

# Distinct response of the hepatic transcriptome to Aflatoxin B<sub>1</sub> induced hepatocellular carcinogenesis and resistance in rats

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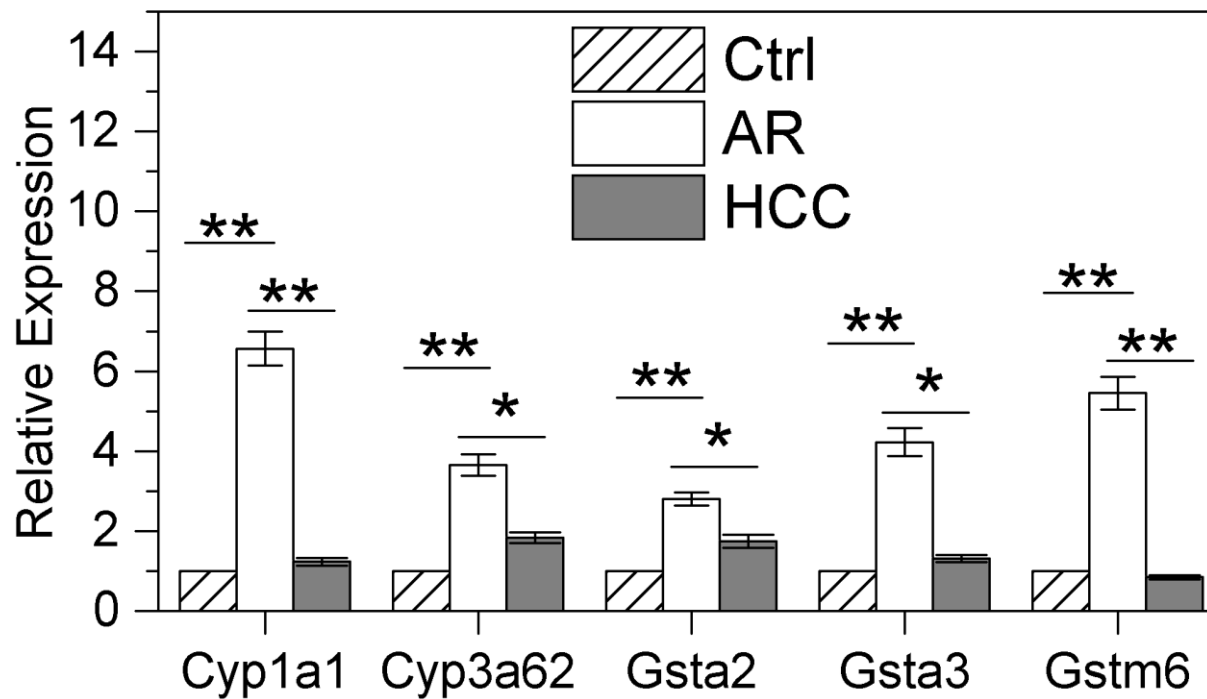
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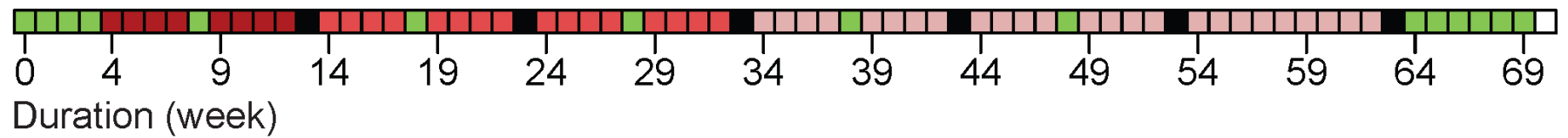
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**Supplementary Figure S1.** qPCR results of AFB1 metabolism related genes.

qPCR results confirm significantly increased expression level of AFB1 metabolism related genes in the AR sample (\* p-value < 0.05, \*\* p-value < 0.01, Student's t-test).



- Control diet(CD) only
- AFB<sub>1</sub>(200 µg/kg body weight, 3 times/week) + CD
- AFB<sub>1</sub>(100 µg/kg body weight, twice/week) + CD
- AFB<sub>1</sub>(100 µg/kg body weight, once/week) + CD
- Liver biopsy for HCC examination + CD
- All rats were sacrificed for sample collection.

Supplementary Figure S2. AFB<sub>1</sub> exposure strategy.

**Supplementary Table S1.** RNA-seq read count and alignment summary.

Samples	Hepatocellular carcinoma				AFB <sub>1</sub> resistance				Control			
	R1 reads		R2 reads		R1 reads		R2 reads		R1 reads		R2 reads	
	Count	%	Count	%	Count	%	Count	%	Count	%	Count	%
Total reads	29,767,084	100	29,767,084	100	33,333,334	100	33,333,334	100	33,333,334	100	33,333,334	100
Total mapped reads	26,983,873	91	26,995,184	91	30,392,438	91	30,410,670	91	30,144,262	90	30,134,074	90
Mapped concordant paired reads	26,022,103	87	26,022,103	87	29,424,474	88	29,424,474	88	28,993,816	87	28,993,816	87

**Supplementary Table S2.** Primers used for qPCR.

<b>Gene</b>	<b>Forward primer</b>	<b>Reverse primer</b>
Bcl2	5'-cagagatgtccagtcagctg-3'	5'-cagtcatccacagagcgatg-3'
MYO18A	5'-caagctcagctcgaagagtc-3'	5'-actggccatgctctccagac-3'
Nfkb1	5'-cagctgacagaagacacgag-3'	5'-tctgtgtagcccatctgtcg-3'
Mapk8	5'-gaagacagccatgagtgtg-3'	5'-ctgggtacctactcactggt-3'
Casp1	5'-tgtctcatggctccaggag-3'	5'-ttctctccacggcatgcctg-3'
Casp4	5'-gggttaatgtctcatggcac-3'	5'-gcaggcctgcacaatgatgac-3'
Il4	5'-gacagccctctgagagagatc-3'	5'-ccgtggtgttcctgttgcc-3'
Mpo	5'-agagctcaagcgctgaatc-3'	5'-ctgagtcattgtaggatcgg-3'
Bnip3	5'-cgtctgacaactccactag-3'	5'-cactaaatcaggaacaccgc-3'
Sstr3	5'-ttcaggtcagctacctagtc-3'	5'-tgctgcaaactgagcttagc-3'
Cyp1a1	5'-caacaggtcatgctaggatc-3'	5'-ggctttgcaaggacaaggag-3'
Cyp3a62	5'-actgatgtggagattgtggc-3'	5'-caggacatcataggtcacag-3'
Gsta2	5'-aaggacatgaaggagagagc-3'	5'-acaaggtagtcttgccatg-3'
Gsta3	5'-gttggcaacaagctgagcag-3'	5'-ttgctgactctggttctcag-3'
Gstm6	5'-cagtggtgctgatgacaaattc-3'	5'-atgtccacacgaatcctctc-3'