## Results

At this stage of the study we compared the differential expression between two aligners; kallisto and salmon based on six samples from 64, three control and three liver cancer patient and we will report the results here;

Salmon has a significant higher number of differentially expressed genes than kallisto as shown in figure (1) and figure (2), both aligners used human transcriptome as a reference. Kindly NOTE we worked on 6 samples out of our proposed 64 paired-end samples, also STAR, Magicblast and HiSat all requested huge memory and time so we were capable of analyzing this section only, although we are committed to analyze the whole study with the other aligners followed by wetlab experiment validation using real-time PCR for the top DEGs.

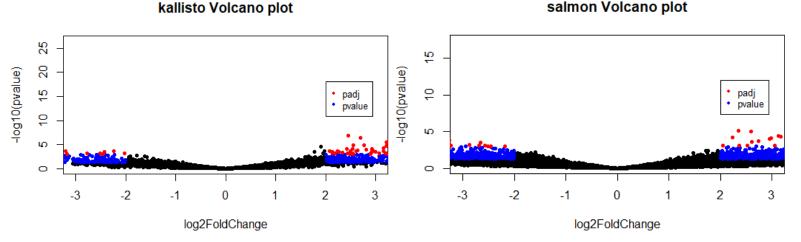


figure (1); volcano plot to visualize the abundance of the differentially expressed genes between kallisto and salmon the colored DEGs are higher than log2 fold change, red for adjusted p value lower than 0.05, and blue for p value lower than 0.05.

^	baseMean <sup>‡</sup>	log2FoldChange	IfcSE <sup>‡</sup>	stat <sup>‡</sup>	pvalue	pad	; ÷					
ENST00000621803.2	8.125057	6.677011	1.6363580	4.080410	4.495636e-05	5.76	0409e-03					
ENST00000633714.1	16.023466	7.659327	1.5389256	4.977061	6.455696e-07	1.90	8900e-04					
ENST00000631466.1	78.903354	8.992410	1.5339303	5.862332	4.564101e-09	4.04	8709e-06					
ENST00000632774.1	11.517318	6.207810	1.5802373	3.928404	8.551160e-05	8.88	3962e-03					
ENST00000626472.2	13.033880	6.384084	1.58		baseMea	baseMean <sup>‡</sup>		Change <sup>‡</sup>	IfcSE <sup>‡</sup>	stat <sup>‡</sup>	pvalue <sup>‡</sup>	padj <sup>‡</sup>
ENST00000390545.3	8.125057	6.677011	1.63 ENST00000372064.8		<b>4.8</b> 39.3	39.311092		9.641372		3.446233	5.684606e-04	3.581302e-02
ENST00000390547.3	16.023466	7.659327	1.53 ENST00000611116.2		64.79	1948	4.909457		1.4630846	3.355553	7.920656e-04	4.354793e-02
ENST00000390549.6	78.903354	8.992410	1.53 ENST00000633705.1		<b>5.1</b> 73.39	9790	0 4.547052		1.2813656	3.548598	3.872873e-04	2.708379e-02
ENST00000390551.6	11.517318	6.207810	1.58 ENST00000436911.6		1.6 12.10	1202	1202 4.984023		1.5117755	3.296801	9.779270e-04	4.813673e-02
ENST00000390559.6	13.033880	6.384084	1.58 ENST	0000061898	1.2 49.98	88447		7.740999	1.5467938	5.004545	5.599418e-07	3.342920e-04
ENST00000390321.2	191.570460	6.129093	1.11 ENST00000621803.2		3.2 397.73	.737426		7.564373	1.1489833	6.583536	4.593890e-11	1.985388e-07
ENST00000390323.2	238.740413	5.748243	0.69 ENST	0000063371	4.1 553.72	553.726667		7.298684	1.4150683	5.157831	2.498264e-07	1.799499e-04
Showing 1 to 12 of 23	2 entries	ENST	0000063146	6.1 3253.76	51687		8.131874	1.0246629	7.936145	2.085618e-15	1.502271e-11	
			ENST	0000062647	2.2 443.76	52266		5.467165	1.6222597	3.370093	7.514291e-04	4.260967e-02
figure (2); kallisto identified 232				ENST00000390539.2		22601		4.450210	1.3460775	3.306058	9.461862e-04	4.776217e-02
differentially expressed genes while				ENST00000390551.6		12787		11.281746	2.6084790	4.325029	1.525121e-05	3.003868e-03
salmon was a	•	ENST	ENST00000390331.3		6437		10.552858	2,4244235	4.352729	1,344535e-05	2.740948e-03	
		, 11122200		Showing 1 to 12 of 446 entries								