

## **Dietary supplementation with hybrid palm oil alters liver function in the common Marmoset**

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## **Supplementary Information**

**Table S1.** GO analysis of `predicted genes modulated by hybrid palm oil-modulated miRNAs

**Supplementary Figure S1:** Body weight progression (in grams) of the experimental groups during the nutritional intervention expressed as the mean of body weight  $\pm$  standard error, (n=10 animals per group). No significant differences were found between the groups according to t test at  $p<0.05$ . AP: African Palm Group; HP: Hybrid Palm Group.

**Supplementary Figure S2:** Lipoprotein cholesterol and triglyceride profiles during dietary supplementation. (n=9 animals per group). AP: African Palm Group; HP: Hybrid Palm Group. A: AP group at 0, 1 and 3 months of dietary treatment, respectively. B: HP group at 0, 1 and 3 months of dietary treatment, respectively. VLDL: very low-density lipoproteins; LDL: low density lipoproteins; HDL: high density lipoproteins.

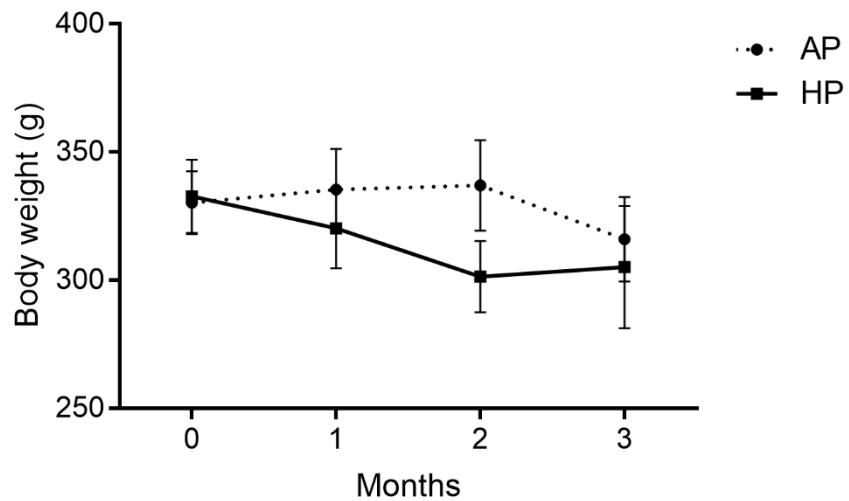
**Supplementary Figure S3.** Liver miRNAs validation by RT-qPCR. Hepatic miRNAs were analysed by RT-qPCR in both sexes. AP: African Palm Group; HP: Hybrid Palm Group. Values are expressed as mean  $\pm$  standard error (n=9 animals per group).

**Table S1.** GO analysis of `predicted genes modulated by hybrid palm oil-modulated miRNAs

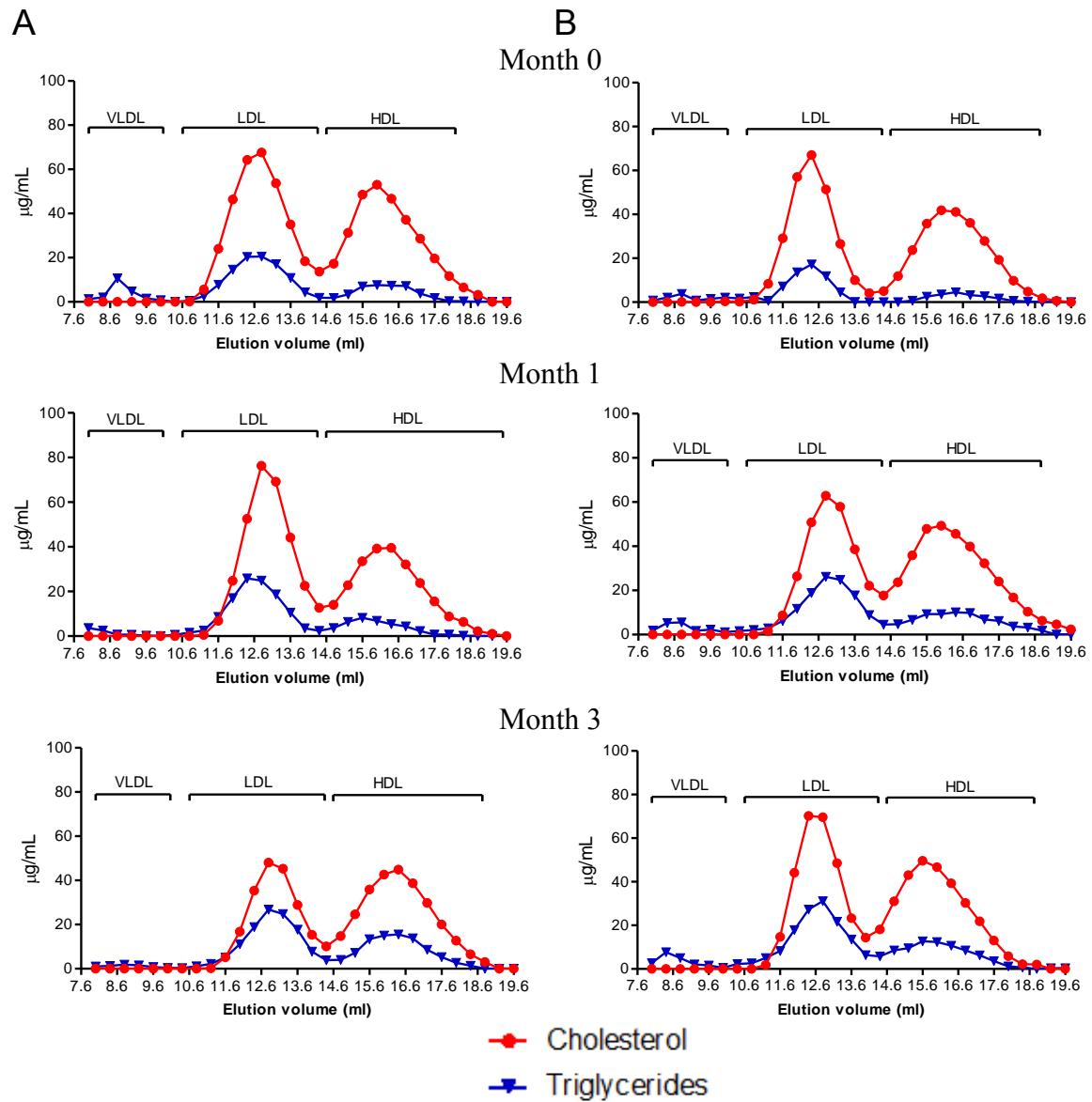
| ID              | Description  | Adjusted P value | Nº of genes | Genes  | microRNA's  |
|-----------------|--|------------------|-------------|--|---|
| Panther: P00044 | Nicotinic acetylcholine receptor signaling pathway | 0.00885829       | 18          | Cacnb1, Chrng, Myh11, Myo5c, Stx1a, Vamp1, Cauna1c, Cauna1d, Chrb3, Myo18a, Myo3a, Acta1, Snap25, Actc1, Actr1b, Chrb2, Vamp2, Vamp8   | calJac-let7b-5p, calJac-miR17b-5p-1, calJac-miR25b-3p, calJac-miR484-1, calJac-miR488-3p1, calJac-miR592, calJac-miR877-5p                                      |
| Kegg: 04514     | Cell adhesion molecules (CAMs)                     | 0.000000917      | 24          | Alcam, Cd8b, Cldn16, Cldn18, Itgb8, Sdc2, Vcan, Cd4, Cldn11, Cldn15, Cldn19, Hla-Doa, Itga9, Itgb7, Ncam1, Sdc1, Cd8a, Cdh3, Itgav, Madcam1, Mpz1l, Ocln, Sell, Siglec1  | calJac-miR17b-5p-1, calJac-miR25b-3p, calJac-miR4745-5p, calJac-miR484-1, calJac-miR488-3p1, calJac-miR877-5p   |
| Kegg: 04146     | Peroxisome   | 0.0000909        | 12          | Acsl4, Acsl6, Crot, Pecr, Pex10, Gstk1, Hacl1, Hsd17b4, Paox, Pex11b, Pex26, Pex5  | calJac-let7b-5p, calJac-miR17b-5p-1, calJac-miR339-3p, calJac-miR488-3p1  |
| GO: 0031175     | neuron projection development                      | 0.00140149       | 10          | Cdk5, Cdk5r1, Cntn2, Igf1r, Lamb1, Lingo1, Tbc1d24, Cdh1, Lamc1, Rb1   | calJac-let7b-5p, calJac-miR17b-5p-1, calJac-miR484-1, calJac-miR488-3p1, calJac-miR592, calJac-miR877-5p  |
| GO: 0030154     | cell differentiation                               | 0.0662095        | 109         | Cbfa2t3, Clic4, Cplx2, Dusp6, Fgf23, Gnptab, Igfs9, Nme1, Ptpru, Rasgrp1, Robo2, Ascl2, Bmpr1b, Cadm1, Dclk1, Dll1, Dll4, Dmrt1, Ereg, Fcrla, Kif2a, L1cam, Mapk7, Pappa, Pkdcc, Racgap1, Rnf17, Robo1, Rufy3, Sema3a, Sema3d, Sh2d2a, Siah1, Srpk1, Stmn1, Ttll7, Usp42, Abhd5, Angpt2, Arhgap24, Bmp1, Bzw2, Camk2g, Caprin1, Caprin2, Col19a1, Csf1, Fgf1, Ggnbp2, Gldn, Grb2, If1a, Ift81, Jmjd6, Mgp, Mkl2, Nfatc4, Piwil2, Sema4b, Sema4g, Sema6d, Sort1, Sqstm1, Strbp, Suv39h2, Utp14c, Acsbg2, Chl1, Osr1, Clptm1, Dbn1, Fev, Hnf4a, Huwe1, Ndel1, Ntng1, Ppard, Slc9a1, Smurf1, Spata20, Tnfrsf12a, Asz1, Bmpr1a, Chrd1, Cit, Dmrt3, Epas1, Gna12, | calJac-let7b-5p, calJac-miR17b-5p-1, calJac-miR25b-3p, calJac-miR339-3p, calJac-miR4745-5p, calJac-miR484-1, calJac-miR488-3p1, calJac-miR592, calJac-miR877-5p |

|  |  |  |  |   |  |
|--|--|--|--|---|--|
|  |  |  |  | Myt1l, Naa15, Nhs, Nrp2,<br>Pafah1b1, Pax5, Pax6,<br>Rbm38, Rnf114, Slc26a8,<br>Tnfsf11, Tp53, Dcx,<br>Dhcr7, Dmrtc2, Hand2,<br>Ngef, Paqr5, Sirt1,<br>Tgfb1i1, Vtn |  |
|--|--|--|--|---|--|

### Supplementary Figure S1



## Supplementary Figure S2



**Supplementary Figure S3**

