

SUPPLEMENTARY TABLES

Supplementary Table 1. The top 10 of the most highly expressed genes in the three postnatal developmental stages of liver.

Group	Ensemble ID	Symbol	NH	YR		
No feeding group	ENSMAMEG00000023436	COX1	9756.00	147337.42		
	ENSMAMEG00000004932	ALB	61178.21	76816.95		
	ENSMAMEG00000023448	ND4	3683.22	81150.51		
	ENSMAMEG00000006318	None	28117.38	55300.09		
	ENSMAMEG00000006871	LOC100472013	5084.93	67570.66		
	ENSMAMEG00000023443	COX3	3508.02	64004.35		
	ENSMAMEG00000023455	CYTB	4054.62	58126.088		
	ENSMAMEG00000023442	ATP6	3188.95	58681.40		
	ENSMAMEG00000023439	COX2	2988.19	55884.78		
	ENSMAMEG00000023430	ND2	2216.17	42200.65		
Group	Ensemble ID	Symbol	AB	LT		
Suckling group	ENSMAMEG00000004932	ALB	136531.04	19953.67		
	ENSMAMEG00000023436	COX1	27751.54	111204.65		
	ENSMAMEG00000006871	LOC100472013	23118.74	91085.25		
	ENSMAMEG00000006318	None	56316.83	19755.60		
	ENSMAMEG00000023448	ND4	13905.71	47491.15		
	ENSMAMEG00000023442	ATP6	14956.36	41074.12		
	ENSMAMEG00000023455	CYTB	11770.93	40879.74		
	ENSMAMEG00000023443	COX3	12716.66	38222.35		
	ENSMAMEG00000023439	COX2	13989.96	36363.50		
	ENSMAMEG00000023430	ND2	10971.47	28860.23		
Group	Ensemble ID	Symbol	YS	HT	CC	PP
Adult group	ENSMAMEG00000004932	ALB	167548.34	76797.42	109881.46	245062.94
	ENSMAMEG00000002287	LOC100482523	88143.42	97980.01	86923.31	107729.87
	ENSMAMEG00000006318	None	27056.67	20188.85	16740.06	38674.56
	ENSMAMEG00000023442	ATP6	38376.79	21968.23	18248.36	12448.58
	ENSMAMEG00000008662	APOA1	27843.45	14730.88	12507.82	34044.19
	ENSMAMEG00000016713	None	21586.80	16322.76	20602.57	10035.15
	ENSMAMEG00000018459	FGB	11069.02	9978.05	21084.40	21685.78
	ENSMAMEG00000018463	FGG	9891.64	10553.21	22160.24	19838.73
	ENSMAMEG00000023448	ND4	26147.82	11195.78	6908.49	13117.63
	ENSMAMEG00000012556	LOC100474871	28393.64	1724.16	17656.30	8186.53