

RNA sequence analysis of rat acute experimental pancreatitis with and without fatty liver: a gene expression profiling comparative study.

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Running title: gene expression profiling of rat acute pancreatitis with fatty liver

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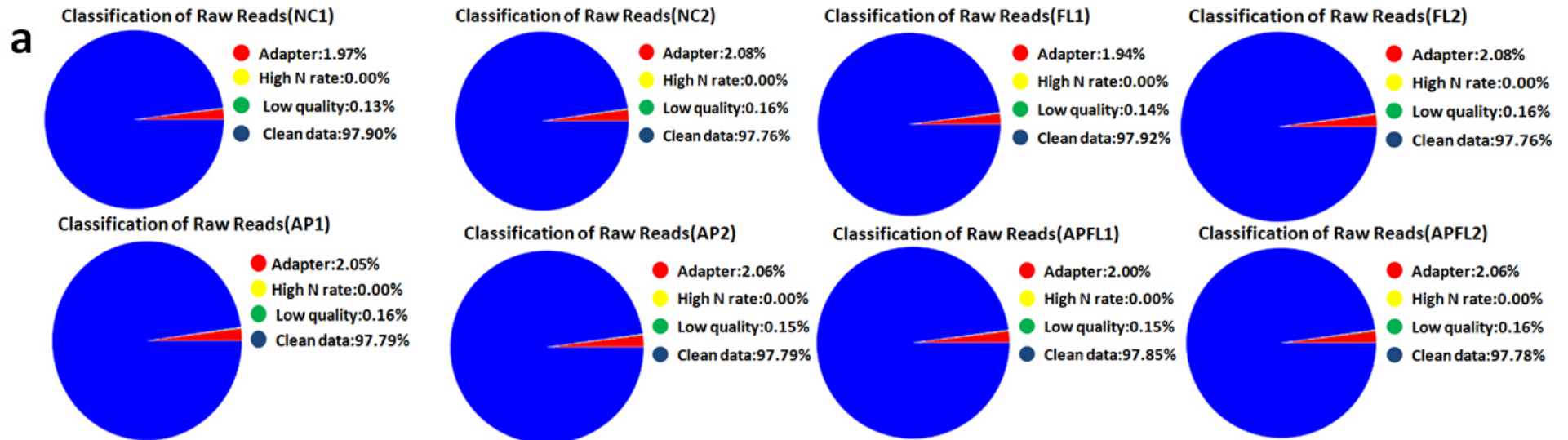
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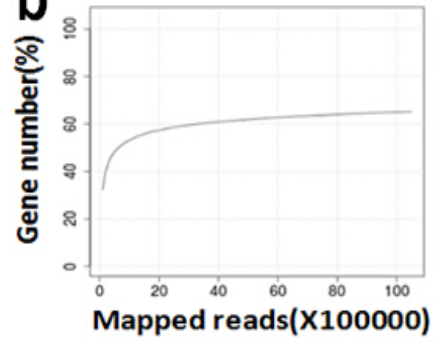
Supplementary Figure S1

RNA sequencing quality assessment. Sequence reads quality assessment (a); Sequencing saturation analysis (b); Distribution of reads on reference genes of sample (c); Statistics of gene coverage (d); Correlations of expression values between replicates (e).

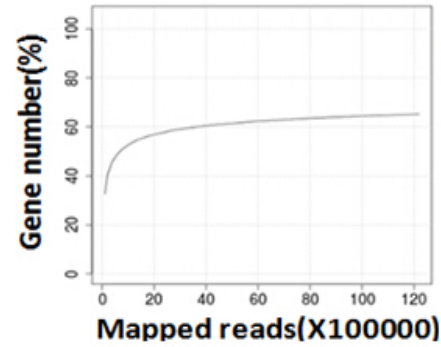


b

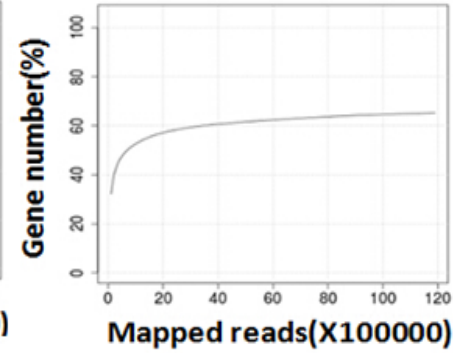
NC1



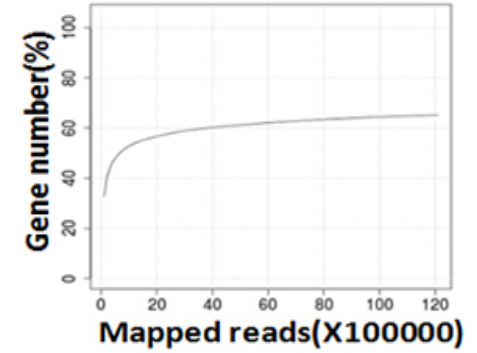
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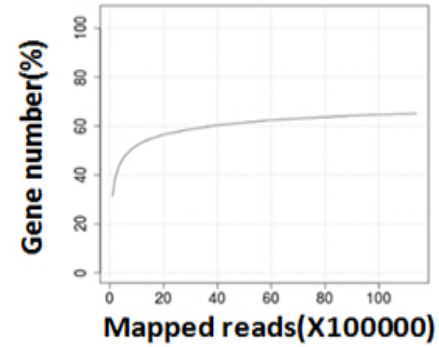
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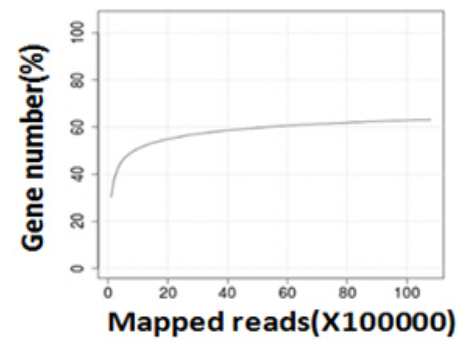
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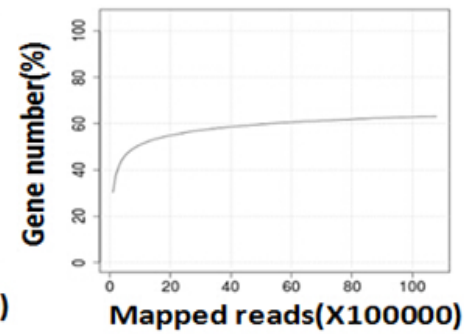
AP1



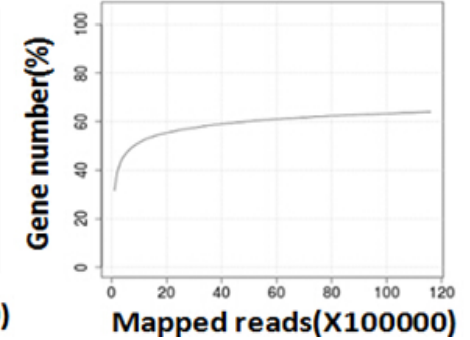
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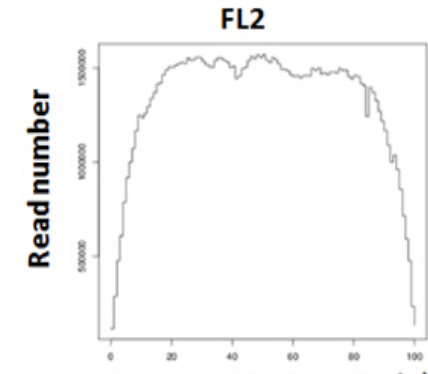
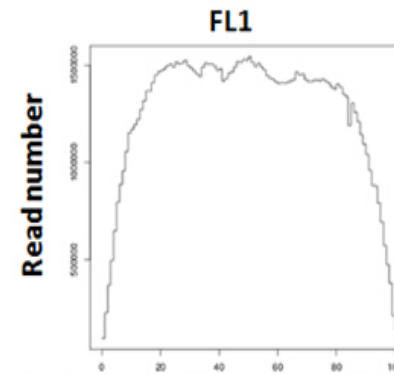
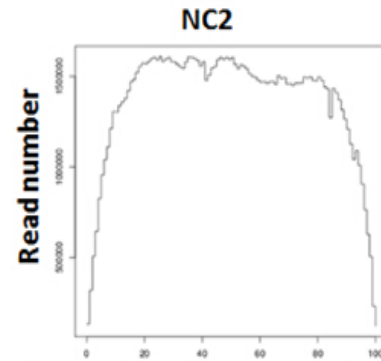
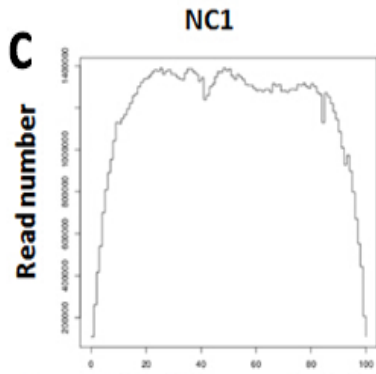


APFL1



APFL2



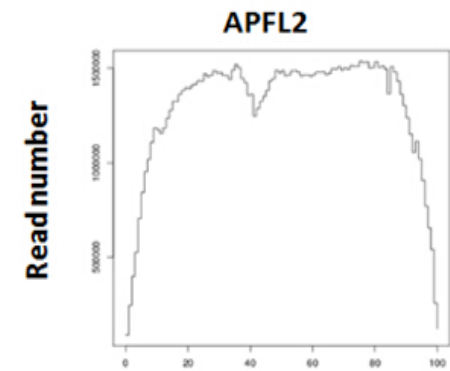
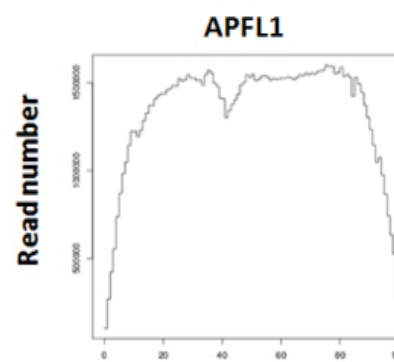
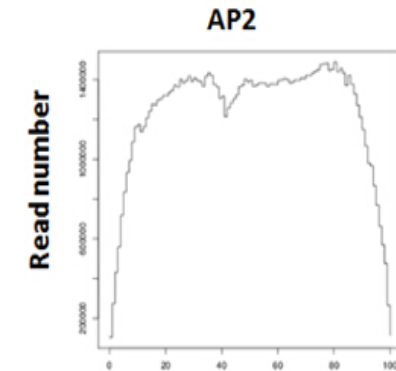
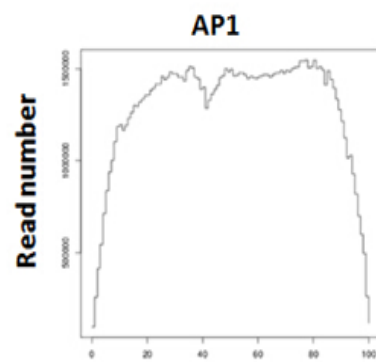
C

Percentile of the gene body(5' ->3')

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Percentile of the gene body(5' ->3')

Percentile of the gene body(5' ->3')



Percentile of the gene body(5' ->3')

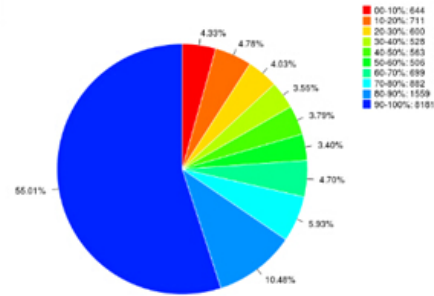
Percentile of the gene body(5' ->3')

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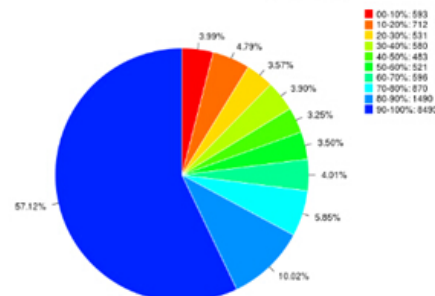
Percentile of the gene body(5' ->3')

d

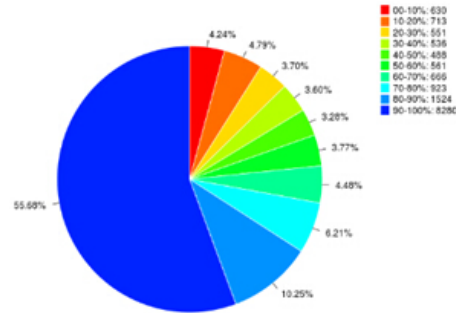
Distribution of Genes' Coverage(NC_1)



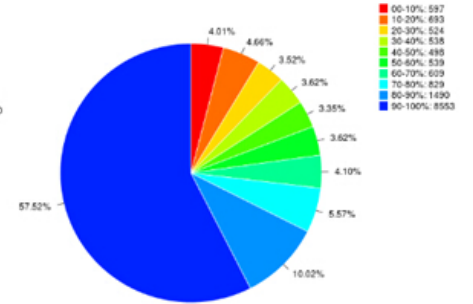
Distribution of Genes' Coverage(NC_2)



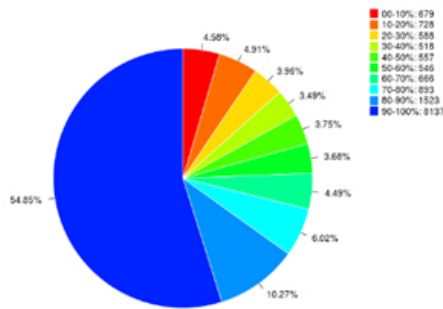
Distribution of Genes' Coverage(FL_1)



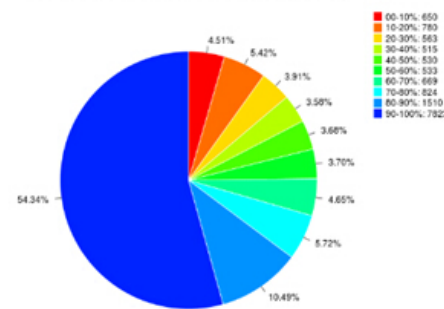
Distribution of Genes' Coverage(FL_2)



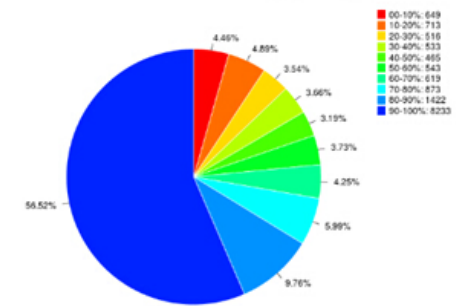
Distribution of Genes' Coverage(AP_1)



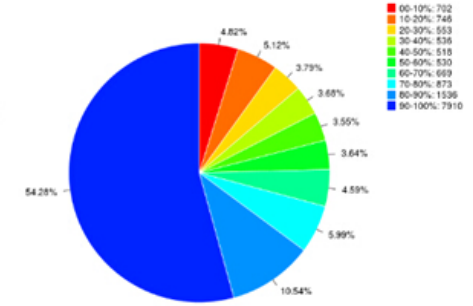
Distribution of Genes' Coverage(AP_2)



Distribution of Genes' Coverage(APFL_1)

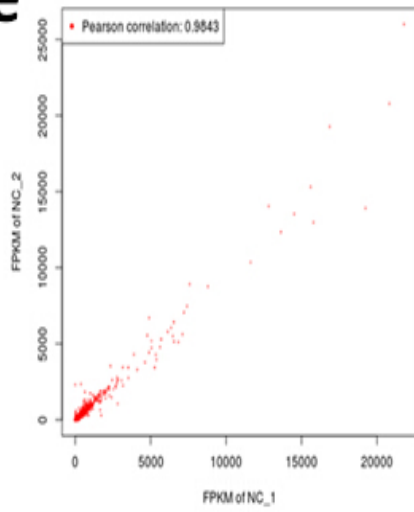


Distribution of Genes' Coverage(APFL_2)

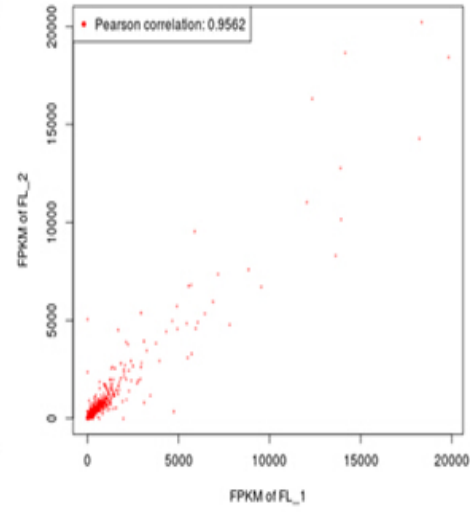


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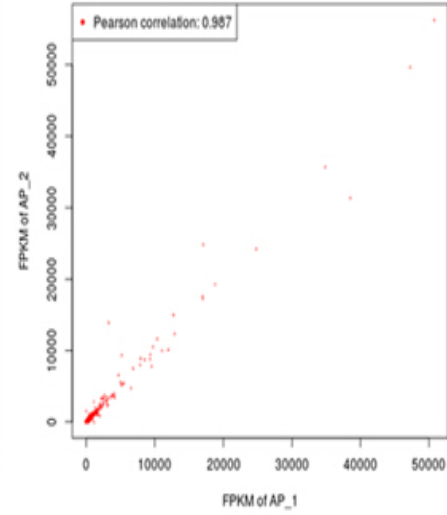
Sample NC1 and NC2 related analysis



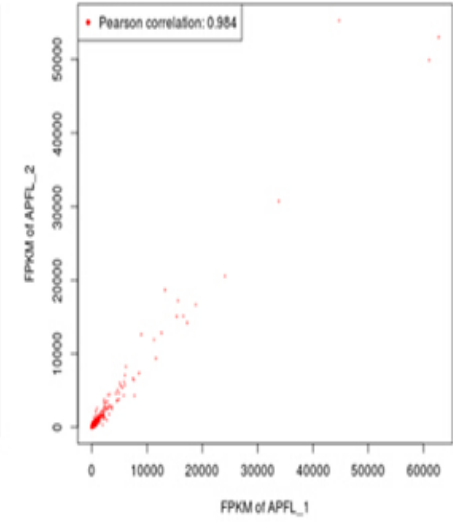
Sample FL1 and FL2 related analysis



Sample AP1 and AP2 related analysis



Sample APFL1 and APFL2 related analysis



Supplementary Table S1

Supplementary Table S1 Primers for commonly dysregulated genes involved in PPAR and lipid metabolism from mRNA expression profiling results for quantitative RT-PCR validation.	
Gene	Primer sequences(5'-3')
PPAR α	Forward GGCAATGCACTGAACATCGAG
	Reverse GCCGAATAGTTCGCCGAAAG
ACSL1	Forward TTCGCAGTGGCATCGTCAG
	Reverse TGTGATCATCAGCCGGACTTTC
CPT1A	Forward AGGTCGGAAGCCCATGTTGTA
	Reverse GCTGTCATGCGCTGGAAGTC
EHHADH	Forward CTCGCCAGAGCACATTGA
	Reverse ACCCAACTGAGGCAGCATAGAAC
ACAA1A	Forward GCATCTCACGGCAGAAGCAA
	Reverse TACGATCTCAGCACGGAAGCAG
ACADM	Forward GCGTGACAGAACCCTCAGCA
	Reverse TCCACGATGAATCCGGTGAA
ACADSB	Forward AATGCCCTATGTTTCGCACCTC
	Reverse TGCTTCAACTTCAATGCCCATC
ALDH1B1	Forward ACATGGGTCAGTGCTGCTGTG
	Reverse ACCTGAGGTCCCTGTTGAGTGTC
HADH	Forward AGCACAGACCTGGTGGTGGA
	Reverse AGACGTGTTGCTGGCAAAGATG