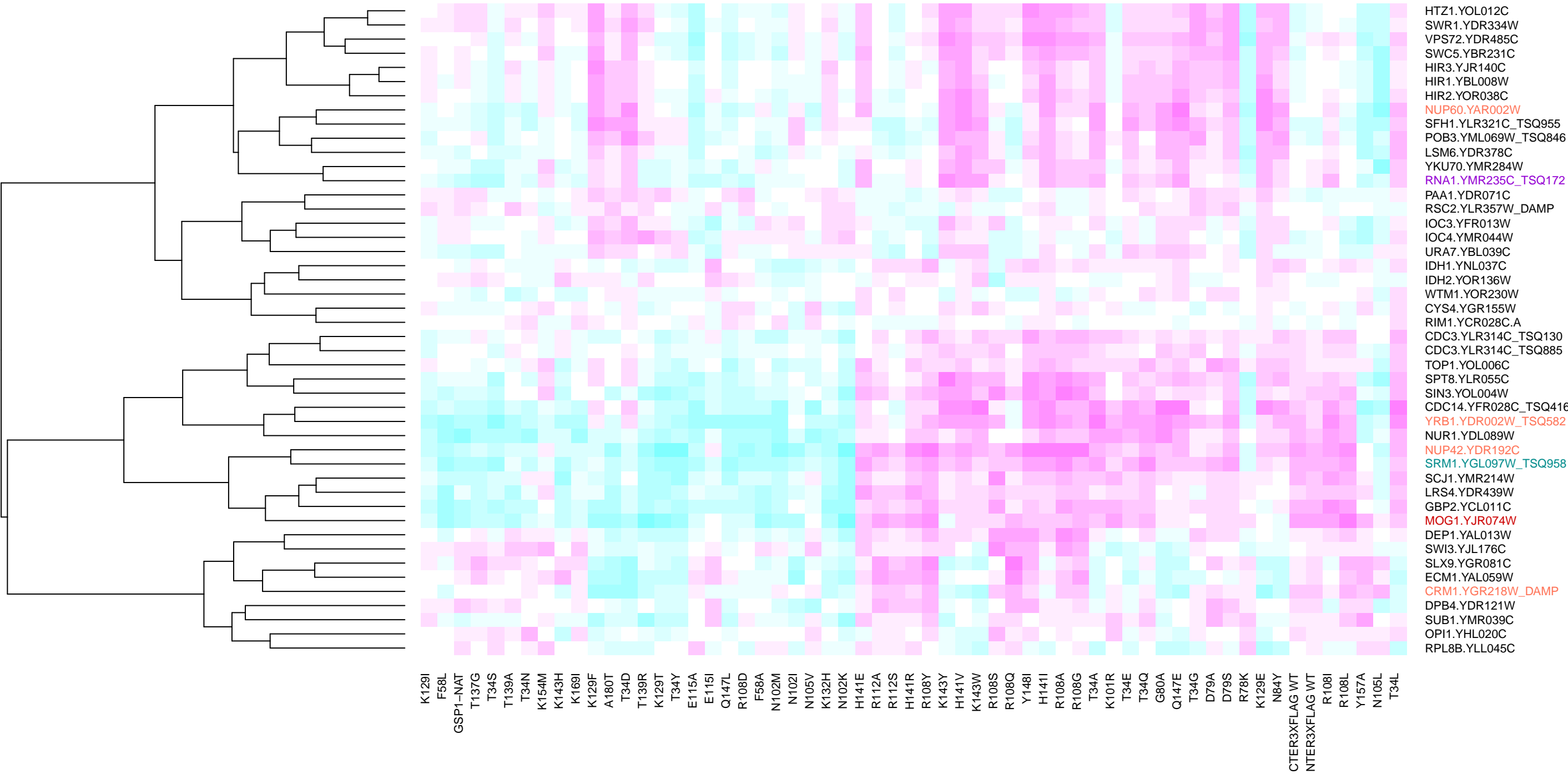
transcription and mRNA processing_GO_15_7_all



sig_Gsp1_Gl_30_1_mut



RNA processing



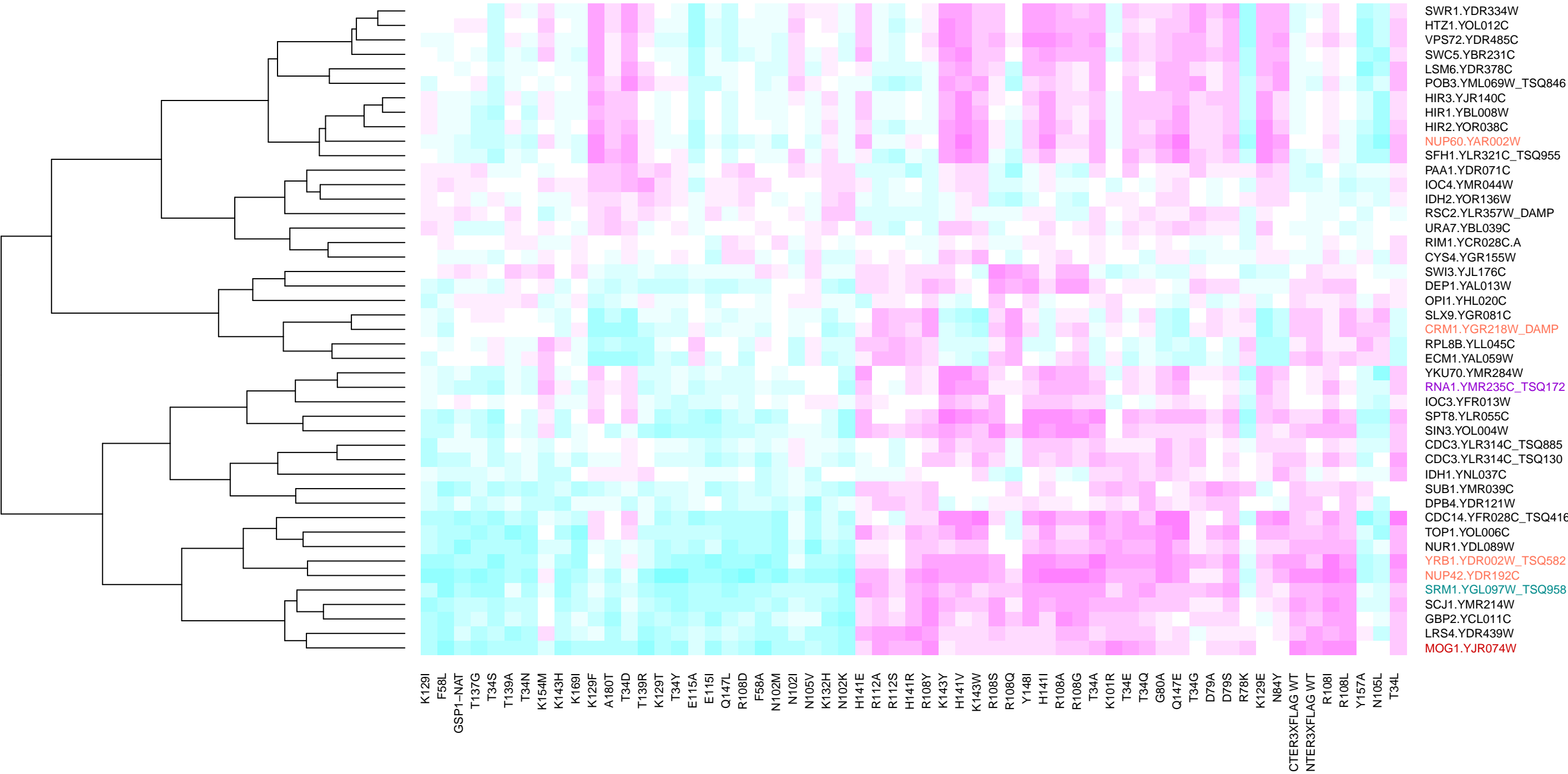
budding_GO_15_1_all



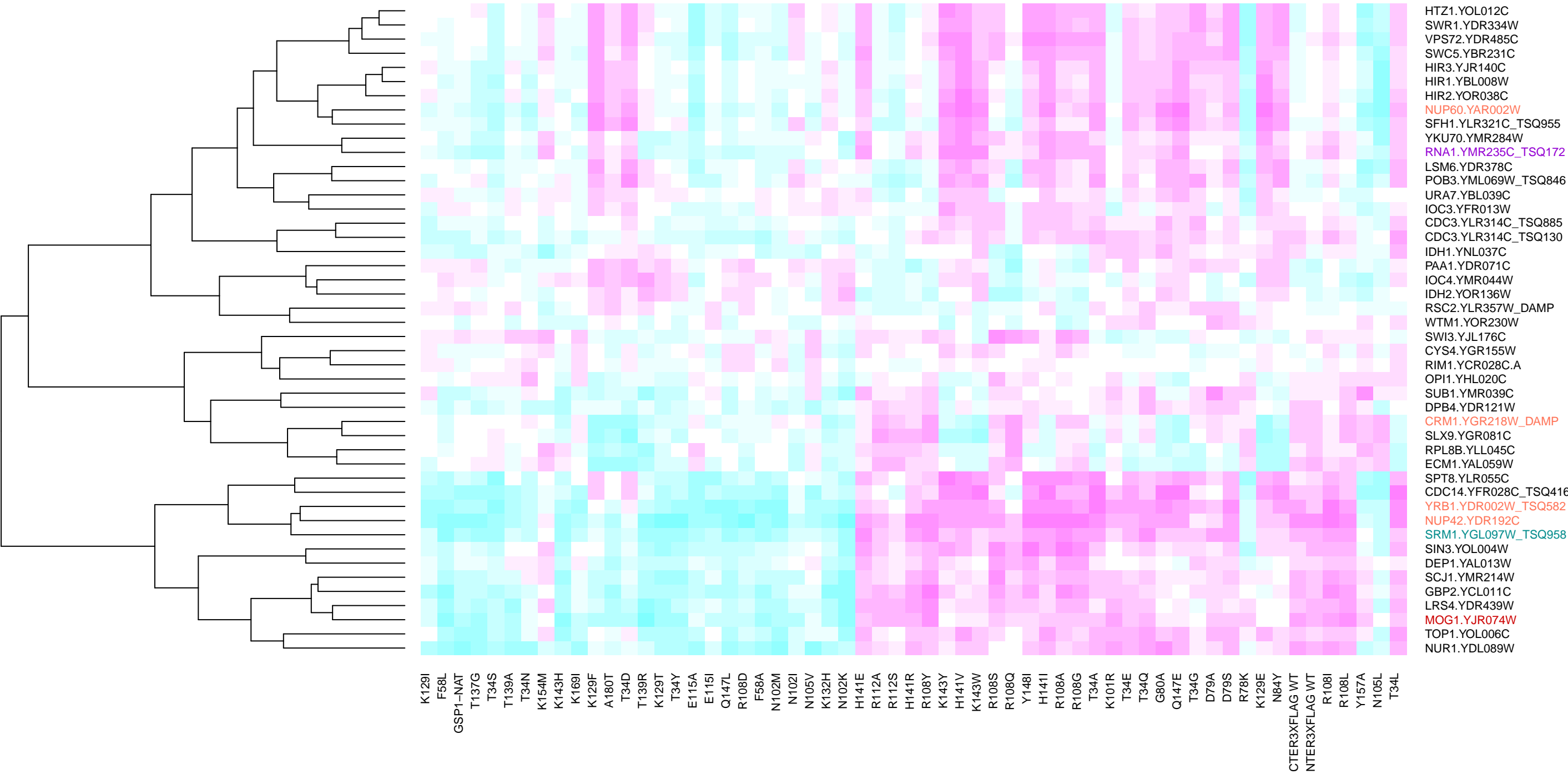
organelle fission_GO_30_1_mut



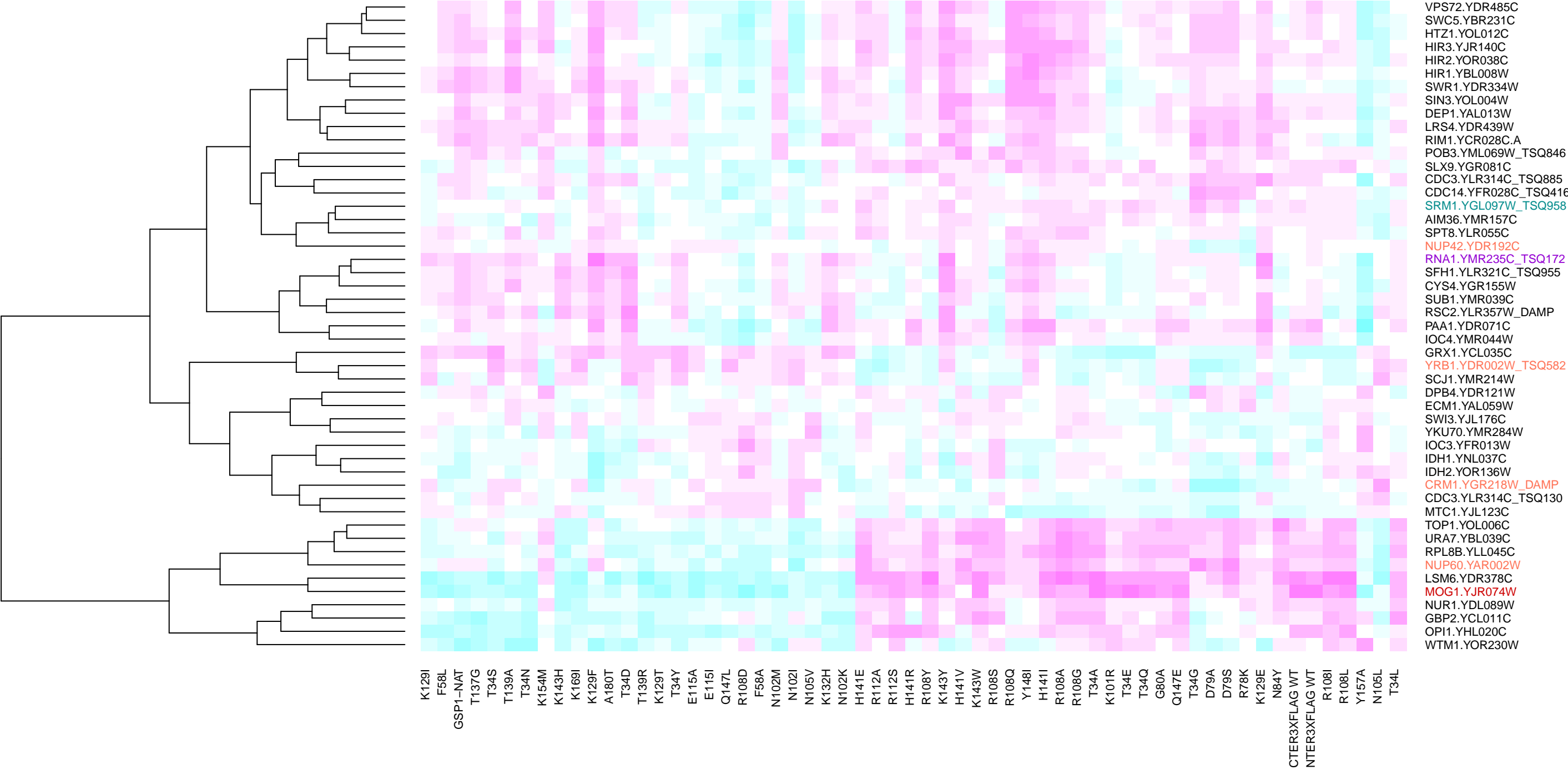
mRNA processing



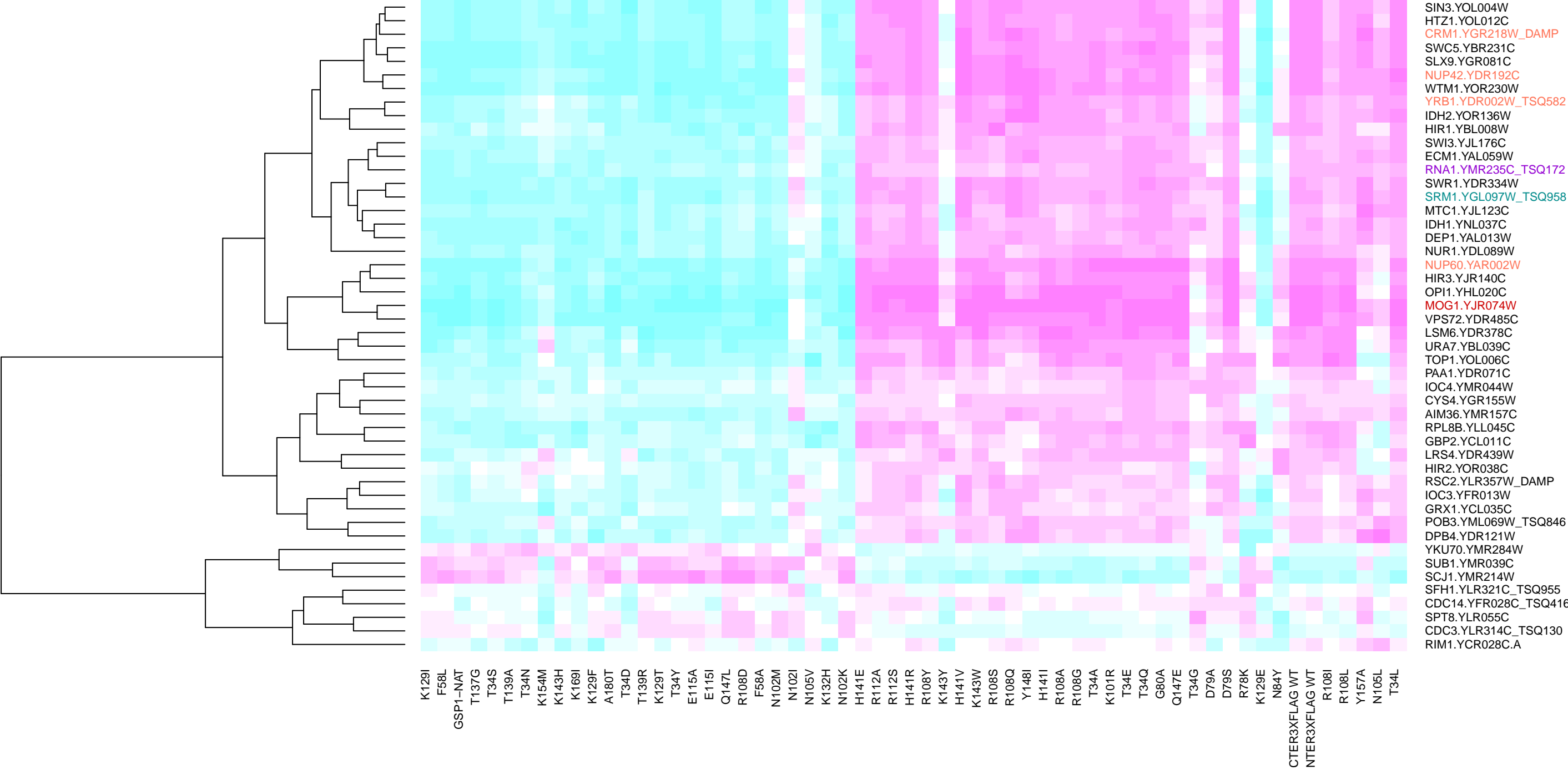
RNA processing_GO_15_1_all



DNA recombination



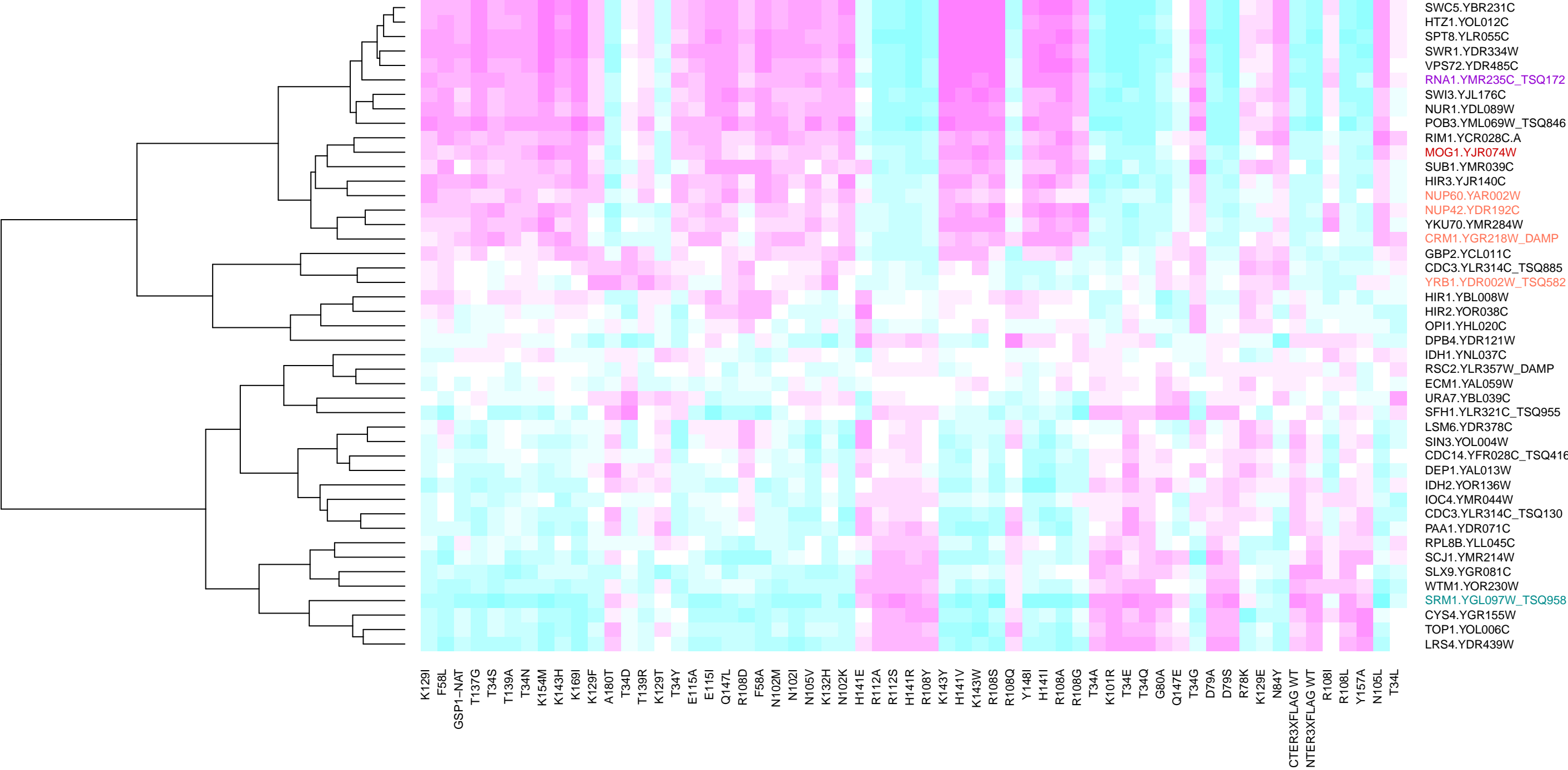
transcription from RNA polymerase II promoter_GO_15_4_all



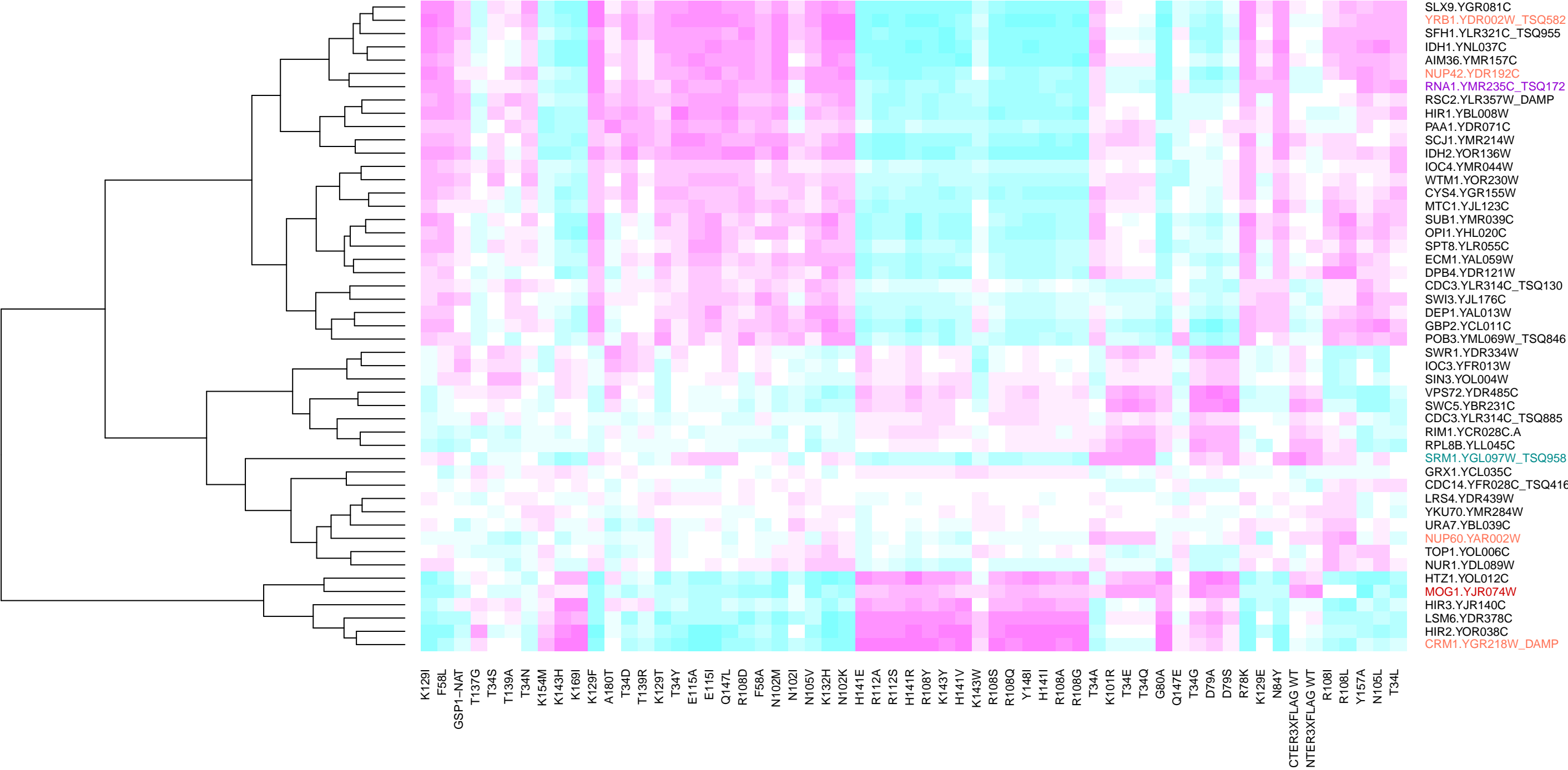
sig_Gsp1_Gl_30_2_all



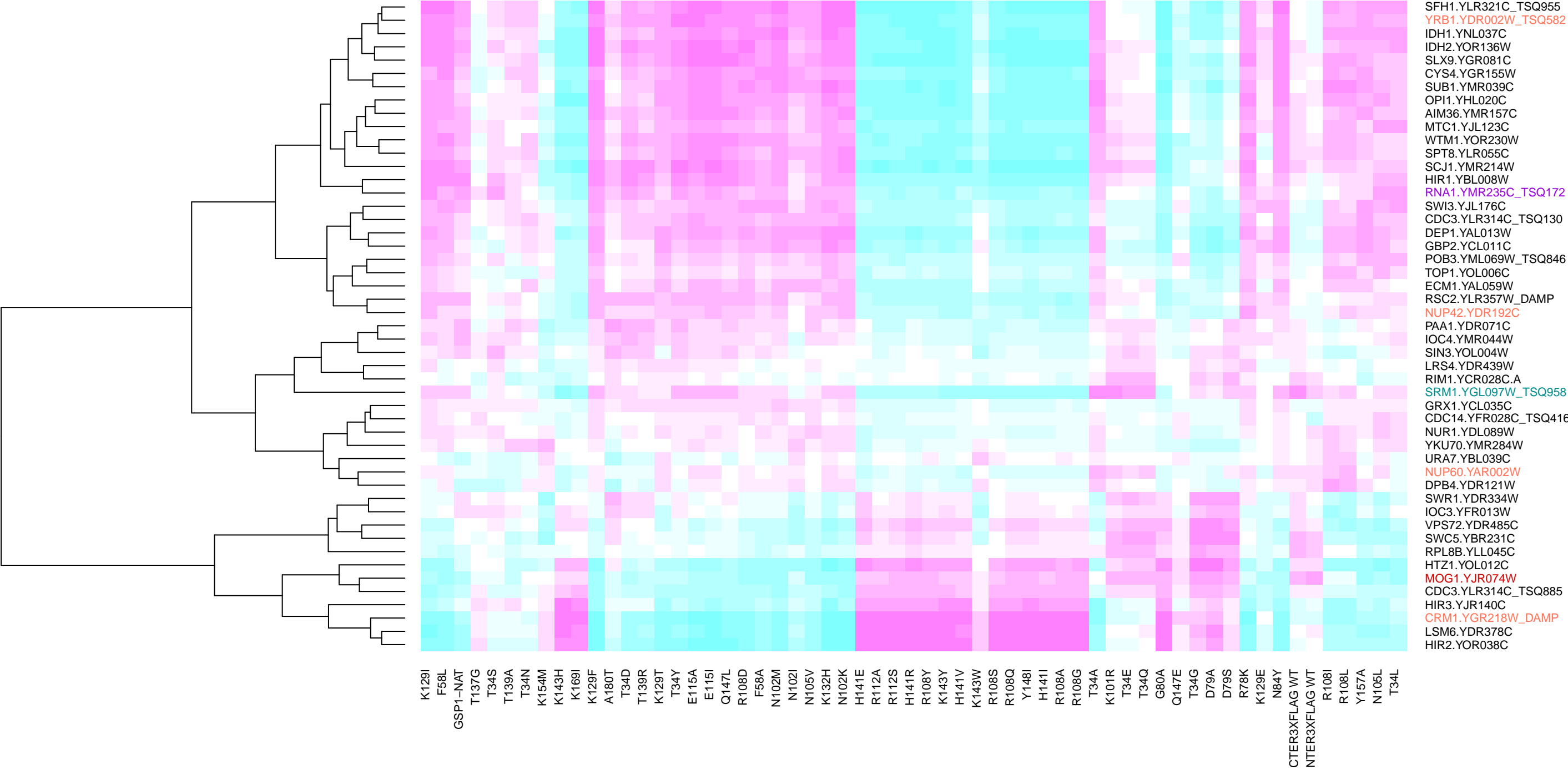
sig_Gsp1_Gl_15_7_mut



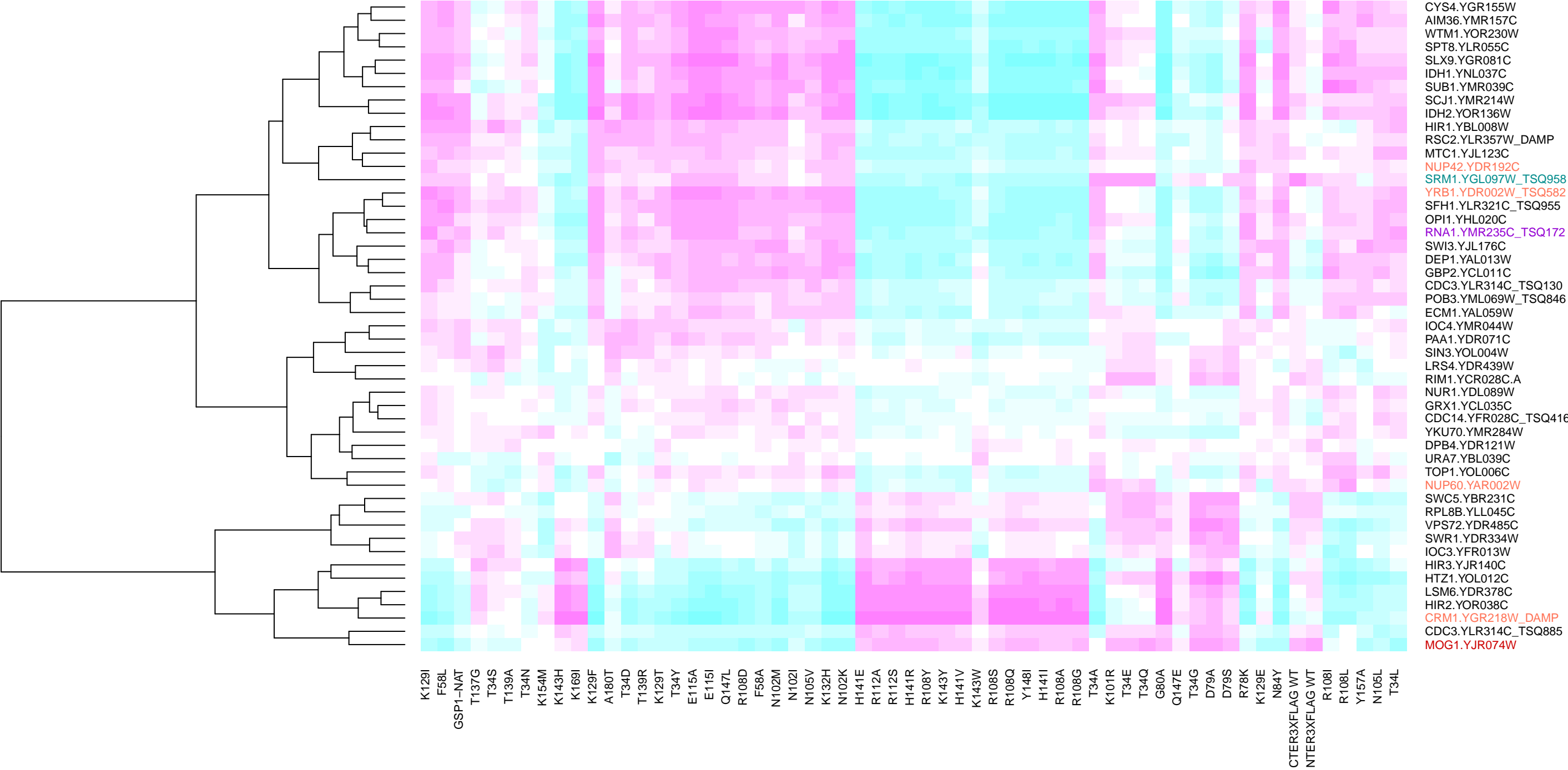
mitotic cell cycle_GO_15_2_all



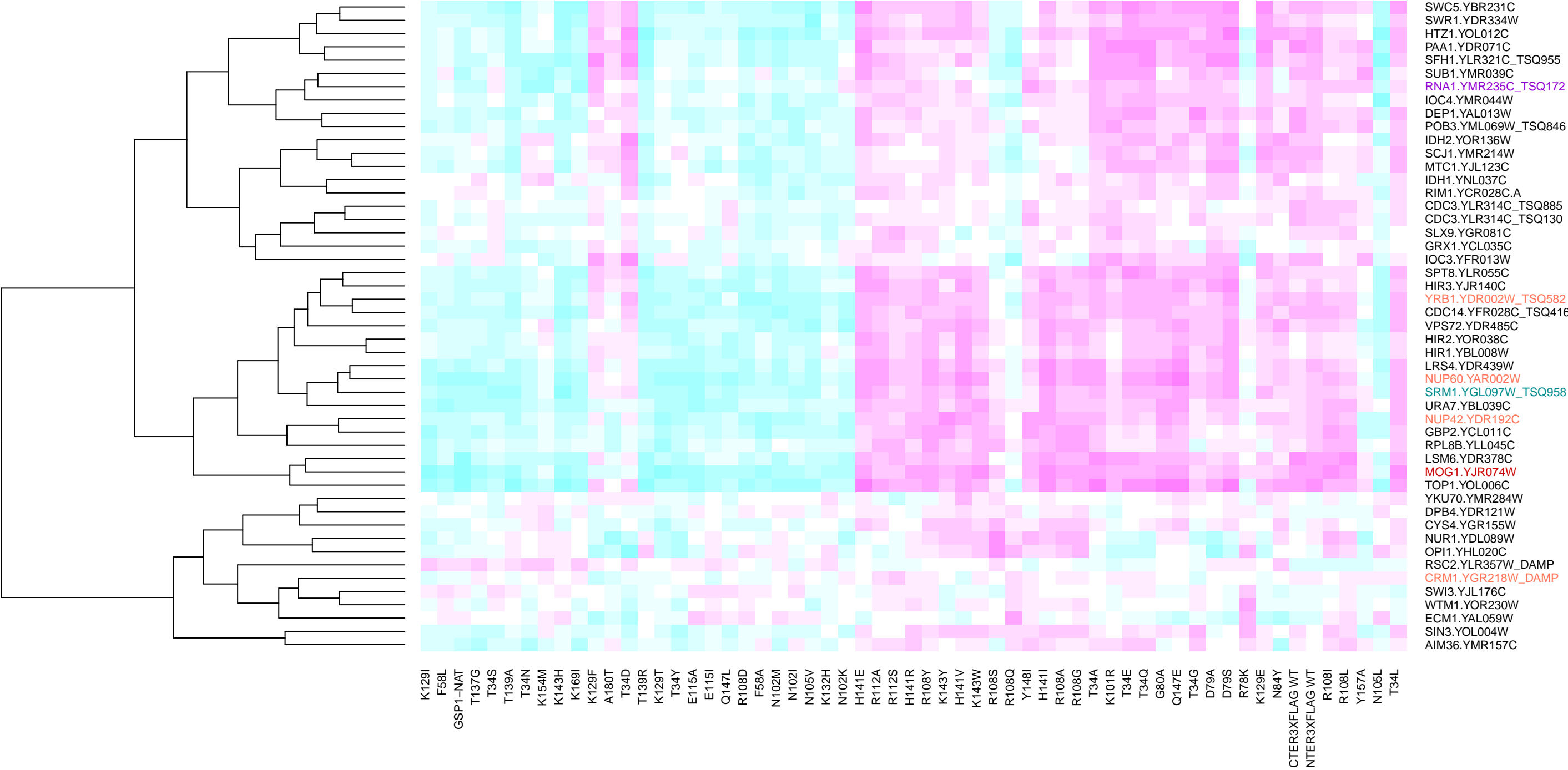
whole_library_15_3_all



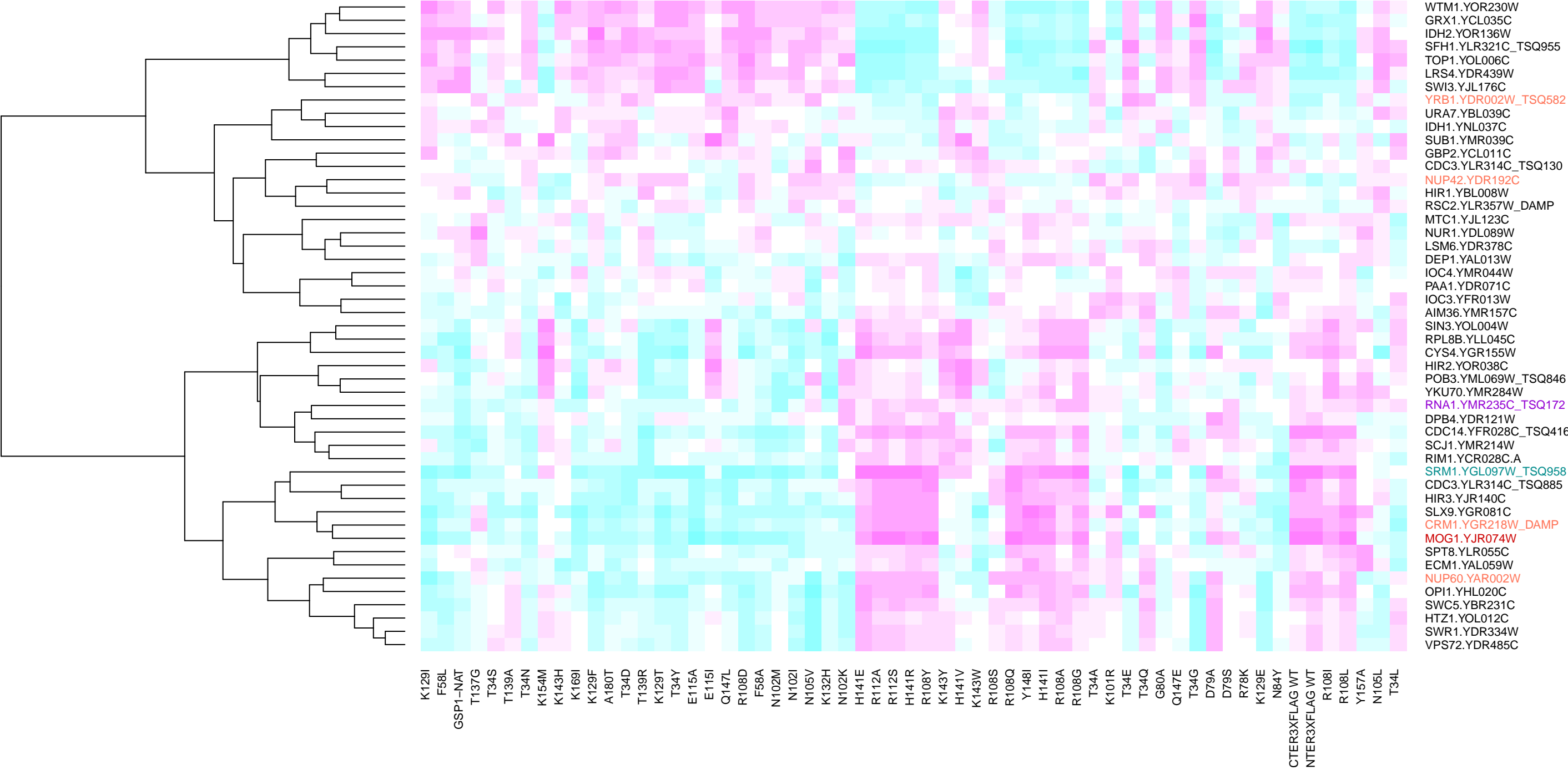
whole_library_30_9_all



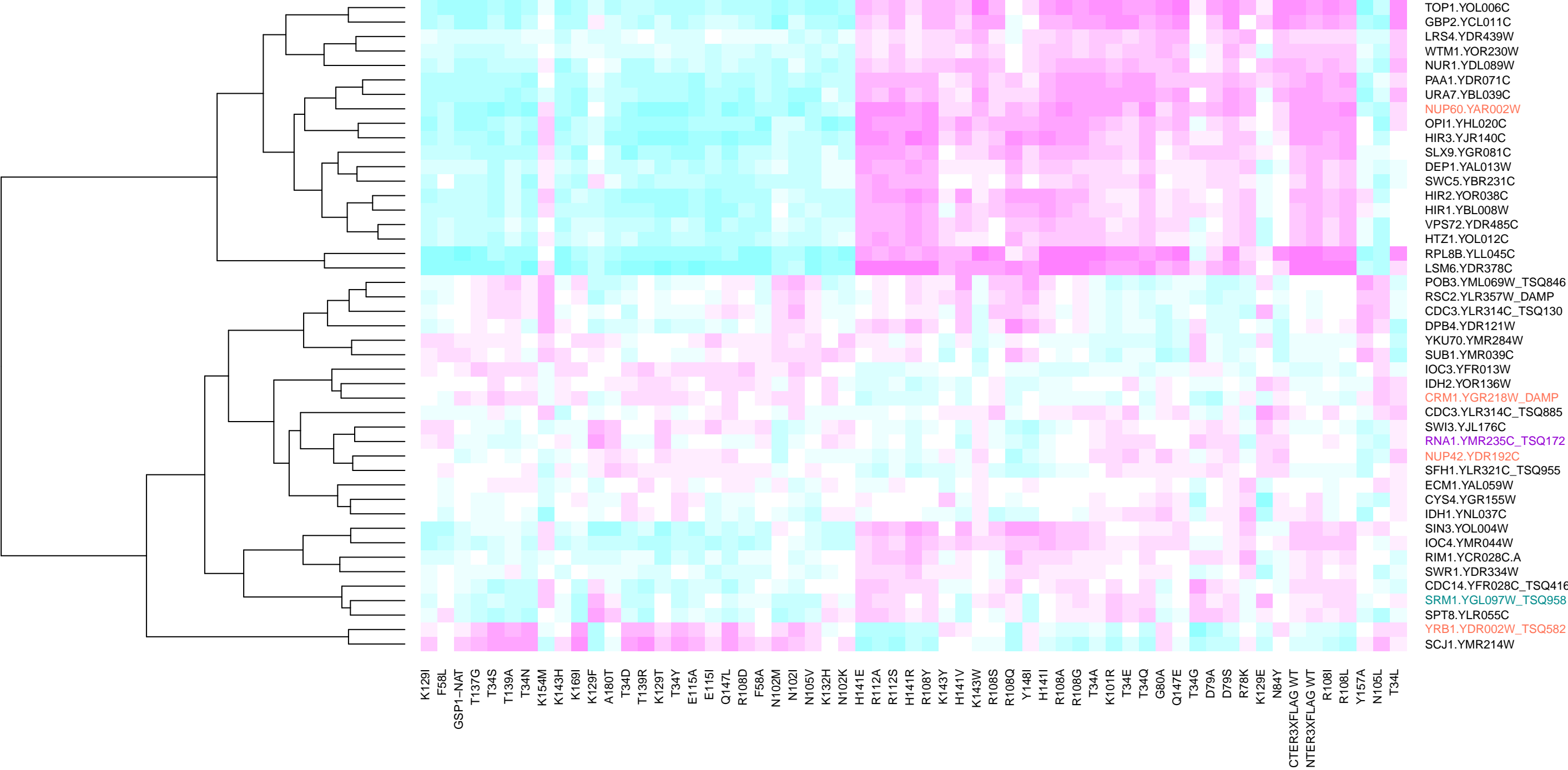
sig_Gsp1_Gl_15_3_all



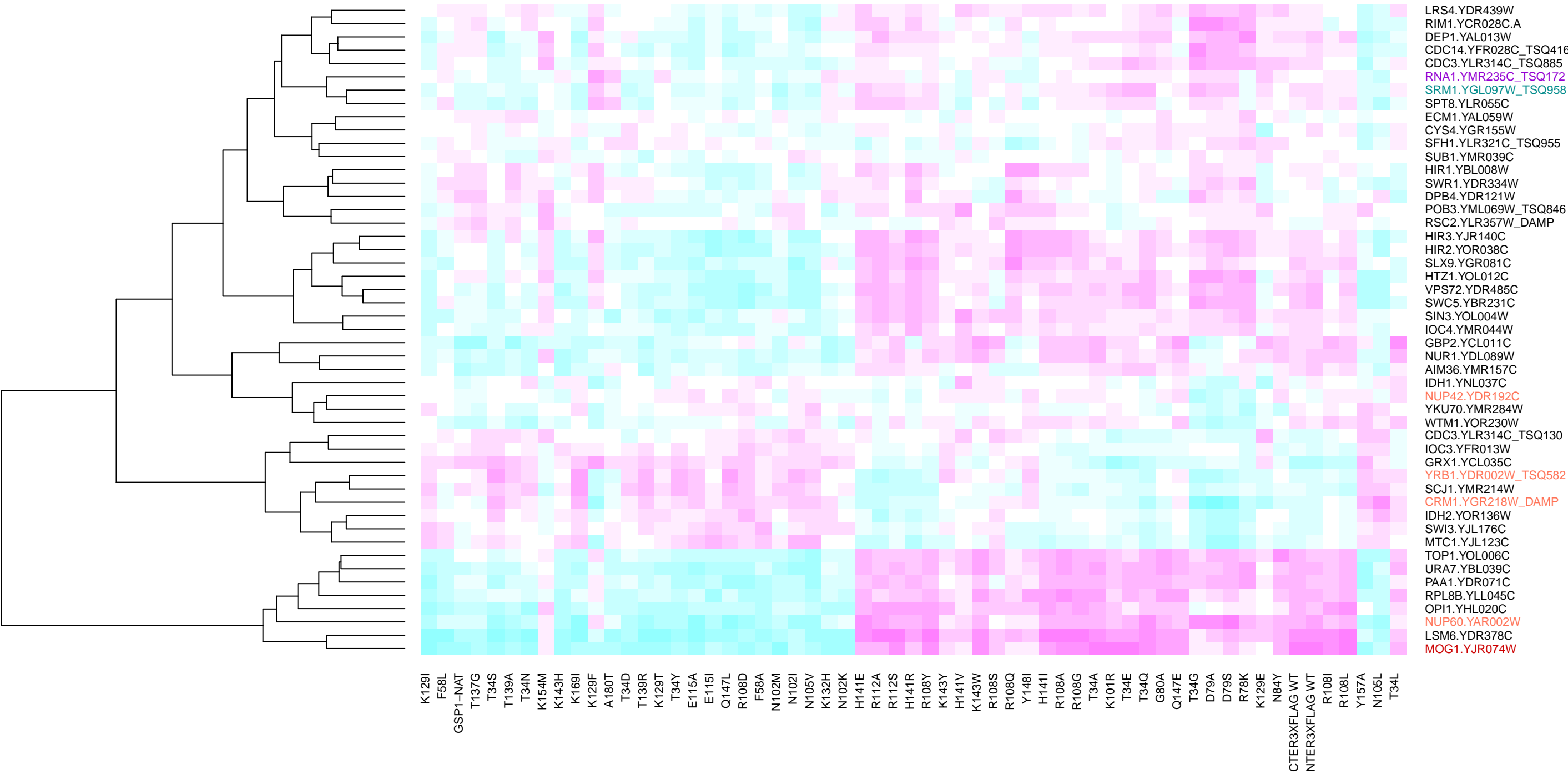
sig_Gsp1_Gl_15_9_mut



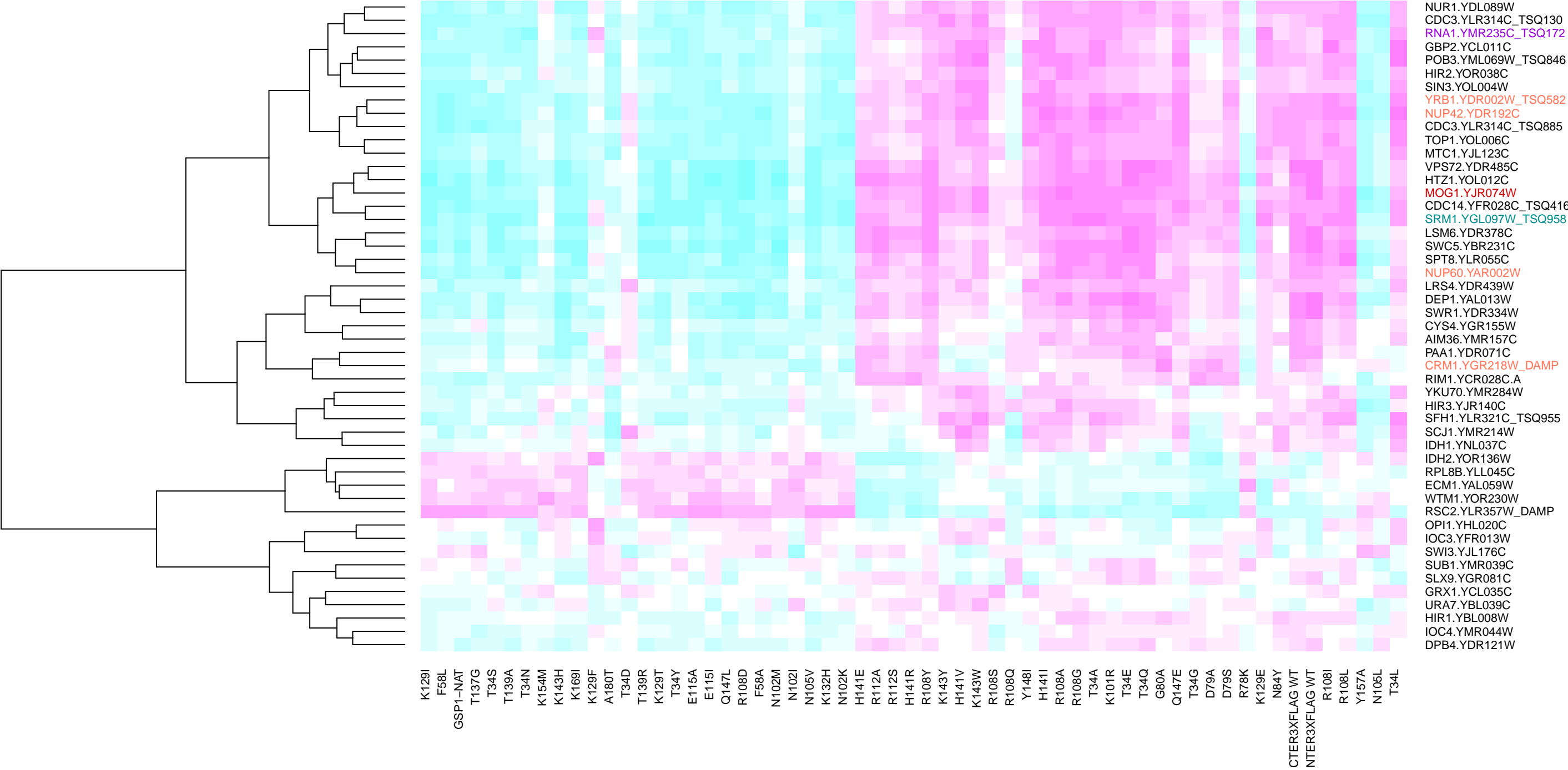
DNA recombination_GO_15_1_mut



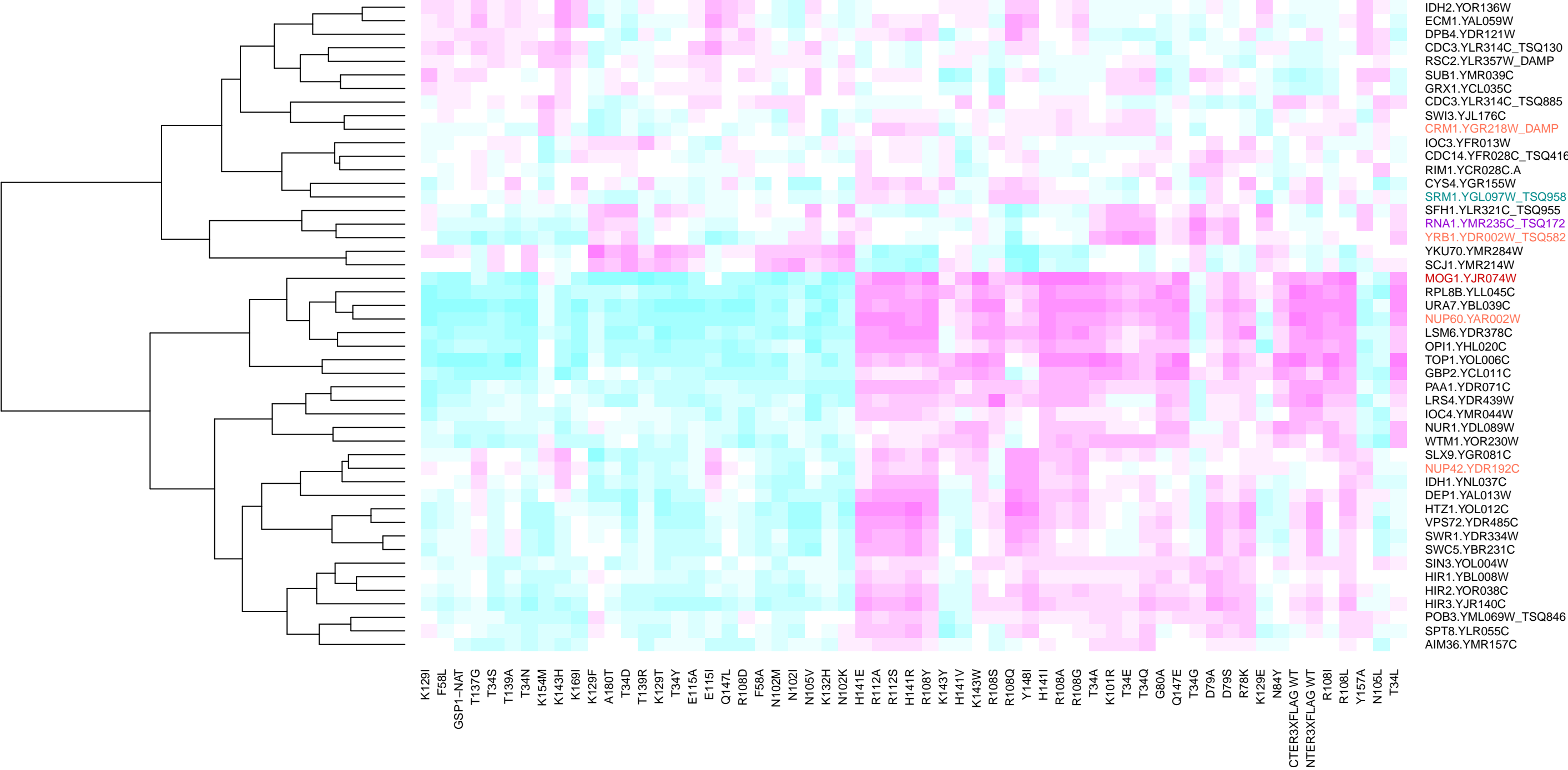
DNA recombination_GO_30_1_mut



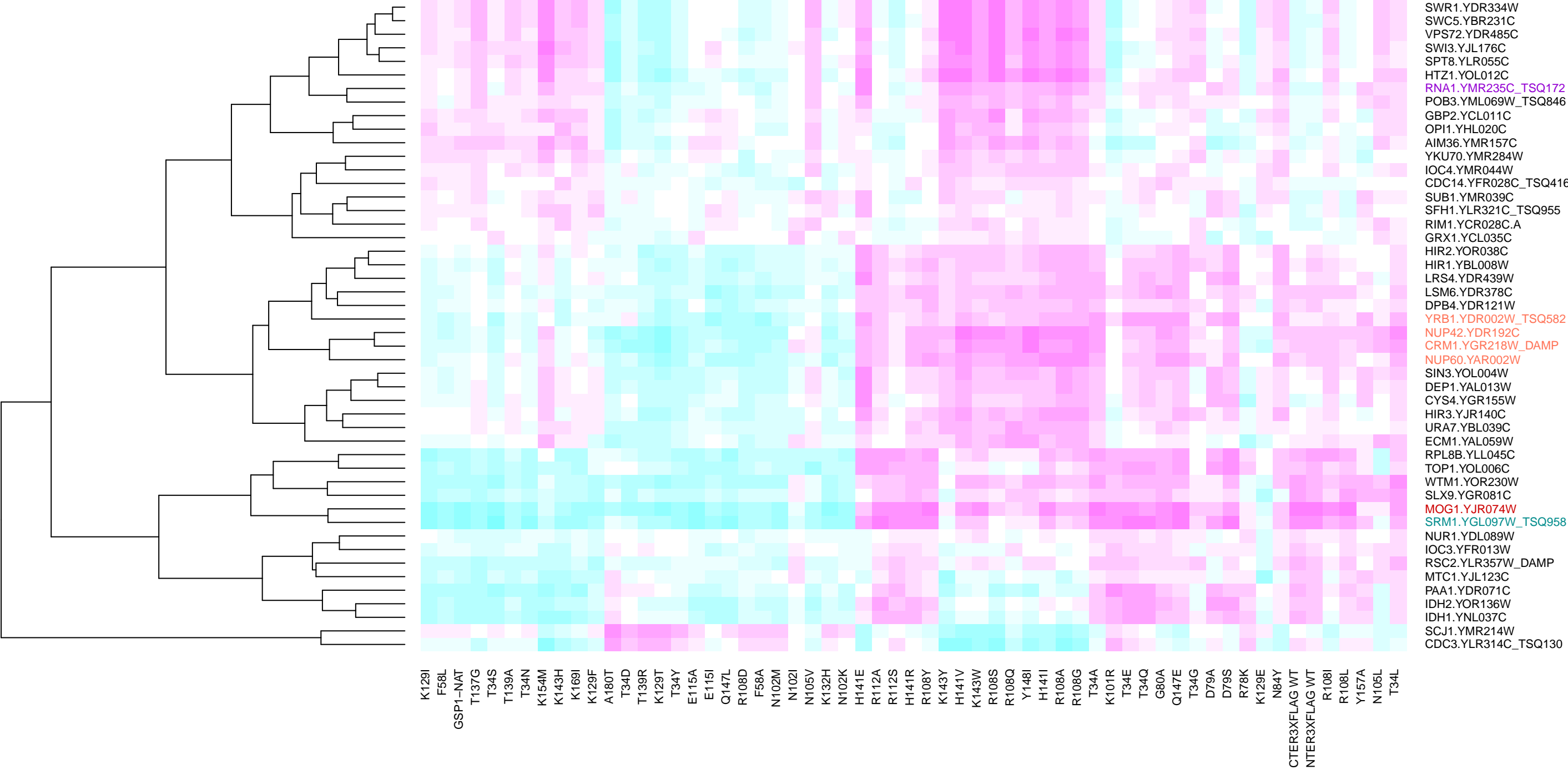
organelle fission_GO_15_3_mut



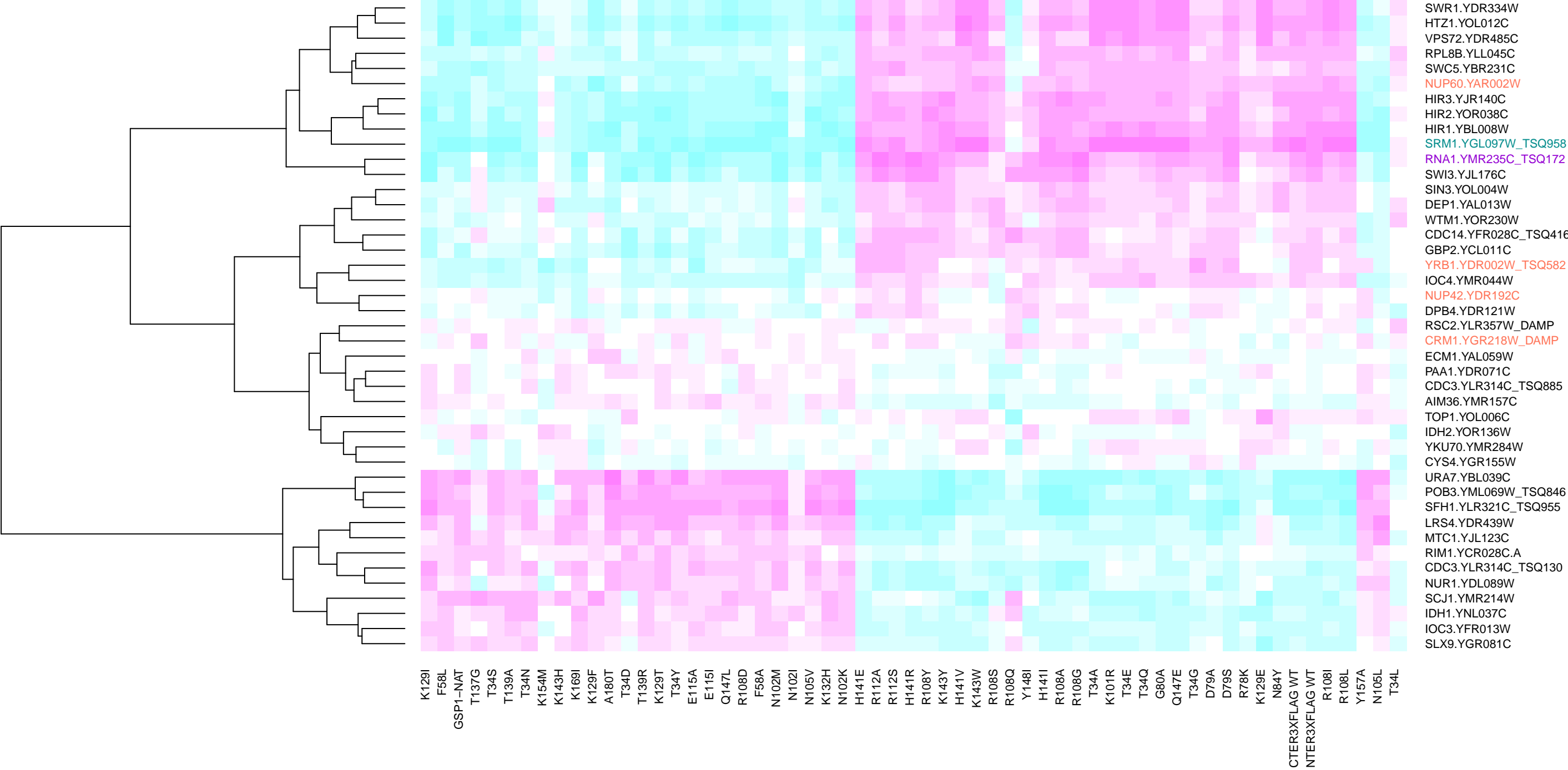
DNA-templated transcription, elongation_GO_15_1_mut



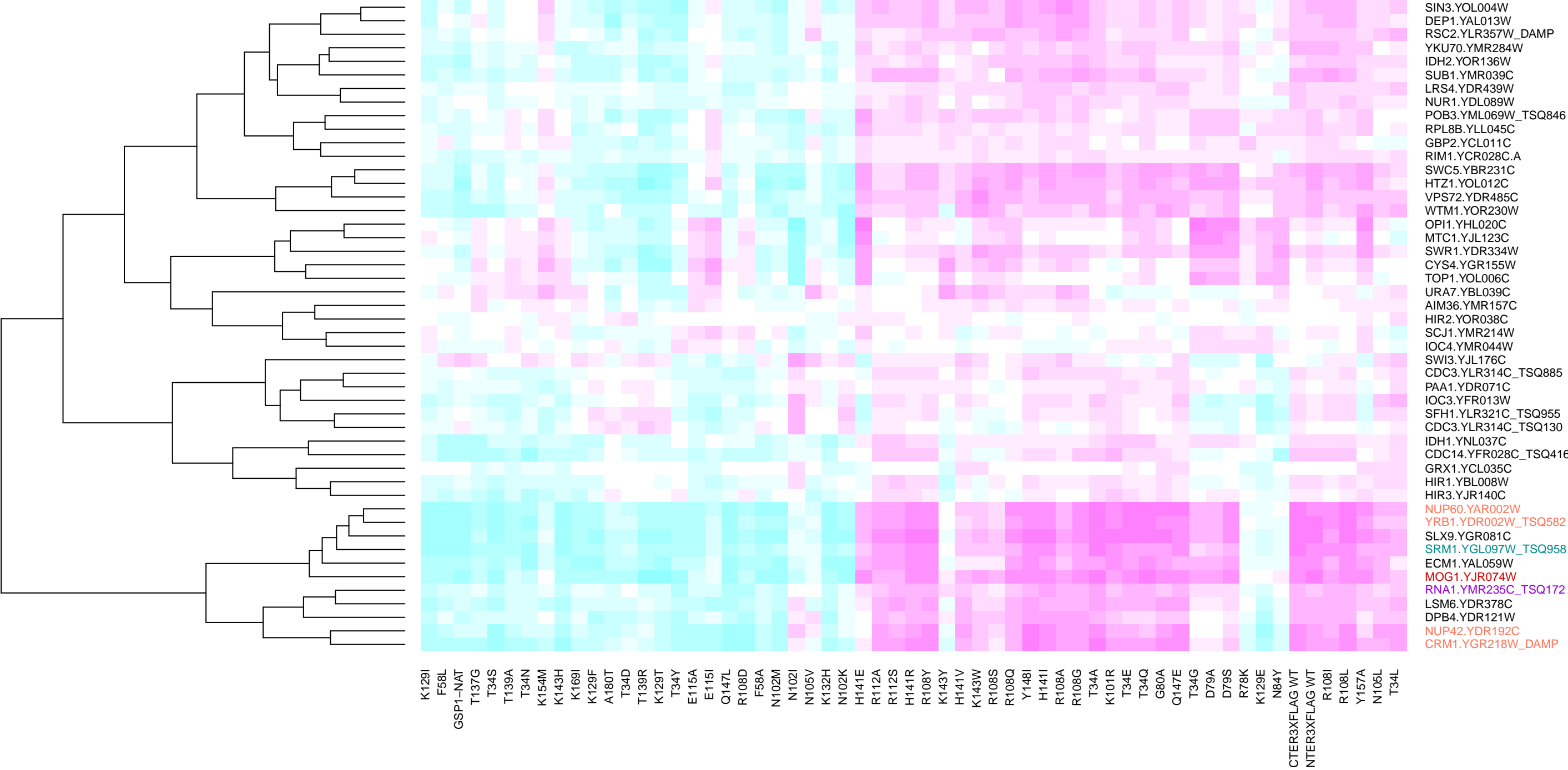
chromatin_GO_30_2_all



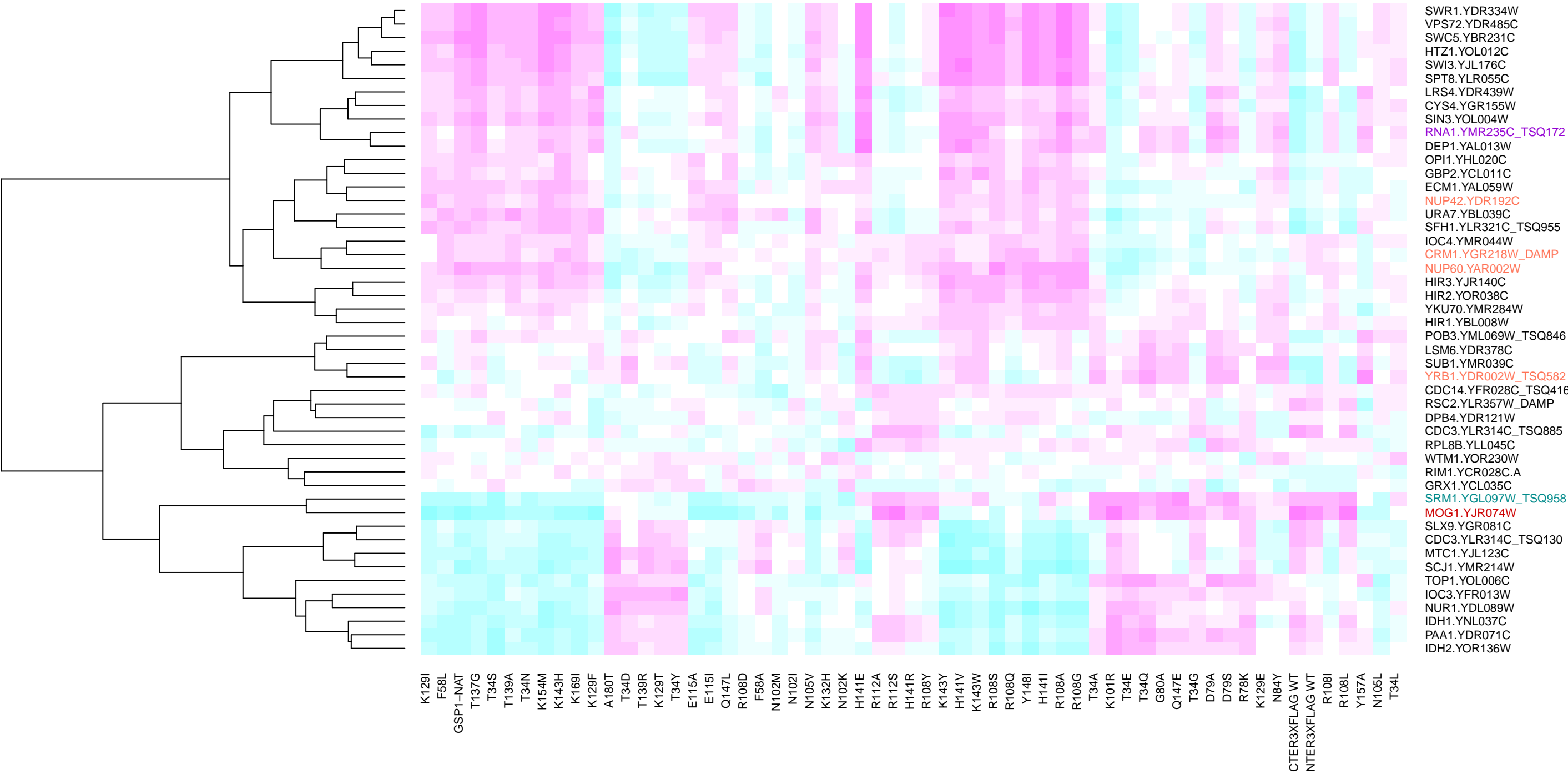
whole_library_15_13_all



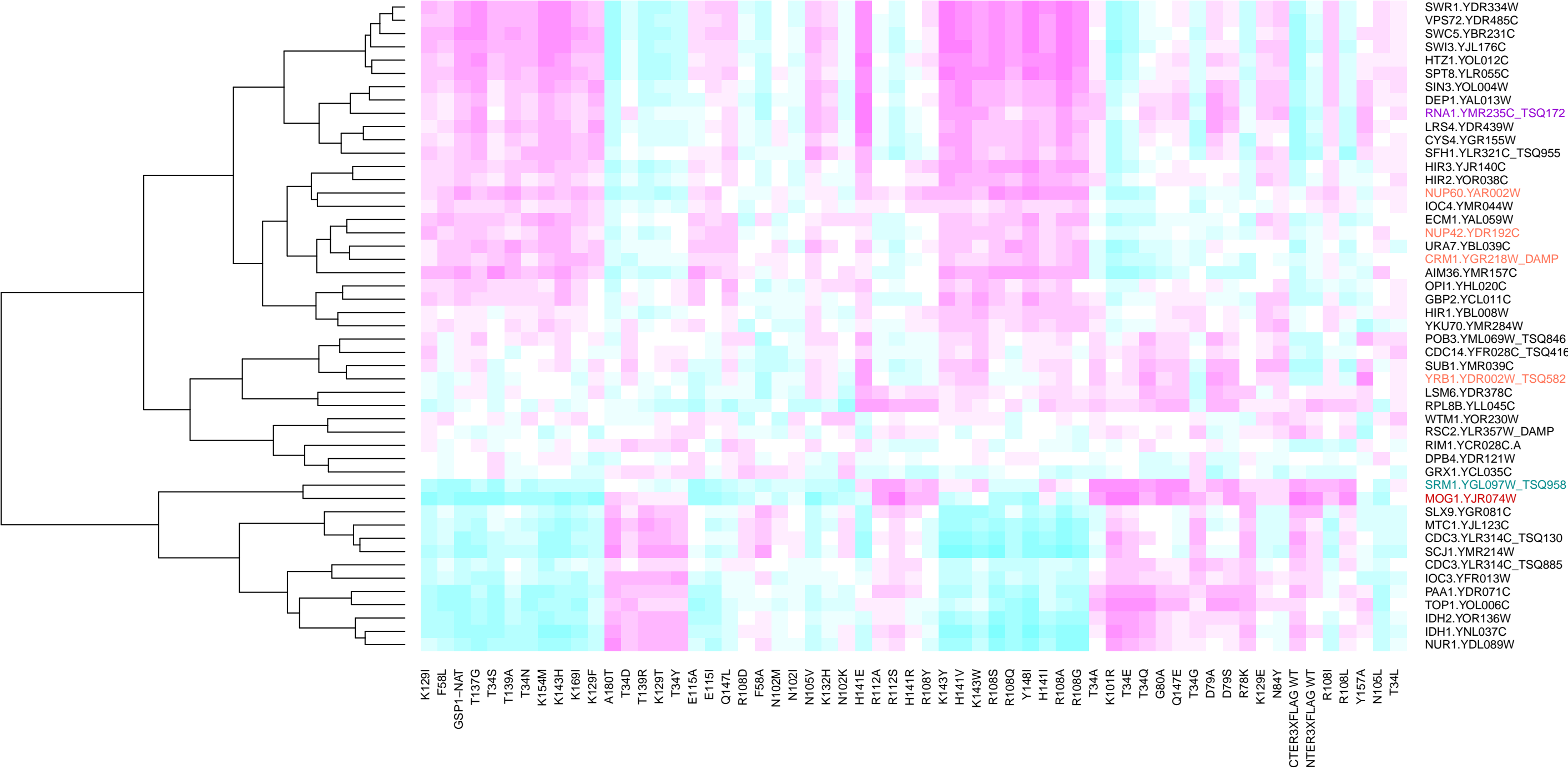
protein targeting



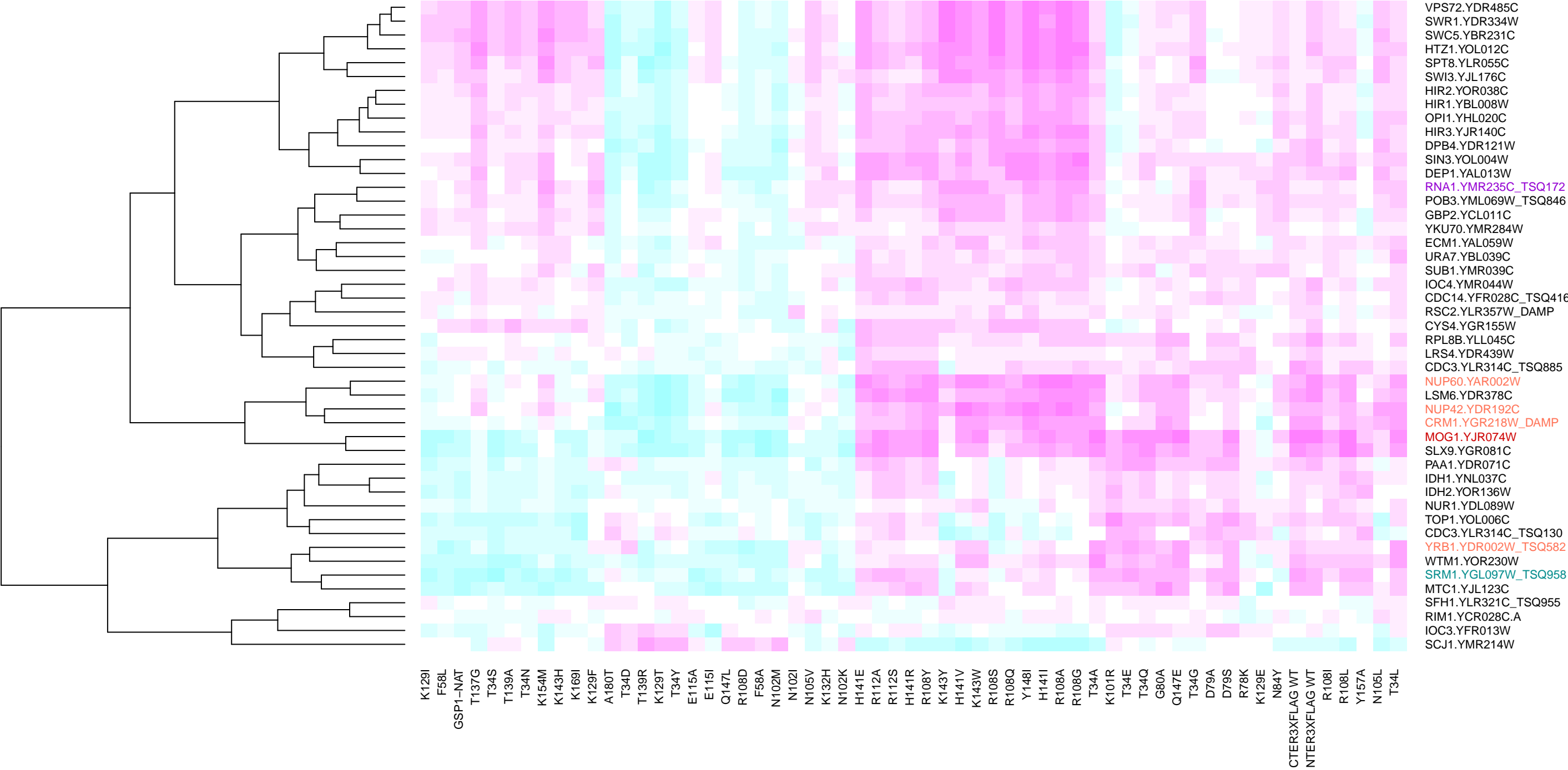
histone modification



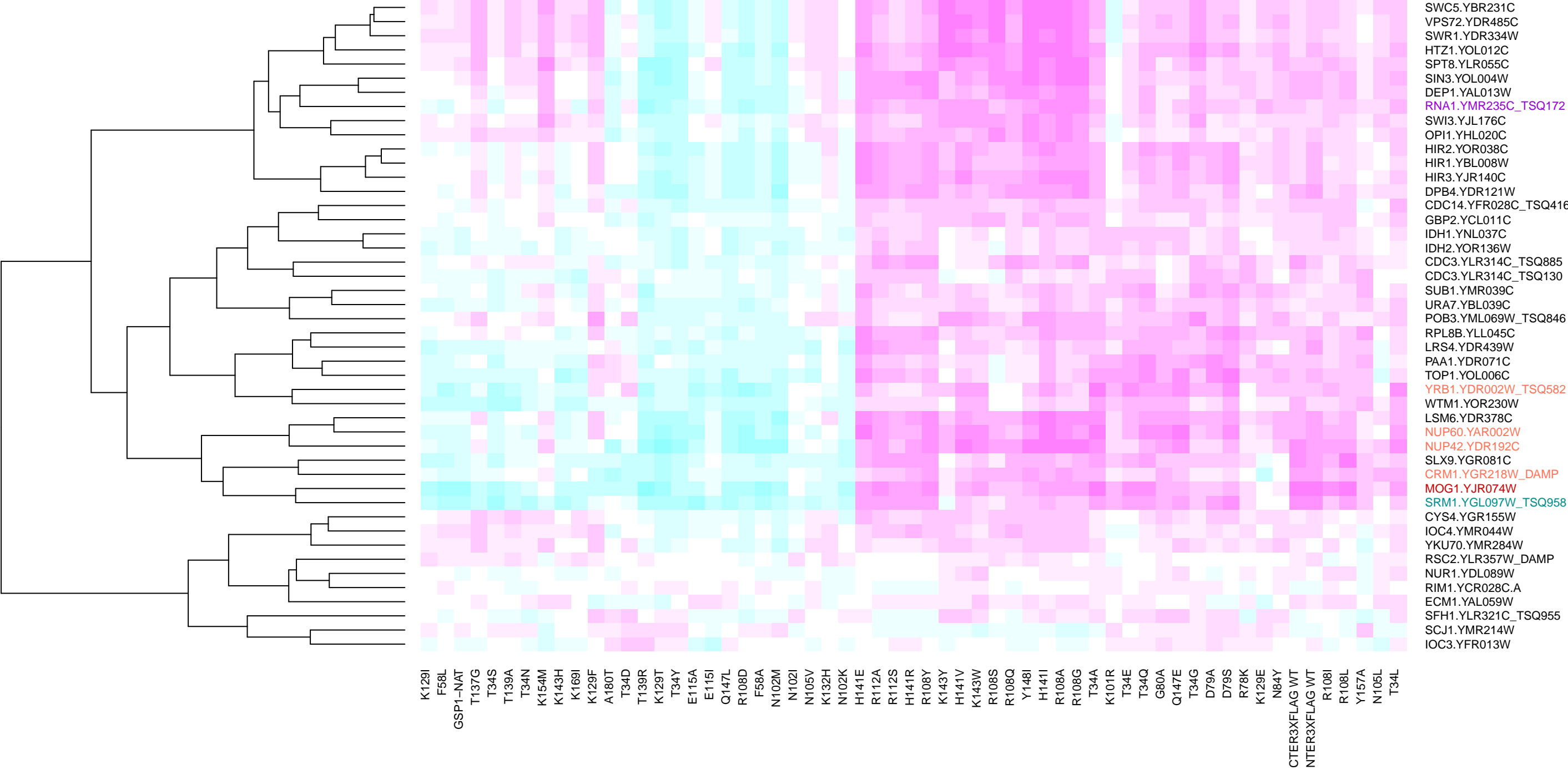
histone modification_GO_30_1_all



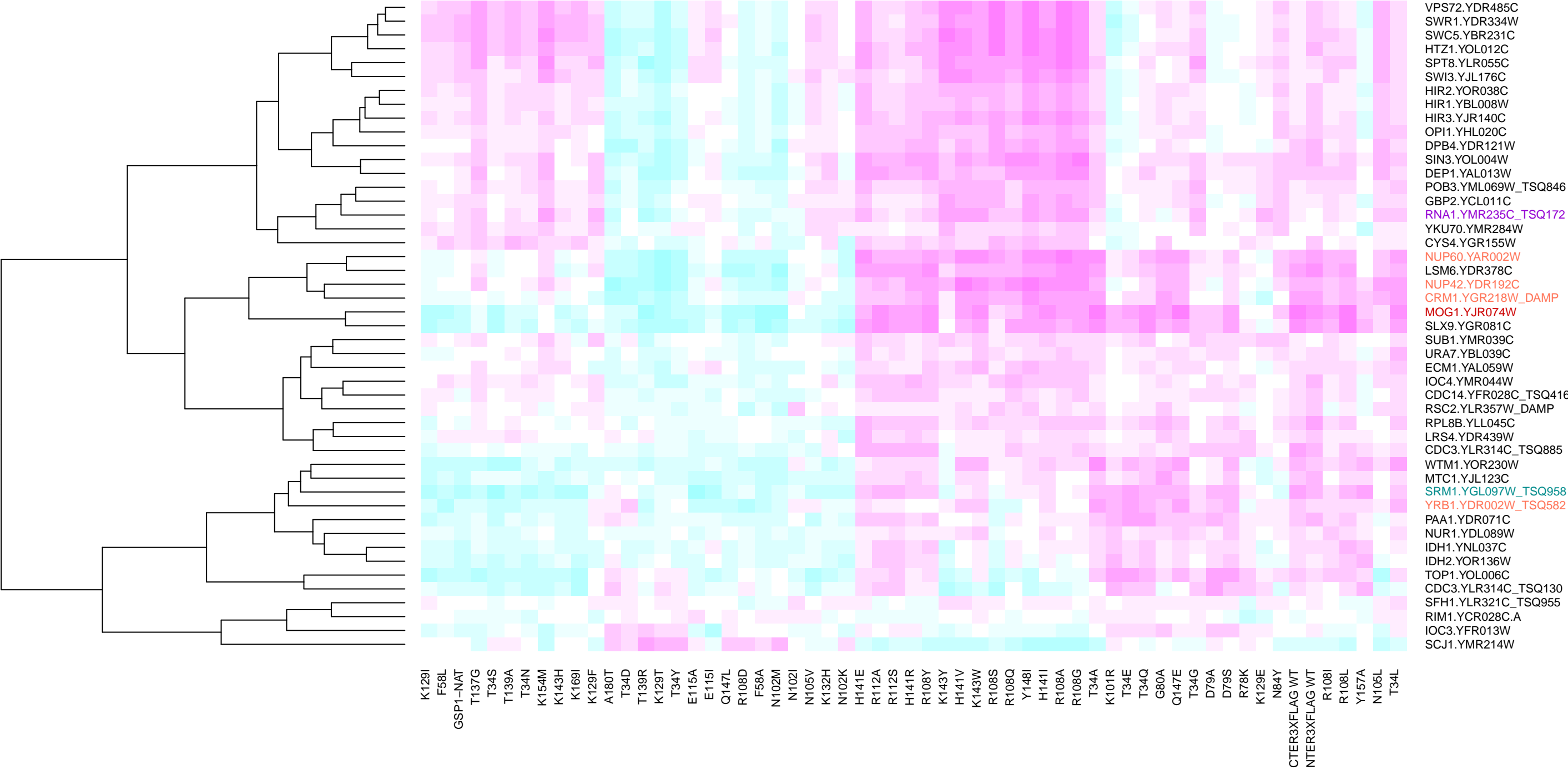
transcription



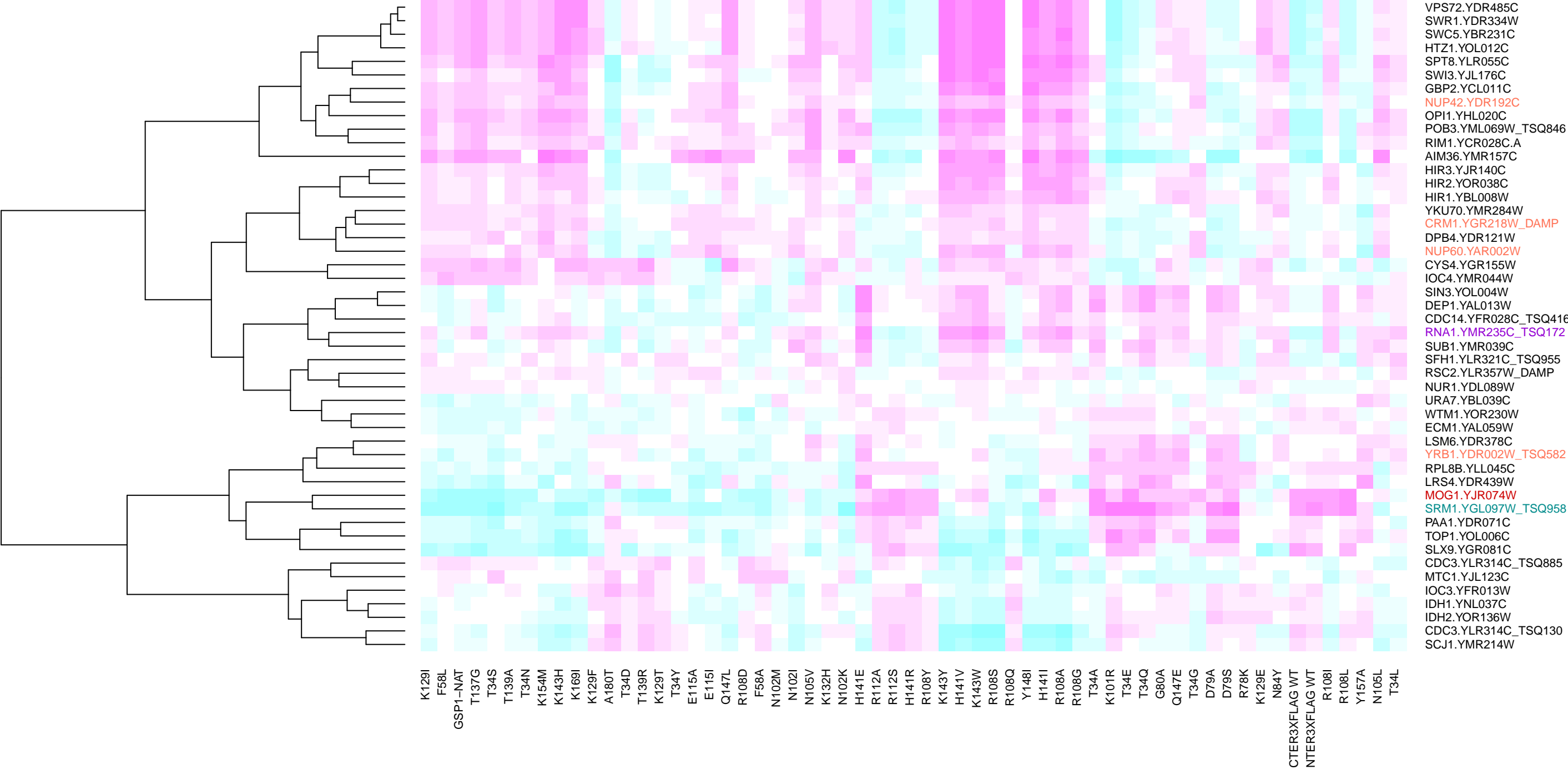
transcription and mRNA processing



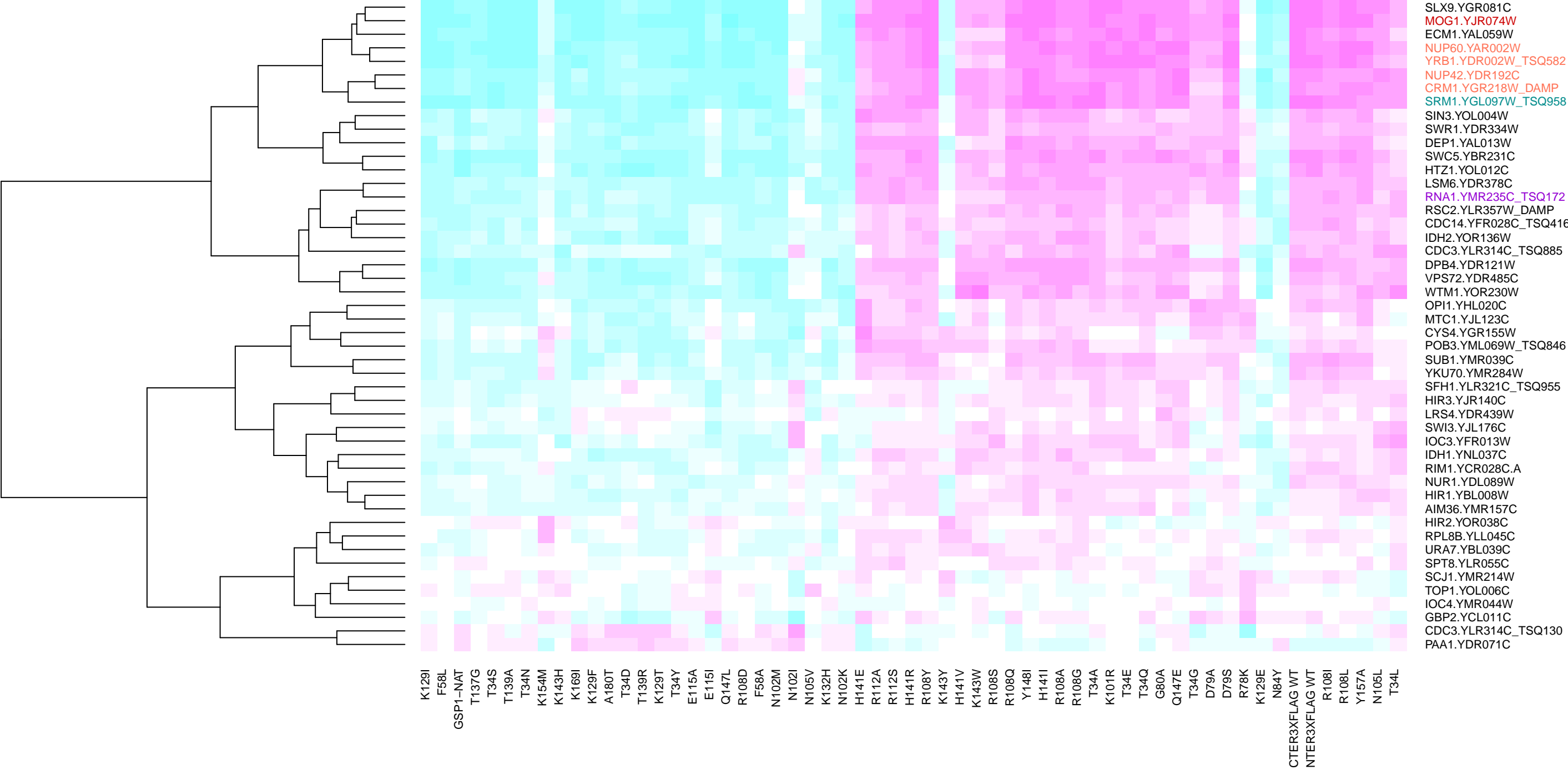
transcription from RNA polymerase II promoter



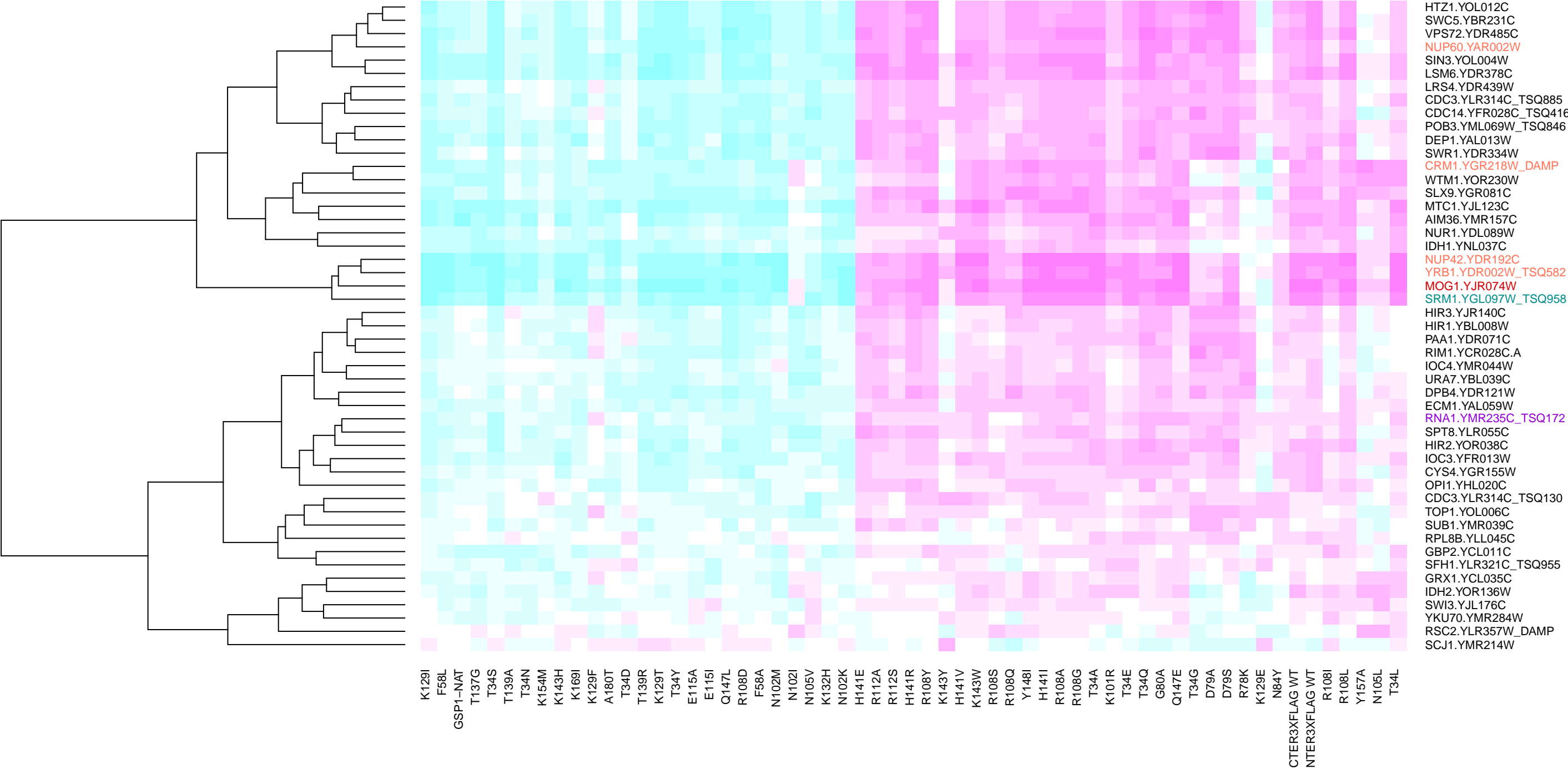
whole_library_30_1_all



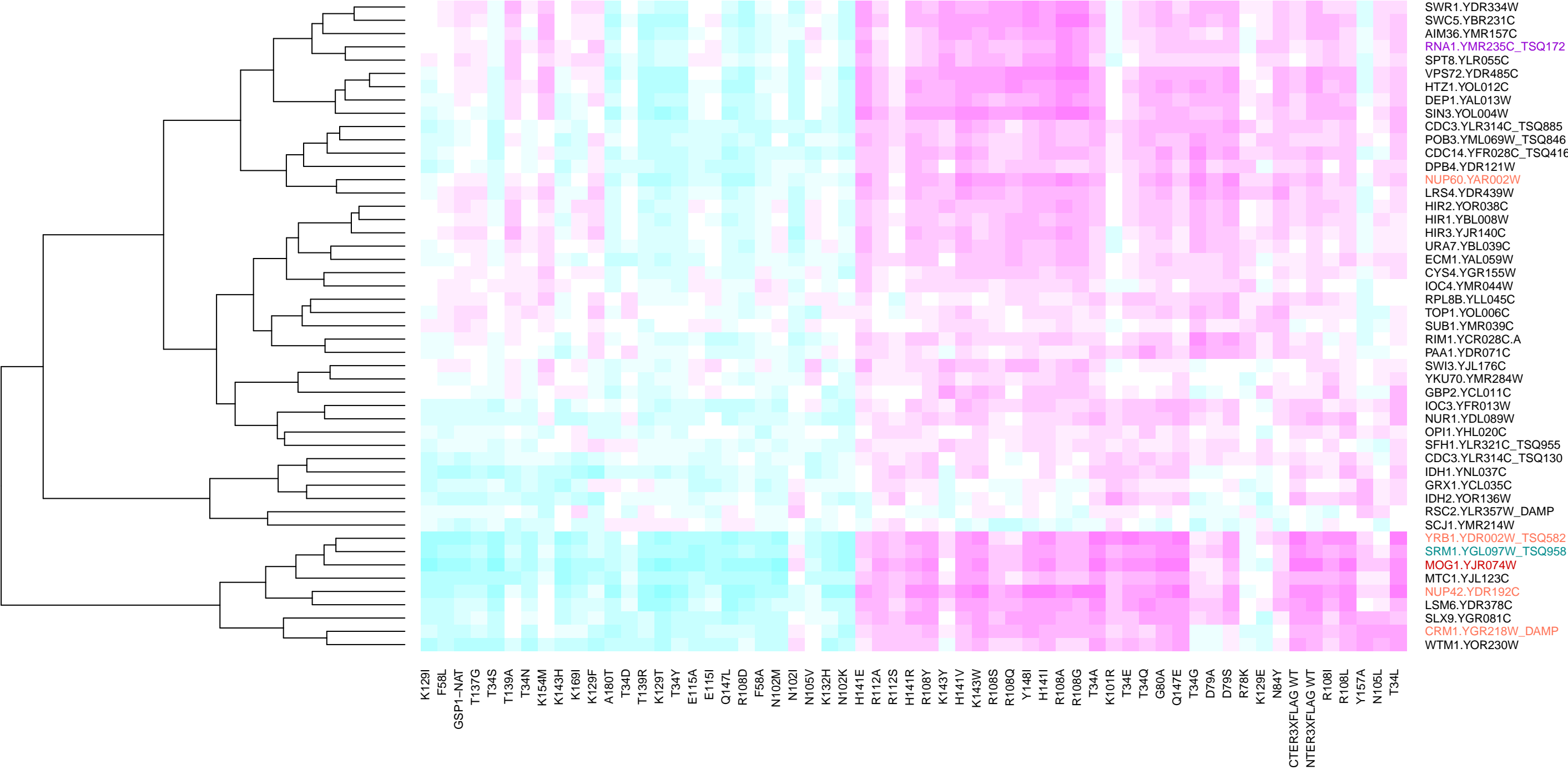
protein targeting_GO_15_2_mut



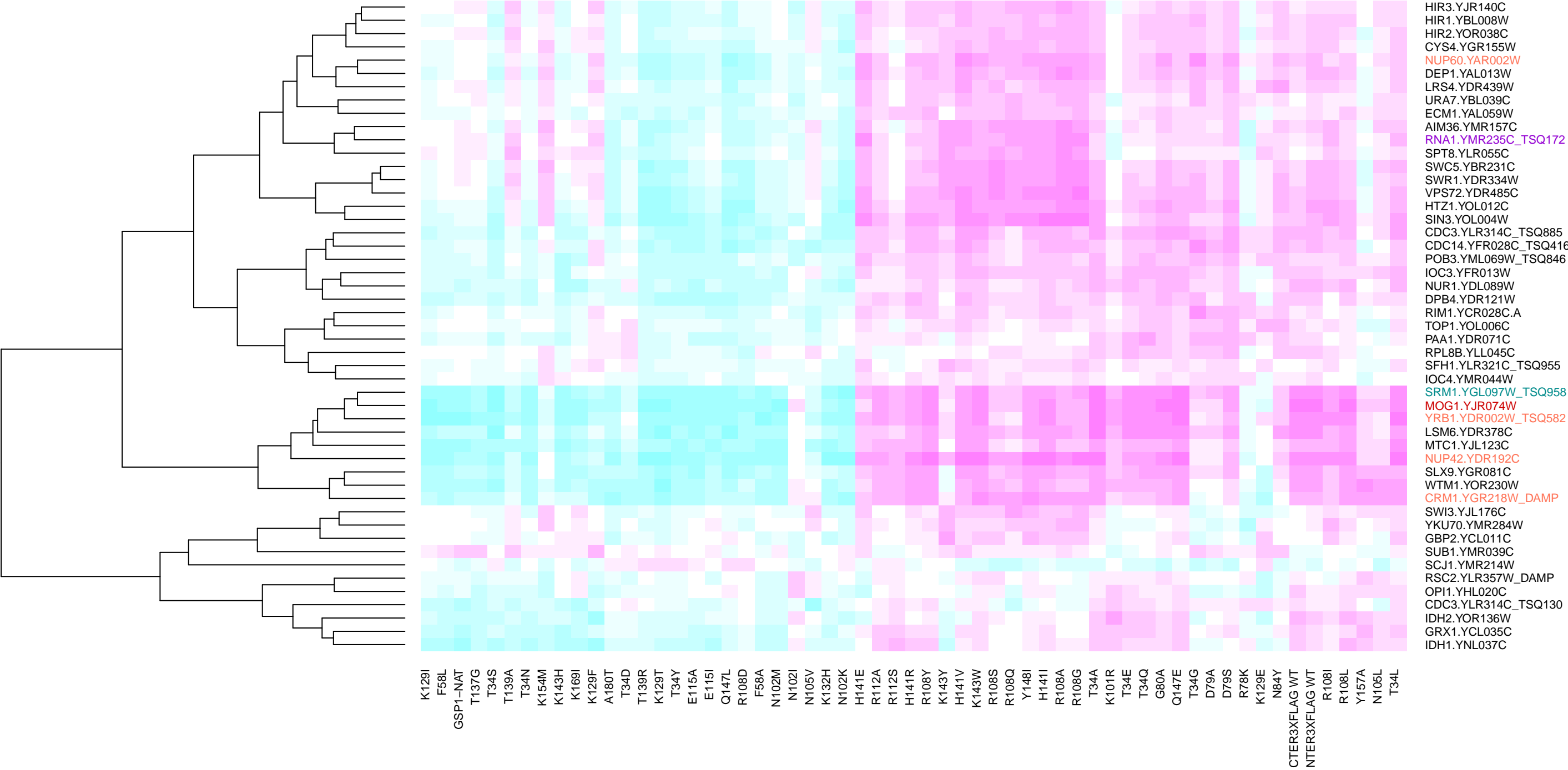
cellular response to DNA damage stimulus_GO_30_1_mut



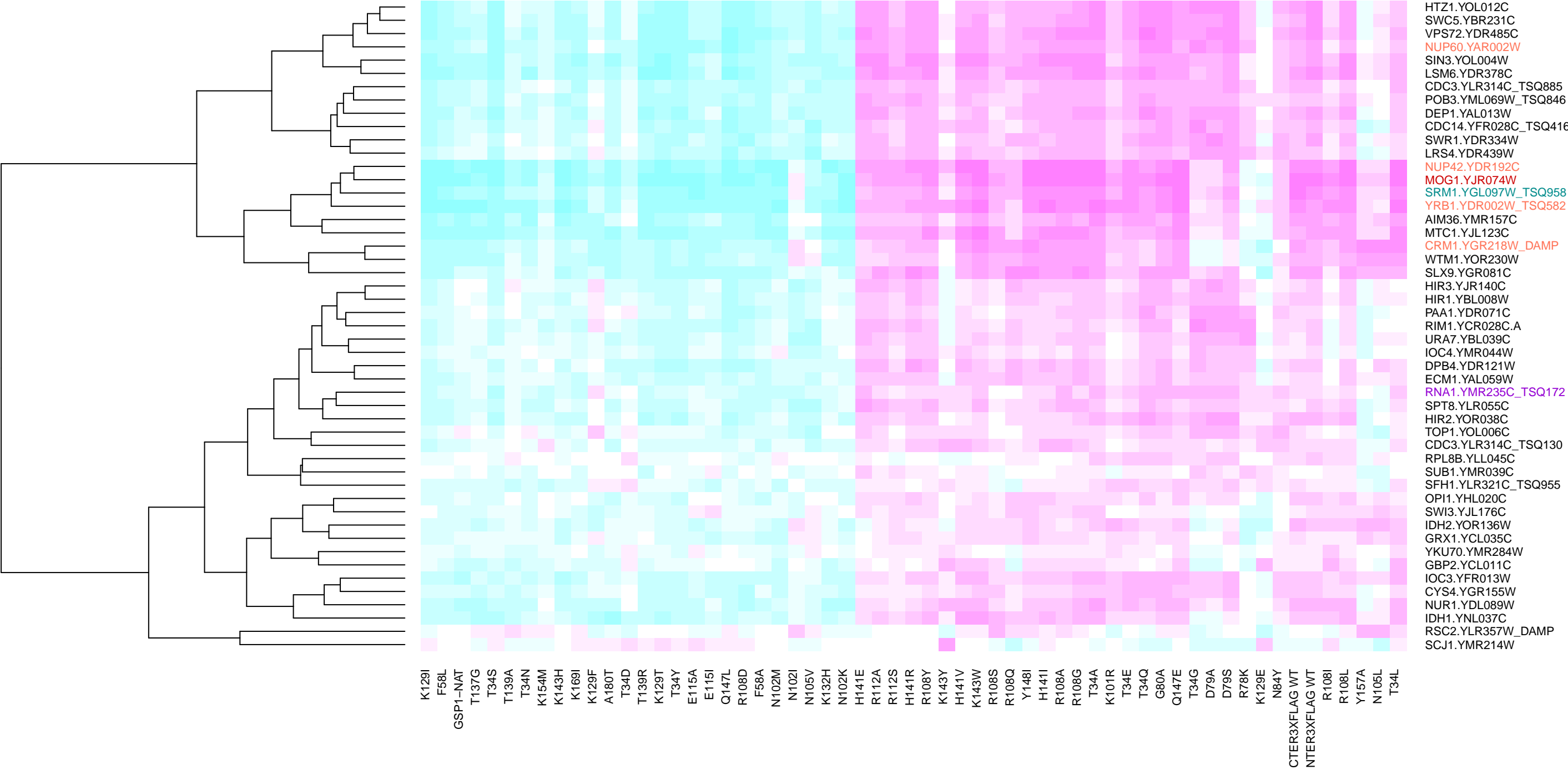
DNA repair



DNA repair_GO_15_1_all



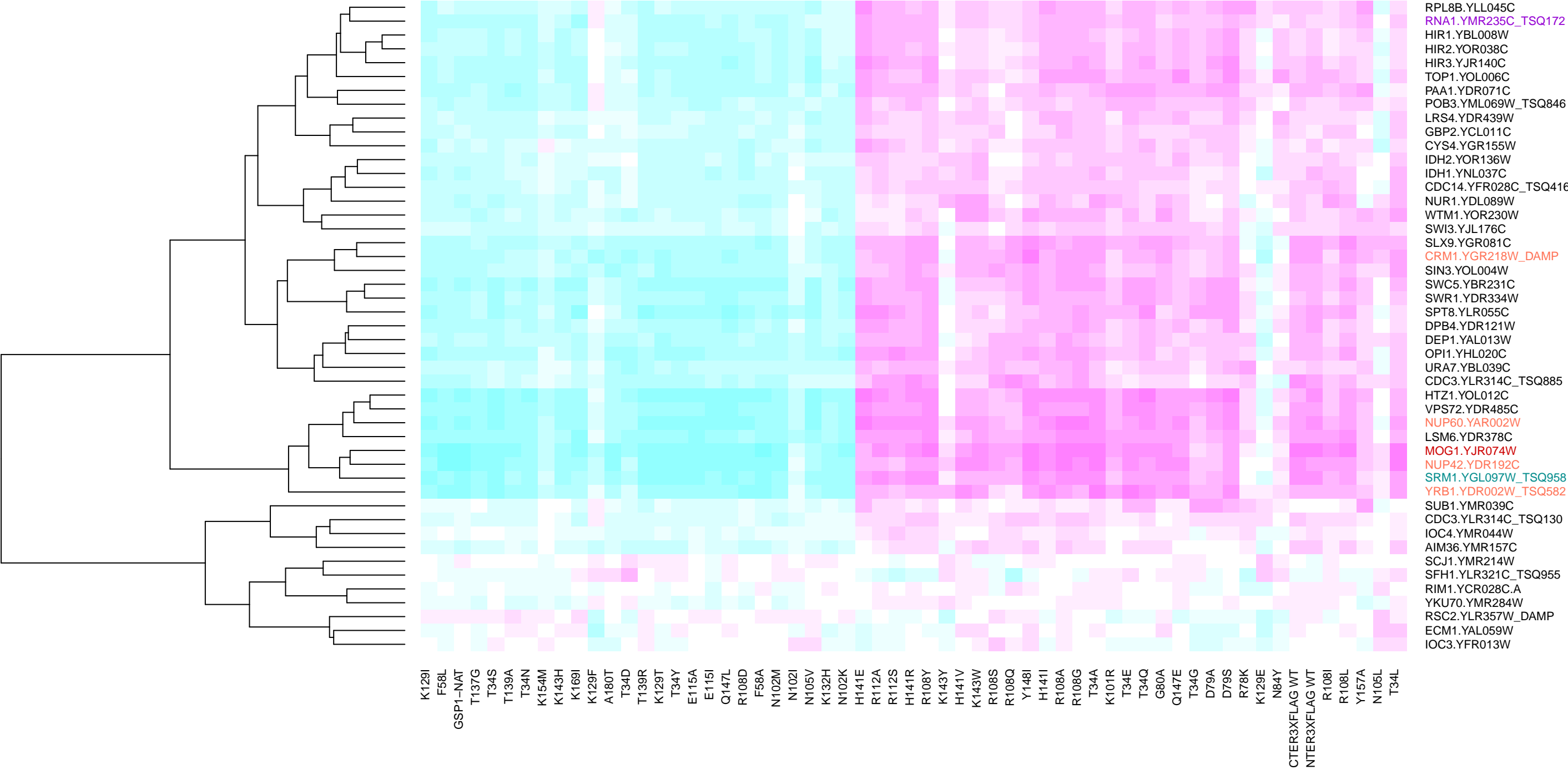
DNA repair_GO_30_1_mut



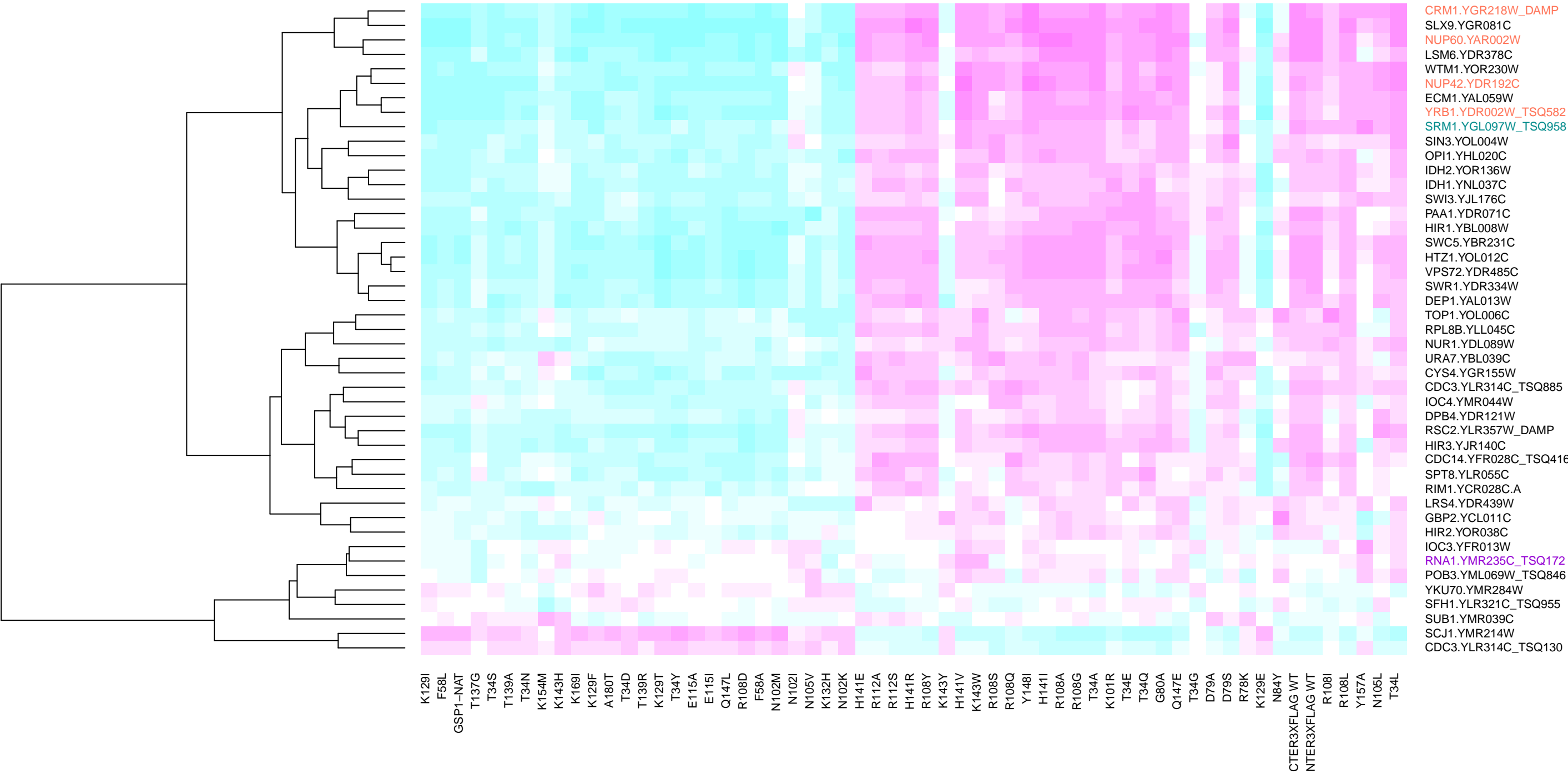
transcription and mRNA processing_GO_15_14_mut



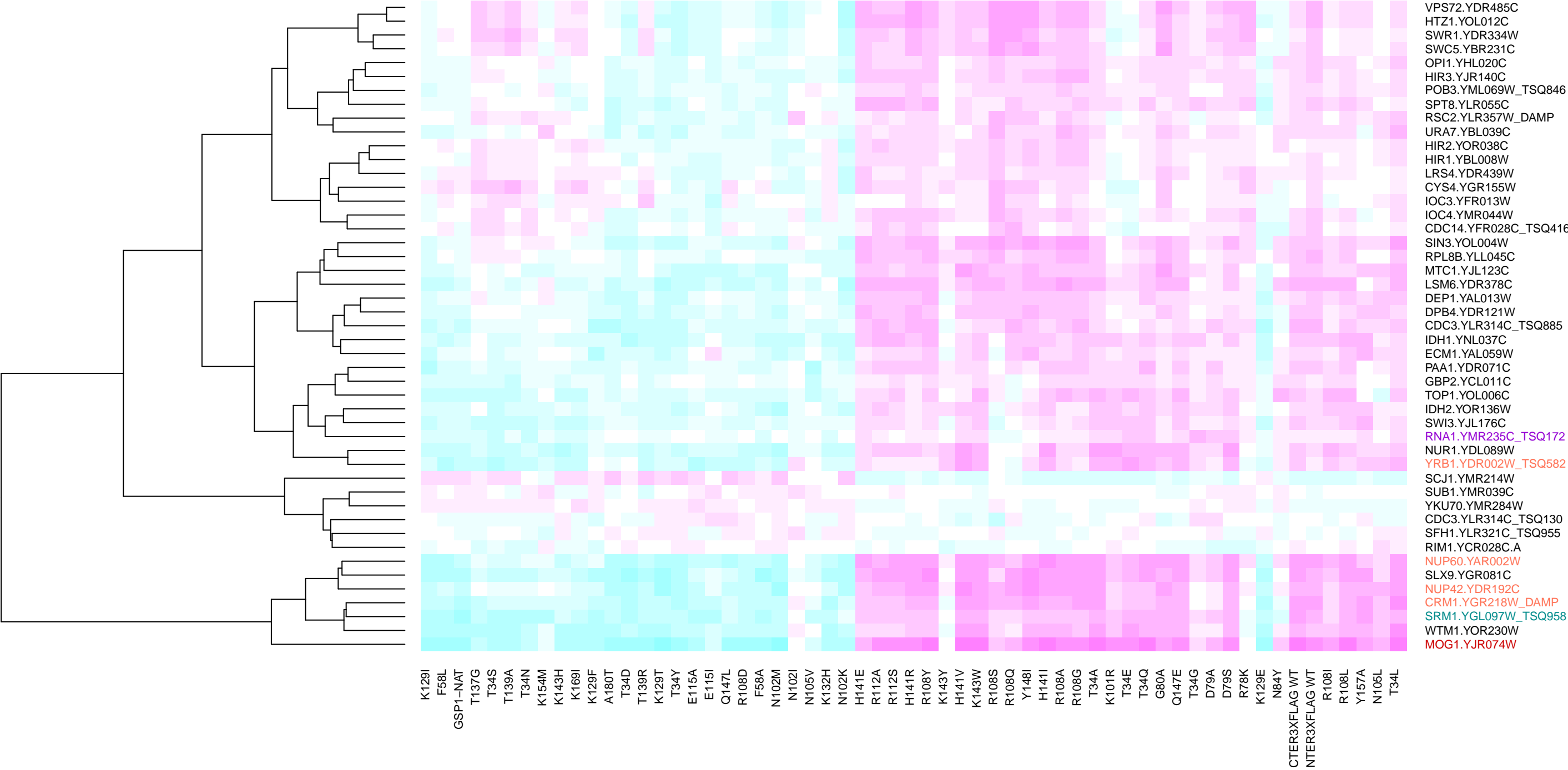
transcription and mRNA processing_GO_30_2_mut



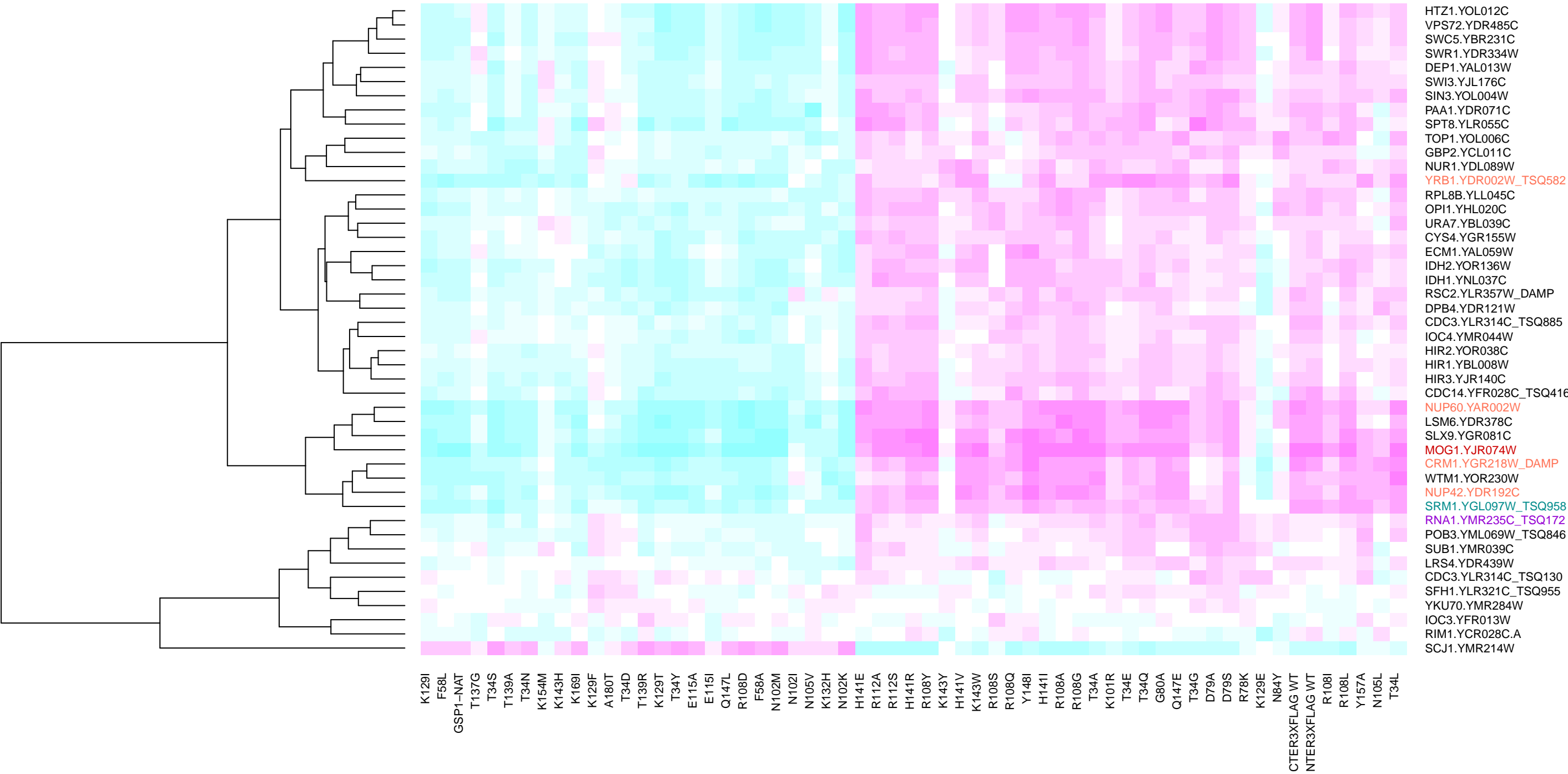
transcription from RNA polymerase II promoter_GO_15_3_mut



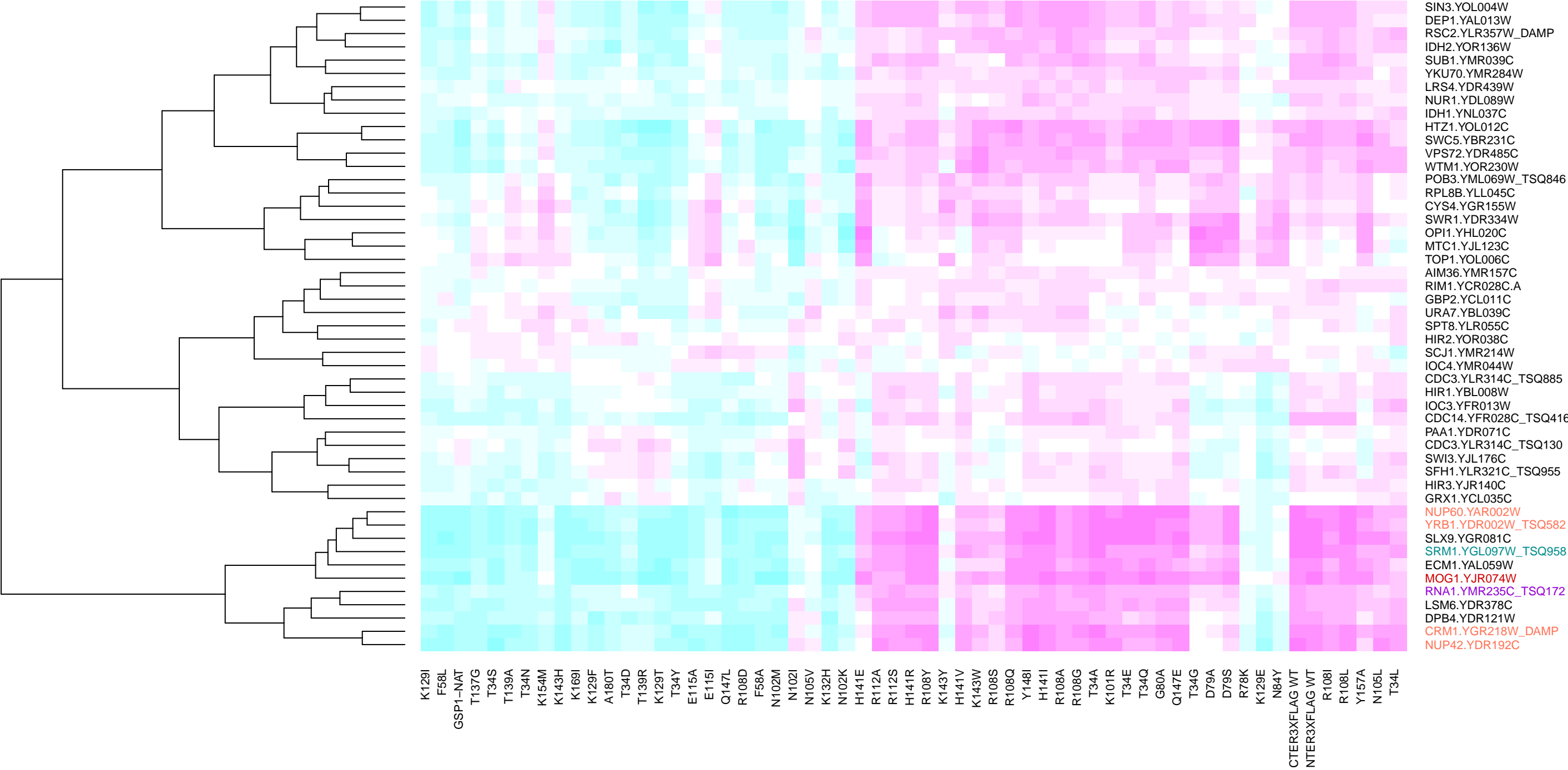
transcription from RNA polymerase II promoter_GO_30_1_all



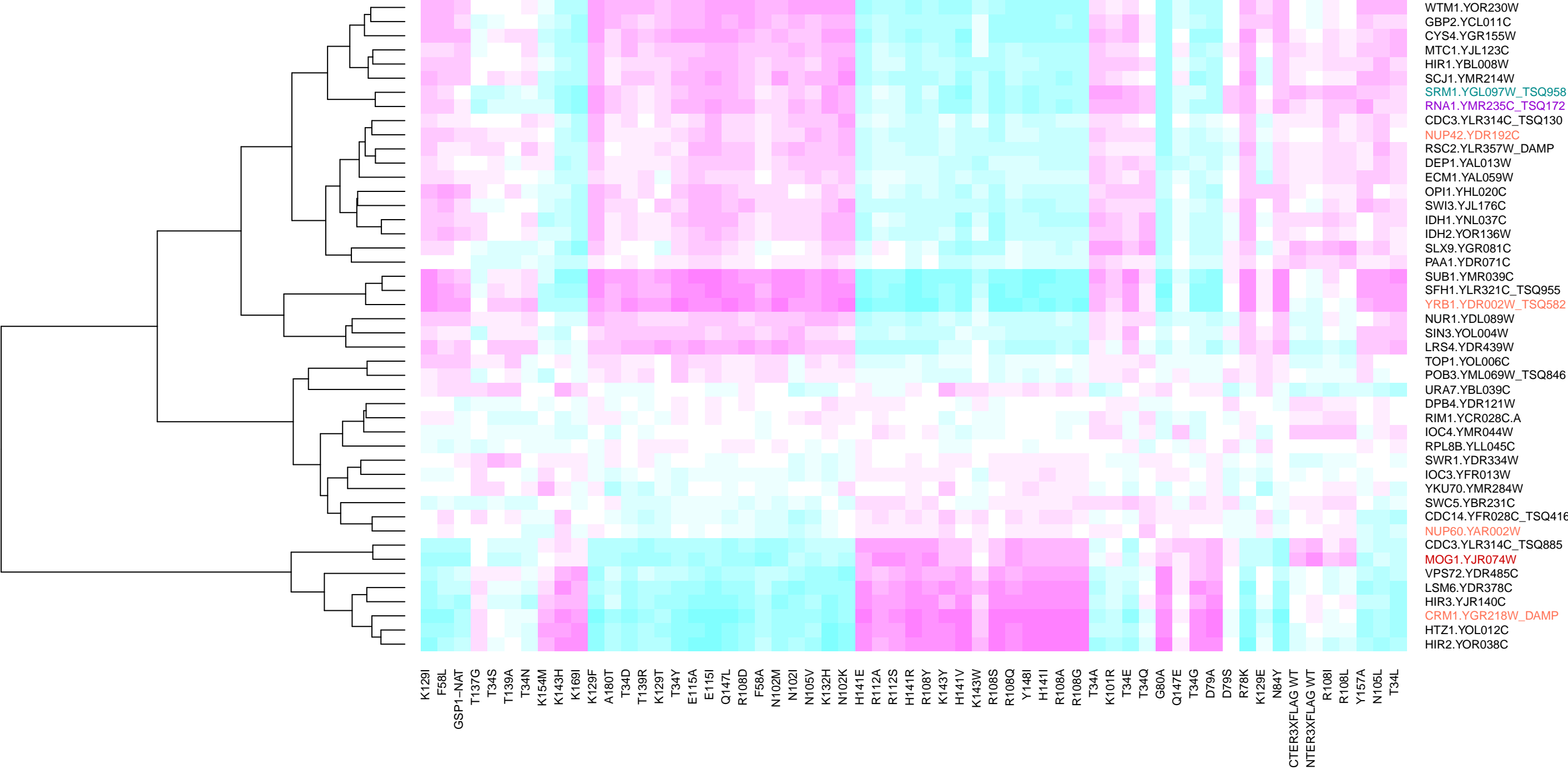
transcription from RNA polymerase II promoter_GO_30_1_mut



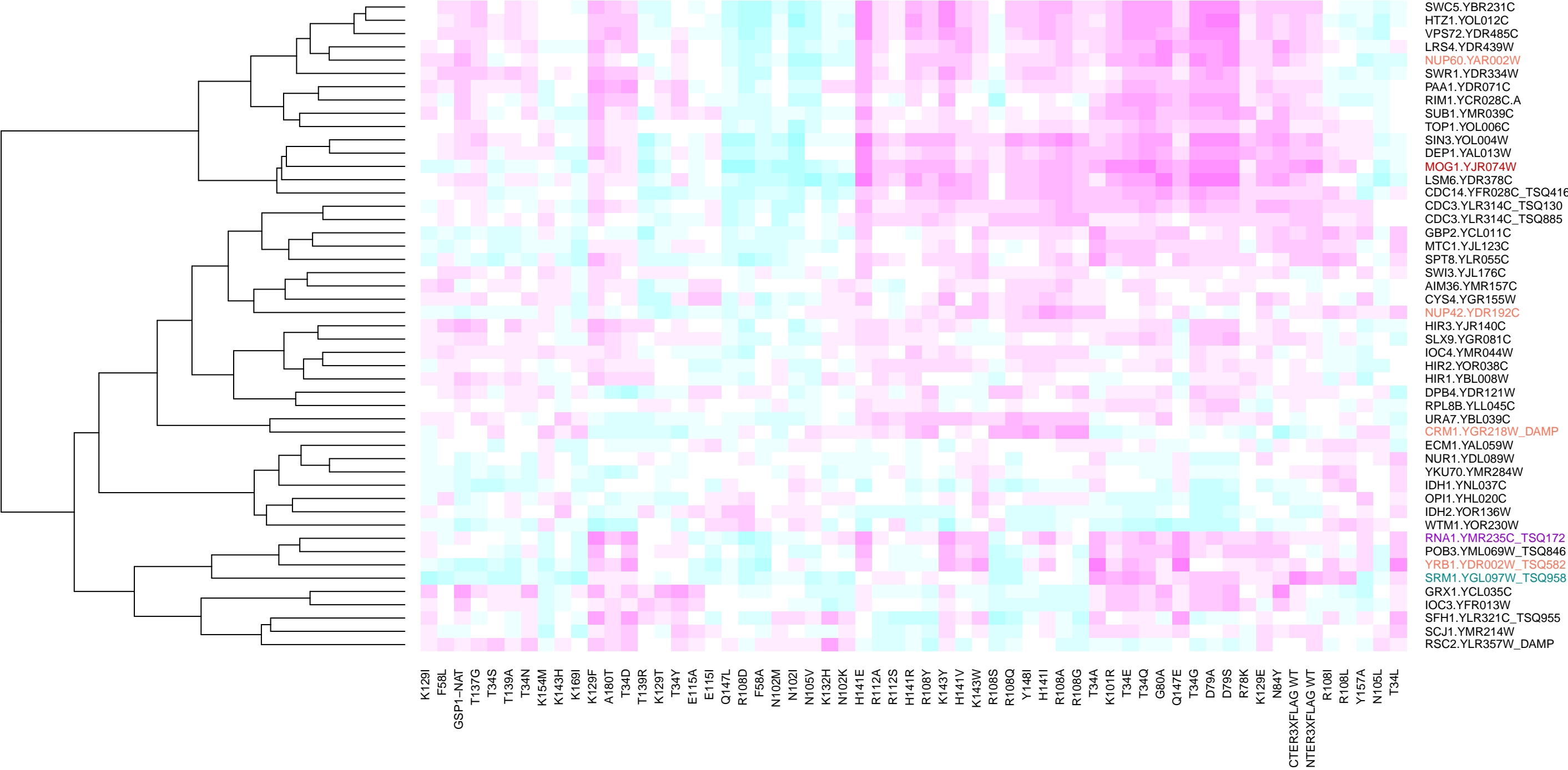
protein targeting_GO_30_1_all



whole_library_30_14_mut



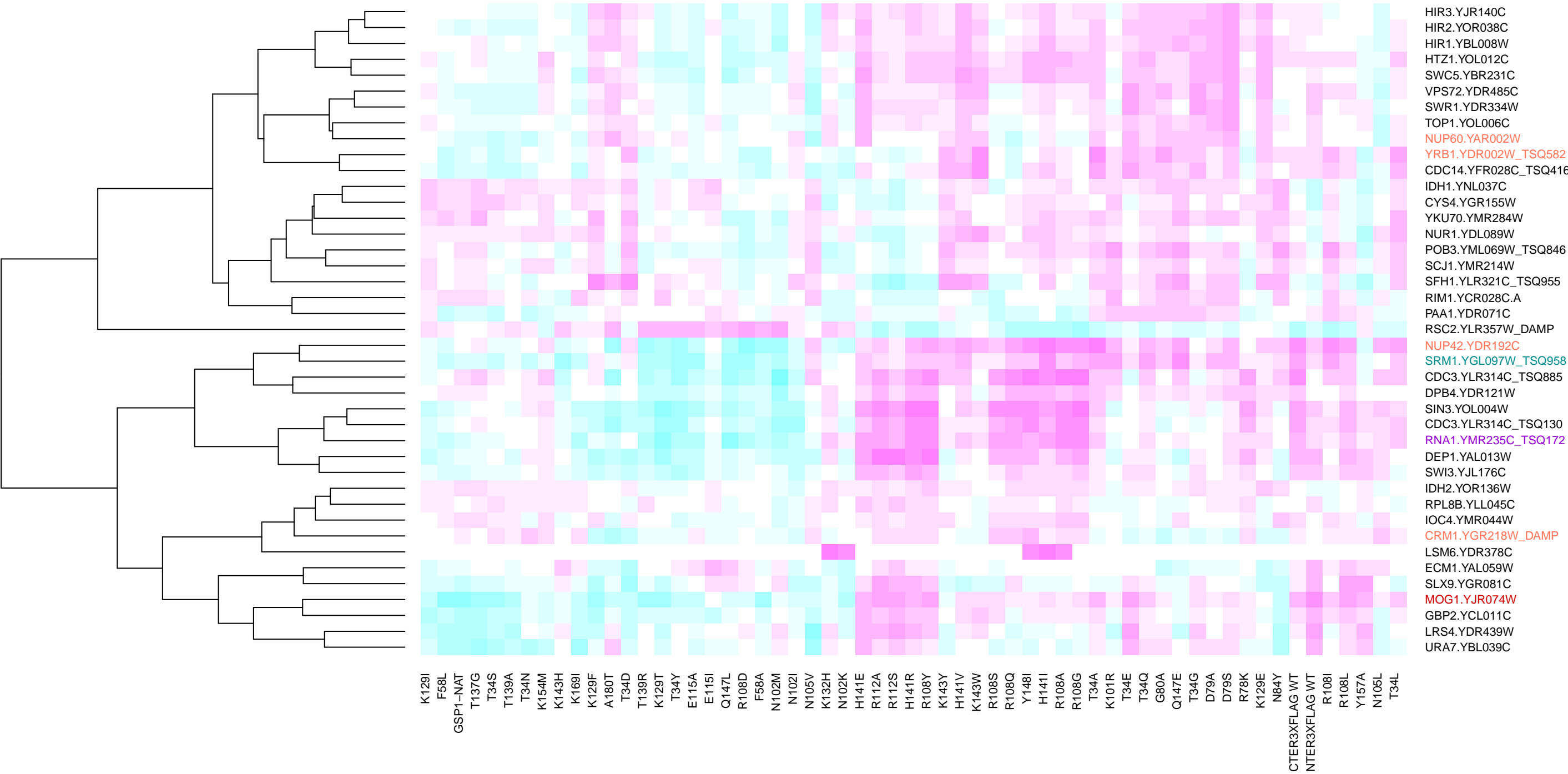
budding



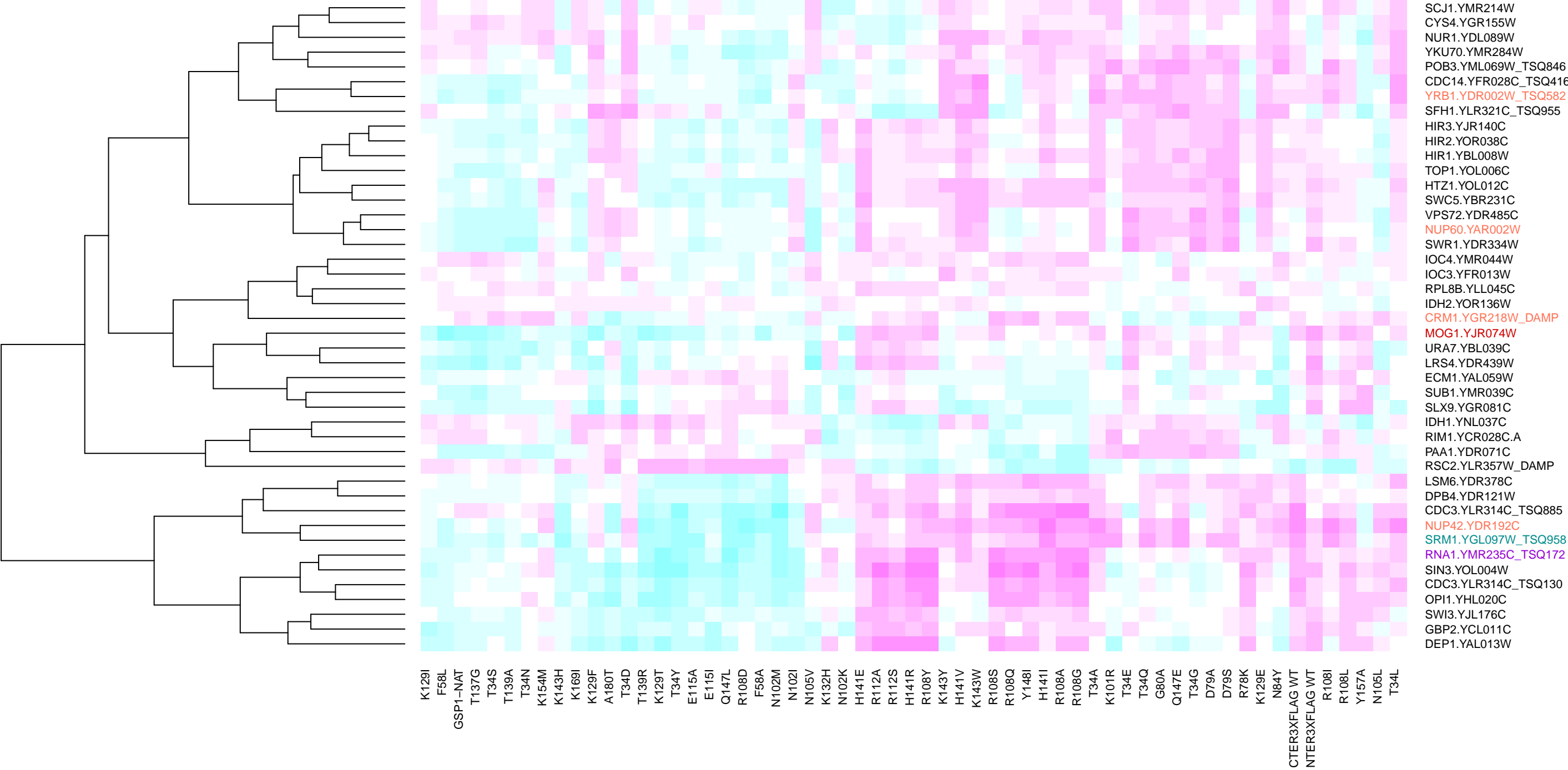
cytoskeleton and microtubules



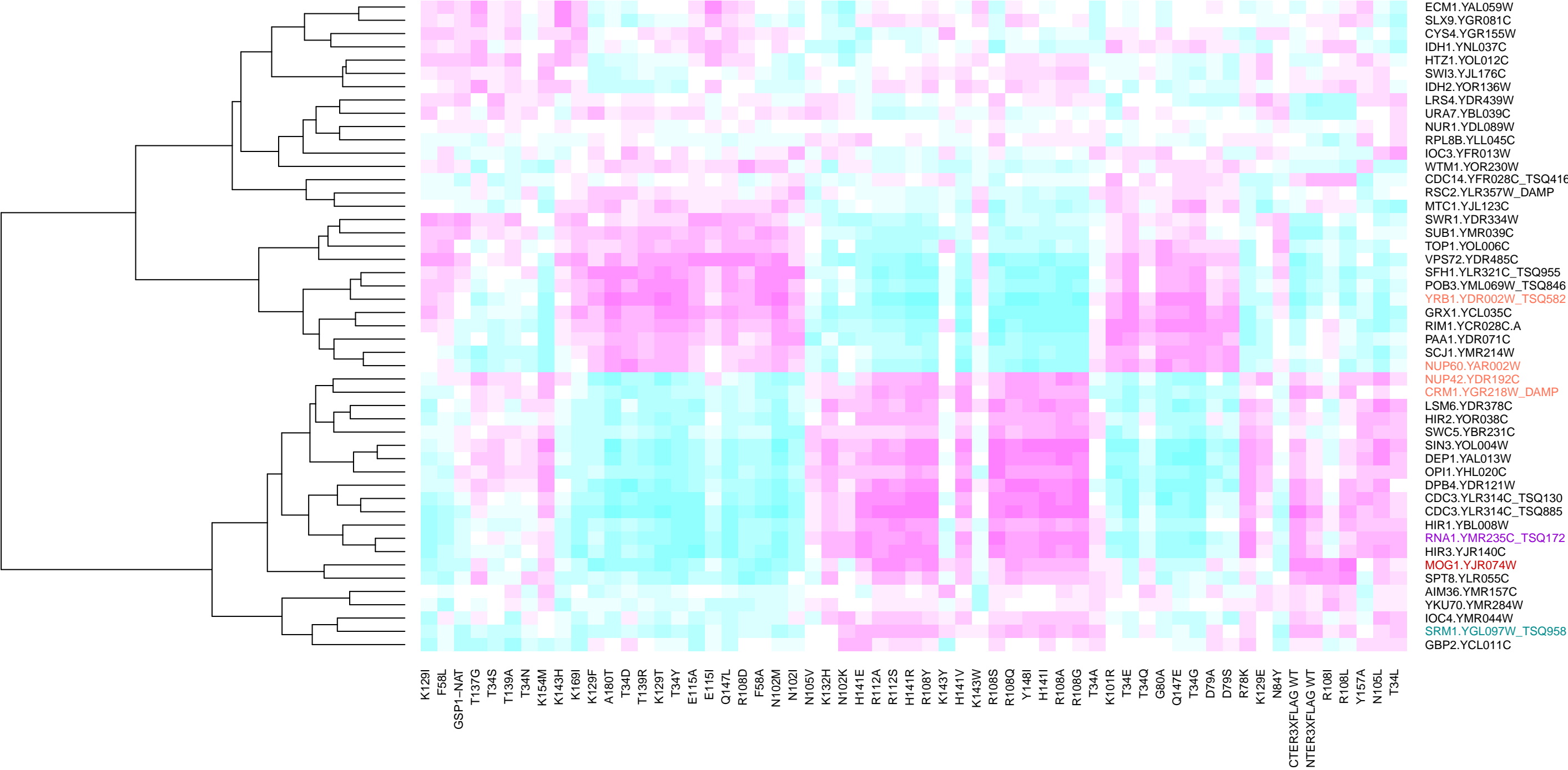
ribosomes and translation



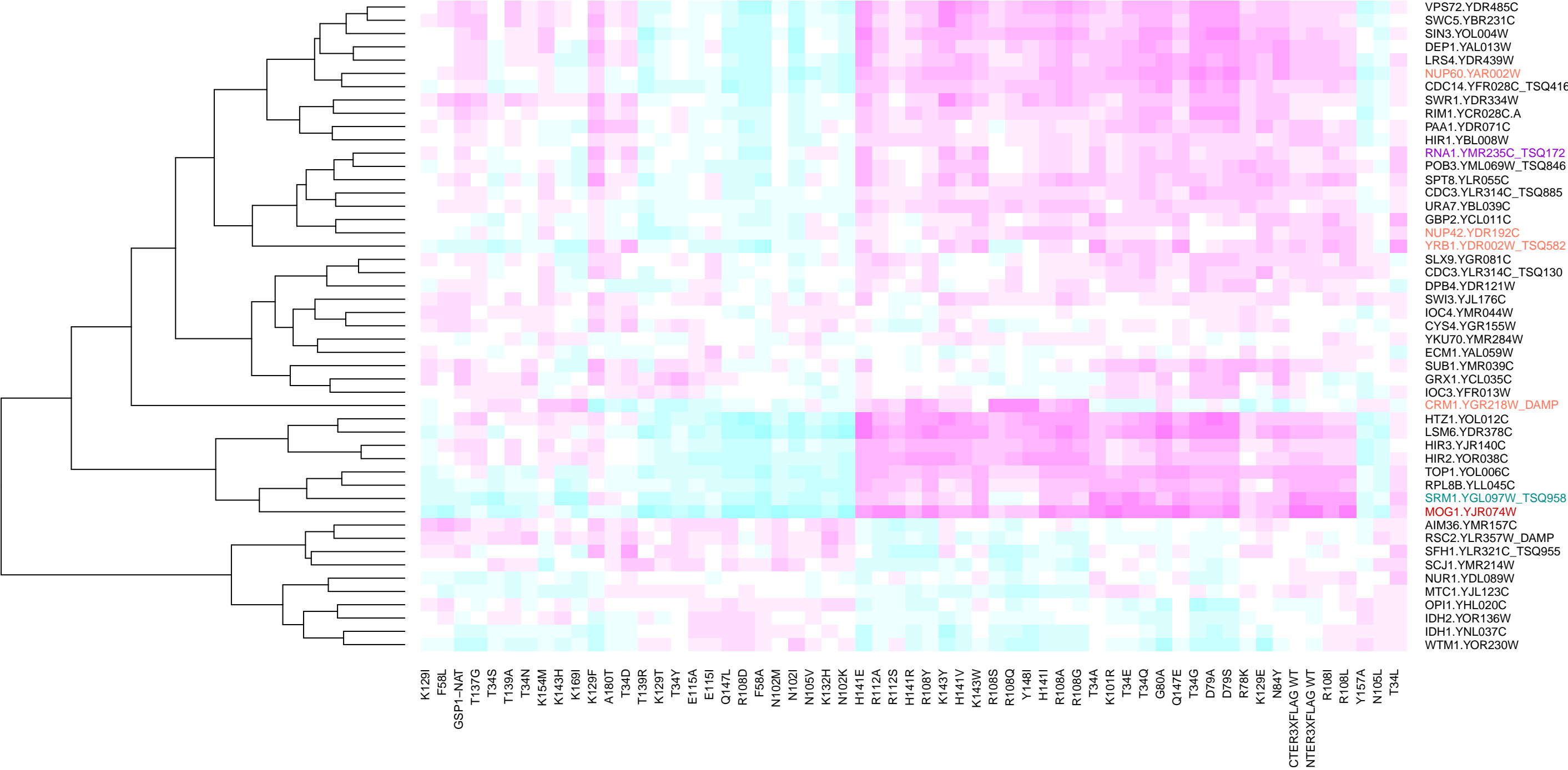
ribosomes and translation_GO_30_1_all



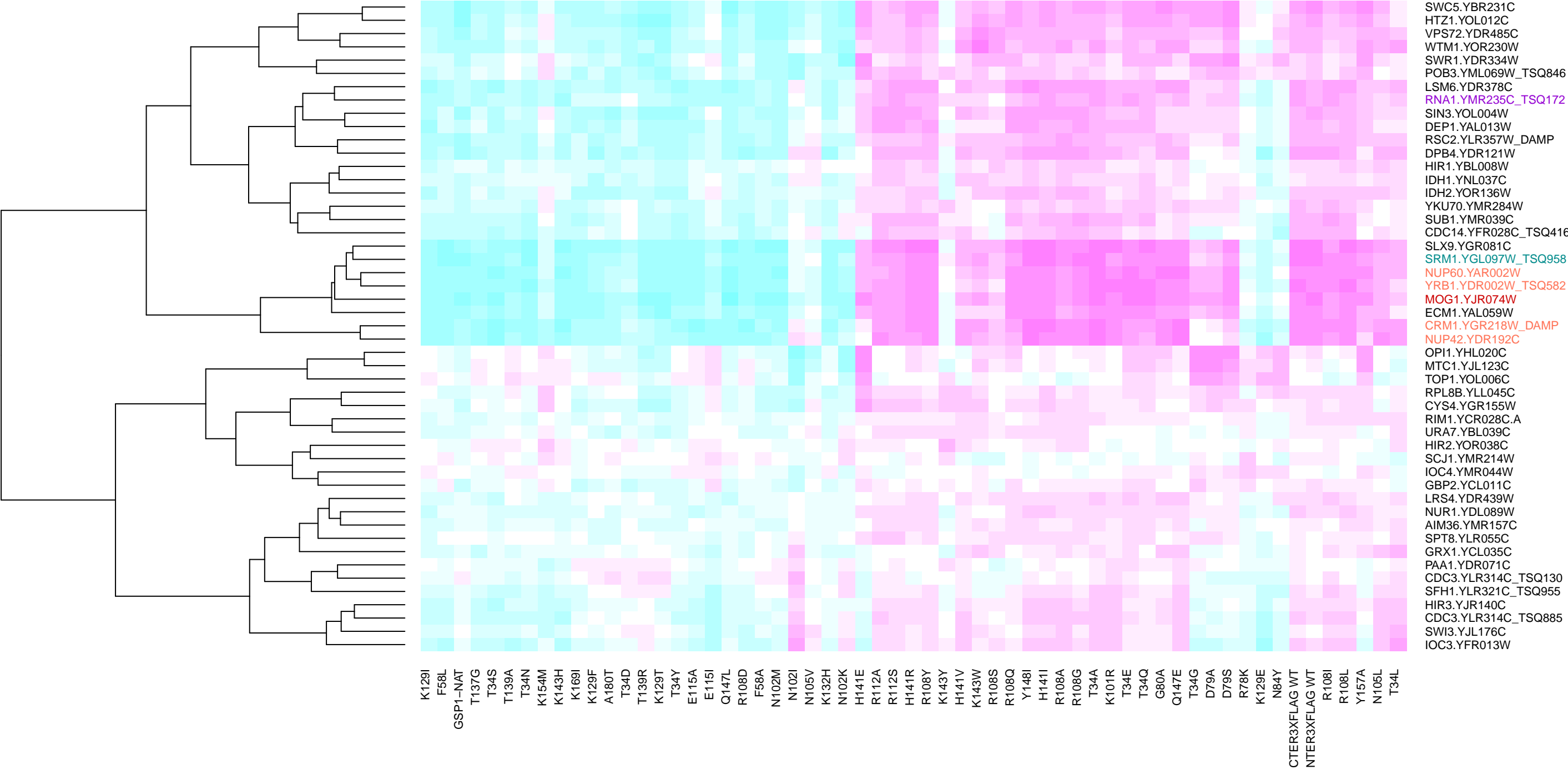
transcription and mRNA processing_GO_15_8_all



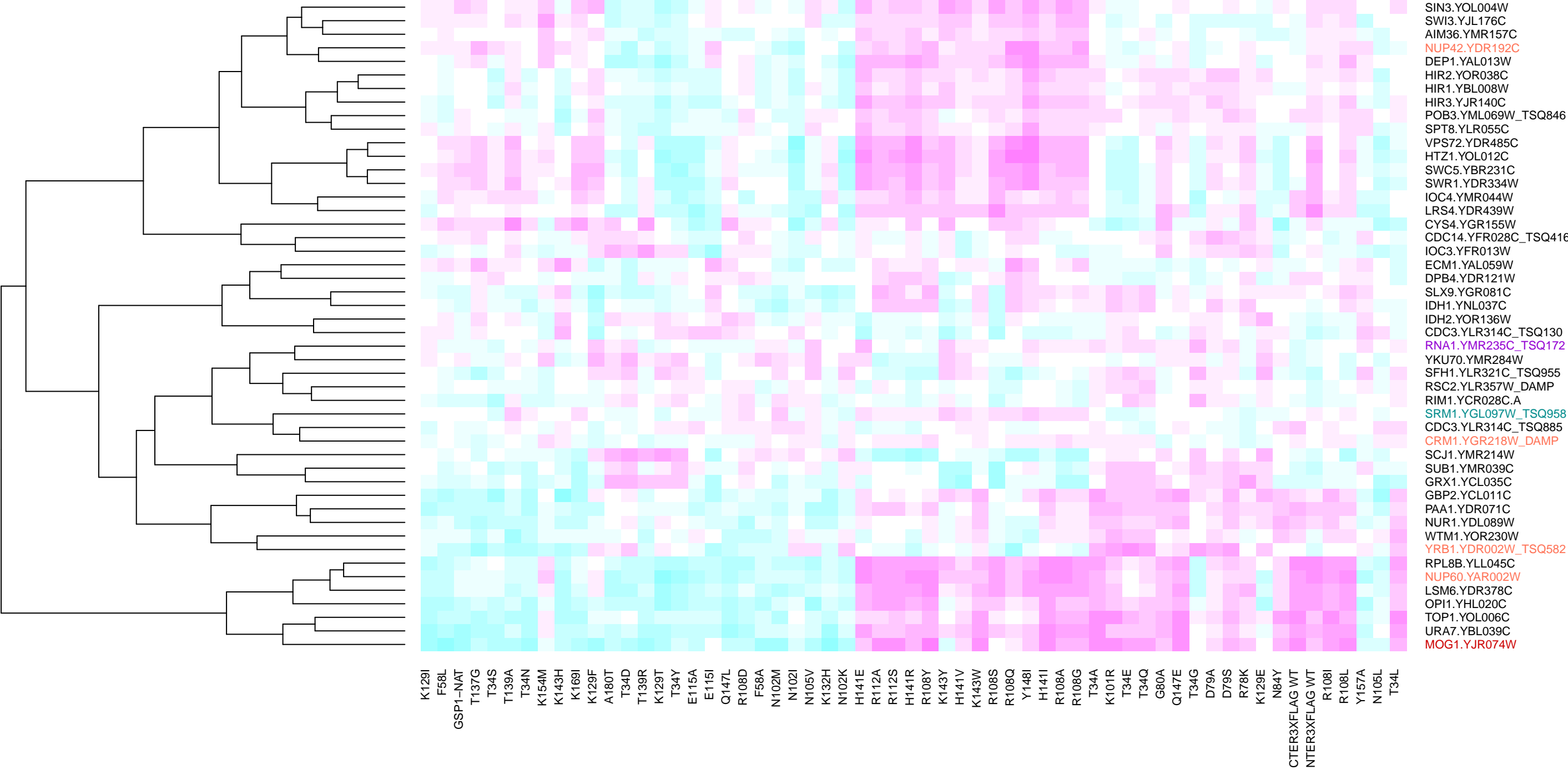
cell cycle



protein targeting_GO_30_1_mut



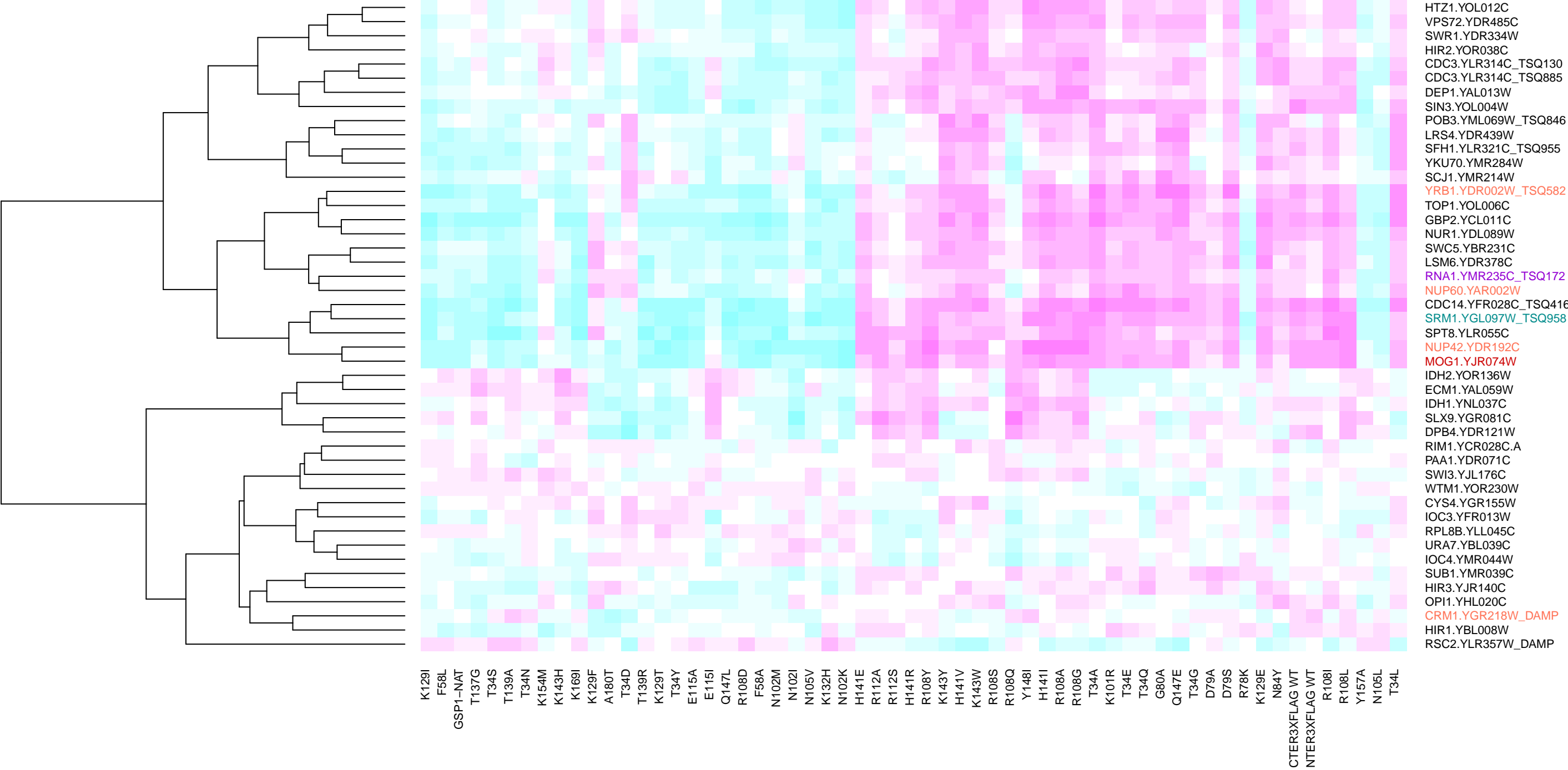
DNA-templated transcription, elongation



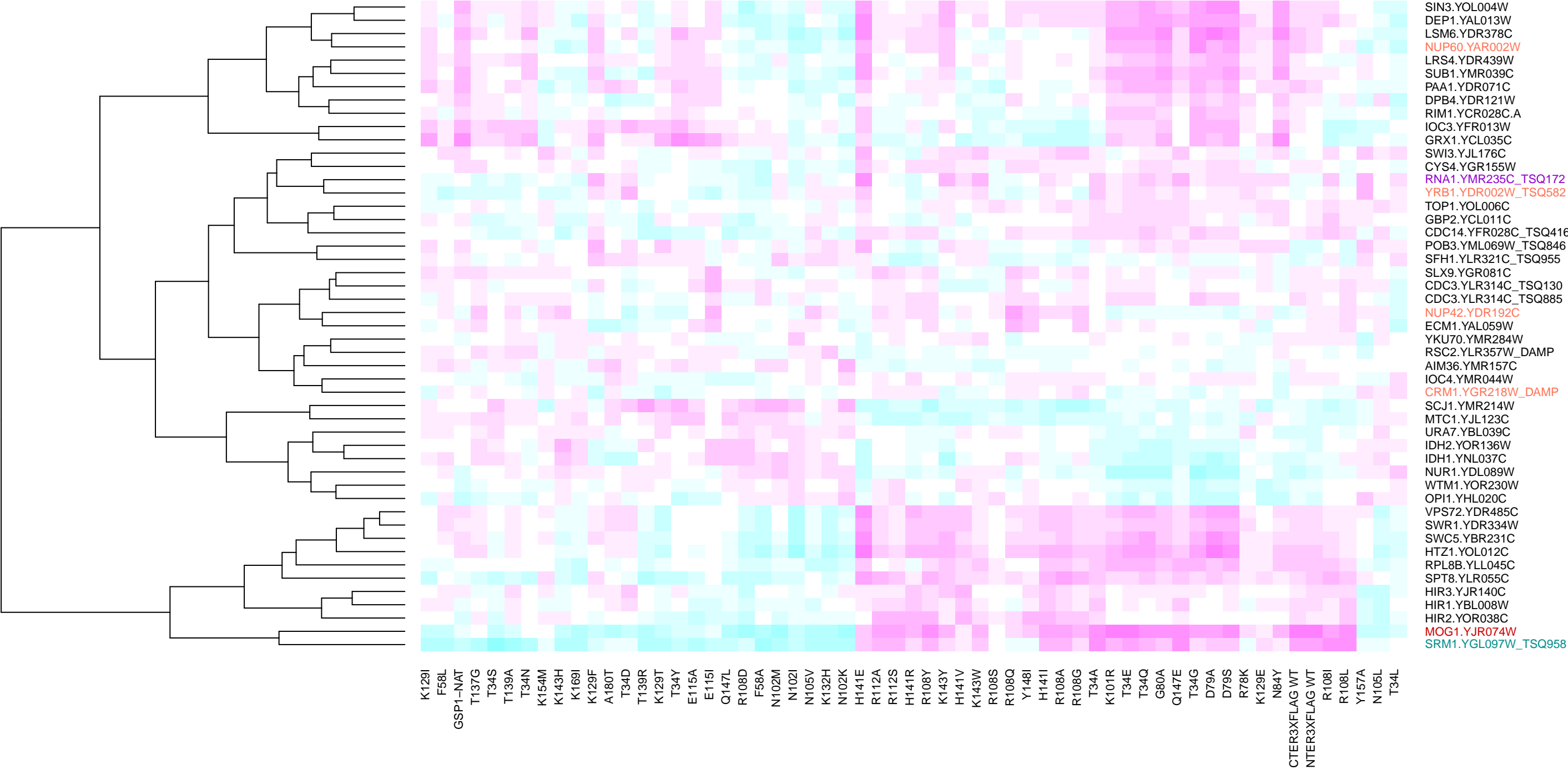
mitotic cell cycle



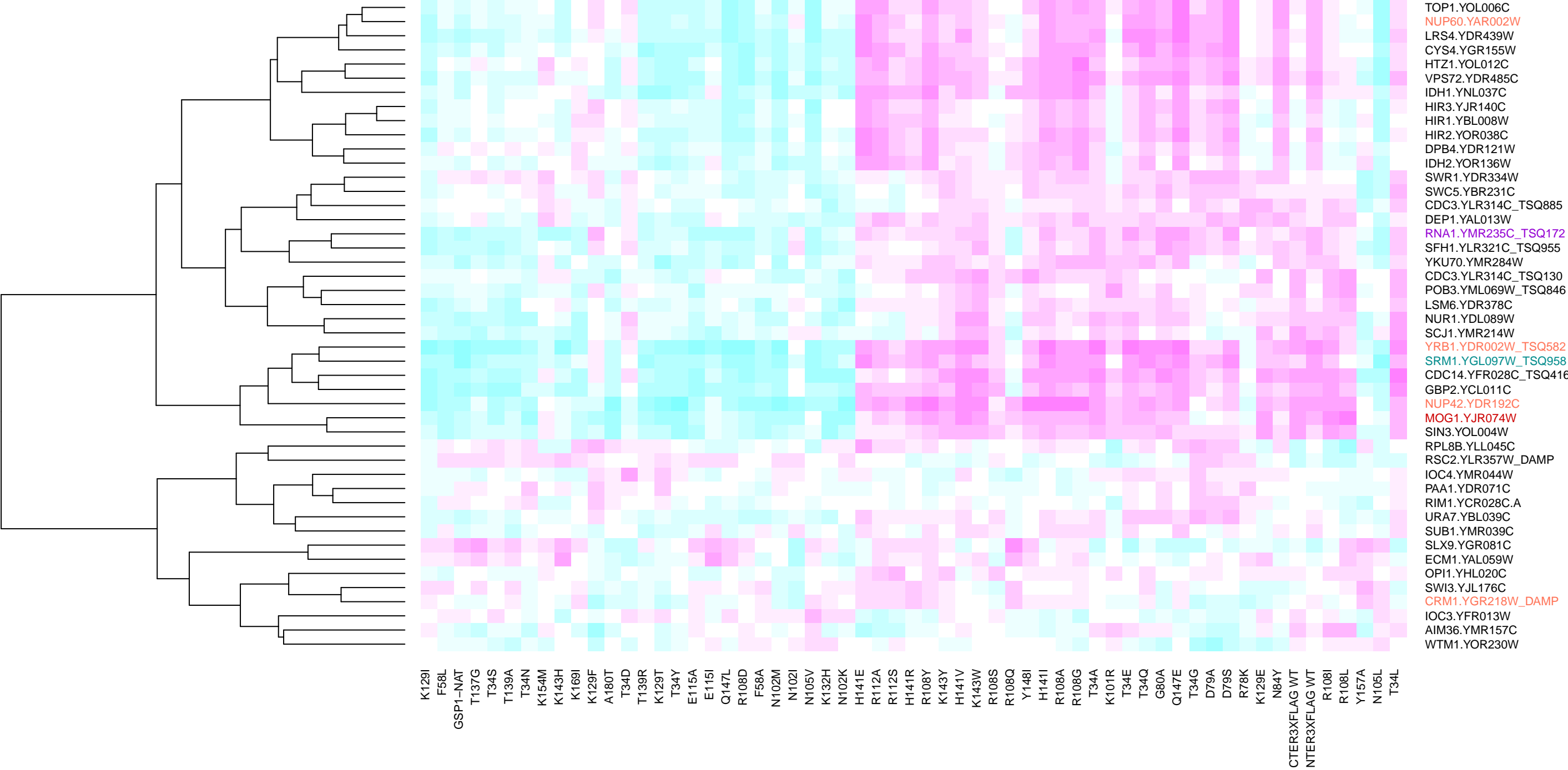
mitotic cell cycle_GO_15_1_all



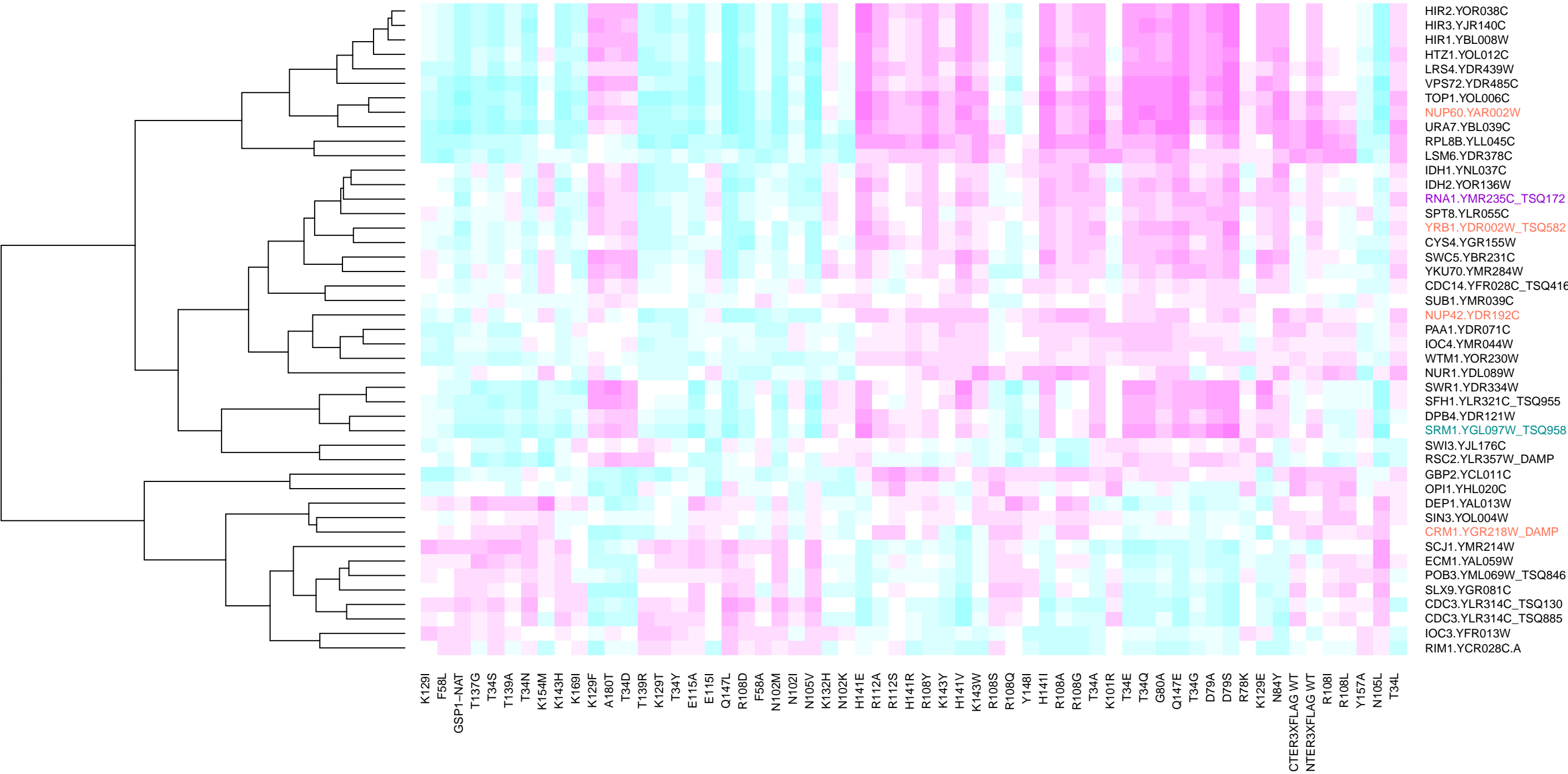
regulation of cell cycle



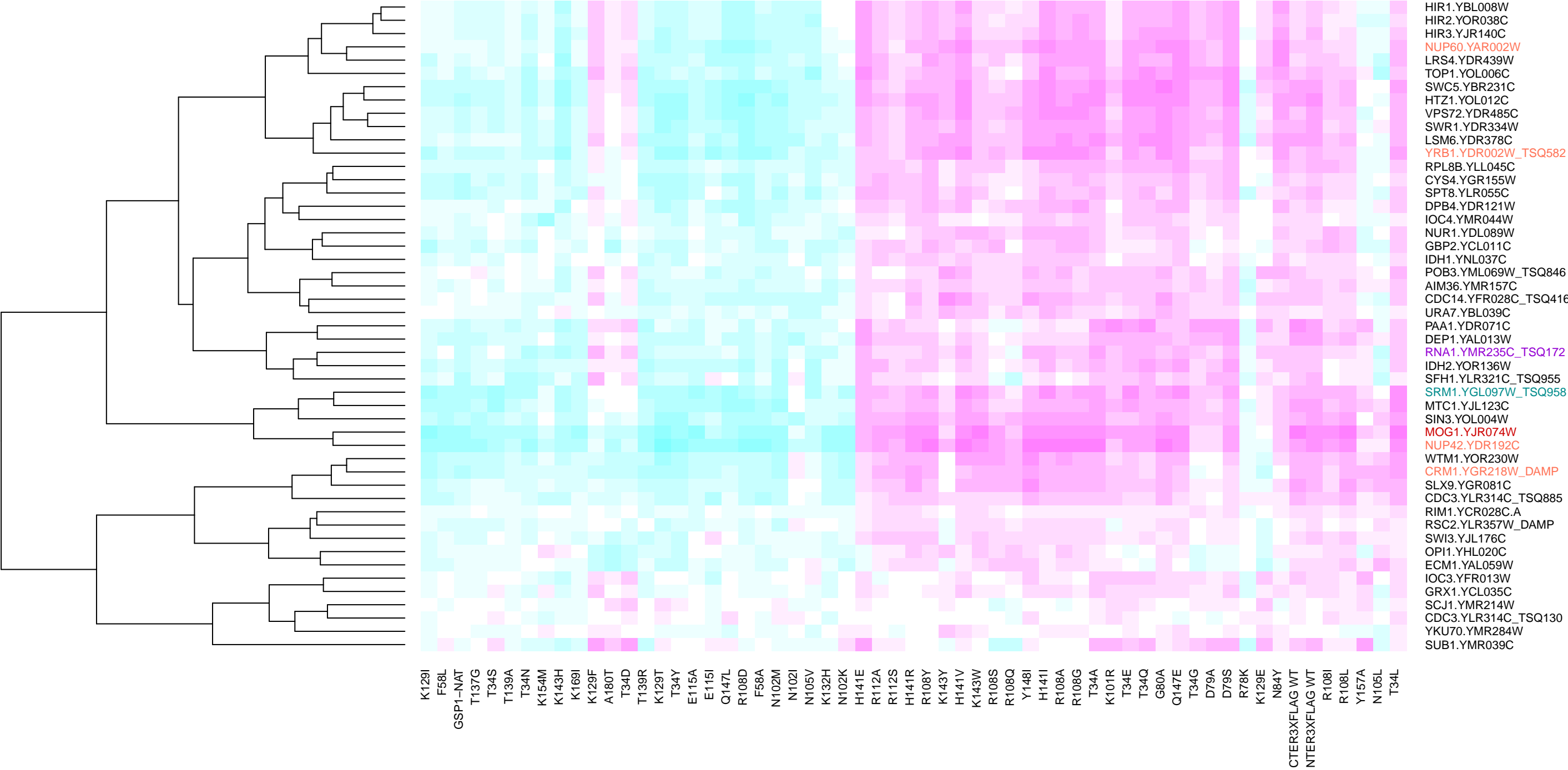
RNA catabolic process



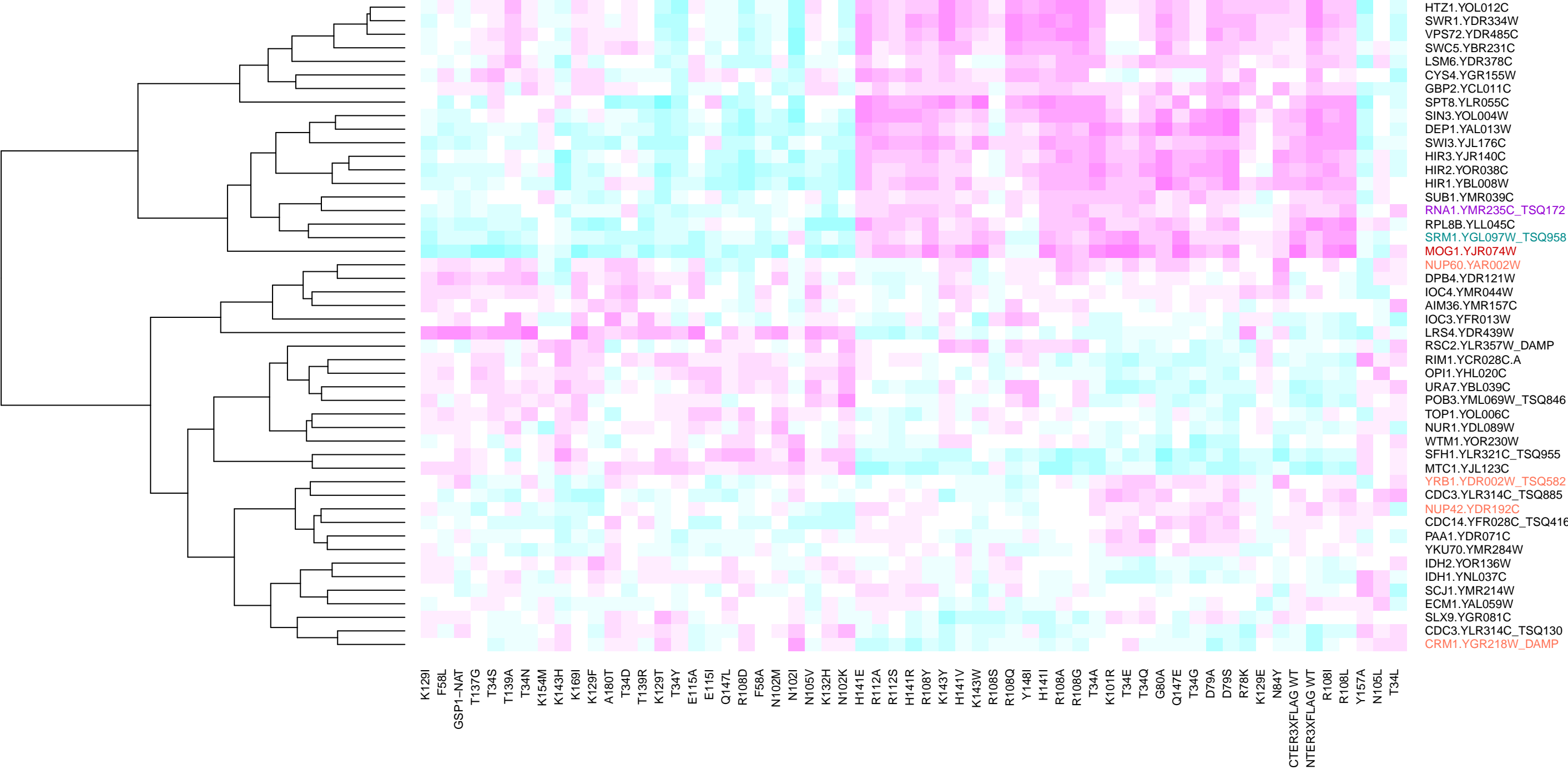
transcription and mRNA processing_GO_30_2_all



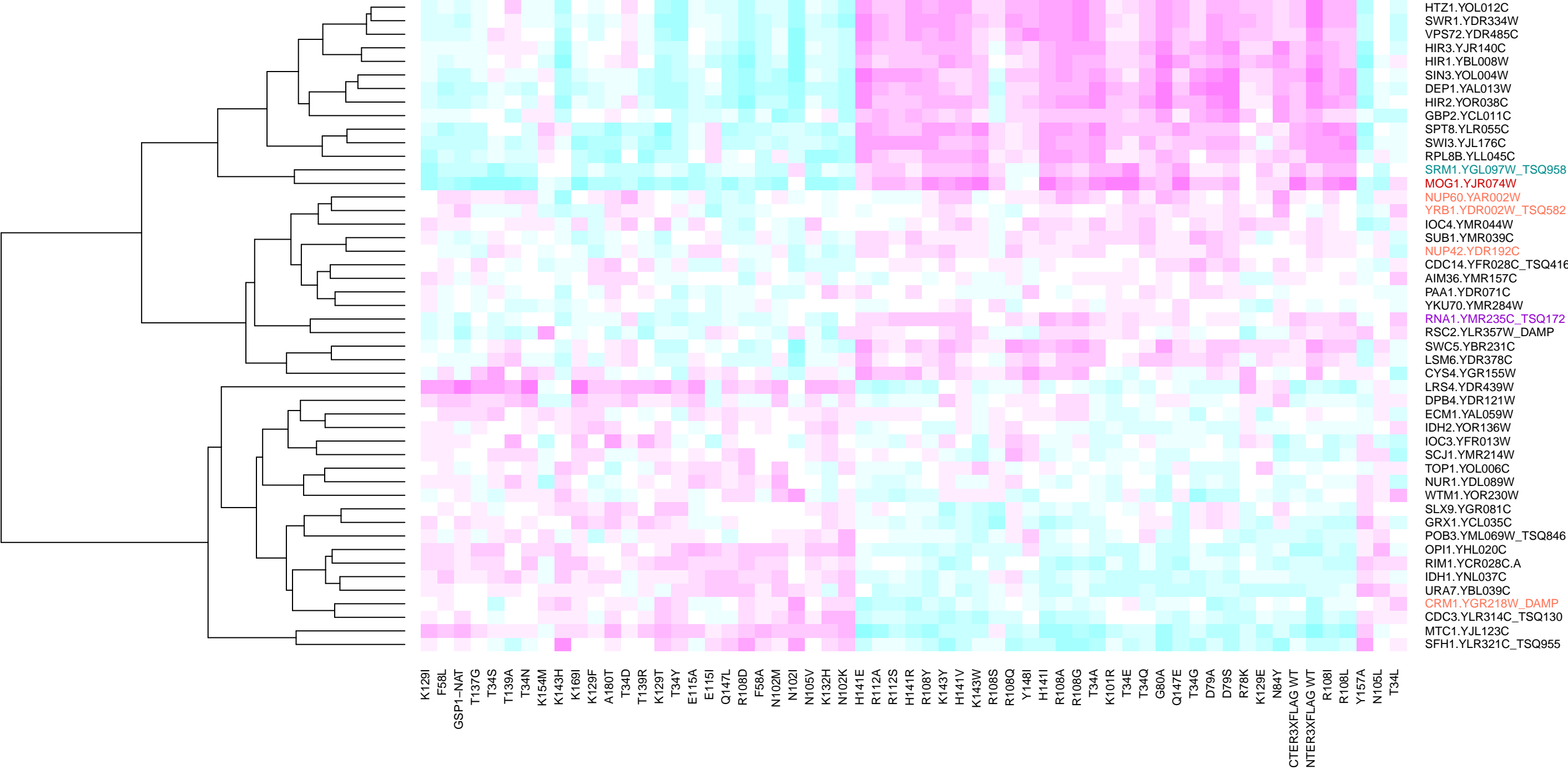
whole_library_30_3_all



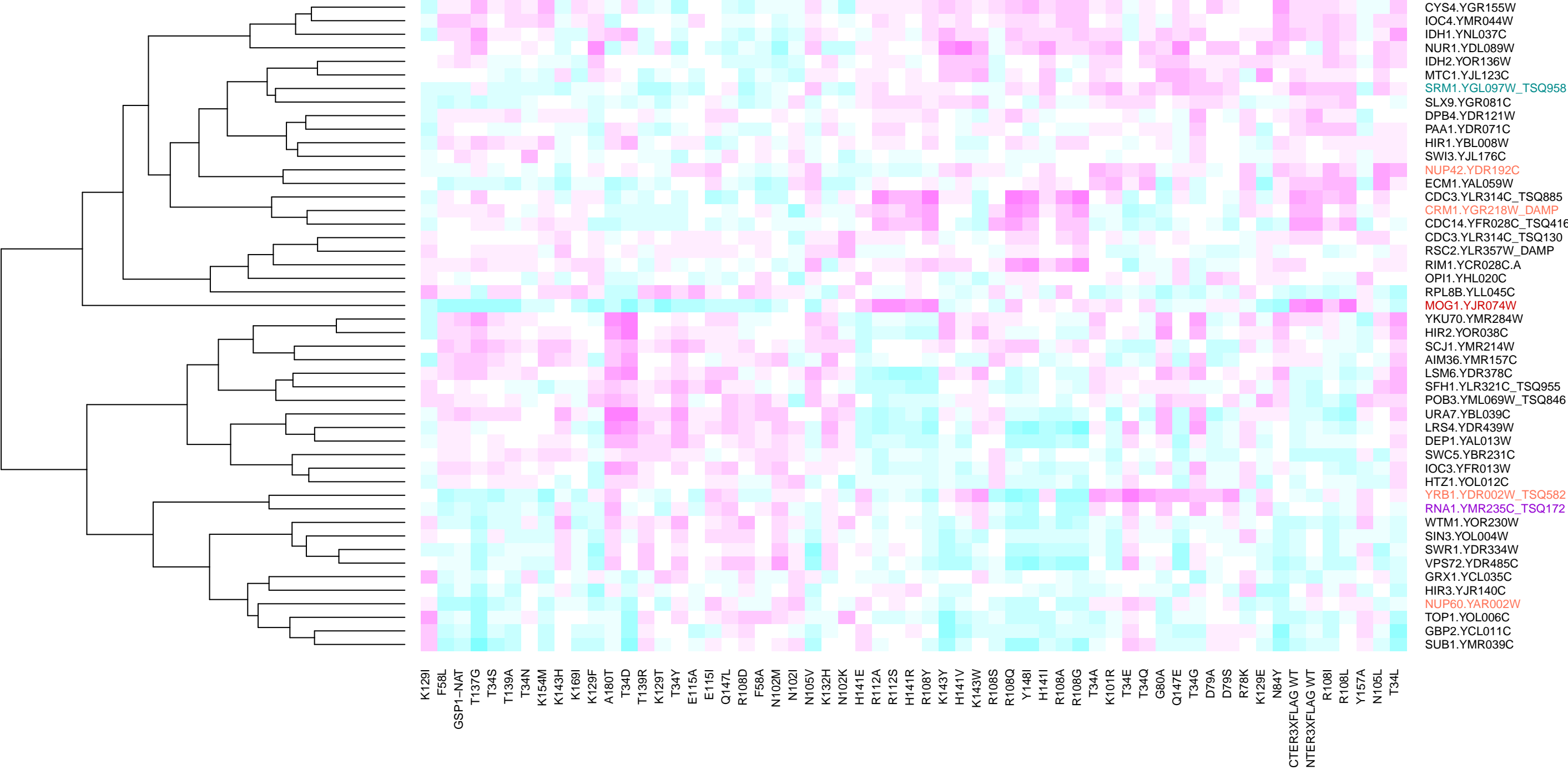
proteolysis involved in cellular protein catabolic process



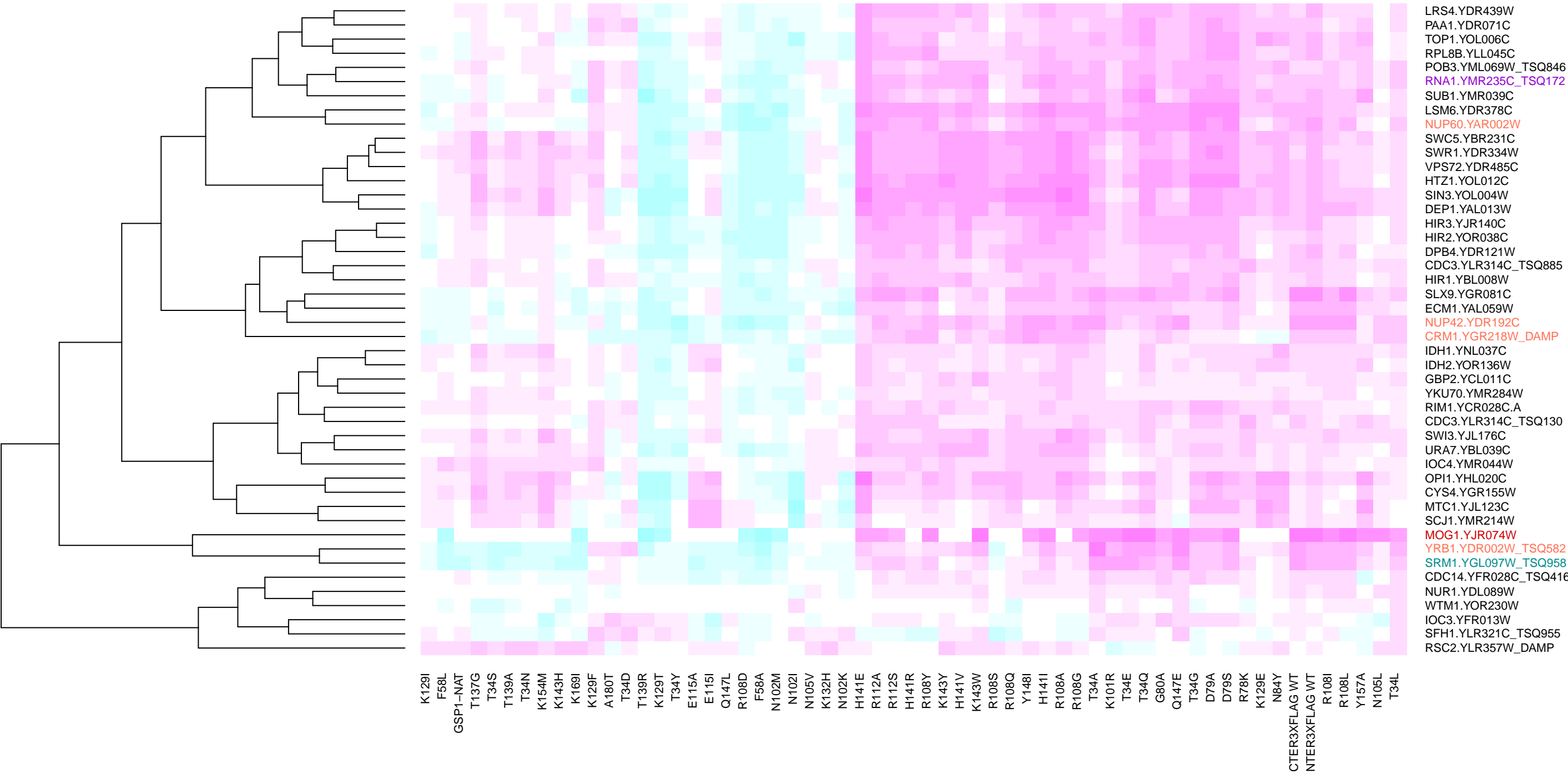
proteolysis involved in cellular protein catabolic process_GO_15_1_all



sig_Gsp1_Gl_30_4_all



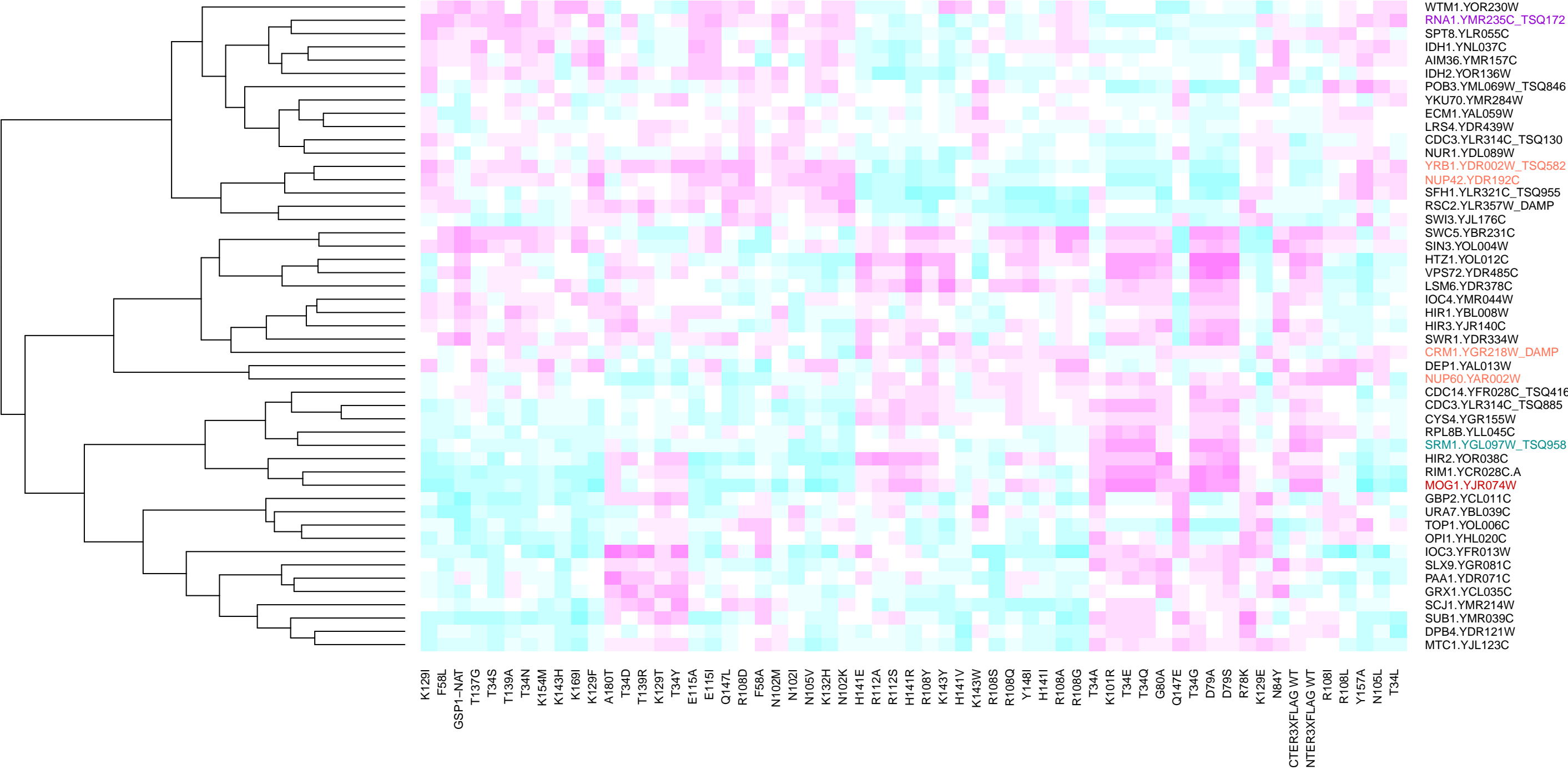
whole_library



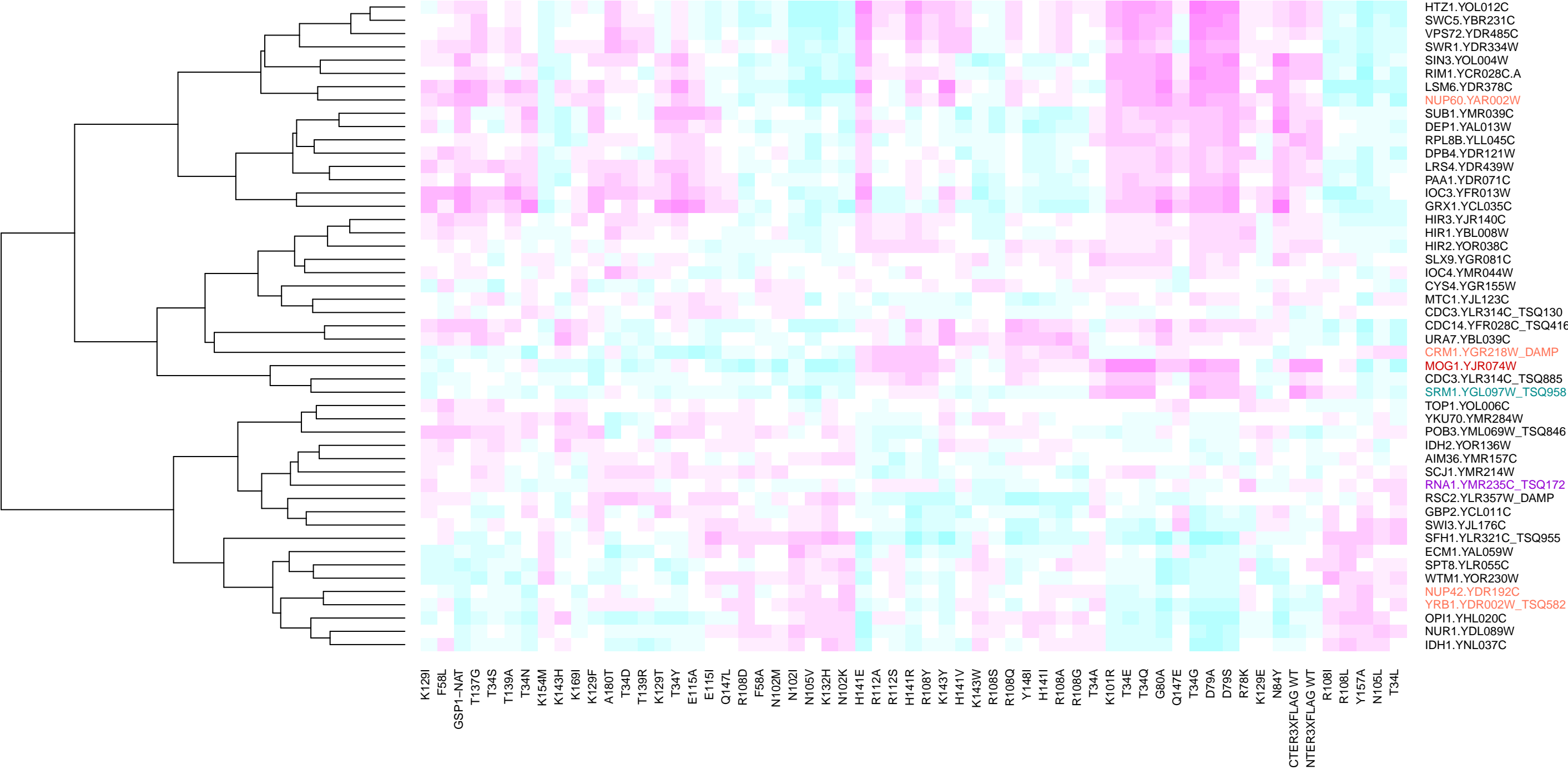
sig_Gsp1_Gl_30_5_mut



chromatin_GO_15_1_all



chromosome segregation



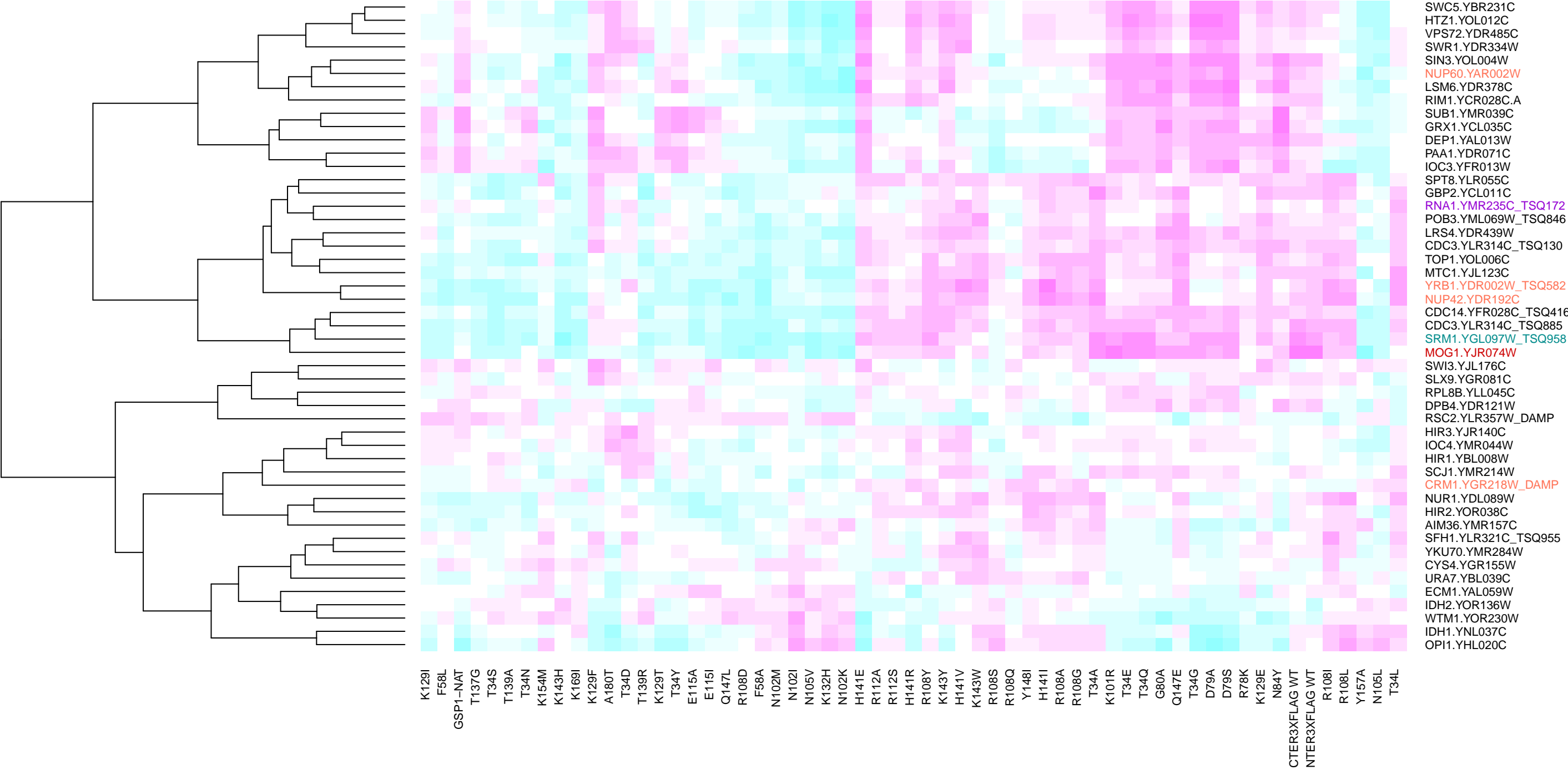
chromosome segregation_GO_15_1_mut



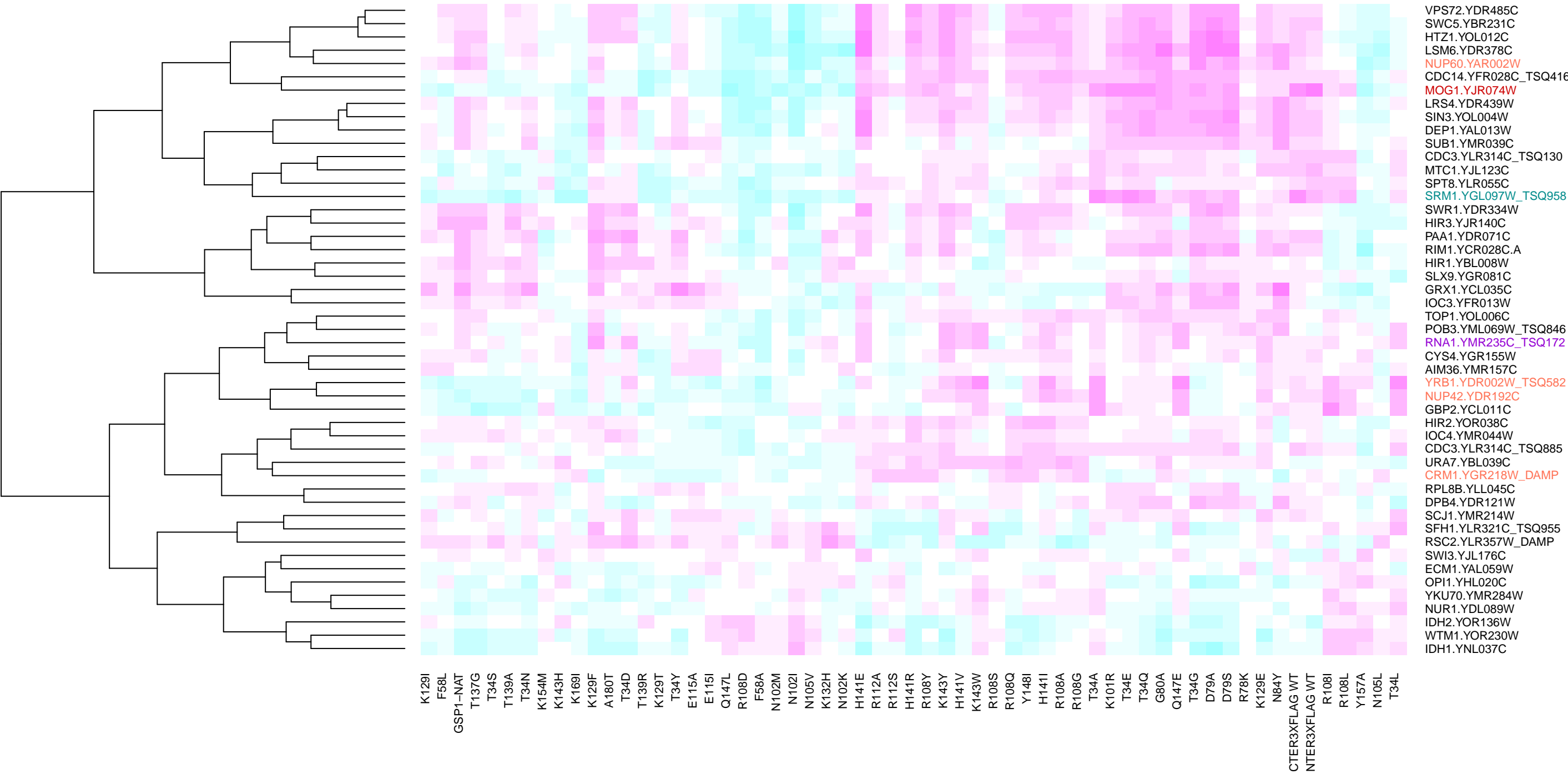
chromosome segregation_GO_30_1_all



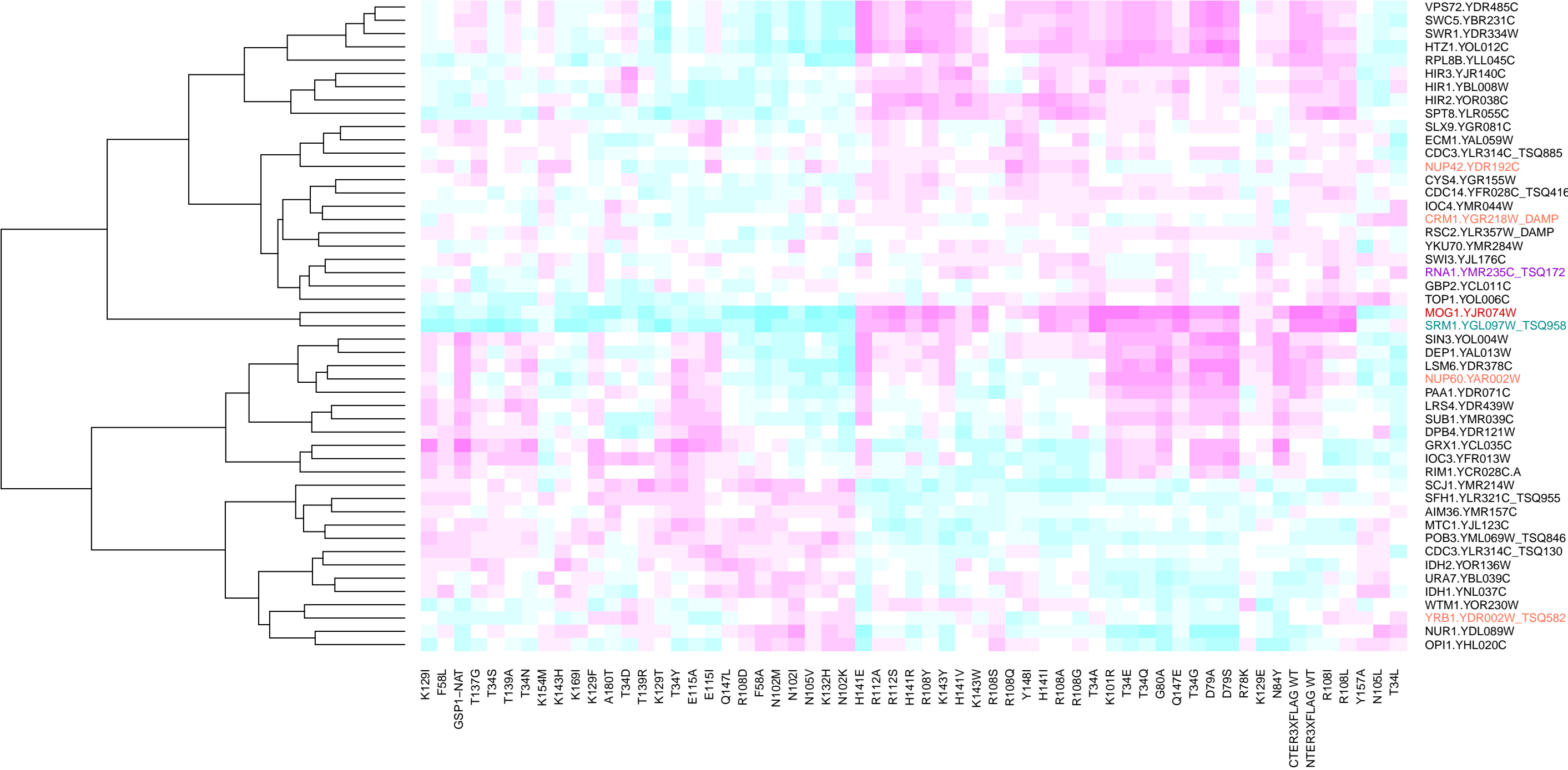
mitotic cell cycle_GO_30_1_mut



organelle fission



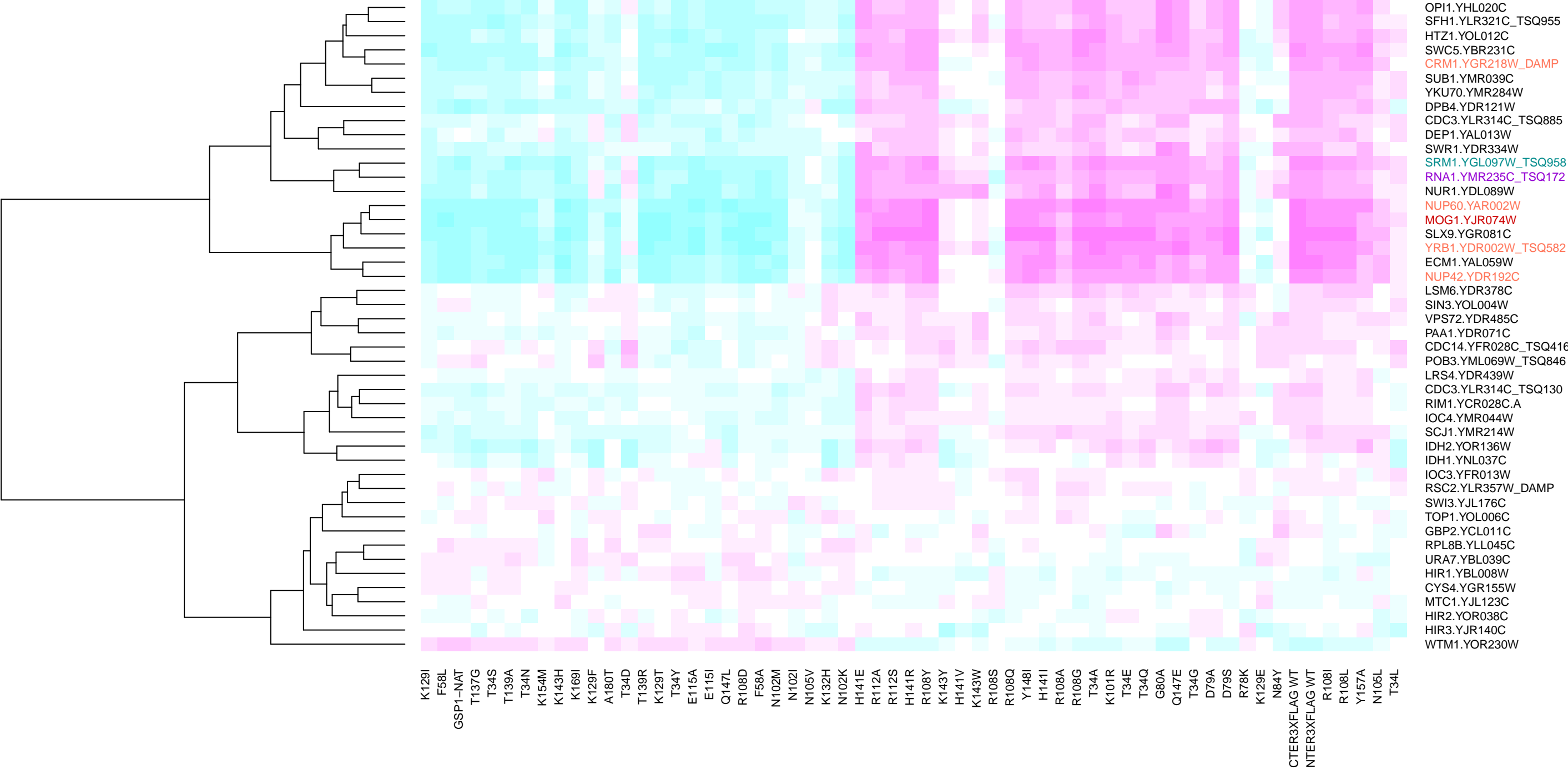
regulation of cell cycle_GO_30_1_all



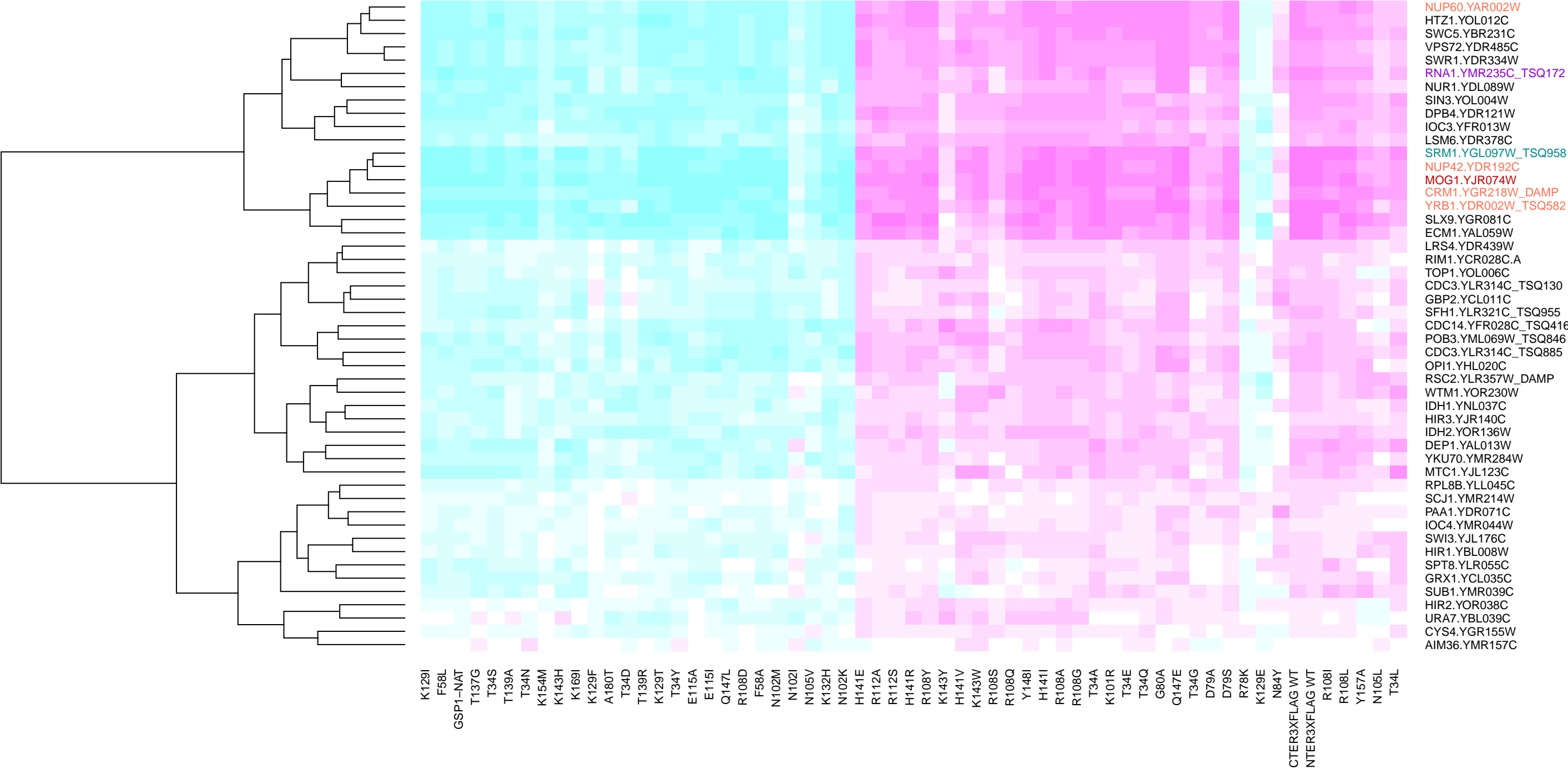
regulation of cell cycle_GO_30_1_mut



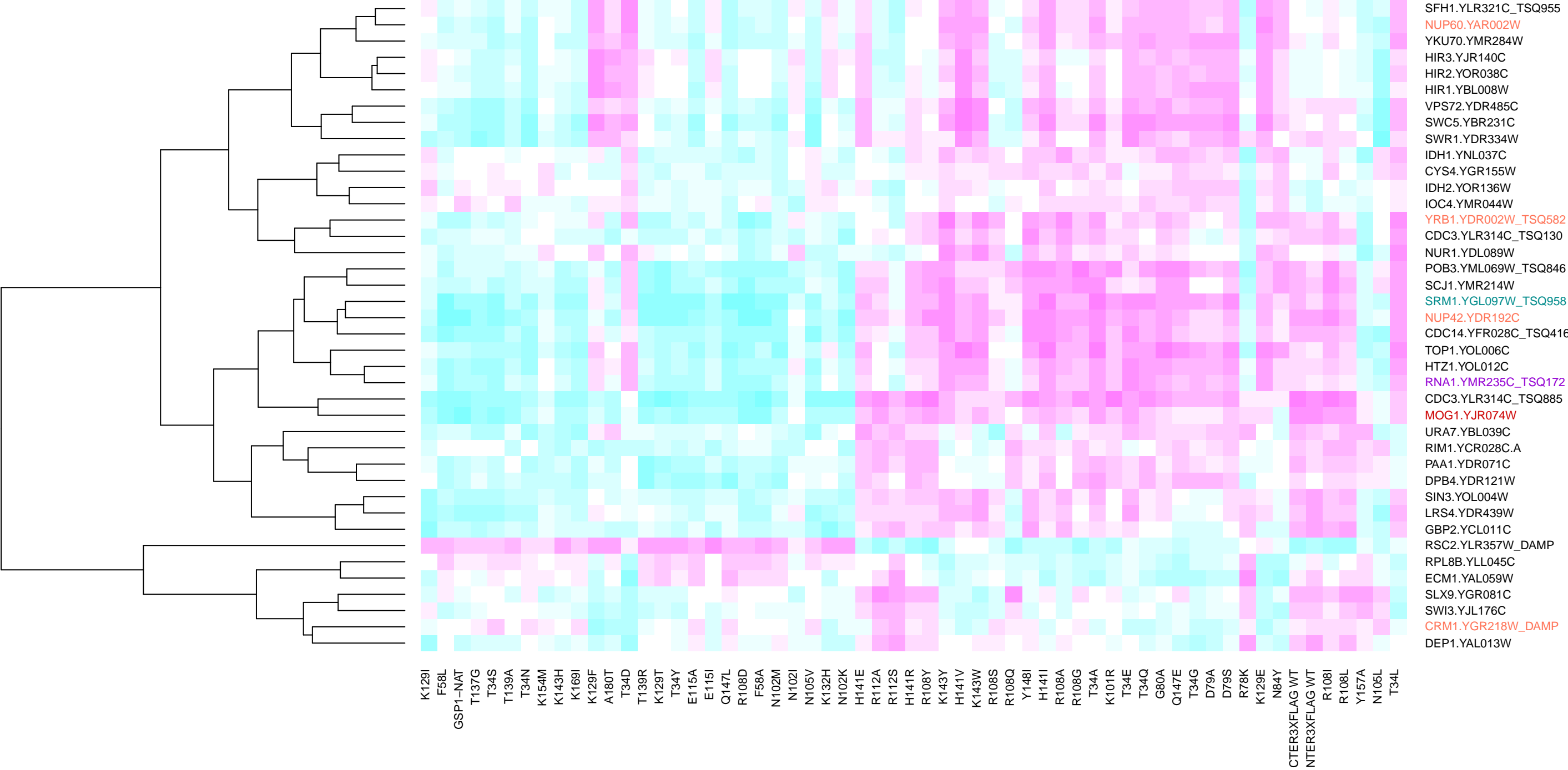
whole_library_30_5_all



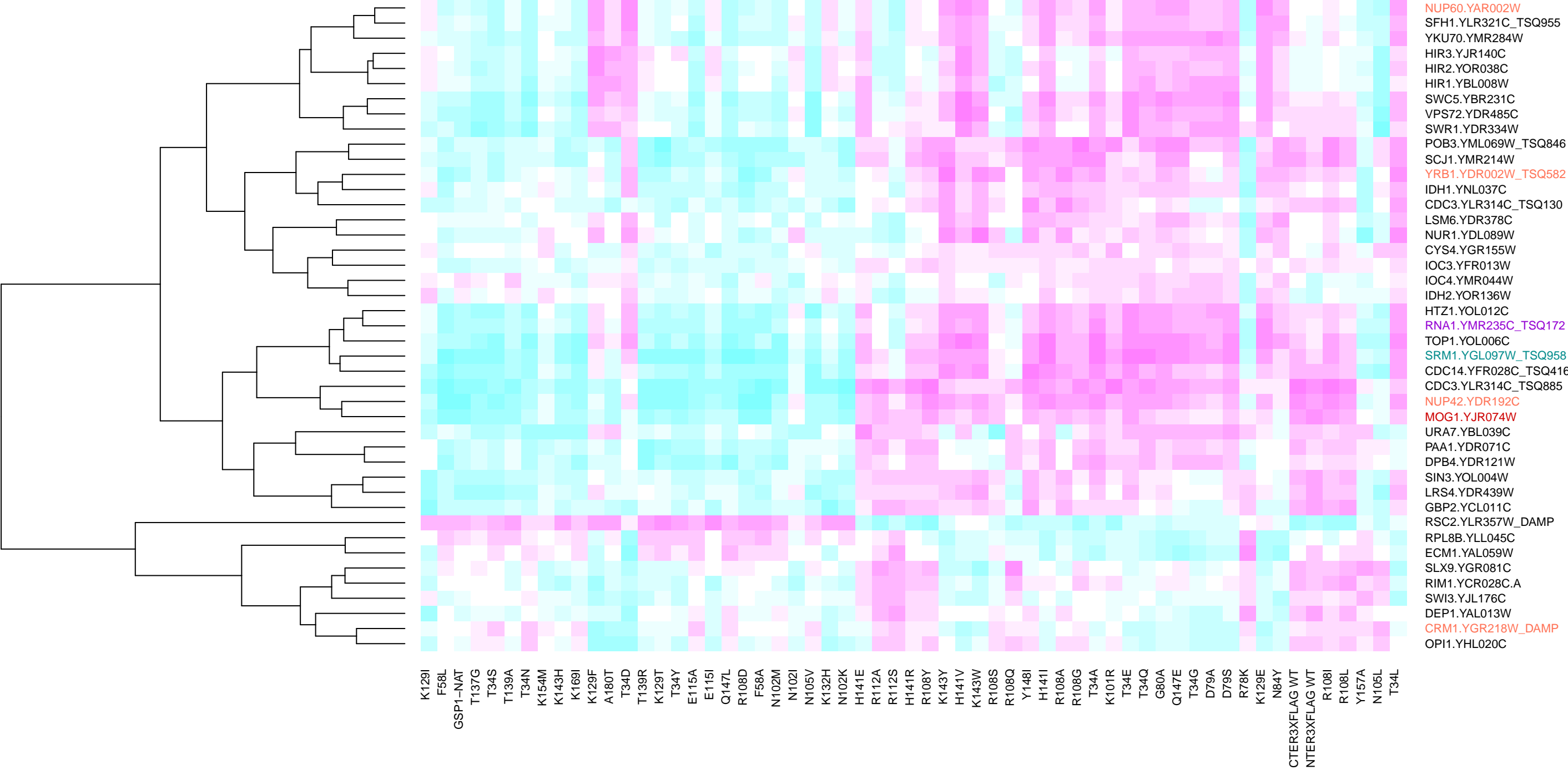
endomembrane system_GO_30_3_all



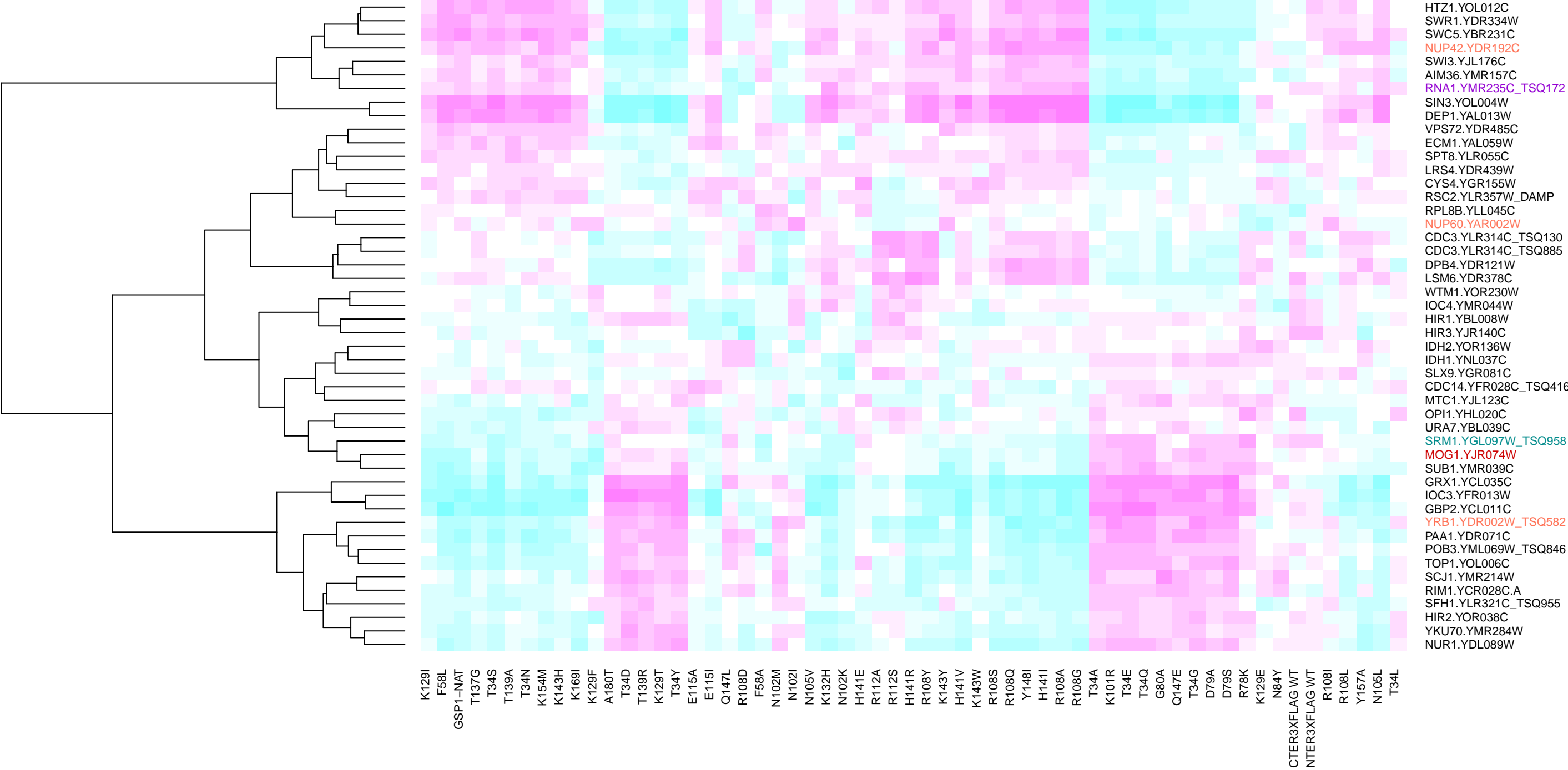
ribosomal small subunit biogenesis



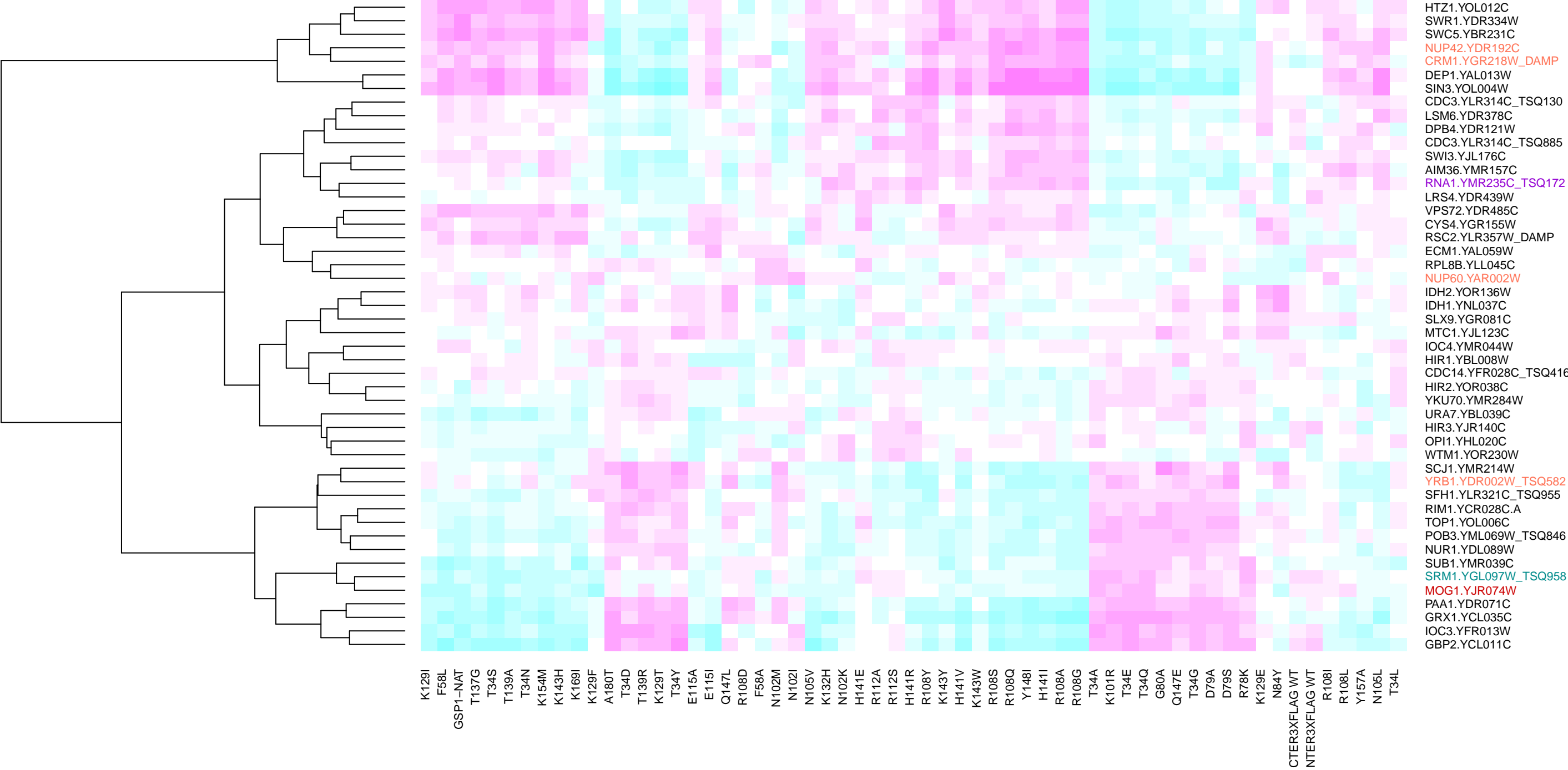
ribosomal small subunit biogenesis_GO_15_1_all



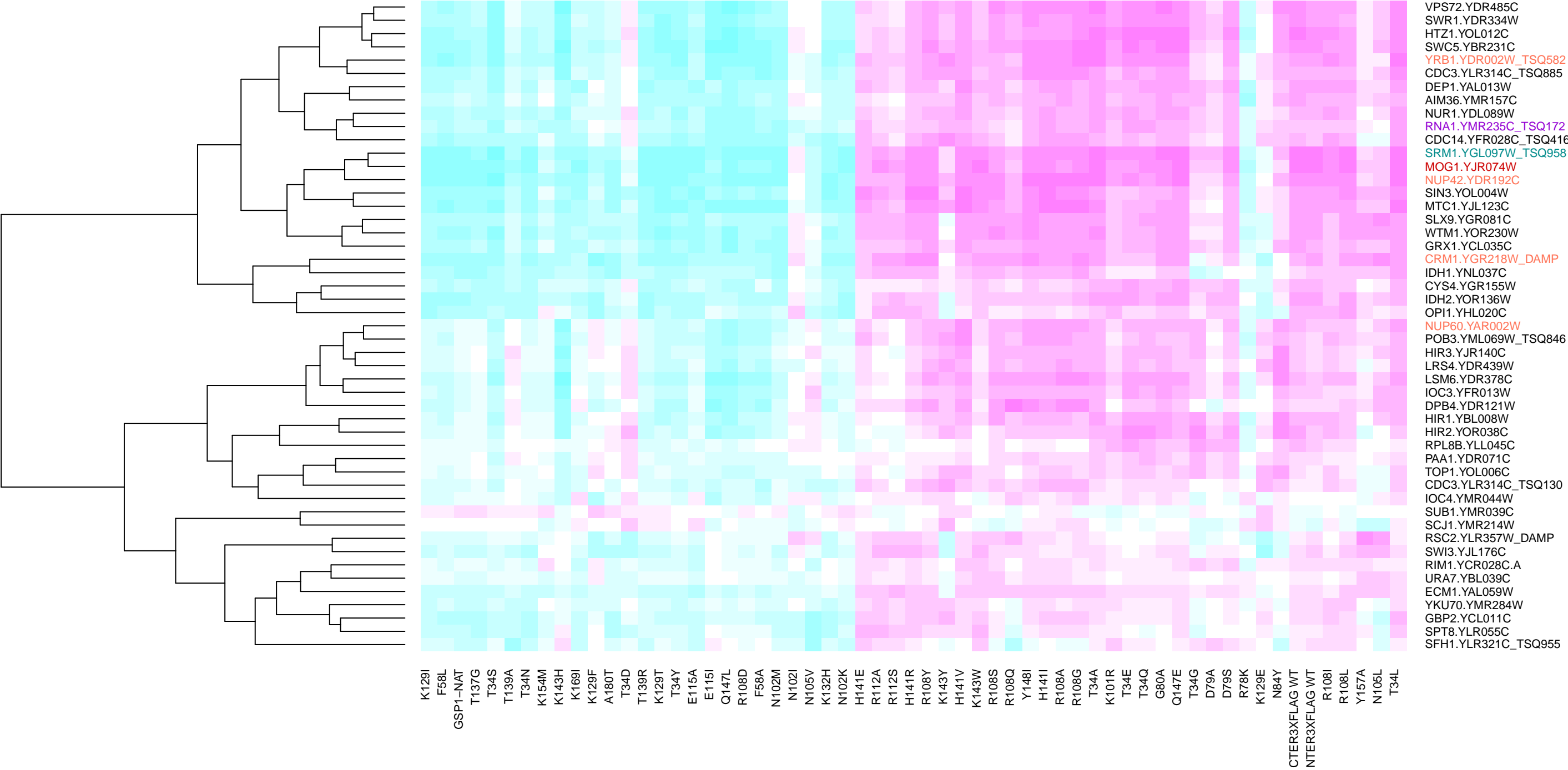
sig_Gsp1_Gl_15_1_mut



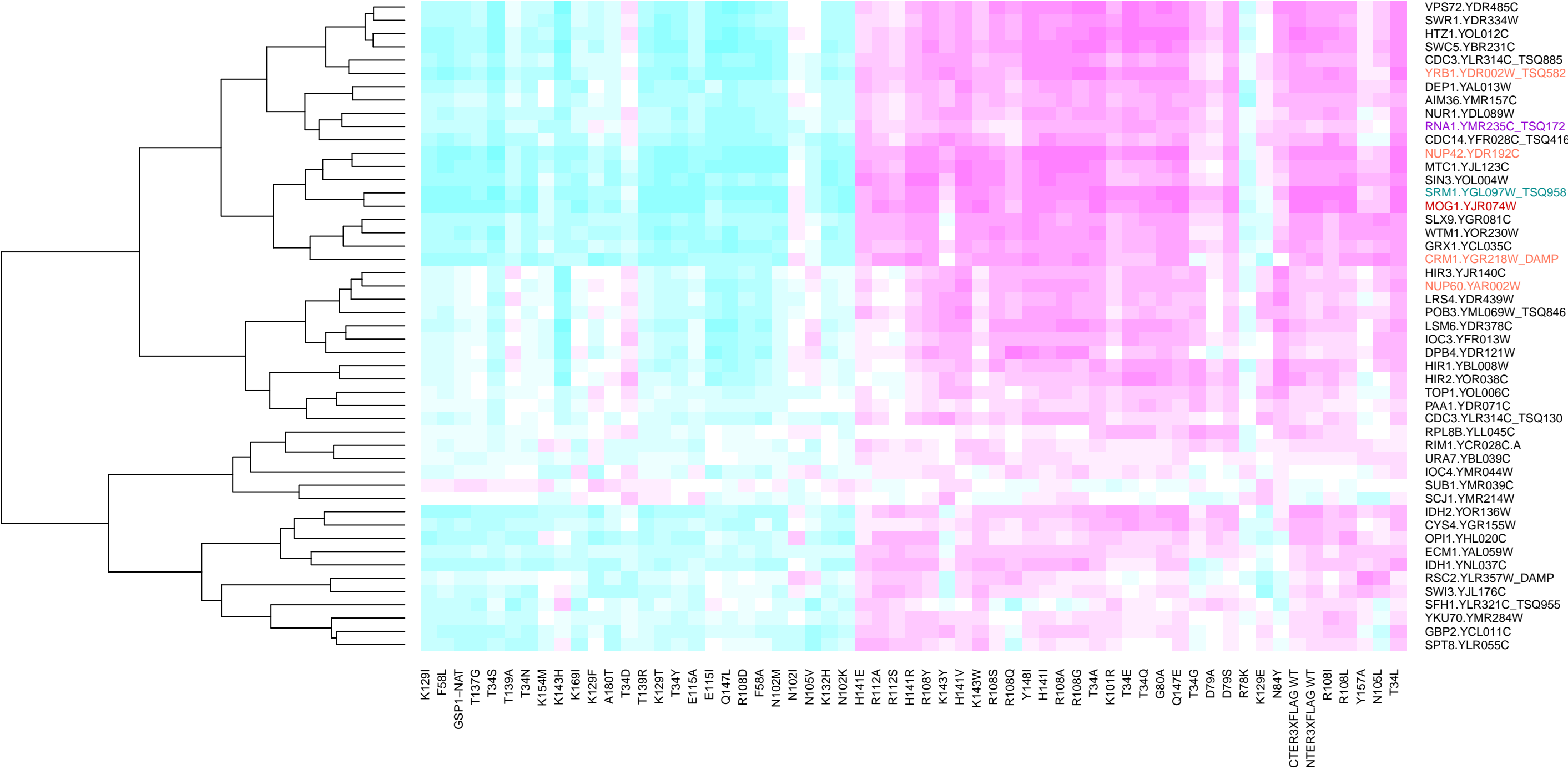
sig_Gsp1_Gl_30_3_mut



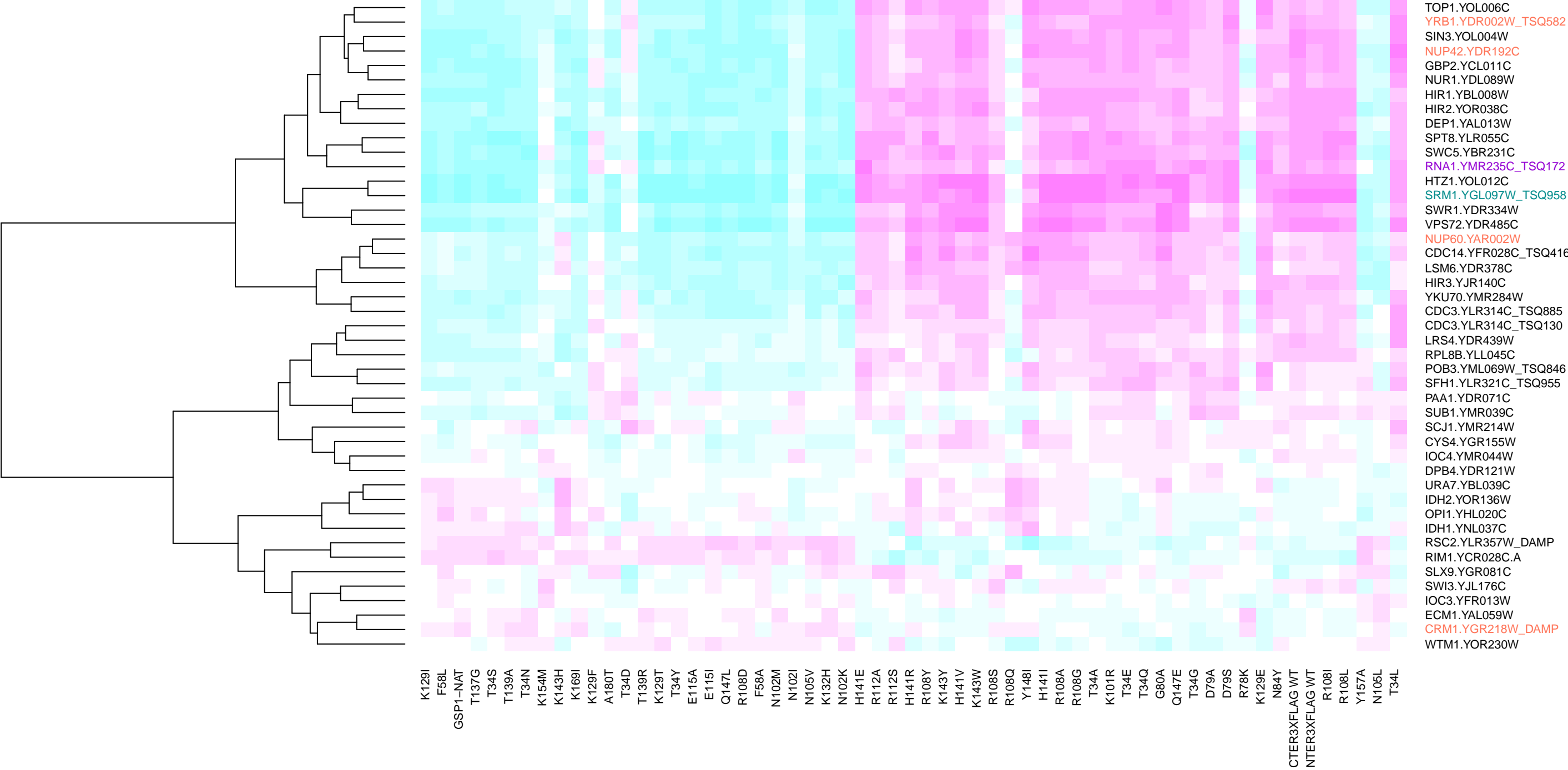
cellular response to DNA damage stimulus_GO_15_1_all



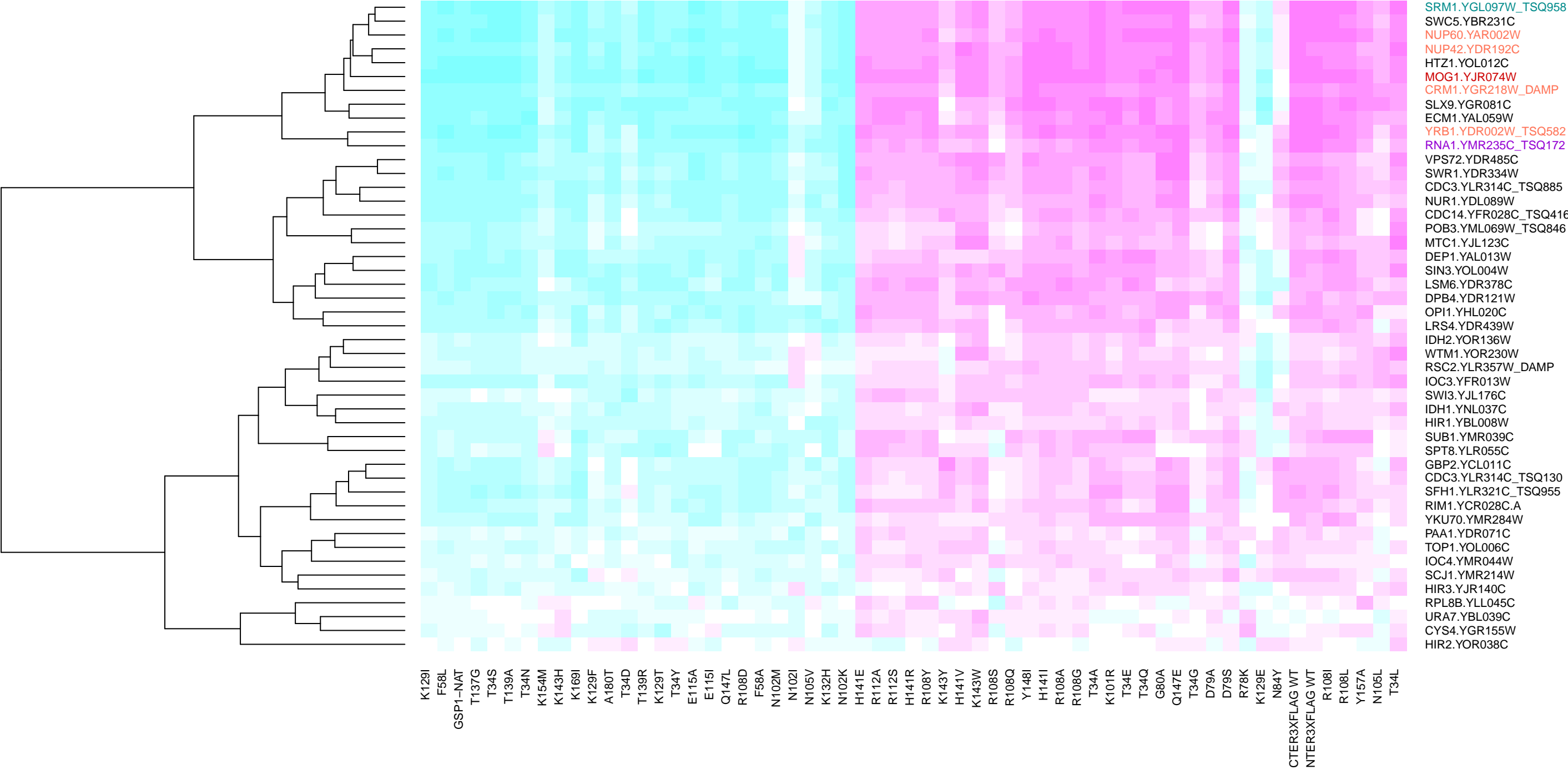
cellular response to DNA damage stimulus_GO_30_3_all



cell cycle_GO_15_3_all



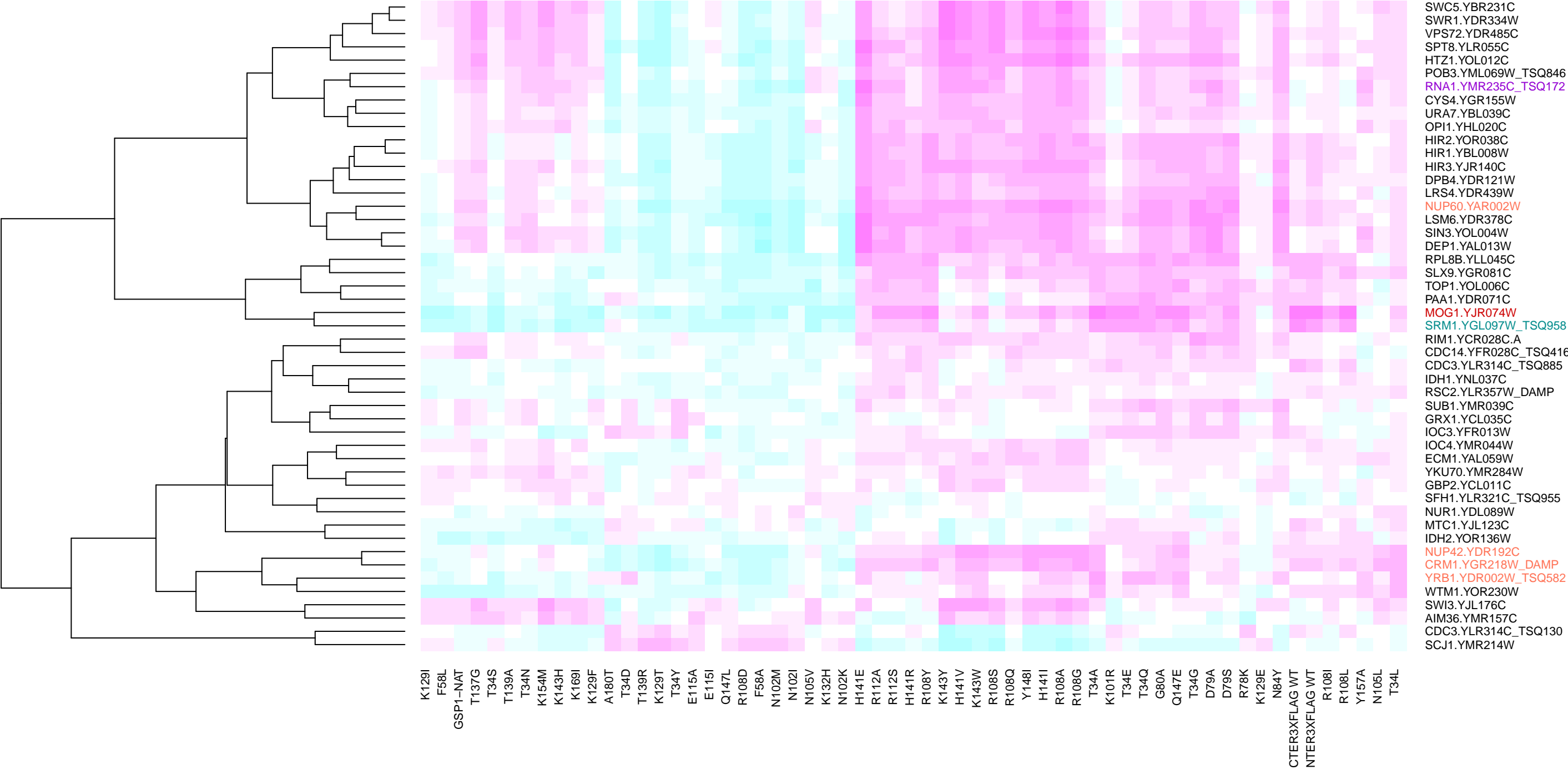
endomembrane system_GO_15_3_mut



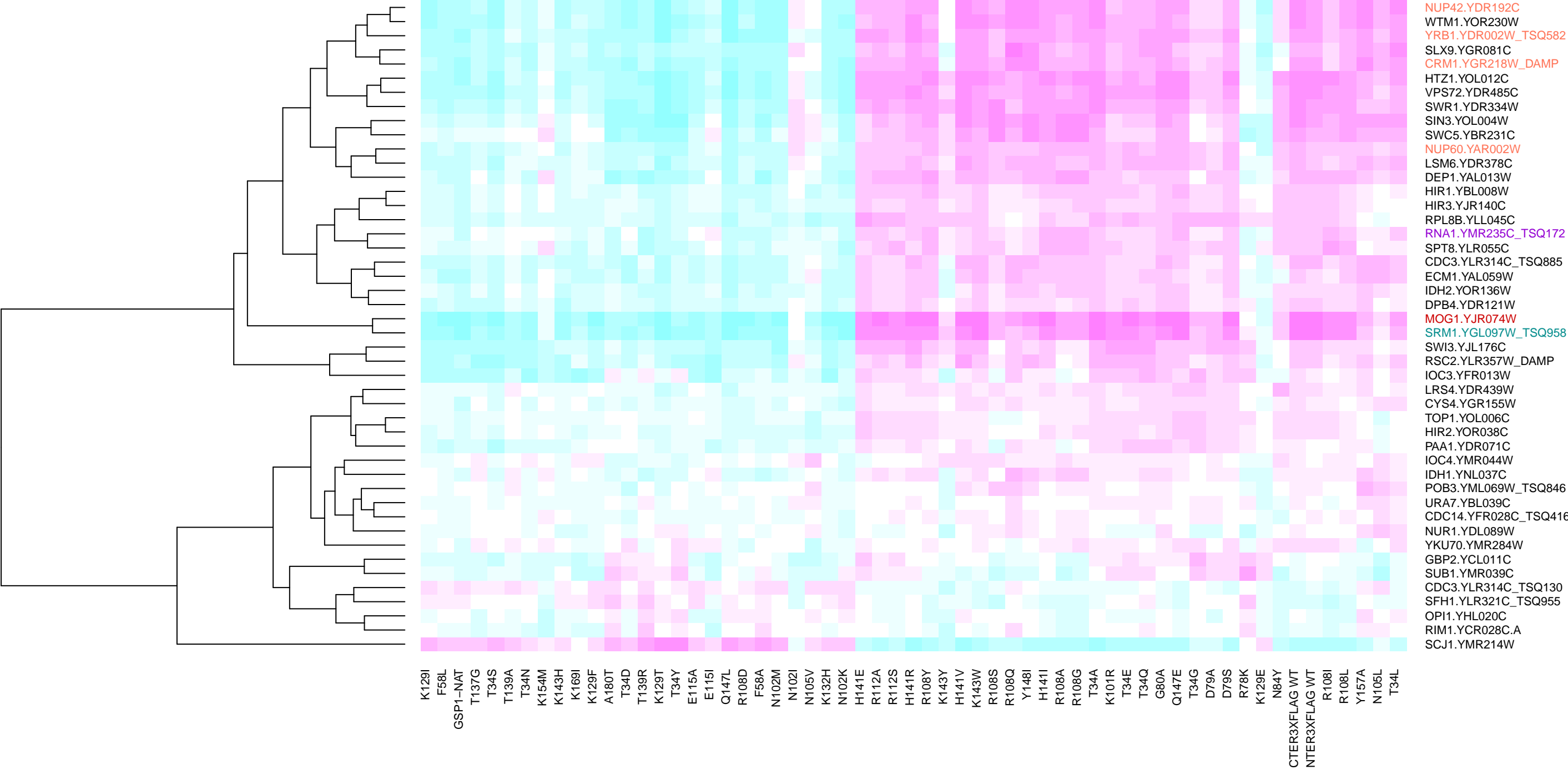
regulation of organelle organization_GO_15_1_all



chromatin



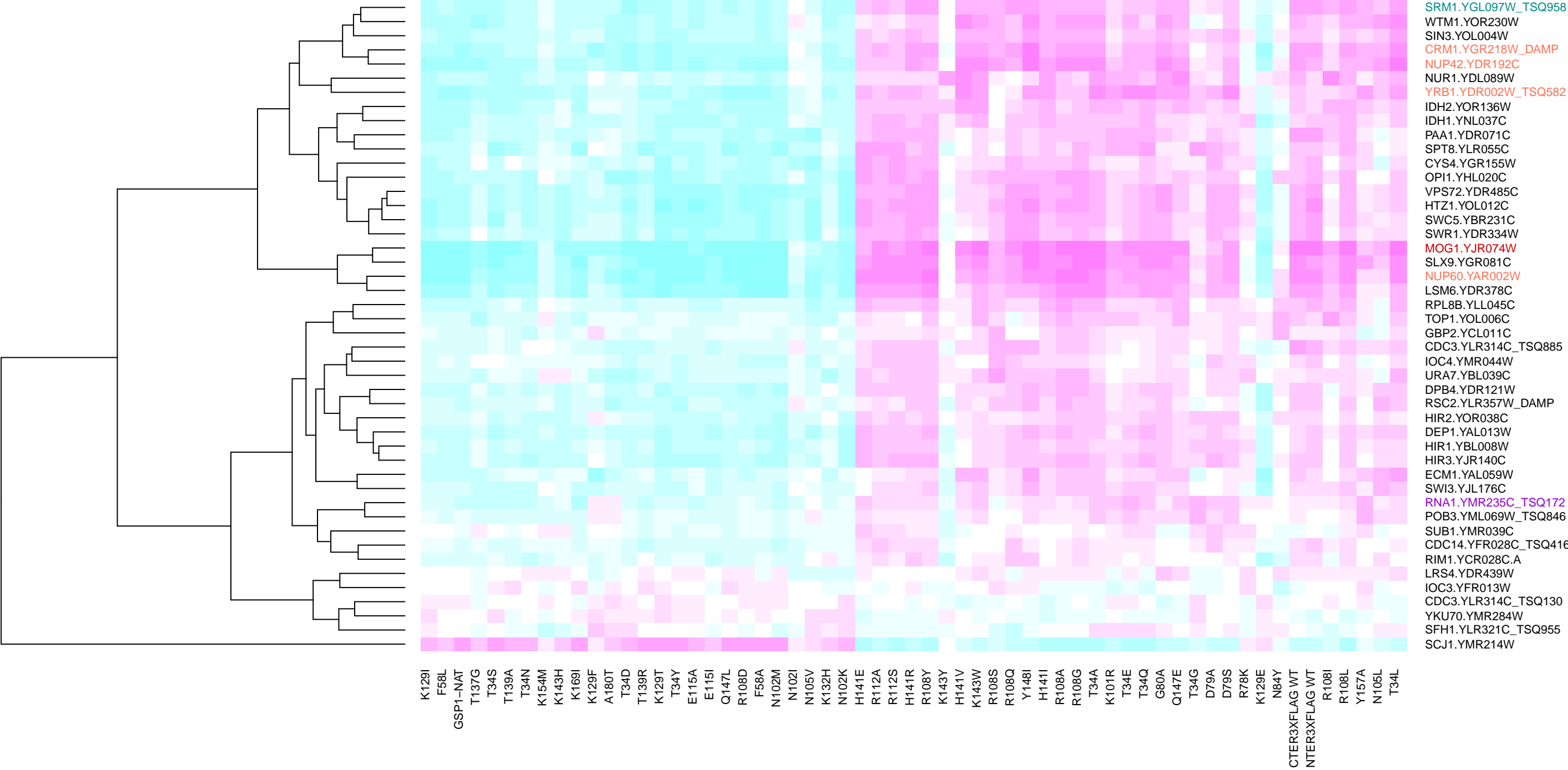
chromatin organization_GO_30_2_all



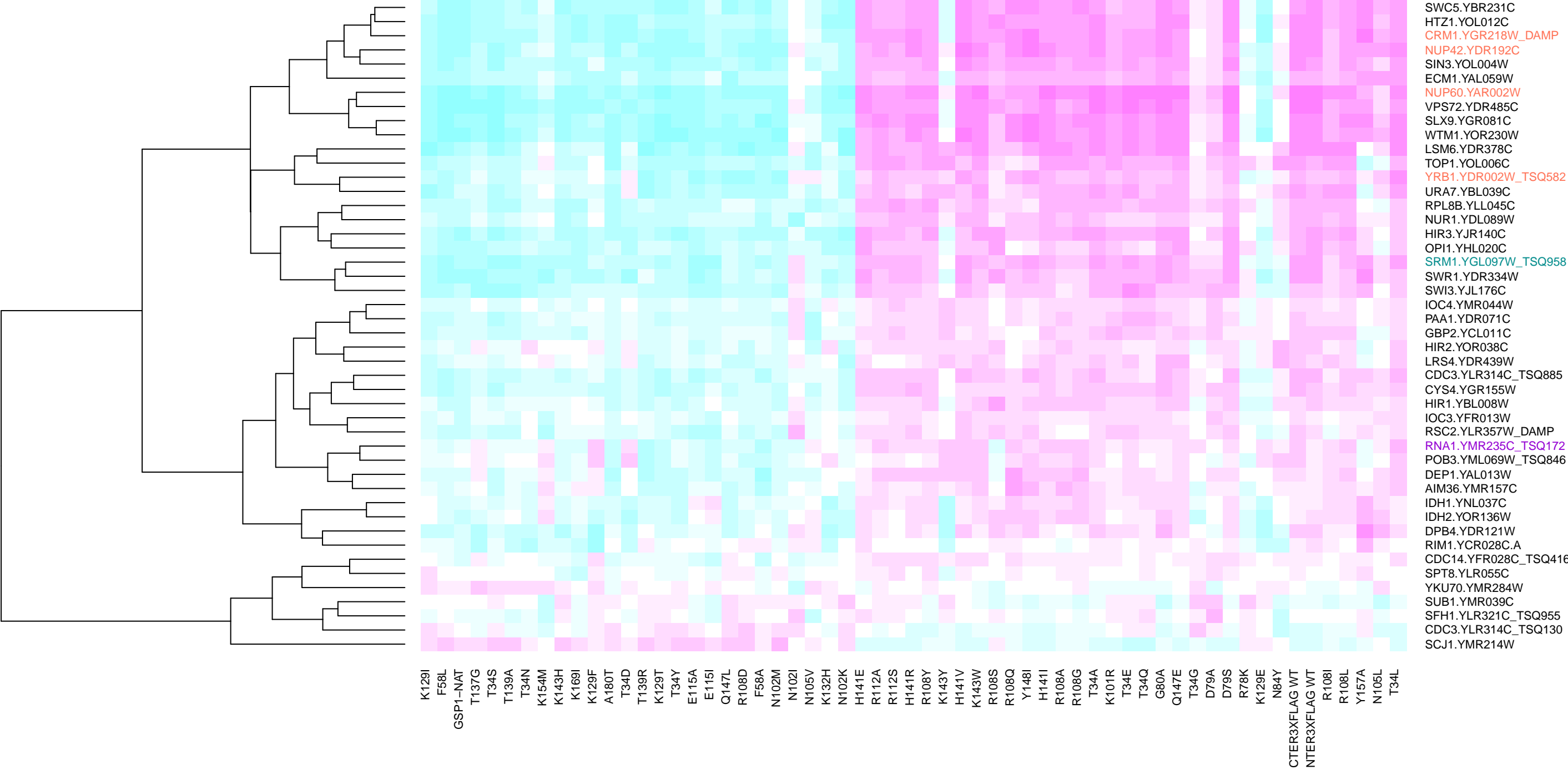
DNA binding_GO_15_3_mut



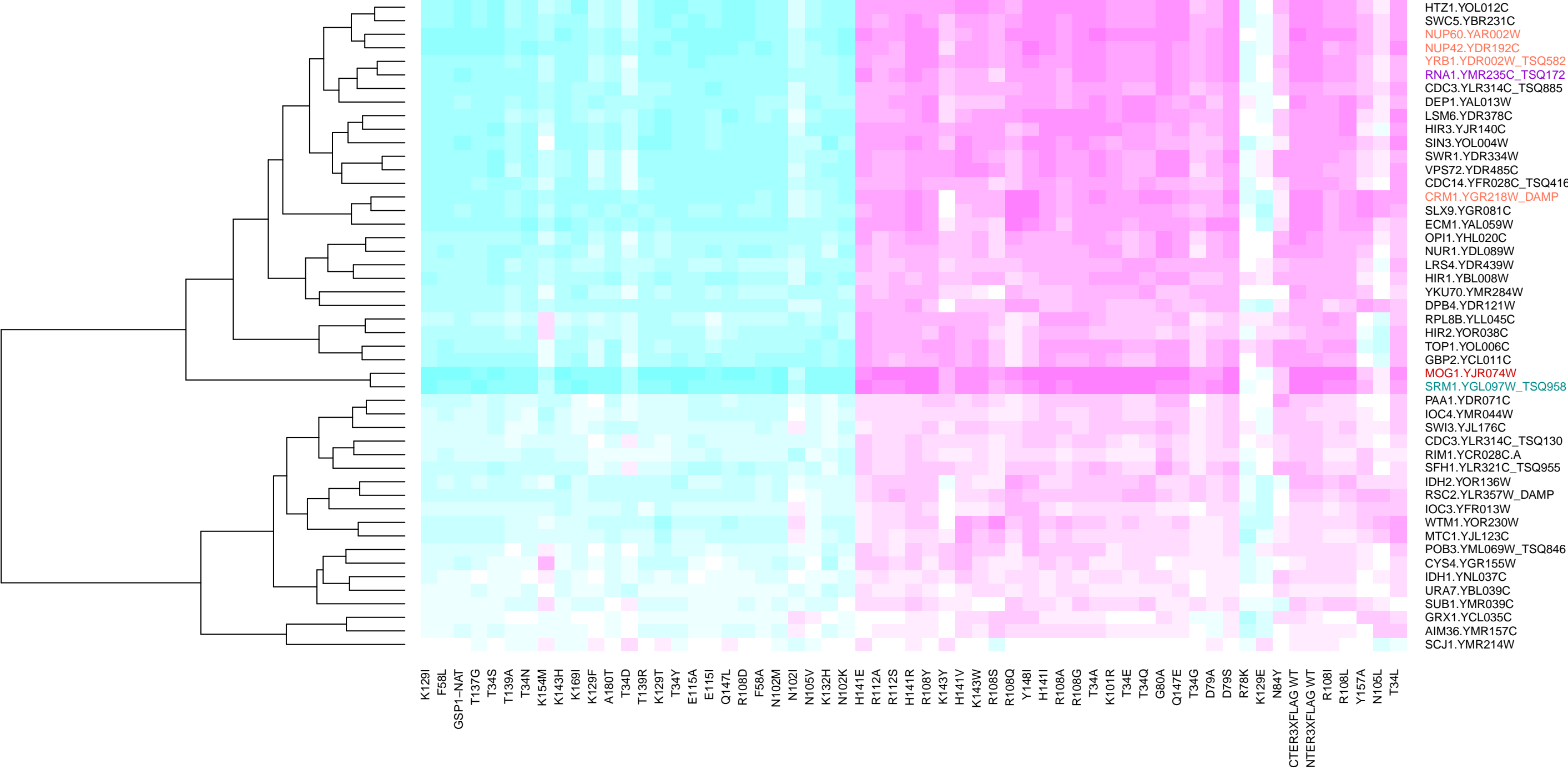
transcription_GO_15_1_mut



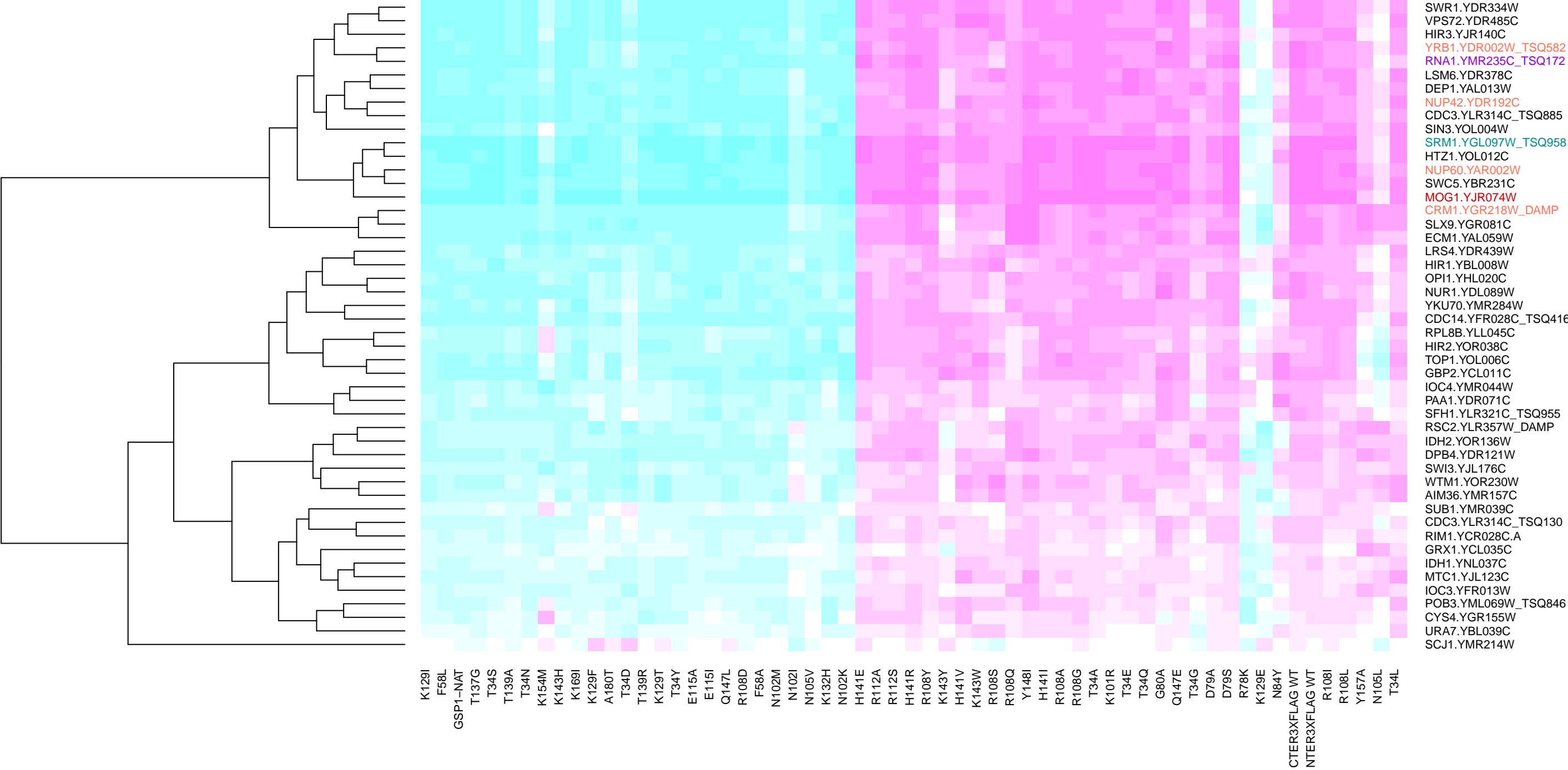
transcription_GO_30_4_all



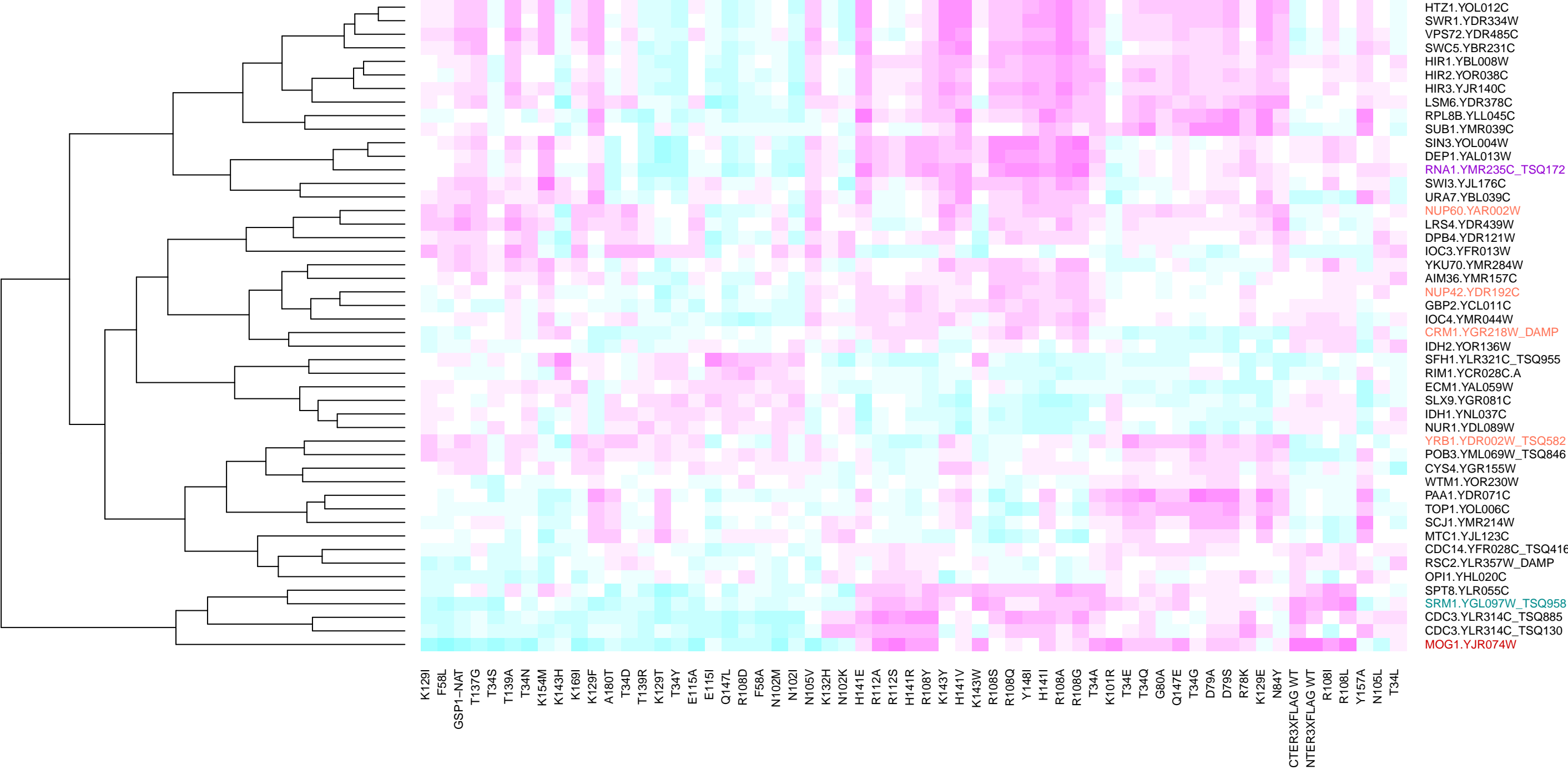
nucleobase-containing compound transport



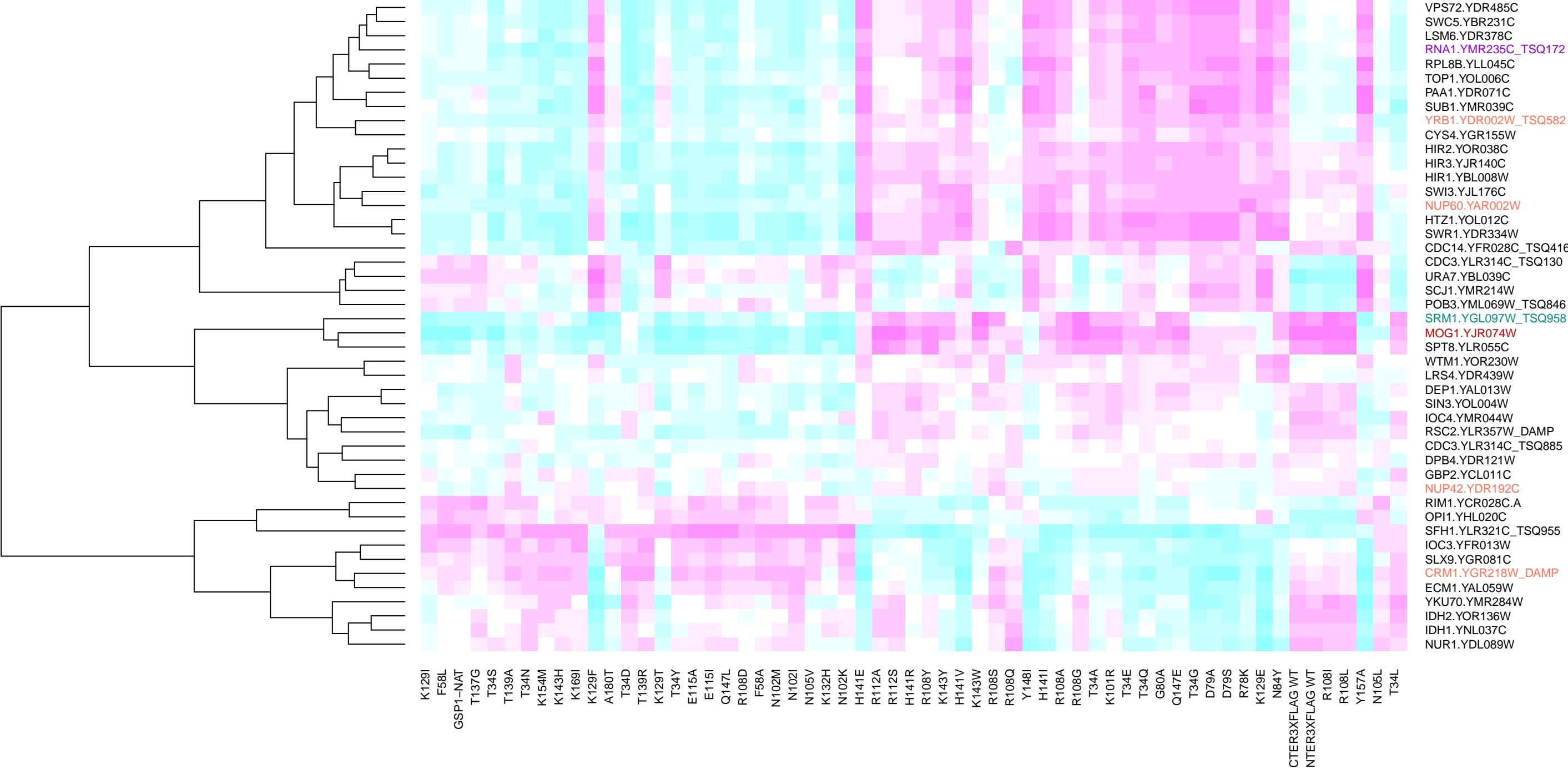
nucleobase-containing compound transport_GO_15_1_mut



protein modification by small protein conjugation or removal



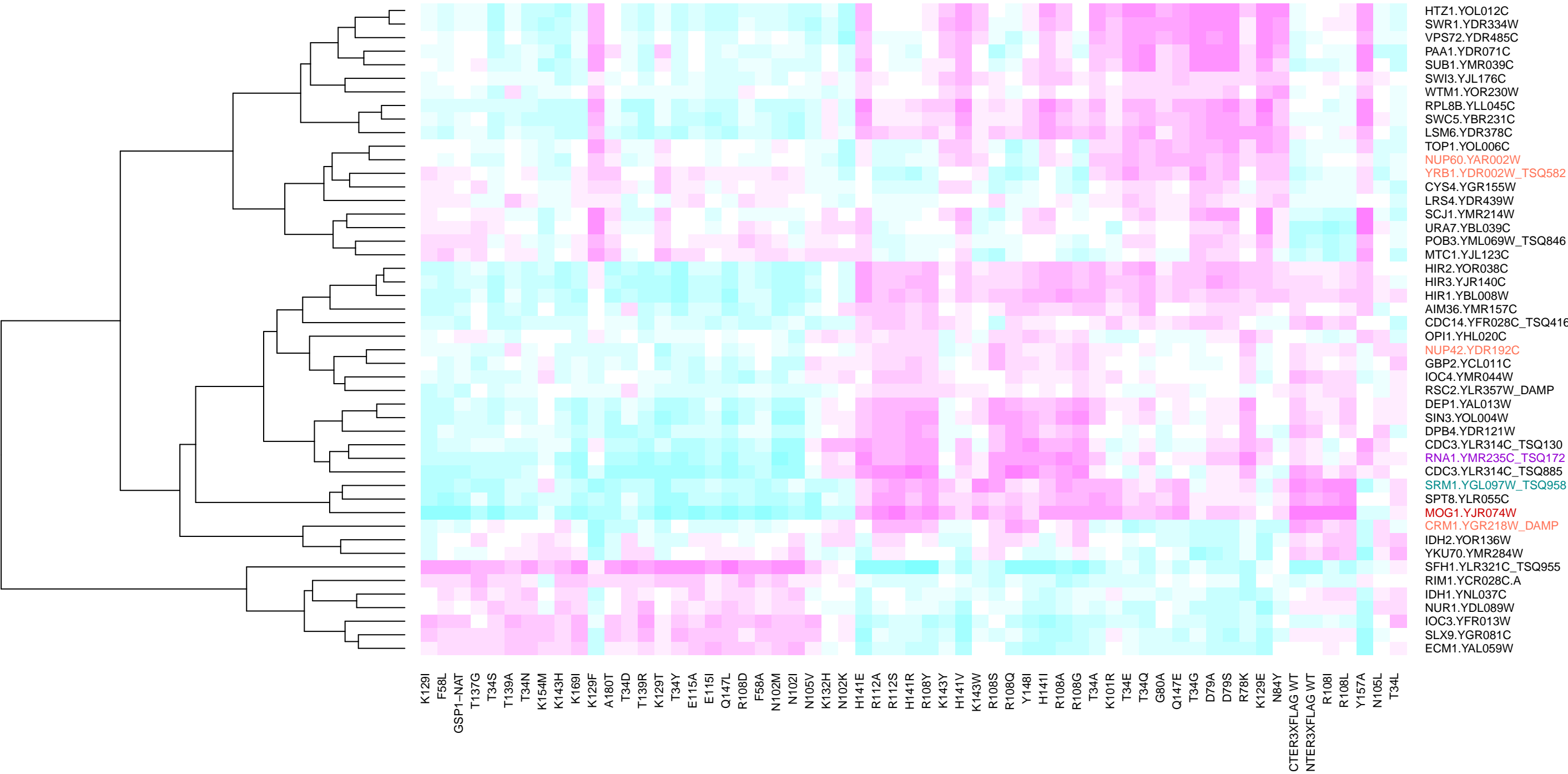
protein modification by small protein conjugation or removal_GO_15_1_mut



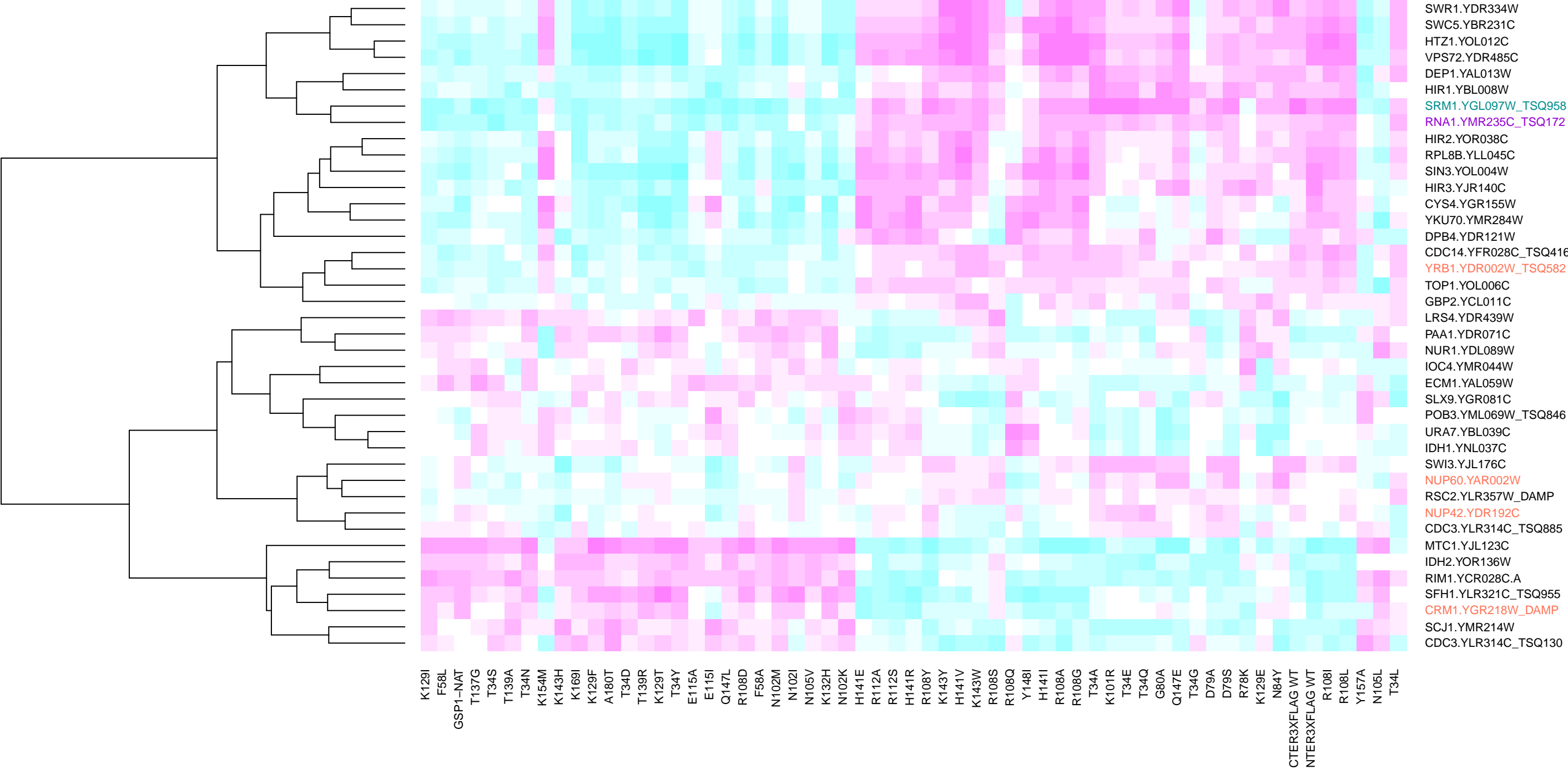
chromatin organization



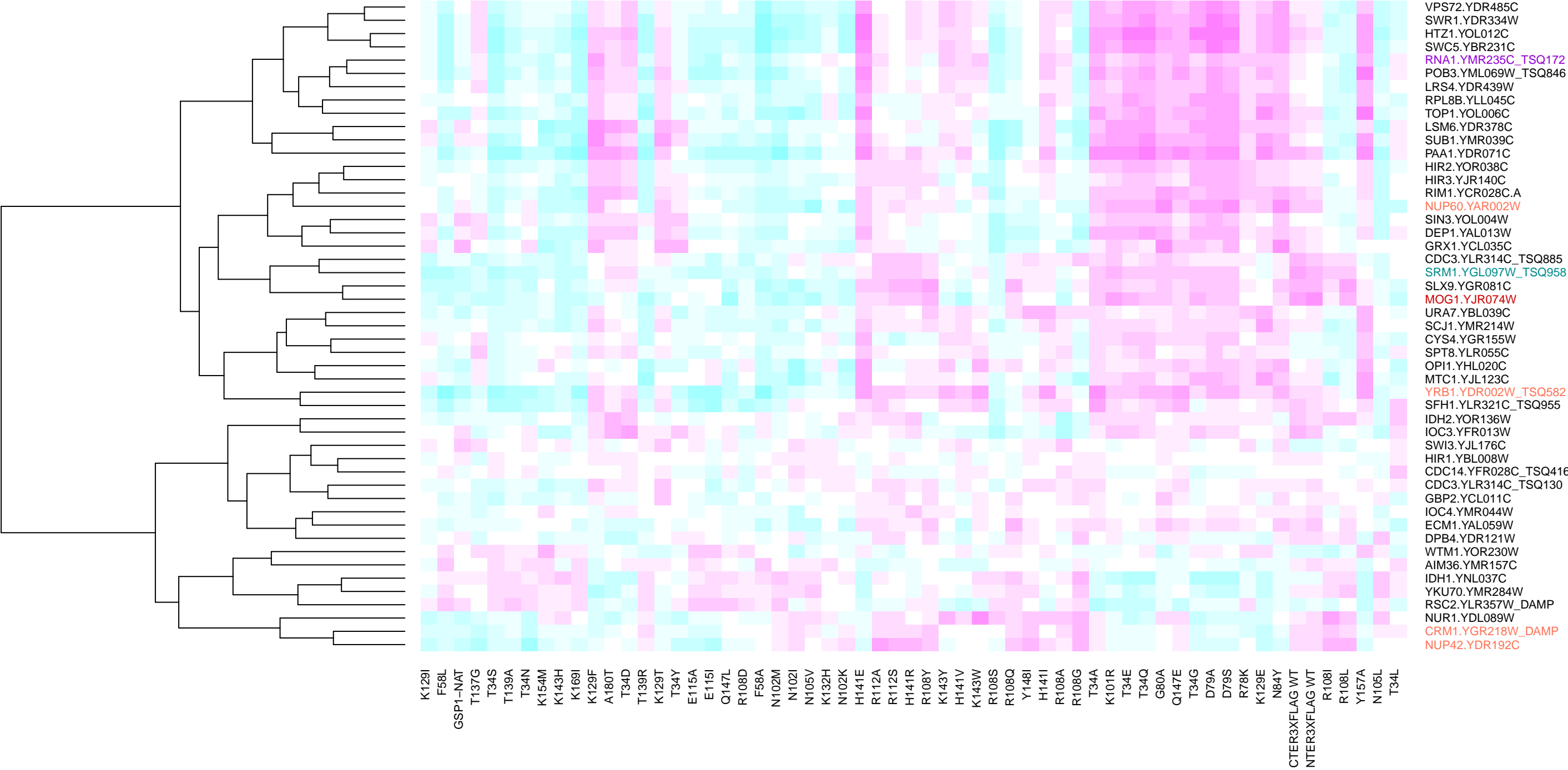
protein modification by small protein conjugation or removal_GO_30_1_mut



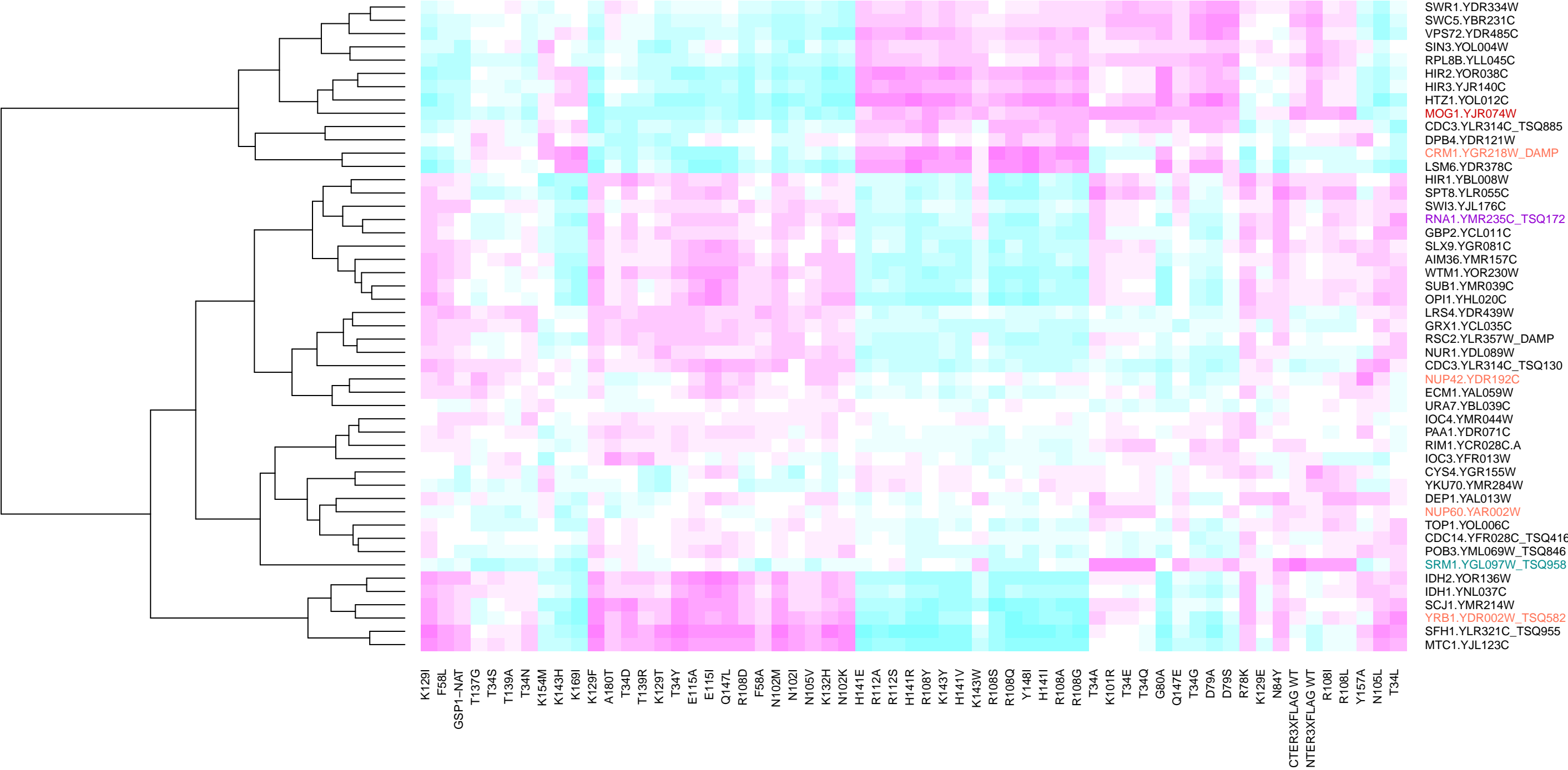
sig_Gsp1_Gl_15_11_all



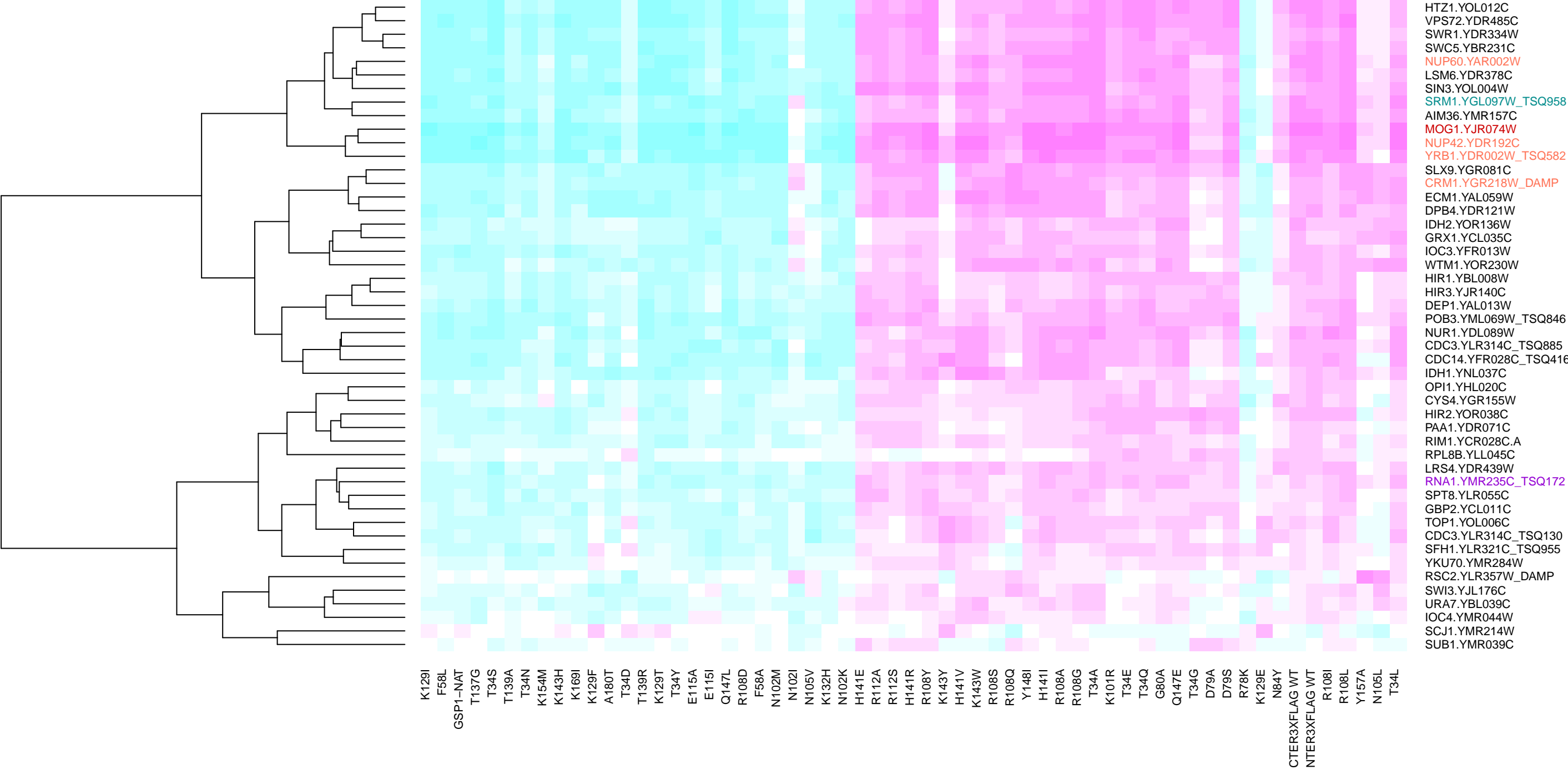
sig_Gsp1_Gl_30_2_mut



sig_Gsp1_Gl_30_3_all



cellular response to DNA damage stimulus_GO_15_3_mut



lipid binding_GO_15_2_all

