

# Single-cell transcriptomes reveal characteristic features of Mouse Hepatocytes with Cholestasis Injury

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Supplemental Table 2: Top10 gene among state (Related to Figure 3)

State	gene	symbol	log2Mean
1	ENSMUSG00000074115	Saa1	8.00
1	ENSMUSG00000057465	Saa2	7.93
1	ENSMUSG00000029368	Alb	6.94
1	ENSMUSG00000005681	Apoa2	6.77
1	ENSMUSG00000066154	Mup3	6.76
1	ENSMUSG00000031722	Hp	6.56
1	ENSMUSG00000030895	Hpx	6.38
1	ENSMUSG00000078672	Mup20	6.22
1	ENSMUSG00000032083	Apoa1	6.17
1	ENSMUSG00000002985	Apoe	6.08
2	ENSMUSG00000049775	Tmsb4x	7.21
2	ENSMUSG00000050708	Ftl1	7.03
2	ENSMUSG00000002985	Apoe	6.41
2	ENSMUSG00000024661	Fth1	6.18
2	ENSMUSG00000092341	Malat1	6.09
2	ENSMUSG00000064351	mt-Co1	5.85
2	ENSMUSG00000064357	mt-Atp6	5.73
2	ENSMUSG00000064358	mt-Co3	5.66
2	ENSMUSG00000029580	Actb	5.16
2	ENSMUSG00000034892	Rps29	5.16