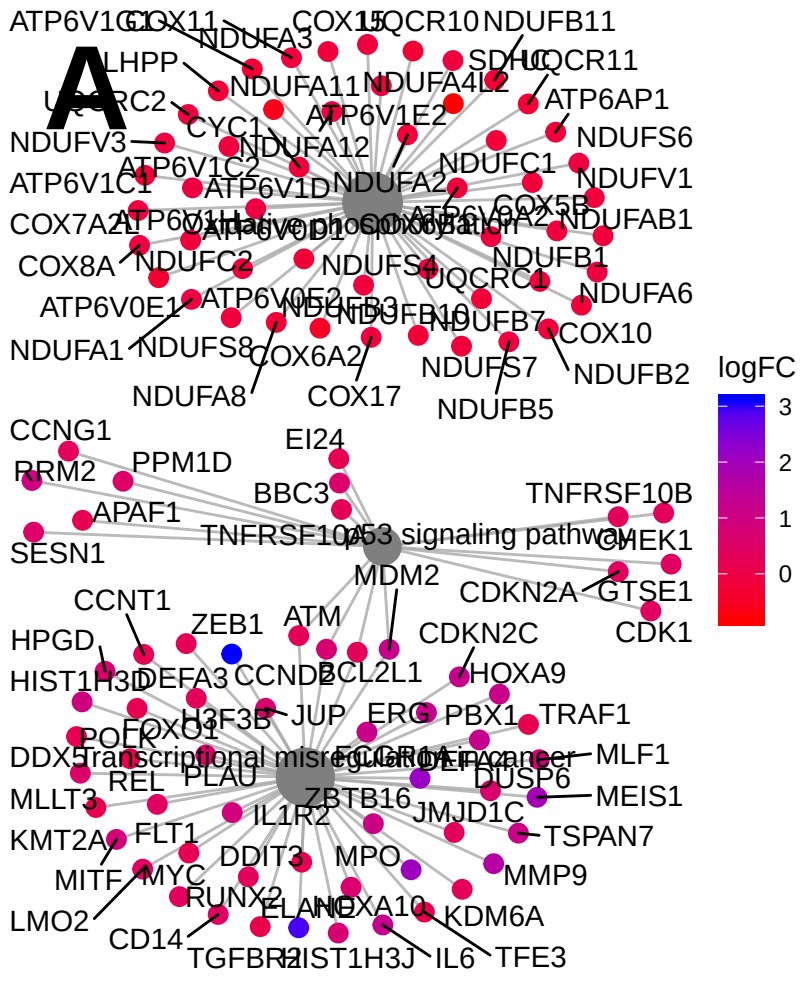
A
ATP6V1G1
COX11
NDUFA3
COX15
UQCR10
NDUFB11
LHPP
NDUFA11
NDUFA4L2
SDHD
UQCR11
UQCRC2
CYC1
ATP6V1E2
ATP6AP1
NDUFV3
NDUFA12
NDUFS6
ATP6V1C2
NDUFC1
ATP6V1C1
ATP6V1D
NDUFA2
NDUFV1
COX7A2L
ATP6V1H
ATP6V0D1
ATP6V0A2
COX5B
NDUFAB1
COX8A
NDUFC2
NDUFS4
NDUFB1
UQCRC1
ATP6V0E1
ATP6V0E2
NDUFB3
NDUFB10
NDUFB7
NDUFA6
NDUFA1
NDUFS8
COX6A2
COX10
NDUFS7
NDUFB2
NDUFA8
COX17
NDUFB5
logFC
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2
1
0
CCNG1
EI24
RRM2
PPM1D
BBC3
TNFRSF10B
APAF1
TNFRSF10A
p53 signaling pathway
CHEK1
SESN1
MDM2
CDKN2A
GTSE1
CCNT1
ZEB1
ATM
CDKN2C
CDK1
HPGD
HIST1H3D
DEFA3
CCND2
BCL2L1
HOXA9
H3F3B
JUP
ERG
PBX1
TRAF1
POFUT
FOXO1
DDX5
Transcriptional misregulation in cancer
FCGR1A
DEFA4
MLF1
REL
PLAU
DUSP6
MLLT3
ZBTB16
MEIS1
IL1R2
JMJD1C
KMT2A
FLT1
TSPAN7
DDIT3
MPO
MITF
MYC
MMP9
RUNX2
LMO2
ELANE
HOXA10
KDM6A
CD14
TGFBR2
HIST1H3J
IL6
TFE3
B
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76
196
TP5
C
CDK12
TP53
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10000
5000
normalized counts
D
PGBD2
1000
500
300
E
HYPK