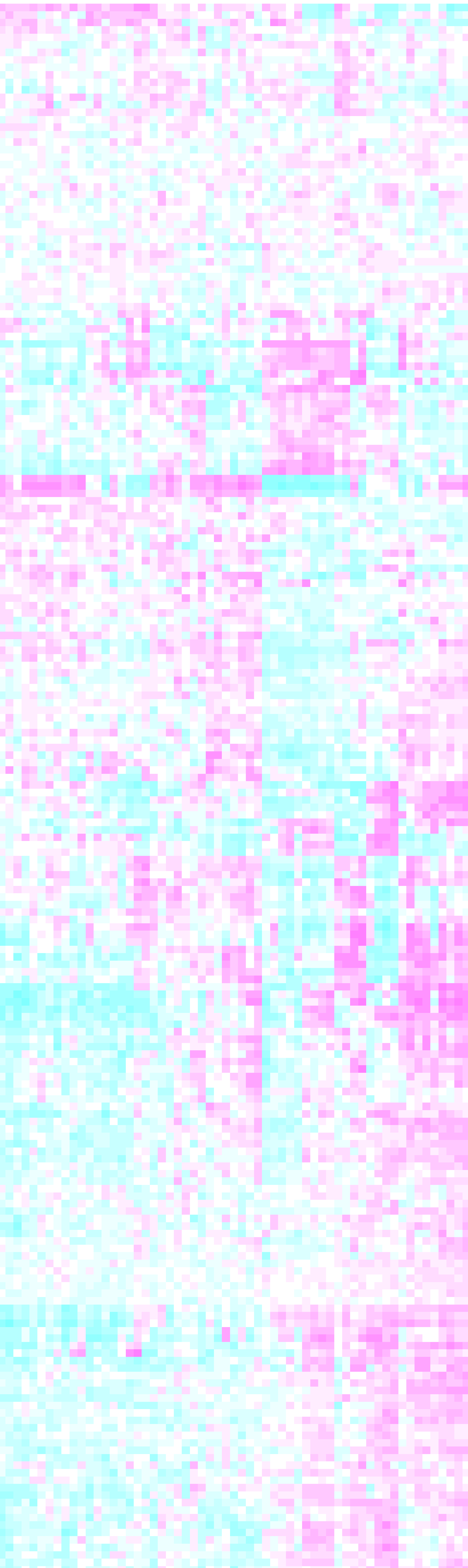
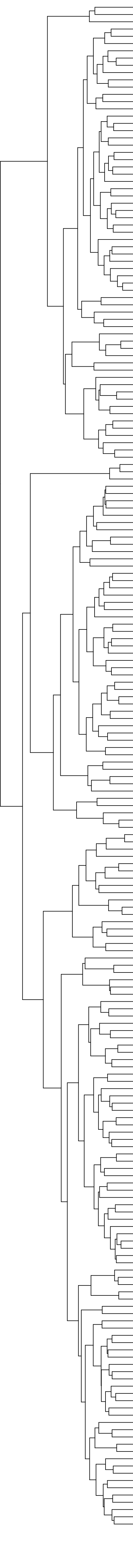
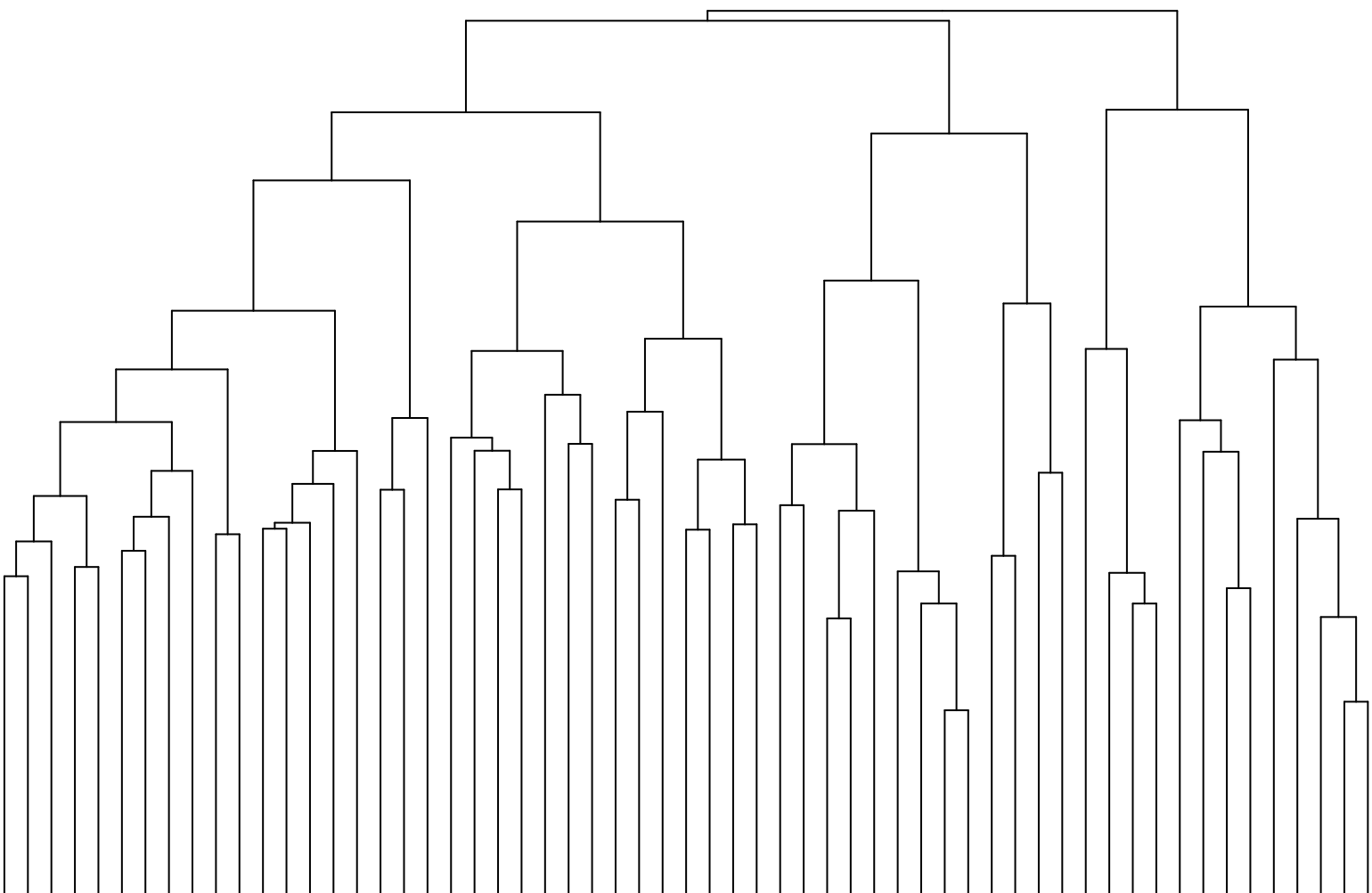
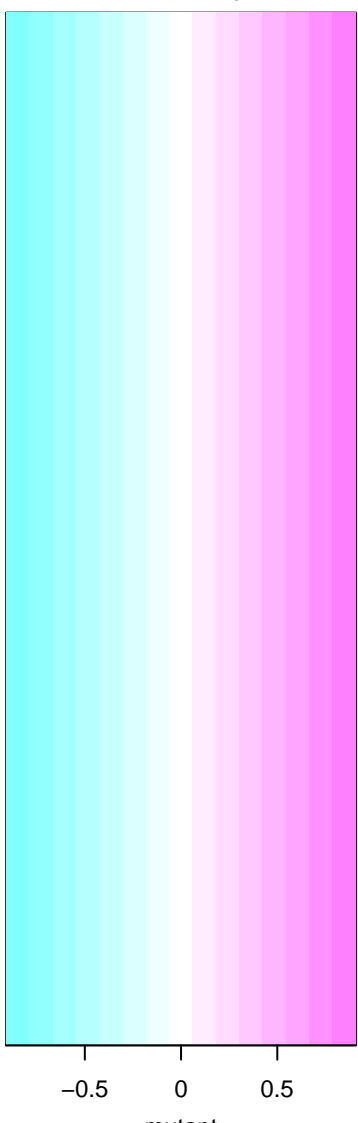


Color Key

all_partners



- RNA1_TSQ172 transcription and mRNA processing_GO
- YRB1_TSQ582 cell cycle_GO_4
- RNA1_TSQ172 transcription and mRNA processing_GO
- PSE1_TSQ683 budding_GO_2
- PSE1_TSQ683 lipids_GO_1
- PSE1_TSQ683 mitochondrion_GO_1
- PSE1_TSQ683 cytoskeleton_GO_2
- PSE1_TSQ683 vacuole_GO_1
- YRB1_TSQ582 vacuole_GO_1
- RNA1_TSQ172 ribosome_GO_2
- YRB1_TSQ582 chromatin_GO_2
- YRB1_TSQ582 cell cycle_GO_1
- RNA1_TSQ172 metabolic_GO_3
- RNA1_TSQ172 nuclear transport_GO_3
- PSE1_TSQ683 transcription and mRNA processing_GO
- YRB1_TSQ582 nuclear transport_GO_2
- PSE1_TSQ683 Golgi and ER_GO_2
- PSE1_TSQ683 lipids_GO_2
- SRM1_TSQ958 mitochondrion_GO_2
- PSE1_TSQ683 nuclear transport_GO_1
- PSE1_TSQ683 ribosome_GO_1
- RNA1_TSQ172 lipids_GO_1
- SRM1_TSQ958 Golgi and ER_GO_1
- RNA1_TSQ172 vacuole_GO_1
- MOG1_chromatin_GO_3
- YRB1_TSQ582 mitochondrion_GO_3
- MOG1_mitochondrion_GO_3
- PSE1_TSQ683 transcription and mRNA processing_GO
- YRB1_TSQ582 nuclear transport_GO_1
- YRB1_TSQ582 budding_GO_1
- PSE1_TSQ683 mitochondrion_GO_2
- RNA1_TSQ172 nuclear transport_GO_2
- MOG1_mitochondrion_GO_1
- SRM1_TSQ958 vacuole_GO_1
- SRM1_TSQ958 lipids_GO_1
- MOG1_nuclear transport_GO_4
- PSE1_TSQ683 transcription and mRNA processing_GO
- YRB1_TSQ582 mitochondrion_GO_2
- RNA1_TSQ172 mitochondrion_GO_2
- PSE1_TSQ683 cell cycle_GO_4
- RNA1_TSQ172 cell cycle_GO_4
- RNA1_TSQ172 mitochondrion_GO_3
- PSE1_TSQ683 nuclear transport_GO_3
- PSE1_TSQ683 chromatin_GO_3
- MOG1_cell cycle_GO_5
- PSE1_TSQ683 budding_GO_1
- PSE1_TSQ683 cytoskeleton_GO_1
- PSE1_TSQ683 cell cycle_GO_3
- RNA1_TSQ172 chromatin_GO_1
- RNA1_TSQ172 nuclear transport_GO_4
- MOG1_Golgi and ER_GO_3
- SRM1_TSQ958 Golgi and ER_GO_3
- YRB1_TSQ582 Golgi and ER_GO_3
- PSE1_TSQ683 Golgi and ER_GO_3
- RNA1_TSQ172 Golgi and ER_GO_3
- MOG1_nuclear transport_GO_5
- RNA1_TSQ172 nuclear transport_GO_5
- PSE1_TSQ683 nuclear transport_GO_5
- SRM1_TSQ958 nuclear transport_GO_5
- RNA1_TSQ172 budding_GO_2
- RNA1_TSQ172 transcription and mRNA processing_GO
- RNA1_TSQ172 ribosome_GO_1
- RNA1_TSQ172 cell cycle_GO_3
- YRB1_TSQ582 cell cycle_GO_3
- SRM1_TSQ958 cell cycle_GO_3
- YRB1_TSQ582 nuclear transport_GO_6
- MOG1_metabolic_GO_1
- RNA1_TSQ172 chromatin_GO_2
- YRB1_TSQ582 lipids_GO_2
- PSE1_TSQ683 nuclear transport_GO_7
- YRB1_TSQ582 transcription and mRNA processing_GO
- YRB1_TSQ582 ribosome_GO_2
- PSE1_TSQ683 mitochondrion_GO_3
- RNA1_TSQ172 cell cycle_GO_1
- SRM1_TSQ958 Golgi and ER_GO_2
- PSE1_TSQ683 nuclear transport_GO_4
- PSE1_TSQ683 chromatin_GO_1
- PSE1_TSQ683 Golgi and ER_GO_1
- PSE1_TSQ683 metabolic_GO_2
- YRB1_TSQ582 budding_GO_2
- RNA1_TSQ172 nuclear transport_GO_6
- PSE1_TSQ683 transcription and mRNA processing_GO
- PSE1_TSQ683 transcription and mRNA processing_GO
- PSE1_TSQ683 nuclear transport_GO_9
- YRB1_TSQ582 cytoskeleton_GO_1
- RNA1_TSQ172 cytoskeleton_GO_1
- YRB1_TSQ582 chromatin_GO_1
- SRM1_TSQ958 mitochondrion_GO_1
- YRB1_TSQ582 transcription and mRNA processing_GO
- RNA1_TSQ172 transcription and mRNA processing_GO
- YRB1_TSQ582 ribosome_GO_1
- PSE1_TSQ683 chromatin_GO_2
- RNA1_TSQ172 metabolic_GO_1
- PSE1_TSQ683 cell cycle_GO_1
- MOG1_cell cycle_GO_1
- MOG1_lipids_GO_2
- RNA1_TSQ172 lipids_GO_2
- YRB1_TSQ582 transcription and mRNA processing_GO
- YRB1_TSQ582 metabolic_GO_3
- PSE1_TSQ683 metabolic_GO_3
- YRB1_TSQ582 nuclear transport_GO_3
- YRB1_TSQ582 chromatin_GO_4
- SRM1_TSQ958 chromatin_GO_1
- MOG1_chromatin_GO_1
- SRM1_TSQ958 cytoskeleton_GO_1
- SRM1_TSQ958 budding_GO_1
- SRM1_TSQ958 cytoskeleton_GO_2
- MOG1_cell cycle_GO_2
- MOG1_nuclear transport_GO_10
- MOG1_Golgi and ER_GO_2
- MOG1_transcription and mRNA processing_GO_1
- SRM1_TSQ958 transcription and mRNA processing_GO
- PSE1_TSQ683 budding_GO_3
- RNA1_TSQ172 budding_GO_3
- YRB1_TSQ582 budding_GO_3
- YRB1_TSQ582 transcription and mRNA processing_GO
- PSE1_TSQ683 cell cycle_GO_2
- RNA1_TSQ172 cell cycle_GO_2
- YRB1_TSQ582 cell cycle_GO_2
- RNA1_TSQ172 cytoskeleton_GO_2
- RNA1_TSQ172 chromatin_GO_3
- RNA1_TSQ172 nuclear transport_GO_10
- PSE1_TSQ683 nuclear transport_GO_10
- YRB1_TSQ582 nuclear transport_GO_10
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- RNA1_TSQ172 Golgi and ER_GO_2
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- RNA1_TSQ172 mitochondrion_GO_1
- RNA1_TSQ172 metabolic_GO_2
- YRB1_TSQ582 metabolic_GO_2
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- SRM1_TSQ958 lipids_GO_2
- YRB1_TSQ582 metabolic_GO_1
- YRB1_TSQ582 cytoskeleton_GO_2
- SRM1_TSQ958 metabolic_GO_2
- RNA1_TSQ172 nuclear transport_GO_7
- SRM1_TSQ958 nuclear transport_GO_7
- MOG1_budding_GO_3
- MOG1_cytoskeleton_GO_1
- MOG1_cytoskeleton_GO_2
- SRM1_TSQ958 nuclear transport_GO_6
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- SRM1_TSQ958 chromatin_GO_4
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- SRM1_TSQ958 nuclear transport_GO_10
- MOG1_transcription and mRNA processing_GO_5
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- PSE1_TSQ683 nuclear transport_GO_6
- PSE1_TSQ683 nuclear transport_GO_2
- PSE1_TSQ683 metabolic_GO_1
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- MOG1_Golgi and ER_GO_1
- RNA1_TSQ172 nuclear transport_GO_1
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- RNA1_TSQ172 transcription and mRNA processing_GO
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- MOG1_nuclear transport_GO_3
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- MOG1_nuclear transport_GO_1
- SRM1_TSQ958 metabolic_GO_3
- MOG1_metabolic_GO_3

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K108I
R108M
N102M
Q147L
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F58A
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K169I
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K143H
K143W
N105V
N105L
K132H
T34L
N102I
R78K
N102K
N84Y
T34Y
K129E
K128E
K128D
A180T
R108S
R108A
R108G
R108Q
Y148I
H141R
R108Y
R112S
R112A
H141V
H141I
Y157A
H141E
R108I
R108L
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CTER3XFLAG WT
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T34G
T34A
K101R
T34E
T34E