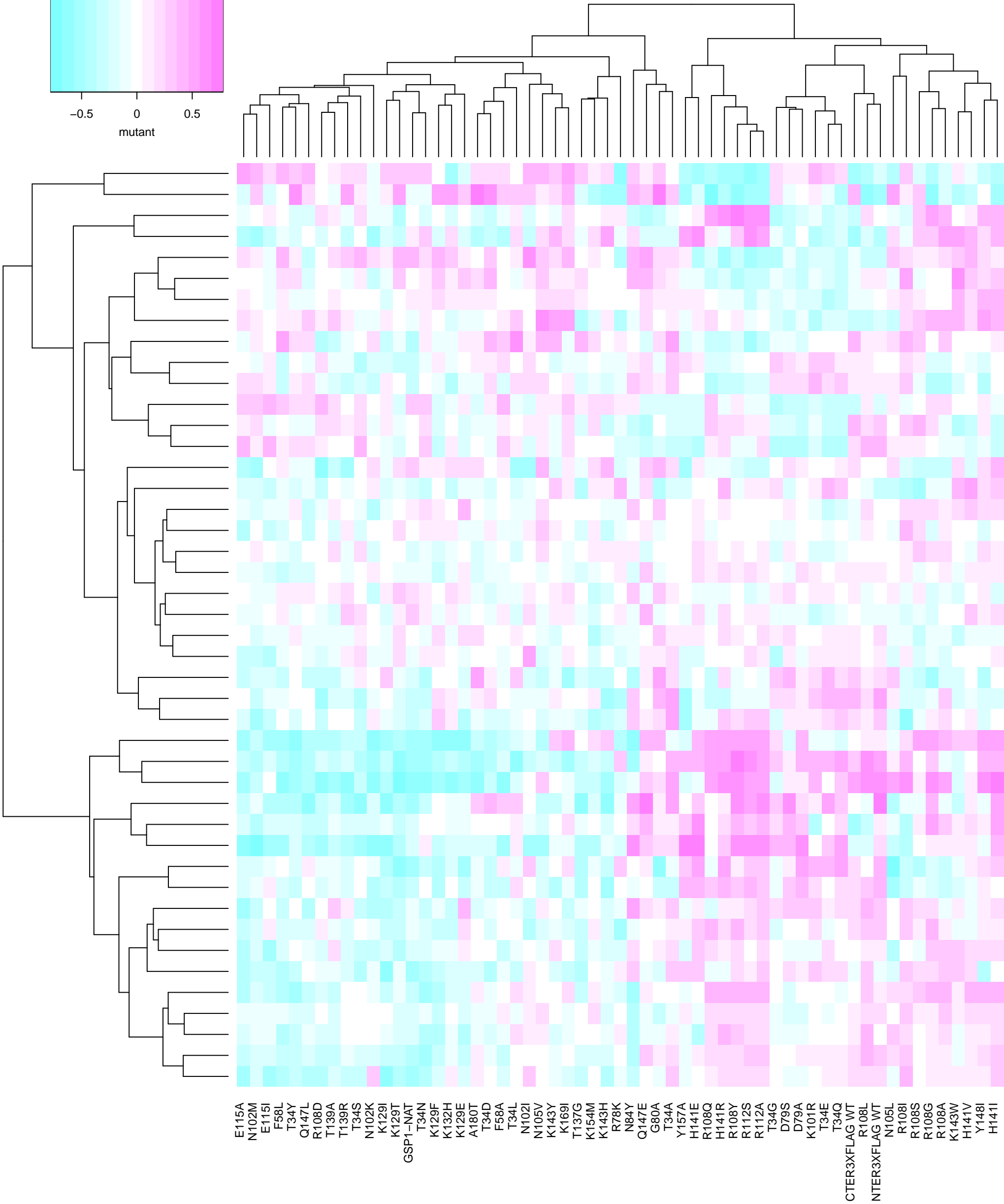
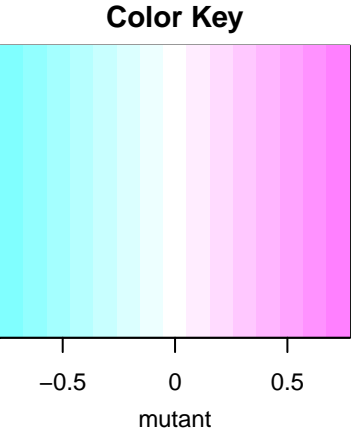


NUP120



- transcription and mRNA processing_GO_3
- peroxisome_GO_0
- chromatin_GO_4
- chromatin_GO_1
- cytoskeleton_GO_2
- budding_GO_2
- vacuole_GO_1
- nuclear transport_GO_10
- Golgi and ER_GO_3
- nuclear transport_GO_3
- ribosome_GO_1
- metabolic_GO_1
- transcription and mRNA processing_GO_1
- Golgi and ER_GO_2
- ribosome_GO_2
- nuclear transport_GO_8
- mitochondrion_GO_1
- cell cycle_GO_2
- Golgi and ER_GO_1
- nuclear transport_GO_2
- metabolic_GO_2
- mitochondrion_GO_2
- lipids_GO_2
- metabolic_GO_3
- nuclear transport_GO_7
- nuclear transport_GO_5
- transcription and mRNA processing_GO_6
- cell cycle_GO_3
- mitochondrion_GO_3
- nuclear transport_GO_9
- chromatin_GO_3
- transcription and mRNA processing_GO_2
- nuclear transport_GO_4
- transcription and mRNA processing_GO_5
- nuclear transport_GO_6
- lipids_GO_1
- transcription and mRNA processing_GO_4
- cell cycle_GO_4
- budding_GO_3
- budding_GO_1
- nuclear transport_GO_1
- cell cycle_GO_1
- cytoskeleton_GO_1
- chromatin_GO_2