## Supplemental Information

## Novel CB1-ligands maintain homeostasis of the endocannabinoid-system in $\omega 3$ - and $\omega 6$ long chain-PUFA deficiency

Ina Hammels*,**, Erika Binczek*, Inga Schmidt-Soltau*, Britta Jenke*, Andreas Thomas $\dagger$, Matthias Vogel $\dagger$, Mario Thevis $\dagger$, Dilyana Filipova $\dagger \dagger$, Symeon Papadopoulos $\dagger \dagger$ and Wilhelm Stoffel ${ }^{*}$, ${ }^{* * 1}$

* Center of Molecular Medicine (CMMC), Laboratory of Molecular Neurosciences, Institute of Biochemistry, University of Cologne, Joseph-Stelzmann-Strasse 52, 50931 Cologne, Germany
** CECAD (Cluster of Excellence, Cellular Stress Response in Aging Related Diseases), University of Cologne, 50931 Cologne, Germany
$\dagger$ Institute of Biochemistry, Deutsche Sporthochschule (DSHS) Cologne, 50933 Cologne, Germany
$\dagger \dagger$ Institute of Vegetative Physiology, Center of Physiology and Pathophysiology, University of Cologne, 50931 Cologne, Germany

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Supplemental Figure S1 Analysis of the phospholipidomes of liver and serum, and kinetics of the PUFA pattern of liver. A, B: GC/MS analysis of the dynamics of the PUFApattern of liver-phospholipids of WT and fads2-/- mice on normal diet (nd). PUFA substitution of phospholipid-classes PC, PS, PI and PE in liver of (A) WT and (B) fads2-/mice. Tracings: 18:2, 20:3, 20:4 and 22:6 at 10, 25, 40, 55, 70 and 120 days. C-E: MS/MS analysis of lipid-classes of serum of cohorts of WT and fads2-\% (4mo) mice on (C) normal diet (nd) (WT: black bars, fads2-/-: red bars), (D) $\omega 6$-AA-diet (WT: black bars, fads2-/-: green bars) and (E) $\omega 3$-DHA-diet (WT: black bars, fads2-/-: blue bars). A pool of $\mathrm{n}=3$ per genotype was used.


Supplemental Figure S2: Novel endocannabinoids 20:3 ${ }^{5,11,14}$-(sciadonoyl)-ethanolamide (SEA) and 2-20:3, ${ }^{5,11,14}$-(sciadonoyl)-glycerol (2-SG) in the endocannabinoid pattern of serum of nd-fads2-/- mice. A-C: Separation and quantification of endocannabinoids in lipid extracts of (A) nd- (WT: black bars, fads2-/-: red bars), (B) $\omega 6$-AA- (WT: black bars, fads2-/: green bars) and (C) $\omega 3$-DHA-WT and -fads $2-/$ mice ( 4 mo ) (WT: black bars, fads2-/-: blue bars) by HPLC-MS/MS. Data represent mean $\pm$ SEM. Two-tailed Student's $t$-test, $p$-values $\leq$ $0.05^{*}, \leq 0.01^{* *}, \leq 0.001 * * *$ were considered significant. N=8 per genotype. Sciadonoylethanolamide (SEA), 2-sciadonoyl-glycerol (2-SG), palmitoyl-ethanolamide (PEA), oleoylethanolamide (OEA), 2-oleoyl-glycerol (2-OG), linoyl-ethanolamide (LEA), 2-linoyl-glycerol (2-LG), arachidonoyl-ethanolamide (AEA), 2-arachidonoyl-glycerol (2-AG), docosahexanoyl-ethanolamide (DHEA), 2-docosahexanoyl-glycerol (2-DG)
A

B

C

D

E

F


H
$100\left[\mathbf{1 -}^{\left.1^{3} \mathrm{C}\right]-\mathbf{2 0 : 3} \mathbf{3}^{5,11,14}}\right.$
J.

Supplemental Figure S3: Structural analysis of novel endocannabinoids isolated from brain of nd-fads2-/- mice. A-J: LC-MS/MS of extracted ion chromatograms (XIC) (A, B) of SEA, transition 350.3-62.06 Da, and TOF product 350.3 at 5.6099 to 5.6831 min ; (C, D) 2SG, transition 381.3-289.3 Da, and TOF product 381.3 at 6.1362 to 6.2217 min . Structures of (E) $\omega 6-20: 3^{5,11,14}$-EA, (F) $\omega 6-20: 4^{5,8,11,14}$-EA, (G) $\omega 6-20: 3^{5,11,14}$-G and (H) $\omega 6-20: 4^{5,8,11,14}$-G. Characterization of synthetic (I) $\left[1-{ }^{13} \mathrm{C}\right]$ - and (J) [D6]-labeled $\omega 6-20: 3^{5,11,14}$.

The chemical structures and analytical data of the novel endocannabinoids, isolated and analyzed by LC-MS/MS from nd-fads2-/- mice (Supplemental Figure S3A-D), proved identical with that of the synthetic unlabeled and $\left[1-\mathrm{C}^{13}\right]$ - and [D6]-labelled $\omega 6-20: 3^{5,11,14}$-EA and $\omega 6-20: 3^{5,11,14}$-G (Supplemental Figure S3E-F).


Supplemental Figure S4: Characterization of the mock and $\boldsymbol{c b 1}$-transfected HEK293cells. A: PCR of cannabinoid receptor 1 (cb1) and glycerinaldehyd-3-phosphatdehydrogenase (gapdh) of cRNA, B: Western blot of anti-CB1 and anti-GAPDH of protein lysates and C: IHC localization of CB1 and EGFP of mock (m) and cbl-transfected (clone \#5) HEK293-cells.


[^0]:    ${ }^{1}$ To whom correspondence should be addressed. E-mail: wilhelm.stoffel@uni-koeln.de

