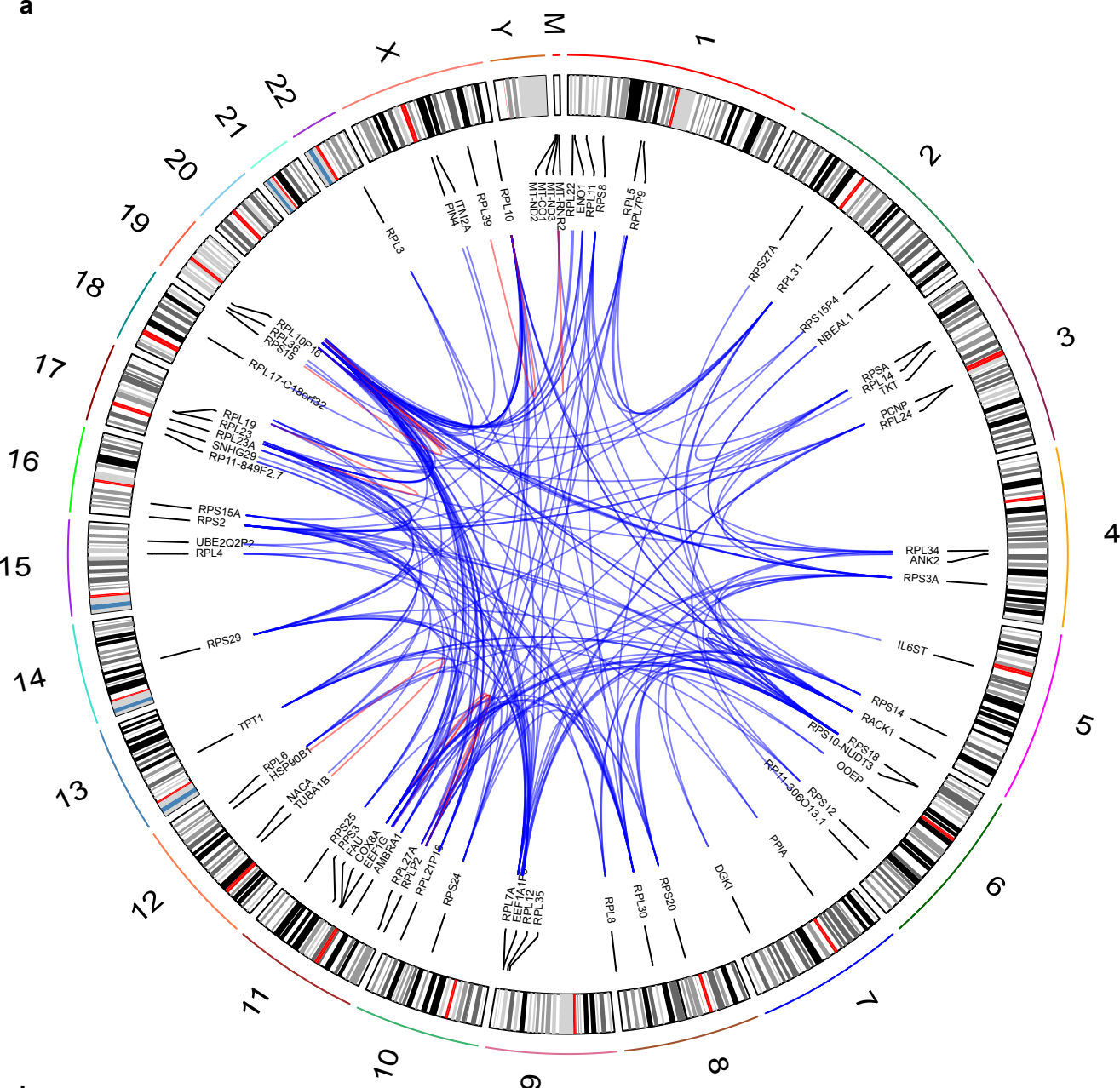
a



b

Sample	GO Term	Genes	P-value
A549	Cytoplasmic translation	RPL34, RPS21	1.467×10^{-2}
A549	Translation	RPL34, RPS21	3.040×10^{-2}
HepG2	Translation	EEF1A1, RPS19, RPL13, RPS12	2.172×10^{-4}
HepG2	Intracellular iron ion homeostasis	MT-RNR2, FTH1, FTL	7.332×10^{-4}
HepG2	Cytoplasmic translation	RPS19, RPL13, RPS12	1.528×10^{-3}
HepG2	Intracellular sequestering of iron ion	FTH1, FTL	4.250×10^{-3}
HepG2	Translational elongation	EEF1A1, EEF1B2	1.150×10^{-2}
HepG2	Iron ion transport	FTH1, FTL	1.630×10^{-2}
HepG2	Ribosomal small subunit biogenesis	RPS19, RPS12	4.526×10^{-2}
HCT116	Regulation of translation	RPS3, RPL38, GAPDH	6.151×10^{-3}
HCT116	Negative regulation of translation	RPS3, RPL13A, GAPDH	6.525×10^{-3}
HCT116	Cytoplasmic translation	RPL4, RPS6, RPL13A, RPSA, RPL7A, RPS29, RPS3, RPL14, RPLP2, RPL13, RPS20, RPL38, RPS2, RPL28	1.008×10^{-24}
HCT116	Translation	RPL4, RPS6, RPL13A, RPSA, EEF1A1, RPL7A, RPS29, RPS3, RPL14, RPLP2, RPL13, RPS20, RPL38, RPS2, RPL28	1.566×10^{-22}
VCaP (SQK-RNA004 kit)	Cytoplasmic translation	RPS16, RPLP2, RPL29	7.15×10^{-5}
VCaP (SQK-RNA004 kit)	Translation	RPS16, EEF1A1P5, RPLP2, RPL29	1.06×10^{-6}