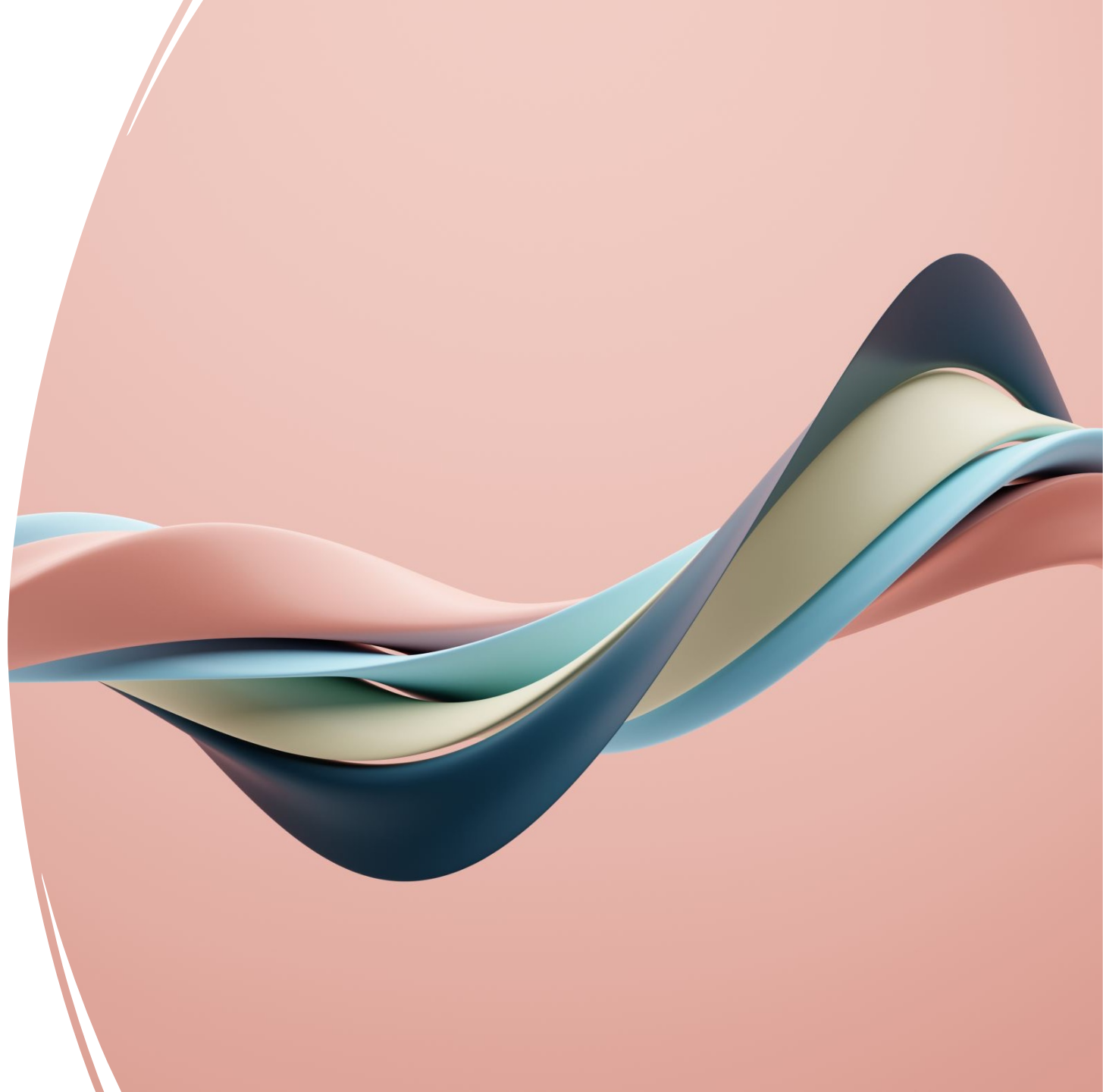


BULK RNA-SEQ ANALYSIS

BCG - GENOMICS AND TRANSCRIPTOMICS

A.A. 2023-2024

SABRINA PERVIN ABU



QUALITY CONTROL

MAT. 30507A

Genes with low/no expression
on all 9 samples are removed:
54042 → 23038

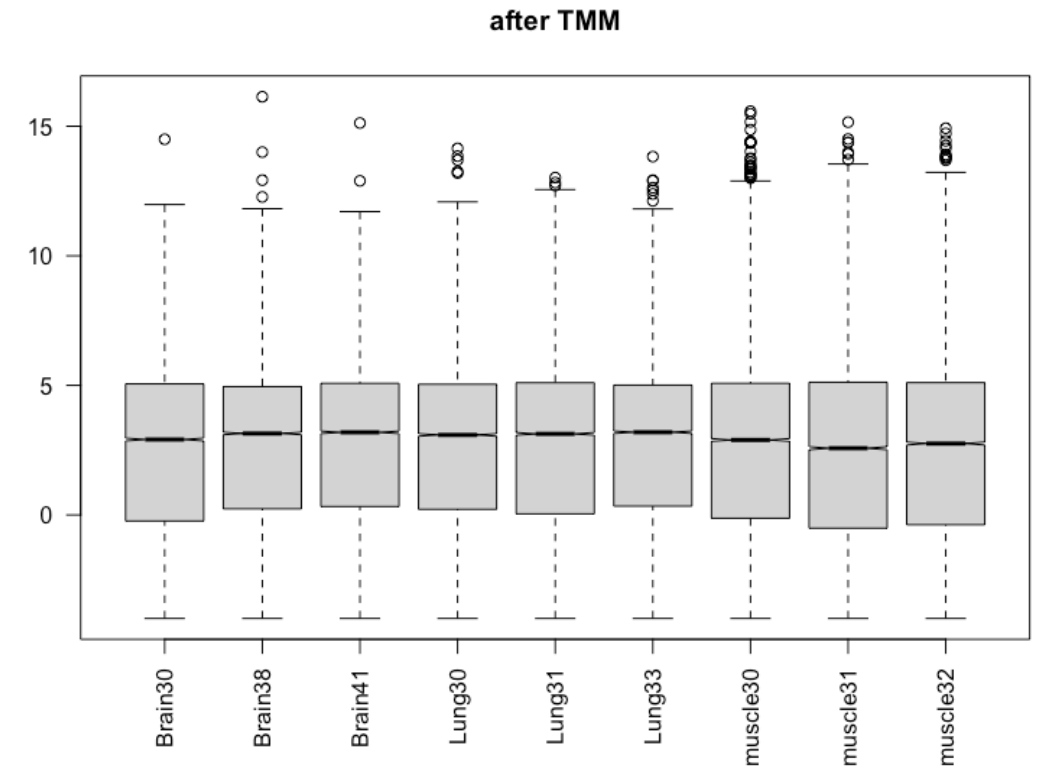
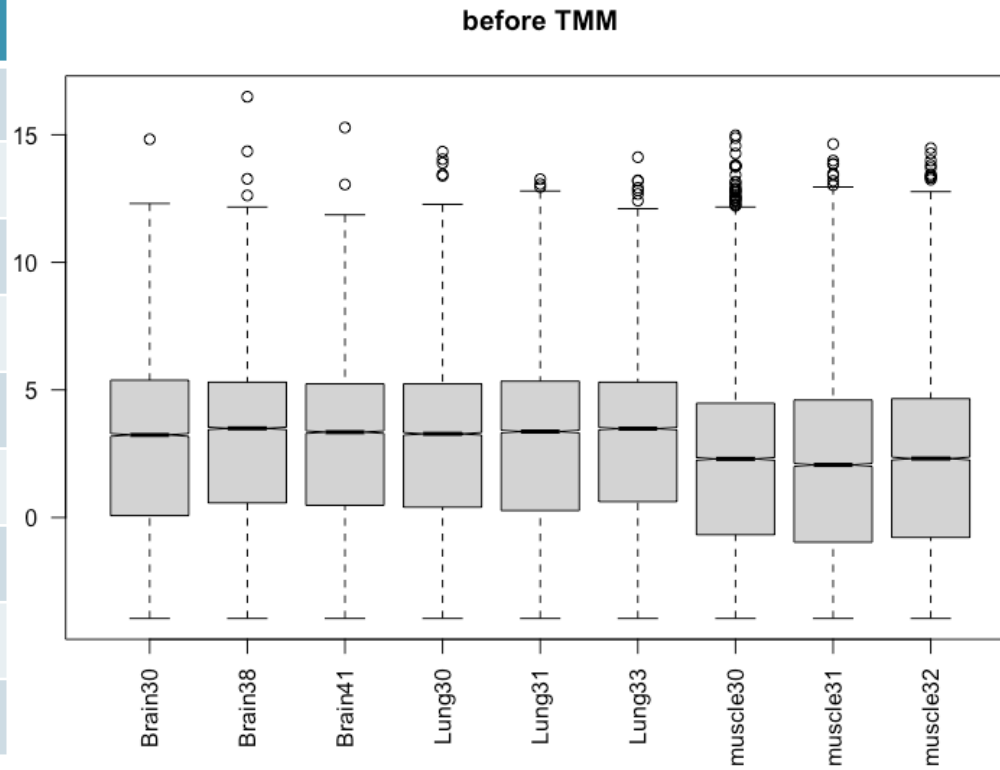
The thresholds set are

- RIN ≥ 6
- rRNA fraction $\leq 10\%$
- uniquely mapped reads $\geq 85\%$

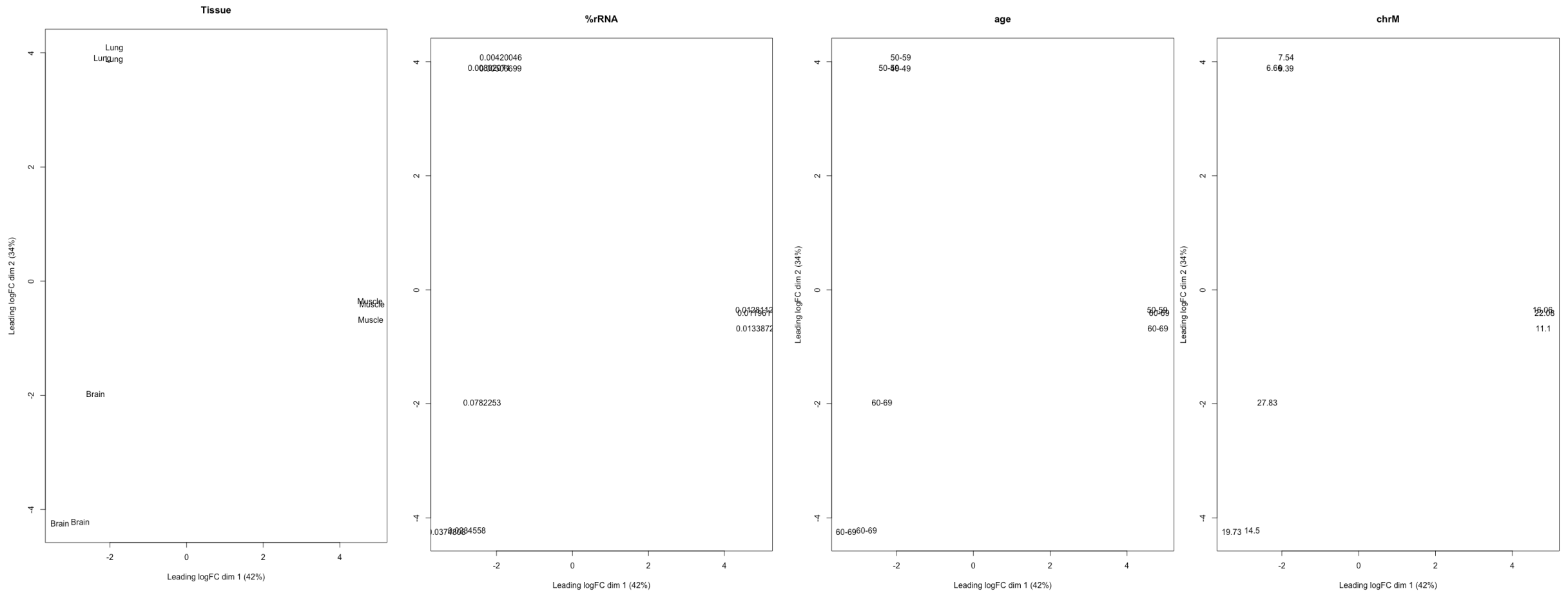
	BRAIN			LUNG			MUSCLE		
SAMPLE	30	38	41	30	31	33	30	31	32
RIN	7.4	7.2	7.3	8.4	8.4	7.7	7	8.4	7.5
%rRNA	0.023	0.078	0.037	0.004	0.008	0.005	0.013	0.012	0.011
%map	87.9	91.3	88	91.2	91.8	89.4	89.2	92.1	89.6

NORMALIZATION

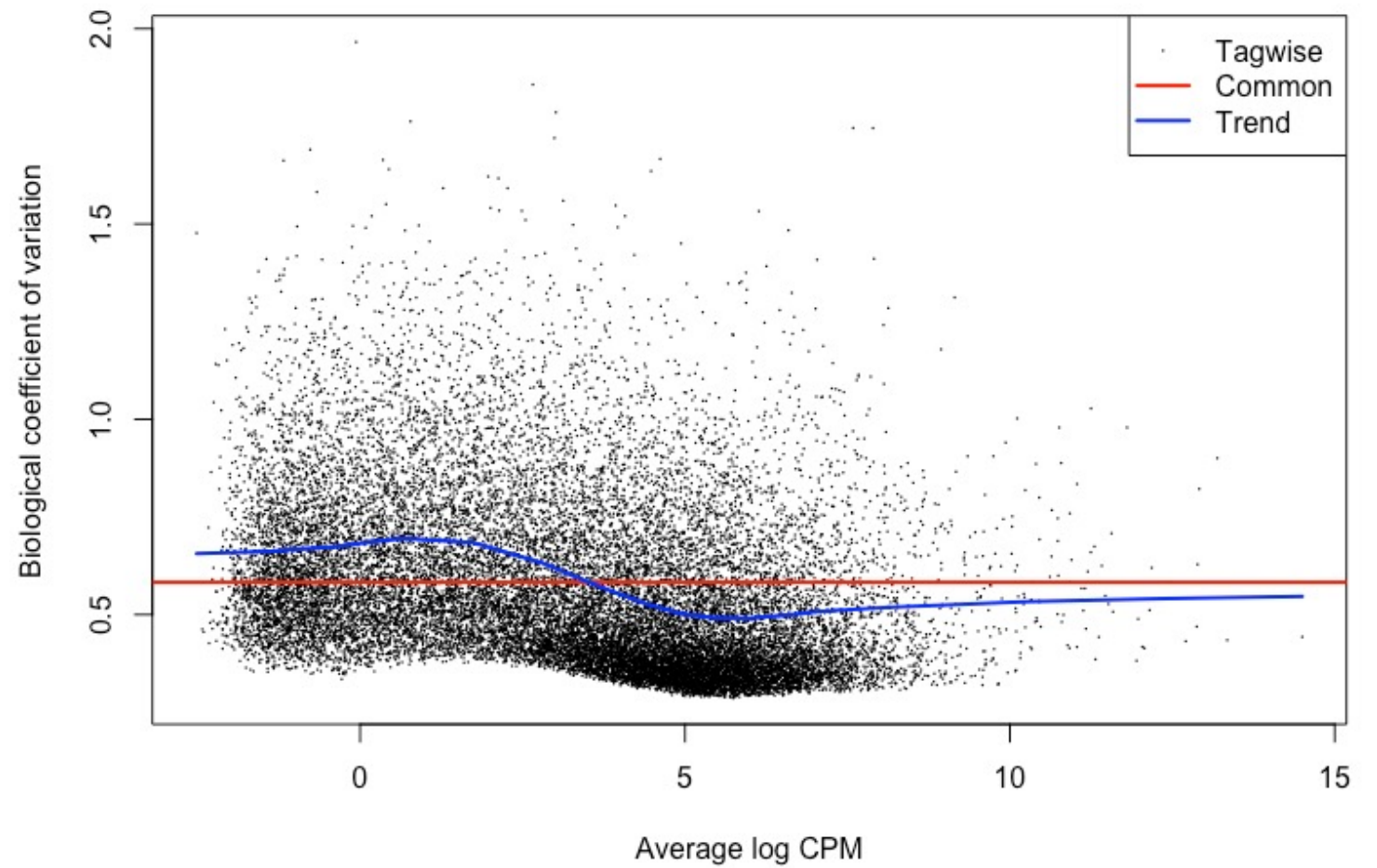
Sample	Norm. factors
BRAIN 30	1.2550309
BARIN 38	1.2764998
BRAIN 41	1.1163924
LUNG 30	1.1445551
LUNG 31	1.1822037
LUNG 33	1.2258074
MUSCLE 30	0.6588416
MUSCLE 31	0.6981477
MUSCLE 32	0.7328724



SAMPLES 2D VISUALIZATION

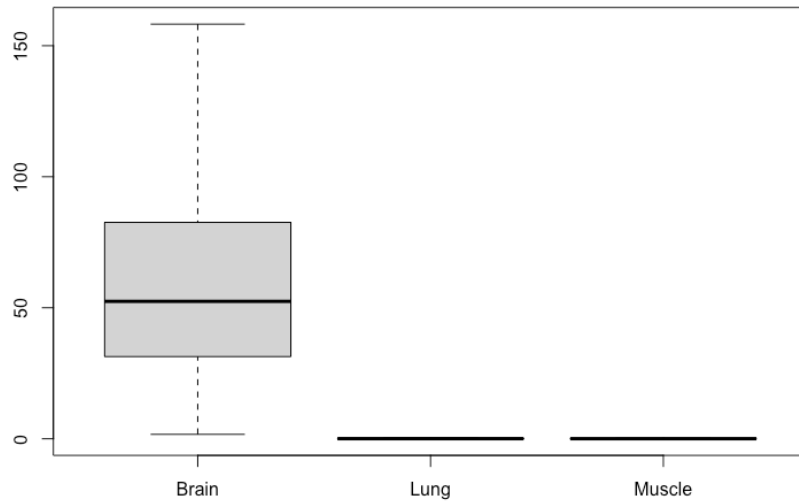


MEAN AND VARIANCE

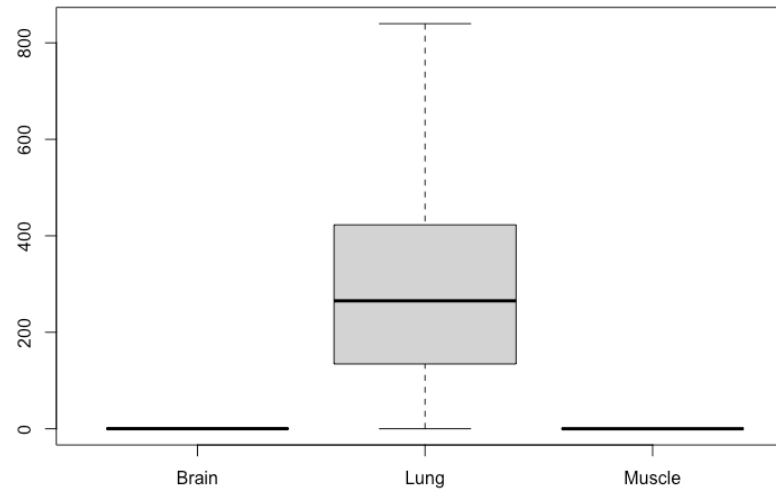


UP-REGULATED GENES

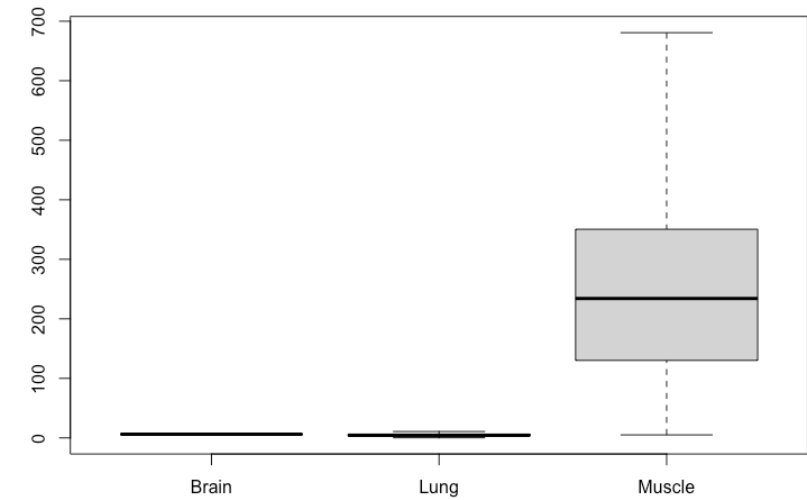
CNTN2



CEACAM6



IDI2



UP-REGULATED GENES IN BRAIN

vs LUNG

895

vs MUSCLE

1732

UP-REGULATED GENES IN LUNG

vs BRAIN

876

vs MUSCLE

1523

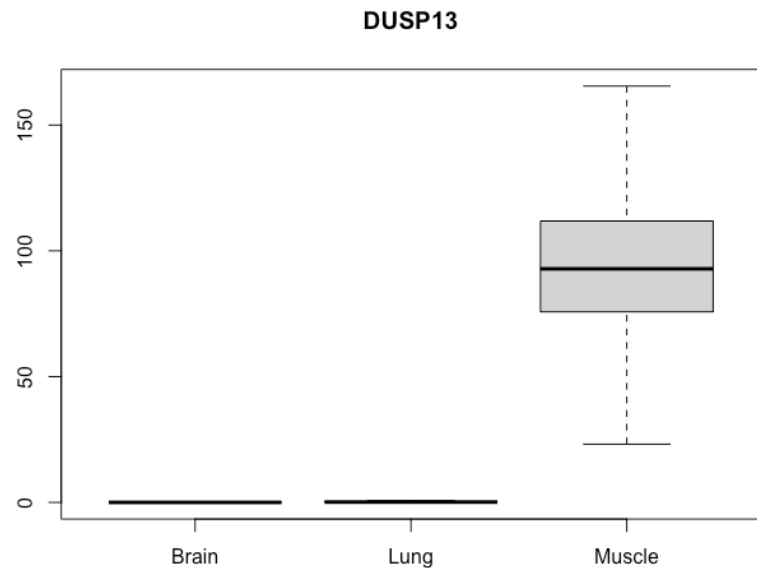
UP-REGULATED GENES IN MUSCLE

vs BRAIN

1691

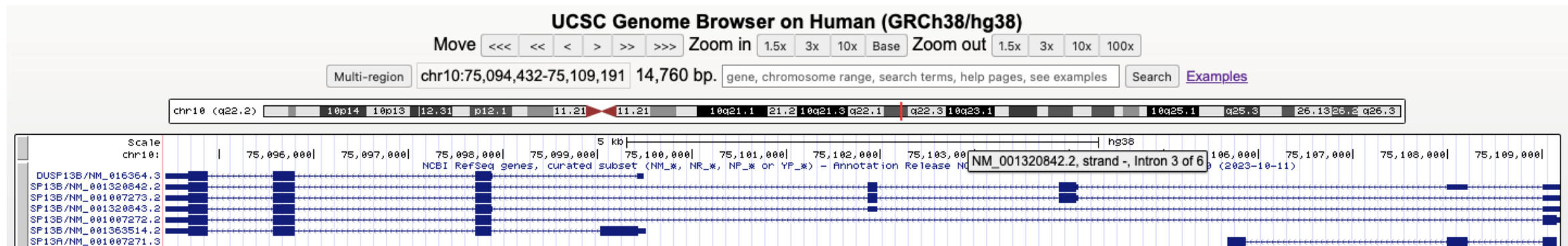
vs LUNG

1665



DUSP13

DUSP13		
<i>Dual Specificity Phosphatase 13</i>		
BIOLOGICAL PROCESS	Negative Regulation Of MAPK Cascade	GO:0043409
	Dephosphorylation	GO:0016311
	Phosphate-Containing Compound Metabolic Process	GO:0006796
	Regulation Of MAPK Cascade	GO:0043408
	Negative Regulation Of Intracellular Signal Transduction	GO:1902532
MOLECULAR FUNCTION	MAP Kinase Phosphatase Activity	GO:0033549
	Protein Tyrosine/Serine/Threonine Phosphatase Activity	GO:0008138
	Phosphoric Ester Hydrolase Activity	GO:0042578
	Phosphatase Activity	GO:0016791



FUNCTIONAL ENRICHMENT (TOP 500 DE GENES)

	BRAIN		LUNG		MUSCLE	
	GO ANNOTATION	FDR	GO ANNOTATION	FDR	GO ANNOTATION	FDR
BIOLOGICAL PROCESS	Nervous System Development (GO:0007399)	1.762e-13	B Cell Receptor Signaling Pathway	0.000001936	Cellular Respiration	1.117e-30
MOLECULAR FUNCTION	Voltage-Gated Monoatomic Cation Channel Activity	0.006338	Immunoglobulin Receptor Binding	0.004326	Oxidoreduction-Driven Active Transmembrane Transporter Activity	1.490e-23
CELLULAR COMPONENT	Neuron Projection	5.765e-21	Lamellar Body	3.328e-8	Mitochondrial Inner Membrane	2.608e-27