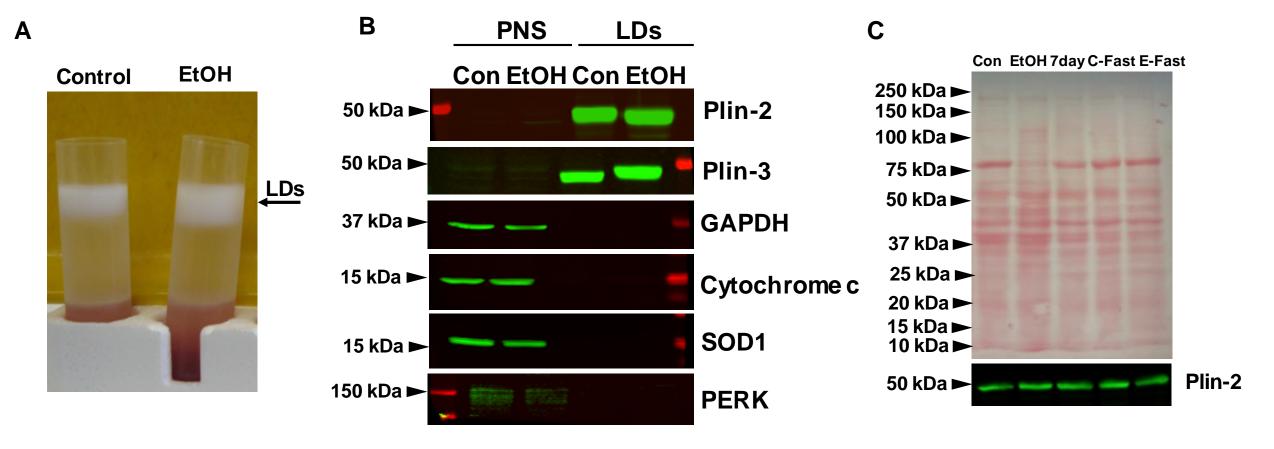
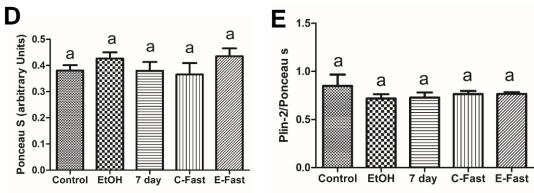
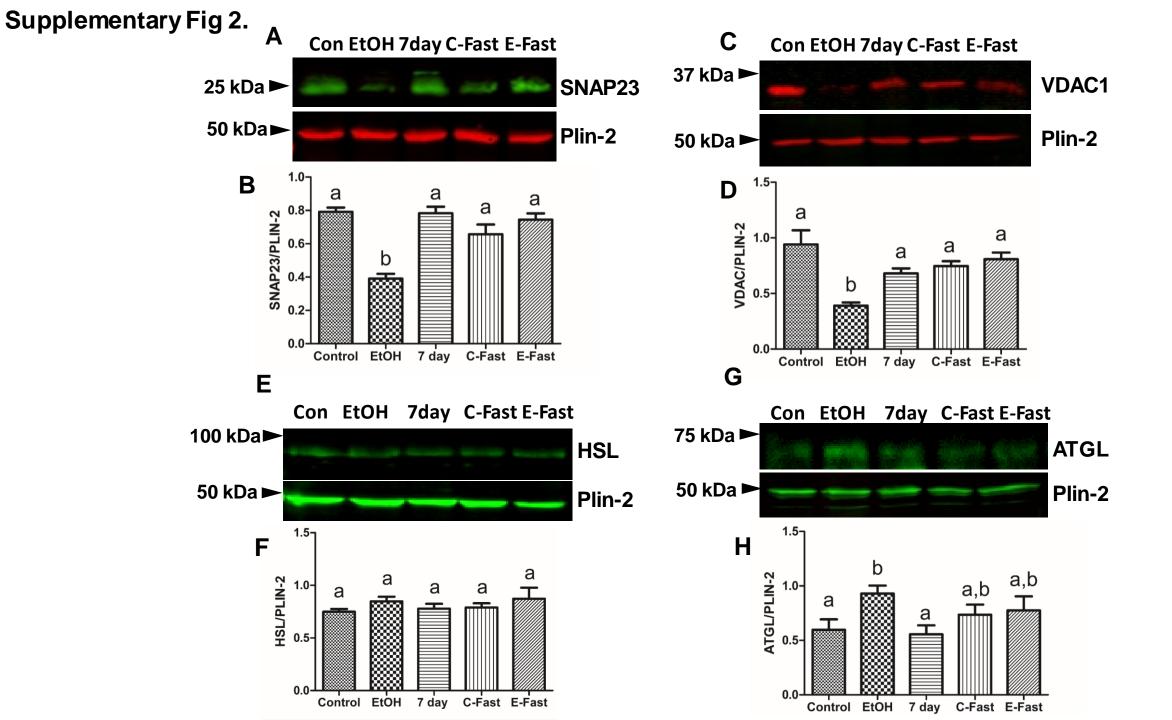
Supplementary Fig 1.





S Fig 1. Purification and quantification of lipid droplet membrane proteins. (A) Representative image showing buoyant lipid droplets (LDs) after gradient centrifugation of post nuclear supernatants (PNS) obtained from the livers of control and ethanol-fed rats (B) Representative Western blot show absence of common contaminating proteins such as cytosol (GAPDH), mitochondria (Cytochrome C and Super Oxide dismutase (SOD1) and endoplasmic reticulum (protein kinase RNA-like endoplasmic reticulum kinase (PERK)) in LDs isolated from control and ethanol-fed groups (C) ponceau S staining and Western blotting show protein bands and PLIN-2 levels, respectively, in equal protein loaded LD fractions from different groups (D) densitometry quantification of ponceau S staining and (E) PLIN-2 levels after normalization with ponceau S staining in each group. Data are mean values of ± S.E.M. of 4 to 6 animals per group. Bars sharing different letters are significantly different. Bars sharing the same letter are not significantly different, P≤0.05.



S Fig 2. Expression of select lipid droplet membrane proteins. (A) Representative Western blot and densitometry quantification of **(A&B)** Synaptosomal-associated protein 23 (SNAP23) **(C&D)** Voltage-dependent anion channel (VDAC1) (**E&F)** Hormone sensitive lipase (HSL) and **(G&H)** Adipose tissue triglyceride lipase (ATGL) in LDs from different experimental groups. Data are mean values of \pm S.E.M. of 4 to 6 animals per group. Bars sharing different letters are significantly different. Bars sharing the same letter are not significantly different, P≤0.05.