

### Supplementary Table 3

Zhao et al., 2009

#### Genomic location and position of hepatic miR-eQTL for 21 miRNAs.

miRNA	genome location				miR-eQTL		
	Chr	start bp	stop bp	strand	Chr	cM	LOD
miR-214	1	164153499	164153608	+	17	2.15	16.26
miR-34a	4	149442563	149442664	+	4	94.01	13.00 *
miR-451	11	77886672	77886743	+	9	74.30	9.96
miR-31	4	88556461	88556566	-	4	51.73	9.56 *
miR-338	11	119876079	119876176	-	6	46.495	7.42
miR-145	18	61807479	61807548	-	17	30.55	6.96
miR-205	1	195333657	195333724	-	7	0.00	6.57
miR-200b	4	155429790	155429859	-	12	7.27	6.36
let-7c-1	16	77599902	77599995	+	8	17.61	6.02 #
let-7c-2	15	85537033	85537127	+			
miR-295	7	3220774	3220842	+	7	0.00	5.25 *
miR-142-3p	11	87570366	87570429	+	9	56.81	6.57
miR-148a	6	51219811	51219909	-	12	17.97	5.04
let-7b	15	85537749	85537833	+	8	17.36	5.74
miR-16-1	14	62250717	62250809	-	12	16.95	5.95 #
miR-16-2	3	68813824	68813918	+			
miR-331	10	93426513	93426608	-	12	7.92	5.82
miR-141	6	124667932	124668003	-	12	17.97	5.68
miR-346	14	35707795	35707892	+	9	79.23	5.46
miR-190	9	67084467	67084533	-	8	57.45	5.44
miR-181c	8	86702772	86702860	-	12	11.14	5.40
miR-195	11	70048544	70048637	+	12	20.00	5.36
miR-143	18	61808850	61808912	-	5	18.47	5.31

Notes:

\* indicates cis-acting regulation; other miRNAs were trans-acting.

# indicates mature miRNAs that potentially derive from two different genomic locations.