

Fig. S1 Scores plots of PLS-DA models generated to maximise the difference between TRAMP and wild-type mice at **(A)** 8 and **(B)** 29 weeks of age regardless treatment. The groups are coloured according to genotype and treatment. The data are scaled to unit variance.

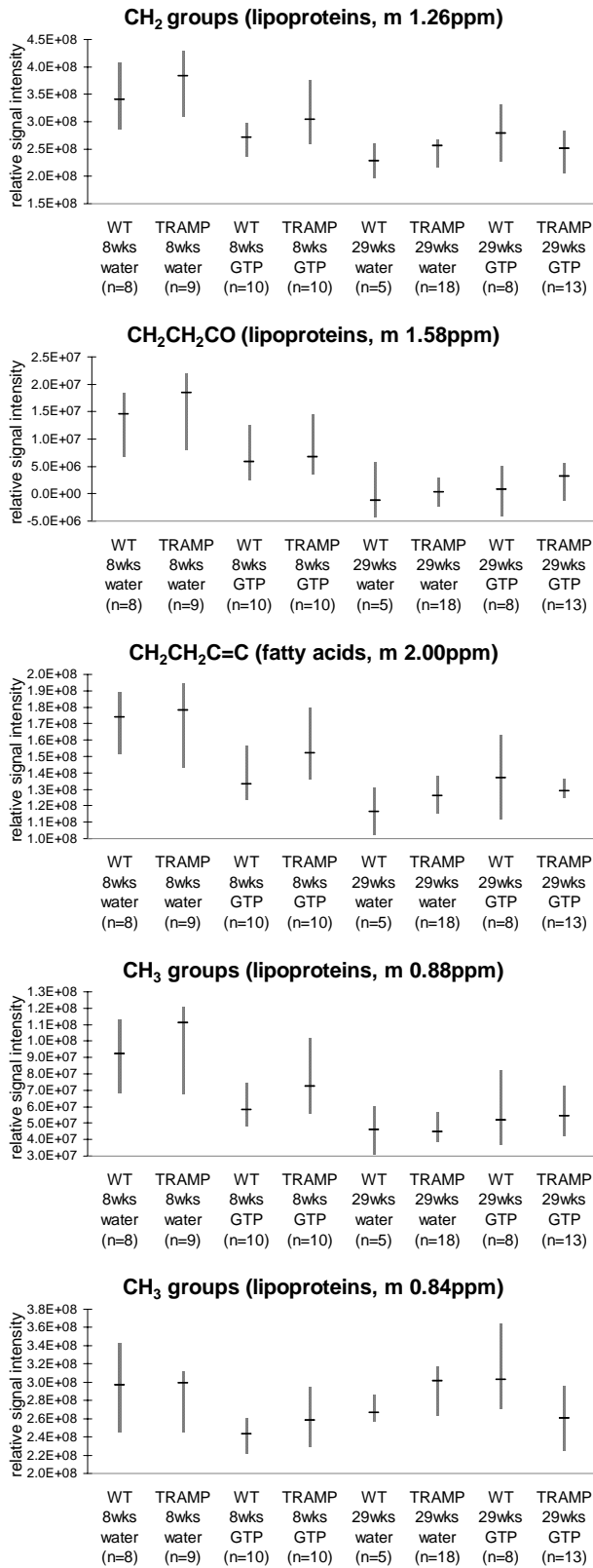


Fig. S2 Median values of lipid signals in individual groups (bars). The columns represent values between the 25. and 75. Percentile; WT = wild-type