

**High-fat diet at adulthood interacts with prenatal stress,
affecting both brain inflammatory and neuroendocrine
markers in male rats**

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Response to Editor

1) Author contributions need to be explicitly included in a statement in the manuscript itself

Done

2) Provide an abbreviations list

Done

3) Provide a clear data sharing statement, and we strongly encourage that you place your data on a public access repository (e.g. Figshare).

Done

4) When revising the manuscript, please **bolden** or underline major changes to the text so they are easily identifiable and DO NOT leave 'track change' formatting marks in your paper.

Done

5) Please ensure that you provide a text and a figure file for the **Graphical Abstract** (as detailed in the instructions below).

Done

6) When carrying out your revisions please refer to the **checklist below**.

Done

Response to reviewers

Reviewer #1

In this paper, Berry et al examine the effects of prenatal stress and the effects of a second-adult challenge (HFD) on the inflammatory and neural plasticity phenotype of adult offspring using gene expression profiles. They conclude that overall PNS decreases the pro-inflammatory phenotype and likewise decreases neuronal plasticity. By contrast, a HFD increases the vulnerability of developing a pro-inflammatory phenotype. In general, I believe that the methods were performed properly and that there was good discussion explaining the possible implications of their findings. However, there were many issues with the paper in its current form, with my biggest hesitation around over-interpretation of the results:

Major concerns:

1) I don't believe many of their conclusions are warranted based on the data they present. For example, they conclude that PNS reduces brain plasticity setting the stage for increased vulnerability for further insults, with a HFD promoting a pro-inflammatory phenotype. However, the data doesn't show this. In the PNS-HFD group, the proinflammatory markers are never significantly different from the Control-CD group. If anything, the HFD doesn't increase the pro-inflammatory phenotype it rather resets it to control levels. I think these conclusions should be toned down to reflect the data more accurately. Also, the title should be changed to reflect this.

We thank the reviewer for pointing out this issue. The title has been changed as follows: “Prenatal stress interacts with an adult metabolic challenge affecting brain inflammatory and neuroendocrine markers in male rats”.

This sentence in the abstract “HFD unveiled a pro-inflammatory phenotype by inducing a specific increase in the above-mentioned cytokines, suggesting that PNS might impinge upon the same mechanisms underlying vulnerability to metabolic challenges at adulthood” has been changed as follows: “Overall, PNS reduced the expression of Bdnf and Tnf- α while HFD administered at adulthood counteracted this effect suggesting that PNS impinges upon the same pathways regulating responses to a metabolic challenge at adulthood”.

The Discussion and the Concluding remarks have been amended by toning down the effect of HFD on the pro-inflammatory phenotype of the PNS rats. See below.

“...HFD feeding, experienced at adulthood, was able to unmask a pro-inflammatory phenotype, inducing an increase in many pro-inflammatory cytokines in those animals that had experienced PNS.” This sentence has been changed as follows: “...moreover, HFD feeding, experienced at adulthood, induced an increase in pro-inflammatory cytokines in those animals that had experienced PNS.”

“Therefore, we can hypothesise that the increase in the pro-inflammatory cytokines observed as assessed in the prefrontal cortex (Il-1b and Tnf-a)...”.

This sentence has been changed as follows: “Here we observed a similar blunted activation upon PNS with HFD triggering a response only in PNS subjects (increased Il-1 β and Tnf- α in the prefrontal cortex and Tnf- α in the dorsal and ventral hippocampus).”

“We have here provided evidence for PNS to reduce brain plasticity (first hit) setting the stage for increased vulnerability to further insults during life (HFD, second hit) promoting a pro-inflammatory phenotype”.

This sentence has been changed as follows: “We have here provided evidence for PNS to reduce brain plasticity (first hit) setting the stage for increased vulnerability to further insults during life (HFD, second hit).”

2) The data relating to ghrelin signaling seems out of place. There is no mention of ghrelin signaling or its significance in the introduction. I think there should be more context/rationale mentioned as to why the authors chose to specifically examine ghrelin in the introduction. Ghrelin is not a unique link connecting mood disorders (HPA axis function) and metabolic disorders. In fact, there are a plethora of neuroendocrine signals connecting the two pathophysiology, including leptin, FGF21, endocannabinoids, etc. It seems out of context to focus in on ghrelin without providing further context.

We thank the reviewer for the thorough suggestion, we have added the following at the end of the Introduction paragraph

“We specifically assessed in these brain regions levels of Bdnf, Tnf- α , Il-6 and Il- β ”

Stress powerfully affects both mood and energy homeostasis. Such effects are achieved not only through the interaction between GC hormones and their receptors (that regulate HPA axis function in addition to glucose metabolism) but also by triggering a multitude of signalling cascades reciprocally modulating one another in a regional-, temporal, and functional-dependent manner (Balsevich et al., 2019). Among these, ghrelin is a peptide hormone produced in the stomach and involved in the signalling of meal initiation. Its action is mediated through the growth hormone secretagogue receptor (Ghs-R) that is found to be highly expressed in several brain regions including the hypothalamus, pituitary gland and hippocampus (see (Zarouna, 2015) and references therein). Both Ghrelin and Ghs-R are modulated by stress (Patterson et al., 2010); there is evidence that ghrelin may decrease anxiety-like and depressive-like behaviours in mice (Lutter et al., 2008) moreover, it appears to be involved in food anticipatory activity and consumption (Blum et al., 2009; Verhagen et al., 2011) as well as in mood disorders (Zarouna, 2015). Interestingly, both genetic variants within ghrelin and GC signalling pathways have been associated with obesity, stress-related mental disorders, or both (see (Balsevich et al., 2019) and references therein). To this regard, we also investigated changes in the expression levels of Ghs-R as well as of Nr3c1 (encoding for the glucocorticoid receptors - GR) and the GC co-chaperon Fkbp5 (encoding for the FK506 binding protein 51) in all the above-mentioned brain regions.”

Moreover we have added in the concluding remarks the following sentence “Further investigation are thus warranted to assess more in detail the role played by these molecules in setting the stage for co-morbidity between metabolic and psychiatric disorders. In addition, the role of additional co-regulators, such as endocannabinoids should be tested (Balsevich, 2019).

3) Although the authors provide sufficient explanation as to why PNS decreases a pro-inflammatory phenotype in the discussion, I still believe that this largely disagrees with previously published data (typically PNS and ELS increase pro-inflammatory phenotypes and risk for mood disorders). I think these conflicting results should be discussed more.

We thank the reviewer for the suggestion and the following has been added to the Discussion paragraph: “Worth to notice, we have previously shown that upon PNS, male rats were characterized by increased corticosterone levels under basal conditions, this effect was associated with a decrease in reactive oxygen species (ROS) as well as a with decreased NF-kB signalling in the hippocampus suggesting a lower set-point under basal conditions in PNS male rats (Anacker et al., 2013). Here we observed a similar blunted activation upon PNS with HFD triggering a response only in PNS subjects (increased Il-1 β and Tnf- α in the prefrontal cortex and Tnf- α in the dorsal and ventral hippocampus). To our knowledge this is one of the first times that such a reduction in inflammatory mediators is described a result of PNS. Because cytokines have been shown to modulate hippocampal development and plasticity (Bourgognon and Cavanagh, 2020; Goshen et al., 2007) a decreased inflammatory profile as a result of a “first hit” (PNS) might indeed set the stage for an increased response to a “second hit” (metabolic challenge).”

4) The study used a candidate approach for both brain regions examined as well as genes examined. Both seemed somewhat arbitrary and not inclusive. For example, why not assess the amygdala, which plays a critical role in the stress response, emotionality, and mood disorders?

We thank the reviewer for the thorough question. Indeed, the aim of this study was to investigate the contribution of prenatal stress to increase the vulnerability to later life metabolic and mood disorders. To this regard, we focused on specific brain regions such as the prefrontal cortex, hippocampus and hypothalamus that are well-known to play a role in mood disorders and that also integrate mood and metabolic signals (hippocampus and hypothalamus). We agree with this reviewer that the amygdala is a very interesting region to study and future research will also try to focus on this area. As far as gene expression is concerned also in this case, we focused on those signalling cascades known to be deeply involved in the modulation of mood and metabolism.

Further, the discussion does mention why GR and Fkbp5 were selected for examination. However, there are major gaps in interpreting these findings. How do changes in either GR or Fkbp5 affect GR sensitivity? The authors should examine circulating CORT levels under baseline or after a challenge or perform a DEX/CRH test to assess whether GR sensitivity is altered. I understand that this may not be possible at this point, however given the direction of changes across different brain regions, it is difficult to determine the significance. With that said, the authors do discuss the possible implications in the discussion. But they don't adequately discuss how FKBP5/GR signaling relates to metabolic vulnerability. think the rationale for selecting these candidates should again be stated in the introduction.

*As suggested by this reviewer, we have introduced the rationale for studying FKBP5/GR signalling in the Introduction paragraph (see above answer to Q2). With regard to HPA axis reactivity in relation to changes in expression levels of Fkbp5 and GR sensitivity, the Discussion paragraph already reported that we have previously shown that PNS male rats were characterised by elevated GC levels when compared to controls (Anacker et al., 2013); this issue has been further strengthened (see also answer to Q3). Moreover, we have now provided evidence that levels of Fkbp5 are positively associated to the time spent in the open arms of the EPM (see Figure 6). **“Interestingly, we also showed that levels of Fkbp5 in the dorsal hippocampus were positively related to the time spent in the open arm of the EPM and that this positive correlation is lost as a result both of PNS as well as of HFD. This result might suggest that under physiological conditions Fkbp5 is required to engage animals in the proper exploration of novel environments through a fine modulation of the GR receptors and that this balance can be greatly affected by both early life stressors as well as by metabolic challenges.”** This sentence has been added to the Discussion paragraph.*

We also added further consideration with regard to this issue in the Concluding remarks (see below)

“Thirdly, genomic and epigenomic regulation of glucocorticoids in the brain may affect mood and metabolism but is based upon the coordinated activity of many chaperone proteins, in addition to proper circadian and ultradian fluctuation of hormone release (Gray et al., 2017).”

5) The correlation graphs between immobility in the EPM and gene expression data should not be included in this study. It is not possible to effectively evaluate the methodology here when the EPM was not included in this study but in a previously published study. Moreover, why did they just correlate to immobility in the EPM? In general time spent in the open and closed arms are the major readouts. Is this the only parameter that significantly correlated? I think this data should be removed.

We thank the reviewer for pointing out this issue. Following this suggestion, and suggestions from reviewer 2, we have briefly added the EPM methodology to the text and carried out a regression analysis between Fkbp5 levels in the dorsal hippocampus and time spent in the open arm of the EPM. We believe that the regression analyses reported in this paper are very important to strengthen data deriving from the analysis of gene expression in the brain area investigated and to characterise the effects of changes in the selected genes. Furthermore, we would like to stress that this paper and Panetta et al. 2017 should be considered as closely related publications since they refer to the same experimental subjects and they complement one-another.

6) The GR and FKBP5 gene expression data should be represented in the form of a graph (not table) like all other data. It is much easier to visualize/interpret graphs than figures

Done

7) The hypothalamus gene expression data should also be included in the graphs, despite many regions showing no change in gene expression from PNS or HFD in the hypothalamus, there were a few cases where there was a change in the hypothalamus (i.e. Fkbp5). Regardless, I think the data should be graphically represented.

Done

8) In the methods, they mention the groups including both males and females. In this study and with the specific research questions, only males are relevant. I think it is confusing to mention females. It only raises questions. I would omit methods related to females. Also, in lines 131-135, they mention methods for a previous paper. This should not be included here. It can be discussed in the discussion in terms of relevance to the current study.

We thank this reviewer for the suggestion; the method section has been modified by removing details about female rats as well as those concerning gene expression levels previously assessed; see below:

“By this time, n=20 CTRL and n=21 PNS males and n=20 CTRL and n=19 PNS females were assigned either to high-fat diet (HFD) or control diet (CD) regimen for 8 weeks. The final number of animals per group was: n=10 CTRL -CD, n=10 CTRL -HFD, n=10 PNS-CD and n=11 PNS-HFD for the male offspring; n=10 CTRL -CD, n=10 CTRL -HFD, n=10 PNS-HFD and n=9 PNS-CD for the female offspring. After 4 weeks on the respective diets, all animals underwent a number of metabolic and behavioural assessments; they were successively sacrificed, brain were dissected out and the ventral hippocampus used to assess Bdnf long 3'UTR, hypothalamus used to assess levels of, LepR, Adipo-R1 and Adipo-R2; total Bdnf expression was assessed in both areas. Results from the above-mentioned studies have been recently published in Panetta et al. 2017. Here we investigated selectively in male animals' brains the expression levels of Il-6, Il-1 β , Tnf- α , Nr3c1, Fkbp5, Ghs-R, in the dorsal hippocampus, ventral hippocampus, hypothalamus and prefrontal cortex; total Bdnf was investigated only in the dorsal hippocampus and prefrontal cortex (since it was already investigated in the other two areas, (Panetta et al., 2017).”

This paragraph has been modified as follows: “By this time, n=20 CTRL and n=21 PNS males were assigned either to high-fat diet (HFD) or control diet (CD) regimen for 8 weeks. The final number of animals per group was: n=10 CTRL -CD, n=10 CTRL -HFD, n=10 PNS-CD and n=11 PNS-HFD. After 4 weeks on the respective diets, all animals underwent a number of metabolic and behavioural assessments; they were successively sacrificed and brains dissected out to investigate the expression levels of Il-6, Il-1 β , Tnf- α , Nr3c1, Fkbp5, Ghs-R, in the dorsal hippocampus, ventral hippocampus, hypothalamus and prefrontal cortex; total Bdnf was investigated only in the dorsal hippocampus and prefrontal cortex (since it was already investigated in the other two areas, (Panetta et al., 2017).”

9) Lines 397-398: Chronic CORT or DEX decreases Fkbp5 methylation and increases FKBP5 expression in mouse models of both HFD and stress (Scharf et al. 2011). Scharf et al. 2011 does NOT show this. They show FKBP5 expression in brain regions after 24h food deprivation or 1x dexamethasone treatment. Correct references should be provided or this statement should be corrected.

We thank this reviewer for pointing out this issue. The sentence, and the associated reference, have been removed from the text.

10) Lines 407-410: “the hippocampal increase in Fkbp5 as a result of HFD might suggest that this metabolic challenge (second hit) might impact cognitive functions in our animal model. This piece of data is further strengthened by a similar increase in the levels of Nr3c1 in the ventral hippocampus of all HFD rats.” Perhaps it is my interpretation, but does this make sense? If you have increase Fkbp5, you in theory decrease GR sensitivity. Therefore, wouldn't the increase in GR offset the increase in FKBP5?

We thank the reviewer for the thorough remarks. Regulation of glucocorticoids is a process relying on the intricate coordination of many chaperone proteins in addition to proper circadian and ultradian fluctuation of hormone release. We have changed the sentence to more clearly indicate that a cross-regulation between a metabolic challenge and Nr3c1/Fkbp5 appears from these data, strengthening the notion that indeed these mediators are involved in “sensing” environmental challenges, be these of psychological nature or metabolic.

“Changes in Nr3c1 and Fkbp5 in the hippocampus a result of HFD indicate a cross-regulation between a metabolic challenge and Nr3c1/Fkbp5 balance, strengthening the notion that indeed these molecules are involved in “sensing” environmental challenges, be these of psychological or metabolic nature”.

We have also added the following sentence to the concluding remarks (see also above Q4). “Thirdly, genomic and epigenomic regulation of glucocorticoids in the brain may affect mood and metabolism but is based upon the coordinated activity of many chaperone proteins, in addition to proper circadian and ultradian fluctuation of hormone release (Gray et al. 2017). Thus, additional analyses should be performed in future studies to assess the interrelationship between psychological and metabolic stressors. In addition, given the observed changes of Ghs-R and Nr3c1 in the dorsal hippocampus following a HFD, future studies should be devoted to better explore the effects of HFD and of PNS, and their interaction, on brain plasticity”.

Minor concerns:

1) For the assignment of pups to CD/HFD and control/PNS groups, how many pups from each litter were represented in each group? This information should be provided given it is best practice to have only 1-2 pups from each litter in each group.

“Groups were composed by 1 pup/litter (CTRL-CD; CTRL-HFD; PNS-CD; PNS-HFD)”.

This sentence has been added to the methods section.

2) Typo, line 108: Thus, the aim of this study.....

Done

3) Typo, line 158: ...offspring were weighed.....and brain were removed.....

Done

4) Line 196: post hoc test just missed statistical significance: ADD P-value

The sentence has been rephrased to make it clearer “A strong PNS x HFD interaction was found ($F(1,34)=4.655$; $p=0.0381$) showing that PNS exposed animals were characterised by decreased levels of this cytokine, while the administration of postnatal HFD counteracted the effects of the PNS (post-hoc comparisons just missed statistical significance).”

5) Lines 199-202: A significant PNSxHFD interaction was found. But this wasn't the case ($p = 0.0580$). Maybe change statistical analysis so that statistical significance is set at 0.05; statistical tendency is set at $p < 0.1$. For interactions at $p < 0.1$, we also examine lower order main effects.

We thank the reviewer for the suggestion; the following sentence has been added to the Statistics paragraph "Statistical tendency was set at $p < 0.1$. For interactions at $p < 0.1$, we also examined lower order effects". Moreover, we have changed the word "significant" into "tendency" or "nearly significant" throughout the text, when appropriate, according with the above mentioned specification in the Statistics paragraph.

6) Line 272: $p = 0.0582$ was described as significant

See above

7) In Figure 3 (for Bdnf), should there not be 2 separate asterisks? One for PNS vs Control, and One for PNS vs PNS-HFD?

Done

8) It is not explained what Nr3c1 or Fkbp5 are/encode in the intro, methods, or results. Then in line 276, they mention GR mRNA. This should be clarified.

Done

9) Typo, line 349:.....level of the IL-1B in the DORSAL hippocampus

Done

10) A lot of the referenced articles in the introduction disagree with the current findings. For example:

a. Bdnf is negatively regulated by pro-inflammatory cytokines. Yet in general there was a decrease of both Bdnf and pro-inflammatory cytokines by PNS, suggesting this is not responsible for the regulation of Bdnf in their study.

We agree with this thorough observation. As suggested by these reviewers, we have toned-down the emphasis on the inflammatory effects observed in order to avoid over-interpretation. Thus, our data rather than pointing directly to changes in the inflammatory phenotype overall suggest a general reduced activation of those pathways possibly involved in brain plasticity. In agreement with this view, we observed a general decrease both in Bdnf and the pro-inflammatory cytokines in PNS male rats.

b. In general they exemplify many studies where PNS leads to a pro-inflammatory phenotype. Are there studies where it reduces the pro-inflammatory phenotype?

See answer to Q3

Yet they did not introduce other key aspects of the paper, such as FKBP5/GR signaling and ghrelin signaling in reference to mood disorders and metabolic regulation. I think the introduction should be re-worked to account for their rationale to look at specific candidates.

See answer to Q2

For Peer Review

Reviewer # 2

Comments to the Author

This nice study by Berry et al. describes the central and site-specific effects of prenatal stress (PNS) in combination with a subsequent dietary challenge (HFD) on the mRNA expression level of inflammatory-, stress response-, mood- and metabolic marker genes. Level of mRNA expression were assessed via qPCR in punches of brain areas known to play key roles in neuro psychopathology and homeostasis. An Elevated Plus Maze test, conducted in a previous study by Panetta et al., set base for the correlation analyses of neuroplasticity marker Bdnf and pro-inflammatory marker Il1- β , both of which were inversely correlated with the duration of immobility in the maze. Further, Berry et al. could clearly show interaction effects between the PNS and subsequent dietary challenge on the expression level of nearly all examined marker genes inducing a pro-inflammatory phenotype. This study further helps to disentangle the complex interplay of stress and metabolic control highlighting the key role of central inflammatory processes in this context. However, I do have a number of minor concerns that should be addressed before this manuscript is suitable for publication.

Comments:

1) In the abstract (lines 38 – 40) and in the beginning of the discussion (327 – 330) the authors state that the changes observed in the expression level of Fkbp5 and Nr3c1 underlie the metabolic vulnerability previously observed in Panetta et al. However, the discussion (lines 379 – 422) focuses on the role of Fkbp5 and Nr3c1 in HPA-axis regulation and states that changes of these two marker genes rather reflect a higher stress vulnerability (lines 401 – 402) and diminished cognitive abilities (lines 408 – 409) without discussing how they induce the observed metabolic vulnerability. Therefore, it is very surprising that there is no linear regression analysis of immobility time in the EPM and Fkbp5/Nr3c1 expression level shown. Please, include these and properly discuss all results, including the role of Fkbp5 and Nr3c1 in metabolic control.

We thank the reviewer for the thorough comment. We have now provided evidence that levels of Fkbp5 are positively associated to the time spent in the open arms of the EPM (see Figure 6). “Interestingly, we also showed that levels of Fkbp5 in the dorsal hippocampus were positively related to the time spent in the open arm of the EPM and that this positive correlation is lost as a result both of PNS as well as of HFD. This result might suggest that under physiological conditions Fkbp5 is required to engage animals in the proper exploration of novel environments through a fine modulation of the GR receptors and that this balance can be greatly affected by both early life stressors as well as by metabolic challenges.” This sentence has been added to the discussion paragraph. Moreover, we have added the following sentence to the Discussion paragraph to stress the involvement of Fkbp5 and of Nr3c1 in metabolic regulations: “Glucocorticoids have also an important function in metabolic regulations as they mobilize glucose to fuel the energy demands of the stress response and furthermore promote energy storage, feeding, and weight gain. Thus, it is possible to hypothesize that both psychological as well as metabolic stress might be able to affect HPA axis functionality (Balsevich et al., 2019)”.

2) The last sentence of the abstract in line 42 - 43: “[...] leading to a greater vulnerability to psychopathology.” is very unexpected as it is not stated in the abstract that the behavioral phenotype was assessed and correlated to mRNA expression level. Further, it is too stringent as the results of this study as well as Panetta et al. show that there are also metabolic complications arising from both PNS and HFD. Please rephrase the last sentence and mention the behavioral assessment already in the abstract.

We thank the reviewer for pointing out this issue. The abstract has been modified as follows: “Furthermore, HFD and PNS affected the expression of both Nr3c1 and Fkbp5, two neuroendocrine mediators involved in the response to stress, metabolic challenges and in the modulation of behavioural plasticity (as shown by

the correlation between Fkbp5 and the time spent in the open arms of the elevated plus-maze).". Results have been reported and discussed. (see methods and Discussion paragraphs).

3) As "first hit" and "second hit" are mentioned a couple of times in terms of prenatal and subsequent adult stressors, please elaborate on the two-hit model including references in the introduction and try to imply it also in the discussion.

We thank the reviewer for the comment. We have added the following sentences to the introduction: "In fact, according to a "two-hit model" of vulnerability to diseases, stress experienced during early developmental phases, might affect brain development leading to a reduced ability to cope with further stressors during life (Daskalakis et al. 2013)"; "Based on the "two hit model" of diseases, we hypothesized that PNS...."

See also graphical abstract and answer to Q3, Reviewer#1

4) Please mention the sort of PNS that was applied already in line 100 of the introduction.

The term "PNS" has been changed with "prenatal restraint stress"

5) Further indicate the motivation/reason for the choice of the specific prenatal stress paradigm used (in the context of this study) in the methods section.

This specification has been added to the methods section "This stress has been selected since it is one of the most well characterised prenatal stressors with well-known effects on developmental trajectories of the offspring".

6) Reference Maccari et al. 1995 which is mentioned in line 147 - 148 is missing in the reference section.

Done

7) In line 135 of the methods section it is stated that experiments were selectively performed in males but it is not explained why. Given the fact that females were used in this cohort (see Panetta et al.) it is even more surprising that they were not included in this data set. It should be clearly explained why females were excluded in this study and stated as a major limitation of this work.

We thank the reviewer for pointing out this issue. Although, this was specified in the introduction, ("In a previous study we have shown that male rats who underwent PNS were more vulnerable than females to the effects of a metabolic challenge experienced at adulthood (high-fat diet feeding - HFD), while females showed an overall greater plasticity, possibly mediated by increased total Bdnf mRNA expression levels both in the hippocampus and in the hypothalamus") we have also added the following sentence for the sake of clarity "A growing body of evidence suggests that the effects of PNS are sex-specific and that males' foetal brain might be more vulnerable to changes in the inflammatory mediators overall showing learning deficits and decreased brain plasticity particularly with regard to the hippocampus prefrontal cortex (McCarthy, 2019; Weinstock, 2007)".

8) Please describe the Elevated Plus Maze (EPM) test briefly in the methods section as this is an important readout of this study and explain why exactly immobility time in the maze was chosen as a “a parameter indicative of increased emotionality” (line 180 – 181) and why this readout “can be related with reduced behavioral plasticity” (line 244). It is important for the overall message of this study to explain why this specific readout of the EPM indicates emotionality and plasticity and to reference this properly.

We thank this reviewer for the suggestion. We have briefly added the EPM methodology to the methods section as follows: “Briefly, the apparatus was made of Plexiglas and consisted of two opposite open arms and two arms closed by transparent walls (50×10×40 cm). Each rat was placed in the central area of the maze and video-recorded for 5 minutes under dim light conditions. Each session was recorded and behavioral analysis was carried out using a commercial software (“The Observer 3.0”, Noldus, The Netherlands) (see (Panetta et al., 2017) for further details”).

In the absence of a significant effect of percent time in open arms (Panetta et al. 2017) we have focuses on “immobility time” as a proxy for emotionality and we have correlated this behavioural outcome with IL-1beta and BDNF. We mentioned behavioral plasticity since we know that both BDNF and IL-1beta are specifically involved in behavioral plasticity especially in the hippocampus. However, following suggestion from reviewer #1 (see Q4) we have also added regression plot of Fkbp5 and time spent in the open arm. The following has been added to the text at the end of the “Animal and experimental design” paragraph: “To better characterise the effects of possible changes in gene expression levels, regression analyses were carried out by taking into account specific outcomes deriving from the Elevated Plus-Maze (EPM), as previously performed in (Panetta et al., 2017). We focused on immobility time since this is a well-known proxy for emotionality and reactivity to novel environment (Fernandes and File, 1996) and correlated it with Bdnf and IL-1 β since both these mediators are involved in behavioural plasticity (Bourgognon and Cavanagh, 2020; Cirulli and Alleva, 2009; Goshen et al., 2007). Moreover, we also assessed the ability of Fkbp5 to affect emotionality in the EPM by correlating expression levels of this gene in the dorsal hippocampus to the time spent in the open arms of the maze.”

9) In figures 1,3 and 5 the significant interaction effects of PNS and HFD are neither visually indicated nor are they described in the figure legends. However, the two-way ANOVA is initially the most important statistical analysis of the data sets to show combined effects of both stressors.

We thank the reviewer for pointing out this issue. We have now specified the significant post-hoc comparisons within the PNS X HFD interaction in each figure legends to increase their readout.

10) In figure 1, there is a main effect of the HFD in dorsal hippocampus indicated (\$\$), however in the figure legend the description of this statistical effect is completely missing. Please indicate this effect in figure legend

Done

Further, adjust graph colour for Il-6.

Done

11) The indicated significance in figure 3 (dorsal hippocampus) is not comprehensible as to my understanding there is a post-hoc effect shown here between CTRL and PNS-HFD. However, in the manuscript lines 255 –

256, it is stated that there's significant effects in PNS vs. CTRL and PNS vs. PNS-HFD, which are not indicated in the figure. Please, clarify this.

We thank this reviewer for pointing out this issue. The figure has been changed also in agreement with Reviewer #1

12) Concerning tables 1 and 2, it is not obvious why the authors would choose tables over graphs here, given the fact that collection and analysis of this data set was the same as for the other markers. Preferably, graphs should be chosen to visualize results for Fkbp5 and Nr3c1. Tables could be included for all shown data sets in supplementary material, including missing results for hypothalamus of figures 1, 3 and 5.

Graphs have been now provided and Tables deleted

13) In lines 322 – 323 it is stated that: “PNS, per se, affected brain plasticity by decreasing the expression levels of the neurotrophin Bdnf and pro-inflammatory cytokines...”. However, in lines 252 – 253 the authors state the exact opposite of this: “[...] did not affect levels of Bdnf per se [...]”. Please be cautious with such phrasings and rewrite either of these two sentences.

We agree with this reviewer and the two sentences have been rephrased to make clear that, in the PNS x HFD interaction, the decrease in Bdnf refers to the comparison PNS-CD vs. CTRL-CD, see below:

“overall PNS and postnatal HFD did not affect levels of Bdnf ($F(1, 36) = 3.678; 1.964; p = 0.0631; 0.1696$), main effect of PNS and HFD respectively)”

“PNS-CD rats were characterised by reduced brain plasticity showing decreased expression levels of the neurotrophin Bdnf and of pro-inflammatory cytokines in a number of brain regions”

14) In figure 5, the word “decreased” is missing in the figure legend.

Done

1 **High-fat diet at adulthood interacts with prenatal stress, affecting both brain**
2 **inflammatory and neuroendocrine markers in male rats**

3

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15 **Short title:** *Prenatal stress programs brain inflammatory and neuroendocrine responses*

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18 **Word count:** 5924 (main text)

19

20 **Figures:** 7

21

22 § These authors contributed equally to the work

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25 **Abstract**

26 Prenatal stress (PNS) affects foetal programming and, through an interaction with subsequent challenges, can
27 increase vulnerability to mood and metabolic disorders. We have previously shown that, following PNS, adult
28 male rats are characterised by increased vulnerability to a metabolic stressor experienced at adulthood (8-
29 week-high-fat diet - HFD). In this study, we specifically assessed whether PNS might interact with an adult
30 metabolic challenge to induce an inflammatory phenotype. Changes in the expression levels of inflammatory
31 (*Il-1 β* , *Tnf- α* , *Il-6*) and of stress response mediators (*Nr3c1*, *Fkbp5*) as well as of mood and metabolic
32 regulators (*Bdnf*, *Ghs-R*) were investigated in the hippocampus, prefrontal cortex and hypothalamus, brain
33 regions involved in the pathogenesis of depression and prone to inflammation in response to stress. **Overall,**
34 **PNS reduced the expression of *Bdnf* and *Tnf- α* , while HFD administered at adulthood counteracted this**
35 **effect suggesting that PNS impinges upon the same pathways regulating responses to a metabolic**
36 **challenge at adulthood. Furthermore, HFD and PNS affected the expression of both *Nr3c1* and *Fkbp5*,**
37 **two neuroendocrine mediators involved in the response to stress to metabolic challenges and in the**
38 **modulation of the emotional profile (as shown by the correlation between *Fkbp5* and the time spent in**
39 **the open arms of the elevated plus-maze). Overall, these results indicate that the same metabolic and**
40 **neuroendocrine effectors engaged by PNS are affected by metabolic challenges at adulthood, providing some**
41 **mechanistic insight into the well-known comorbidity between mood and metabolic disorders.**

42

43 **Keywords:** Animal model, Prenatal stress, High-fat diet, Neuroinflammation, Mood disorders, Co-morbidity.

44

45 LIST OF ABBREVIATIONS

- 46 BDNF= Brain-Derived Neurotrophic Factor
- 47 CD= Control Diet
- 48 CRP= C-Reactive Protein
- 49 CTRL= control
- 50 EPM= Elevated Plus-Maze
- 51 EU= European
- 52 GC= Glucocorticoids
- 53 GD= Gestational Day
- 54 GHS-R= Ghrelin Receptor
- 55 GR= Glucocorticoid Receptor
- 56 HFD= High-Fat Diet
- 57 HPA= Hypothalamic–Pituitary–Adrenal
- 58 IL= interleukin
- 59 LTP= Long-Term Potentiation
- 60 NF- κ B= Nuclear Factor Kappa-Light-Chain-Enhancer of Activated B Cells
- 61 PND= Post-Natal Day
- 62 PNS= Prenatal Stress
- 63 ROS= Reactive Oxygen Species
- 64 SEM= Standard Error Mean
- 65

66

67 **1. INTRODUCTION**

68 Mood disorders are common conditions with major public health implications. It is estimated that each year
69 around 40% of the EU population suffers from a mental disorder. Adjusted for age and comorbidity, this
70 corresponds to 164.8 million people affected (Health Organization Regional Office for Europe, 2015). Among
71 the risk factors that might trigger the onset of psychiatric conditions, or an exacerbation of symptoms, stressful
72 life events play a pivotal role. **In fact, according to a “two-hit model” of vulnerability to diseases, stress
73 experienced during early life phases, might affect brain development leading to a reduced ability to cope
74 with further stressors during life (Daskalakis et al., 2013).** To this regard, a suboptimal intrauterine
75 environment, prompted by maternal stress (ranging from poor socio-economic status and maternal obesity to
76 depression or maltreatments) may predispose the offspring to lifelong negative health outcomes, including
77 neuropsychiatric disorders as well as metabolic and immune dysregulations (Barker, 1995; Berry et al., 2015;
78 Boersma et al., 2013; Cattane et al., 2020; Cirulli et al., 2020; Krontira et al., 2020).

79 Consistent evidence suggests that depressed patients are characterised by alterations in the functional activity
80 of the immune system, affecting both peripheral and central tissues and that early life stress represents an
81 important factor influencing the inflammatory status (Cattaneo et al., 2015; Dantzer et al., 2008; Dowlati et
82 al., 2010; Liu et al., 2012; Valkanova et al., 2013). As an example, Danese and colleagues reported increased
83 blood C-reactive protein (CRP) levels in maltreated children, an effect that was larger in those subjects that
84 also developed depression later in life (Danese et al., 2009, 2008). Likewise, Slopen and co-workers reported
85 increased CRP and interleukin-6 (IL-6) in adolescents who experienced early life adversities (Slopen et al.,
86 2014). Offspring of women who experienced stressful life events during pregnancy show higher levels of IL-
87 1 β , IL-4, IL-5, IL-6, and IL-8 in umbilical cord blood at delivery (Andersson et al., 2016). Interestingly, it has
88 been recently proposed that inflammation may act as a key mediator linking exposure to prenatal stress (PNS)
89 and enhanced vulnerability to psychopathology in the offspring (Hantsoo et al., 2019).

90 Preclinical studies in rodent models provide strong support to the above mentioned clinical findings, showing
91 for example that PNS exposure causes extended inflammation in the foetal brain (Ślusarczyk et al., 2015). To
92 this regard, Gur and colleagues found increased levels of IL-1 β in the placenta of mouse dams undergoing
93 stress during pregnancy and in the foetal brain of their offspring; this was also associated with decreased levels
94 of the neurotrophin Brain-Derived Neurotrophic Factor (BDNF) specifically in the amygdala (Chen et al.,
95 2020; Gur et al., 2017) suggesting that increased expression of placental immune responsive genes upon
96 maternal stress might be considered as a potential mechanism (Bronson and Bale, 2014; Mueller and Bale,
97 2008).

98 Despite the large body of clinical and preclinical evidence showing associations among PNS, the
99 hyperactivation of the immune system and an enhanced vulnerability to mood disorders, cause-effect
100 mechanisms are still largely unknown, and the prenatal environment may not act alone to set the stage for long-
101 term onset of mood disorders. Interestingly, psychiatric conditions and metabolic pathologies are often found

102 to co-occur within the same individual showing a feed-forward pattern overall suggesting the activation of
103 shared mechanisms/pathways (Cattane et al., 2020; Cirulli et al., 2020; Milaneschi et al., 2019). Among these,
104 BDNF and glucocorticoids (GC) hormones have been identified as main effectors of brain plasticity and
105 metabolic function in response to stressful events. Thus, it is possible that these actors may represent both
106 effectors as well as targets of stress - during sensitive developmental periods - leading, in the developing
107 organism, to a remodelling of the mechanisms associated with stress responsiveness (Cattaneo et al., 2015;
108 Cirulli and Alleva, 2009; McEwen, 2000). The neurotrophin BDNF plays a pivotal role in brain and
109 behavioural plasticity as well as in the control of energy homeostasis (Cirulli and Alleva, 2009; Marosi and
110 Mattson, 2014); its expression is finely tuned by GC in response to acute or chronic stressful stimuli (Cirulli,
111 2017; Cirulli & Alleva, 2009) and it has been also suggested to be negatively regulated by pro-inflammatory
112 cytokines (Barrientos et al., 2004; Bilbo et al., 2008). Thus, BDNF appears as a sensitive target as well as a
113 biomarker of stress response as it could actively contribute to maintain brain health/homeostasis in different
114 challenging conditions. To this regard, the interplay between the activation of stress-related pathways by
115 adverse experiences during development and adult metabolic challenges might be crucial in shaping individual
116 vulnerability to stress (Nederhof and Schmidt, 2012).

117 **A growing body of evidence suggests that the effects of PNS are sex-specific and that males' foetal brain**
118 **might be more vulnerable to changes in the inflammatory mediators overall showing learning deficits**
119 **and decreased brain plasticity particularly with regard to the hippocampus prefrontal cortex**
120 **(McCarthy, 2019; Weinstock, 2007).** Indeed, in a previous study we have shown that male rats who
121 underwent PNS were more vulnerable than females to the effects of a metabolic challenge experienced at
122 adulthood (high-fat diet feeding - HFD), while females showed an overall greater plasticity, possibly mediated
123 by increased total *Bdnf* mRNA expression levels both in the hippocampus and in the hypothalamus. **Based on**
124 **the "two hit model" of diseases,** we hypothesized that PNS might disrupt the intrauterine environment
125 affecting brain developmental trajectories leading, in turn, to increased brain inflammation and reduced
126 plasticity in the offspring (first hit). We also hypothesized that a metabolic challenge such as a HFD,
127 experienced at adulthood, might interact with inflammatory pathways to exacerbate the PNS-induced pro-
128 inflammatory phenotype. Thus, **the** aim of this study was to investigate the effect of a second hit - represented
129 by an HFD - specifically in male rats. We focused on selected brain regions - characterised by high degree of
130 sexual dimorphism (Handa et al., 1994) - such as the dorsal and the ventral hippocampus, the prefrontal cortex
131 and the hypothalamus known to be involved in the pathogenesis of depression and to be prone to inflammatory
132 status in response to external challenges. **We specifically assessed in these brain regions levels of *Bdnf*, *Tnf-***
133 **α , *Il-6* and *Il- β .***

134 **Stress powerfully affects both mood and energy homeostasis. Such effects are achieved not only through**
135 **the interaction between GC hormones and their receptors (that regulate HPA axis function in addition**
136 **to glucose metabolism) but also by triggering a multitude of signalling cascades reciprocally modulating**
137 **one another in a regional-, temporal, and functional-dependent manner (Balsevich et al., 2019). Among**
138 **these, ghrelin is a peptide hormone produced in the stomach and involved in the signalling of meal**

139 initiation. Its action is mediated through the growth hormone secretagogue receptor (*Ghs-R*) that is
140 found to be highly expressed in several brain regions including the hypothalamus, pituitary gland and
141 hippocampus (see Zarouna, 2015 and references therein). Both Ghrelin and *Ghs-R* are modulated by
142 stress (Patterson et al., 2010); there is evidence that ghrelin may decrease anxiety-like and depressive-
143 like behaviours in mice (Lutter et al., 2008) moreover, it appears to be involved in food anticipatory
144 activity and consumption (Blum et al., 2009; Verhagen et al., 2011) as well as in mood disorders
145 (Zarouna, 2015). Interestingly, both genetic variants within ghrelin and GC signalling pathways have
146 been associated with obesity, stress-related mental disorders, or both (see Balsevich et al., 2019 and
147 references therein). To this regard, we also investigated changes in the expression levels of *Ghs-R* as well
148 as of *Nr3c1* (encoding for the glucocorticoid receptors - GR) and the GC co-chaperon *Fkbp5* (encoding
149 for the FK506 binding protein 51) in all the above-mentioned brain regions.

150

151 2. MATERIALS AND METHODS

152 2.1 Animals and experimental design

153 Twelve adult nulliparous female (230–260 g) and 6 male Sprague-Dawley rats (400 g) were purchased from a
154 commercial breeder (Charles River, Calco, Italy). Upon arrival, animals were pair-housed with same sex
155 conspecifics under standard laboratory conditions (see Panetta et al. 2015 for further details). Pellet food
156 (Altromin-R, Rieper, Italy) and tap water were continuously available. Following one week of adaptation, two
157 females and one male were mated for 24 hours. To assess pregnancy, changes in body weight were monitored.
158 Pregnant females were randomly assigned to either the control (CTRL) or prenatal stress (PNS) groups. CTRL
159 rats were left undisturbed throughout gestation, while PNS females underwent a repeated restraint stress
160 procedure during the last third of gestation until the day delivery day (postnatal day 0 - PND-0). On PND-1 all
161 pups were weighted and litters culled to an average of 5 male and 5 female pups. On PND-21 pups were
162 weaned and housed in groups of 2 or 3 same-sex littermates until 2 months of age. **By this time, n=20 CTRL**
163 **and n=21 PNS males were assigned either to high-fat diet (HFD) or control diet (CD) regimen for 8**
164 **weeks. The final number of animals per group was: n=10 CTRL-CD, n=10 CTRL -HFD, n=10 PNS-CD**
165 **and n=11 PNS-HFD. Groups were composed by 1 pup/litter (CTRL-CD; CTRL-HFD; PNS-CD; PNS-**
166 **HFD. After 4 weeks on the respective diets, all animals underwent a number of metabolic and**
167 **behavioural assessments; they were successively sacrificed and brains dissected out to investigate the**
168 **expression levels of *Il-6*, *Il-1 β* , *Tnf- α* , *Nr3c1*, *Fkbp5*, *Ghs-R*, in the dorsal hippocampus, ventral**
169 **hippocampus, hypothalamus and prefrontal cortex; total *Bdnf* was investigated only in the dorsal**
170 **hippocampus and prefrontal cortex (since it was already assessed in the other two areas, (Panetta et al.,**
171 **2017).**

172 To better characterise the effects of possible changes in gene expression levels, regression analyses were
173 carried out by taking into account specific outcomes deriving from the Elevated Plus-Maze (EPM), as

174 **previously performed in (Panetta et al., 2017). We focused on immobility time since this is a well-known**
175 **proxy for emotionality and reactivity to novel environment (Fernandes and File, 1996) and correlated it**
176 **with *Bdnf* and *Il-1 β* since both these mediators are involved in behavioural plasticity (Bourgognon and**
177 **Cavanagh, 2020; Cirulli and Alleva, 2009; Goshen et al., 2007). Moreover, we also assessed the ability**
178 **of *Fkbp5* to affect emotionality in the EPM by correlating expression levels of this gene in the dorsal**
179 **hippocampus to the time spent in the open arms of the maze. As for the EPM protocol used, briefly, the**
180 **apparatus was made of Plexiglas and consisted of two opposite open arms and two arms closed by**
181 **transparent walls (50×10×40 cm). Each rat was placed in the center of the maze and video-recorded for**
182 **5 minutes under dim light conditions. Each session was recorded and behavioral analysis was carried**
183 **out using a commercial software (“The Observer 3.0”, Noldus, The Netherlands) (see (Panetta et al.,**
184 **2017) for further details).**

185 All experimental procedures were reviewed by the ethical body of the Istituto Superiore di Sanità for animal
186 welfare and conducted in conformity with the European Directive 2010/63/EU and the Italian legislation on
187 animal experimentation, D.Lgs. 26/2014. They were authorized by the Italian Ministry of Health.

188

189 **2.2 Dams’ stress procedure**

190 Pregnant females (at gestational day - GD - 14) were restrained in a transparent Plexiglas cylinder (7.5×19 cm)
191 under a bright light (6.500 lux) for 45 minutes three times daily at random times (between 9:00 a.m.-5:00 p.m.)
192 during the dark phase until the expected delivery day (GD-21, see Maccari et al., 1995 for further details).
193 **This stress has been selected since it is one of the most well characterised prenatal stressors with well-**
194 **known effects on developmental trajectories of the offspring.**

195

196 **2.3 High-fat diet administration**

197 CTRL and PNS offspring were fed *ad libitum* either with HFD (energy: 5.24 kcal/g; composition: fat 60%,
198 carbohydrate 20% and protein 20%) or CD (energy: 3.3 kcal/g; composition: fat 17%, carbohydrate 60% and
199 protein 23%) starting from two months of age. The HFD diet (D12492) was purchased from Research Diets,
200 Inc., New Brunswick, NJ, USA; the CD was purchased from Altromin-R, Rieper, Italy.

201

202 **2.4 Tissue collection**

203 At 4 months of age offspring **was** weighted and sacrificed. **Brains were** removed and the dorsal and ventral
204 hippocampus, hypothalamus and prefrontal cortex were dissected out and immediately frozen at -80°C (Panetta
205 et al., 2017).

206

207 2.7 Molecular analysis

208 2.7.1 RNA isolation and Real Time PCR Analyses

209 Total RNA was isolated using PureZol RNA isolation reagent (Bio-Rad Laboratories, Italy), treated with
210 DNase to avoid DNA contamination and quantified by spectrophotometric analysis. The quantified RNA was
211 analyzed by TaqMan q-RT PCR Instrument (CFX384 real time system, Bio-Rad Laboratories) using the
212 iScript™ one-step RT-PCR kit for probes (Bio-Rad Laboratories) and Applied BioSystem Assays (Gene
213 Expression Assays: *Il6*, *Il-β*, *Tnf-α*, *Bdnf*, *Nr3c1*, *Fkbp5*, and *Ghs-R*). Samples were run in triplicate and each
214 target gene analyzed has been normalized to the expression of the housekeeping (HK) gene β-actin (ActB).
215 The expression of target genes was calculated by using to the Ct method (-ΔΔCt method) (Schmittgen and
216 Livak, 2008), where CTRL-CD rats have been used as a reference group.

217

218 2.8 Statistical analysis

219 Data were evaluated by a two-way ANOVA with Diet (HFD vs CD) and Prenatal condition (PNS vs CTRL)
220 as between-subject factors. *Post-hoc* comparisons between groups were performed using the Tukey's test. A
221 linear regression model was used to assess the ability of *Bdnf* levels and of *Il-1β* in the dorsal hippocampus to
222 affect immobility duration (a parameter indicative of increased emotionality) in the EPM test. A level of
223 probability set at $p < 0.05$ was used as statistically significant. **Statistical tendency was set at $p < 0.1$. For**
224 **interactions at $p < 0.1$, we also examined lower order effects.** Data are presented graphically as means ±
225 SEM box plot (observations outside the ranges are represented with dots outside the boxes). **The raw data**
226 **supporting the conclusions of this article will be made available by the authors upon request without**
227 **undue reservation.**

228

229 3. RESULTS

230

231 3.1 Inflammatory mediators *Il-1β*, *Tnf-α* and *Il-6* mRNA expression

232 *Il-1β* - We first analyzed the expression of *Il-1β*, a prototype pro-inflammatory cytokine. We observed that
233 PNS resulted in a nearly significant decrease in the dorsal hippocampus while neither a main effect of HFD
234 nor a PNS x HFD interaction effect were found ($F(1,35)=3.818, 0.004, 2.184; p=0.0587, 0.9518, 0.1484$).

235 No main effects of PNS nor of HFD were observed in the ventral hippocampus ($F(1,34)=0.152, 0.335;$
236 $p=0.6995, 0.5665$). **A strong PNS x HFD interaction was found ($F(1,34)=4.655; p=0.0381$) PNS exposed**
237 **animals being characterised by decreased levels of this cytokine, while the administration of postnatal**
238 **HFD counteracted the effects of the PNS (post-hoc comparisons just missed statistical significance).**

239 In the prefrontal cortex while no main effects of PNS was observed ($F(1,33)=0.969; p=0.3321$) postnatal HFD
240 overall increased levels of *Il-1β* (main effect of HFD: $F(1,33)=5.112, p=0.0305$). Similarly to what was
241 observed in the ventral hippocampus, a **nearly significant** PNS x HFD interaction was found with decreased

242 levels of *Il-1 β* in PNS animals, this effect being reverted by postnatal administration of HFD ($F(1,33)=3.858$;
243 $p=0.0580$, post hoc comparisons: $p<0.01$, PNS vs. PNS-HFD); (see Figure 1A).

244 In the hypothalamus, levels of *Il-1 β* were not affected by PNS, HFD nor an interaction was found
245 ($F(1,36)=0.042, 0.0001633, 0.070$; $p=0.8386, 0.9899, 0.7934$ respectively for PNS, HFD and PNS x HFD, see
246 Figure 1A).

247

248 *Tnf- α* - In the dorsal hippocampus no main effects of PNS nor of HFD were observed ($F(1,37)=1.288, 0.209$;
249 $p=0.2638, 0.6506$). By contrast, a significant interaction was found between PNS and HFD ($F(1,37)=5.772$;
250 $p=0.0214$). In particular, PNS animals were characterised by decreased levels of *Tnf- α* , although this effect
251 was no longer observed in PNS-HFD animals that showed levels of this cytokine comparable to those observed
252 in CTRL animals (post hoc comparisons: $p<0.05$, PNS vs. CTRL).

253 No main effects of PNS nor of HFD were found in the ventral hippocampus ($F(1,34)=0.033, 2.941$; $p=0.8573,$
254 0.0955 , respectively for PNS and HFD), while a PNS x HFD interaction was found showing that administration
255 of postnatal HFD in the PNS group selectively increased levels of this cytokine ($F(1,34)=6.747$; $p=0.0138$);
256 post hoc comparisons: PNS vs. HFD-PNS, $p<0.01$).

257 No main effects of PNS nor of HFD were observed in the prefrontal cortex ($F(1,33)=0.079, 1.648$; $p=0.7809,$
258 0.2082 respectively for PNS and HFD) however, as also observed for the ventral hippocampus a PNS x HFD
259 interaction effect was found showing that the administration of postnatal HFD in the PNS group selectively
260 increased levels of this cytokine ($F(1,33)=7.946$; $p=0.0081$); post hoc comparisons: PNS vs. HFD-PNS,
261 $p<0.01$); (see Figure 1B).

262 Levels of *Tnf- α* in the hypothalamus (see Figure 1B) were not affected neither by PNS nor by postnatal HFD
263 ($F(1, 35)=0.286, 0.702, 2.111$; $p=0.5964; 0.4079; 0.1551$ respectively for PNS, HFD and PNS x HFD).

264

265 *Il-6* - Postnatal HFD decreased levels of *Il-6* in the dorsal hippocampus (main effect of HFD: $F(1,37)=0.814$;
266 $p=0.0066$), whereas no effect of the PNS nor of the PNS x HFD interaction effect were found (PNS main
267 effect: $F(1,37)=1.387$; $p=0.2465$; PNS x HFD interaction: $F(1,37)=0.164$; $p=0.6881$).

268 Levels of *Il-6* did not change as a result of PNS, HFD or their interaction neither in the ventral hippocampus
269 ($F(1,36)=0.759, 2.631, 0.029$; $p=0.3893, 0.1135, 0.8647$ respectively for PNS, HFD and PNS x HFD) nor in
270 the prefrontal cortex ($F(1,35)=0.144, 0.053, 0.202$ respectively for PNS, HFD and PNS x HFD); (see Figure
271 1C). PNS decreased nearly significantly the levels of *Il-6* also in the hypothalamus ($F(1,36)=3.889$; $p=0.0563$),
272 but no effect of HFD nor an interaction PNS x HFD were found to be significant ($F(1,36)=0.903, 0.757$;
273 $p=0.3482, 0.3899$, respectively for HFD and PNS x HFD).

274

275 -----FIGURE 1 about here-----

276

277 Interestingly, when expression levels of *Il-1 β* in the dorsal hippocampus were related to immobility duration
278 in the EPM (a test run in a companion paper on the same experimental animals, see (Panetta et al., 2017)), we

279 observed a significant strong negative association linking these two parameters only in the PNS-CD animals,
 280 overall suggesting that reduced levels of this cytokine can be related with reduced behavioral plasticity (see
 281 discussion; CTRL-CD: $F(1,8)=0.007$; $p=0.9367$; $R^2=0.01$; CTRL-HFD: $F(1,8)=3.461$; $p=0.1051$; $R^2=33.1$;
 282 PNS-CD: $F(1,8)=7.538$; $p=0.0287$; $R^2=51.8$; PNS-HFD: $F(1,9)=1.739$; $p=0.2237$, $R^2=17.9$; see Figure 2).

283

284 -----FIGURE 2 about here-----

285

286 3.2 Neuronal plasticity marker - *Bdnf* mRNA expression

287 We first investigated the expression of *Bdnf*, a neurotrophin that is considered a prototype marker of plasticity.
 288 Within the dorsal hippocampus, **overall PNS and postnatal HFD did not affect levels of *Bdnf*** ($F(1, 36)=$
 289 3.678 ; 1.964 ; $p=0.0631$; 0.1696), main effect of PNS and HFD respectively), although we observed a
 290 significant interaction between PNS and postnatal HFD ($F(1, 36)=5.606$; $p=0.0234$). More in detail, PNS
 291 decreased *Bdnf* expression, while the postnatal administration of HFD was able to counteract this effect (PNS
 292 vs. CTRL, $p<0.05$; PNS vs. PNS-HFD, $p<0.05$). In the prefrontal cortex no effect of PNS, HFD or their
 293 interaction was observed ($F(1, 36)=0.200$, 1.377 , 0.174 ; $p=0.6572$; 0.2482 ; 0.6794 , respectively for PNS, HFD
 294 and PNS x HFD); (see Figure 3).

295

296 -----FIGURE 3 about here-----

297

298 Interestingly, levels of *Bdnf* in the dorsal hippocampus were negatively associated with the time spent
 299 immobile in the EPM (a test run in a companion paper on same experimental animals, see (Panetta et al.,
 300 2017)) only in the PNS-CD group (CTRL-CD: $F(1,9)=0.730$; $p=0.4261$; $R^2=0.081$; CTRL-HFD: $F(1,9)=$
 301 0.090 ; $p=0.7713$; $R^2=0.011$; PNS-CD: $F(1,8)=11.084$; $p=0.0127$; $R^2=0.612$; PNS-HFD: $F(1,9)=0.716$;
 302 $p=0.4221$; $R^2=0.082$), supporting a role of this neurotrophin in the emotional phenotype (see Figure 4).

303

304 -----FIGURE 4 about here-----

305

306 3.3 Stress response mediators - *Nr3c1*, *Fkbp5* mRNA expression

307 *Nr3c1* - In the dorsal hippocampus the postnatal HFD decreased the levels of *Nr3c1* when compared to controls
 308 and a similar **nearly significant** effect was observed for PNS (main effect: $F(1,36)=3.827$, 7.063 , $p=0.0582$;
 309 0.0117 respectively for PNS, HFD). Moreover, a significant PNS x HFD interaction was found showing that
 310 both PNS-CD and CTRL-HFD as well as the combination PNS-HFD were characterised by a decrease of GR
 311 mRNA levels ($F(1,36)=6.228$; $p=0.0173$ and PNS x HFD; post hoc comparisons $p<0.05$).

312 In the ventral hippocampus levels of *Nr3c1* were increased a result of postnatal HFD (main effect:
 313 $F(1,36)=10.405$; $p=0.0027$), while no effects of PNS nor of the PNS x HFD interaction were found ($F(1,$
 314 $36)=0.011$, 0.258 ; $p=0.9167$; 6146). In the prefrontal cortex the postnatal HFD overall increased the levels of
 315 *Nr3c1* (main effect of HFD: $F(1, 37)=12.453$; $p=0.0011$); no effects of PNS nor of the PNS x HFD interaction

316 were observed ($F(1, 37)=1.001, 0.702; p=0.3236; 0.4076$). No effects of PNS, HFD nor of their interaction
 317 were found in the hypothalamus ($F(1, 37)=0.157, 0.009, 1.067; p= 0.6943; 0.9257; 0.3084$); (see **Figure 5**).

318

319 *Fkbp5* - In the dorsal hippocampus the postnatal HFD increased the levels of *Fkbp5*
 320 ($F(1,35)=10.331;p=0.0028$), while no effect of PNS nor of the PNS x HFD interaction was observed
 321 ($F(1,35)=1.502, 0.958; p= 0.2285; 0.3343$). No effect of PNS, HFD or their interaction was found in the ventral
 322 hippocampus ($F(1,36)=0.781, 3.222, 0.007; p=0.3828; 0.0811; 0.9359$). In the prefrontal cortex, PNS increased
 323 the levels of *Fkbp5* ($F(1,37)=5.244; p=0.0278$), while no effect of HFD nor of the PNS x HFD interaction was
 324 observed ($F(1,37)=1.219, 2.019; p=0.2767; 0.1637$). A main effect of the postnatal HFD was also observed In
 325 the hypothalamus, the Postnatal HFD decreased levels of *Fkbp5* ($F(1,37)=9.156; p=0.0045$); no effect of PNS
 326 nor of the PNS x HFD interaction was observed ($F(1,37)=0.094, 3.353; p= 0.7608; 0.0751$); (see **Figure 5**).

327

328 -----FIGURES 5 about here-----

329

330 **Interestingly, levels of *Fkbp5* in the dorsal hippocampus were positively related with increased time**
 331 **spent in the open arm of the EPM (a test run in a companion paper on same experimental animals, see**
 332 **(Panetta et al., 2017)) only in the CTRL-CD group (CTRL-CD: $F(1,8)=26.617; p=0.0013; R^2= 0.792$;**
 333 **CTRL-HFD: $F(1,9)= 0.104; p=0.7713; R^2=0.013$; PNS-CD: $F(1,8)=0.253; p=0.6307; R^2=0.035$; PNS-**
 334 **HFD: $F(1,9)=0.503; p=0.6307; R^2= 0.4982$). This result might suggest that *Fkbp5* is physiologically**
 335 **required to engage animals in the proper exploration of novel environments through a fine modulation**
 336 **of GR receptors (see Figure 6).**

337

338 -----FIGURES 6 about here-----

339

340 **3.4 Metabolic/mood regulator marker - *Ghs-R* mRNA expression**

341 *Ghs-R* - In the dorsal hippocampus the postnatal HFD overall decreased levels of *Ghs-R* ($F(1,33)=1,948$;
 342 $p=0.0007$). By contrast, PNS increased *Ghs-R* levels ($F(1,33)=5.192; p=0.0293$), but no interaction between
 343 PNS x HFD was observed ($F(1,33)=0.290; p= 0.5941$). In the ventral hippocampus HFD overall increased the
 344 levels of *Ghs-R* ($F(1,32)=6.278; p=0.0175$), while no main effect of PNS was observed ($F(1,32)=0.593$;
 345 $p=0.4468$). However, *Ghs-R* was increased in the CTRL-HFD and in the PNS-CD groups (interaction between
 346 PNS and HFD: $F(1,32)=14.778; p=0.0005$, post hoc: CTRL vs. CTRL-HFD, $p<0.01$; CTRL vs. PNS, $p<0.01$).
 347 In the prefrontal cortex, PNS increased the levels of *Ghs-R* ($F(1,29)=5.865; p=0.0219$) and no effect of HFD
 348 was observed ($F(1,29)=2.191; p=9.1496$). However, in this brain area, *Ghs-R* was increased in the CTRL-HFD
 349 and in the PNS-CD groups (interaction between PNS and HFD: $F(1,29)=6.870; p= 0.0138$, post hoc: CTRL
 350 vs. CTRL-HFD, $p<0.05$; CTRL vs. PNS, $p<0.01$); (see **Figure 7**). As for the hypothalamus, neither PNS nor
 351 HFD affected levels of *Ghs-R* ($F(1,34)=3.155, 3.033, 0.907; p=0.0846; 0.0906; 0.3477$).

352

353 -----FIGURE 7 about here-----

354

355 4. DISCUSSION

356 In the present study we assessed the long-term effects of exposure to PNS (a first hit) on foetal brain
357 programming with regard to neuronal plasticity and brain inflammation. Moreover, we used a metabolic
358 challenge (HFD) at adulthood (second hit) to assess potential interaction effects with the PNS-induced
359 phenotype. We have previously shown (Panetta et al., 2017) that although PNS resulted in a strong metabolic
360 liability in male rats fed with HFD at adulthood it also dampened the negative effects driven by such a
361 metabolic challenge on the emotional phenotype as assessed in the EPM. Here we extended such body of
362 evidence by providing information on the possible mechanisms underlying this interaction. **PNS-CD rats were**
363 **characterised by reduced brain plasticity showing decreased expression levels of the neurotrophin *Bdnf***
364 **and of pro-inflammatory cytokines in a number of brain regions; moreover, HFD feeding, experienced**
365 **at adulthood, induced an increase in pro-inflammatory cytokines in those animals that had experienced**
366 **PNS.** This suggests that PNS might impinge upon the same mechanisms underlying vulnerability to metabolic
367 challenges at adulthood. Furthermore, HFD greatly affected the expression levels of the main effectors of HPA
368 axis function (the glucocorticoid receptor - GR - gene *Nr3c1* and of the GR's co-chaperone *Fkbp5*) an effect
369 possibly underlying the metabolic vulnerability previously observed in male rats.

370 In our study, we found that PNS animals were characterised by differential expression of mRNA levels of *Il-*
371 *1β* and *Tnf-α* in almost all the brain regions investigated except for the hypothalamus, an area that, in our
372 experimental condition, appeared to be resilient to inflammatory changes driven by both pre- and postnatal
373 stressors. While *Tnf-α* was tightly modulated upon pre- and postnatal stressors, changes in *Il-1β* were observed
374 only when considering the interaction with the second hit (HFD), suggesting that the former cytokine (*Tnf-α*)
375 might be considered as a reliable marker of stress adaptation in response to a prenatal challenge. As for *Il-6*,
376 no effect was observed upon PNS in neither of the brain areas considered, though HFD overall decreased its
377 expression levels in the dorsal hippocampus. TNF-α, IL-1β and IL-6 are the main activators of the HPA and
378 are in turn modulated (inhibited) by GC hormones. As described by O'Connors and colleagues these cytokines
379 hold differential sensitivity for adrenal steroids with TNF-α being the most sensitive to such inhibition in the
380 range of physiological levels, IL-1β being second and IL-6 being the most resistant (O'Connor et al., 2000).
381 There is general consensus that elevated inflammation can exacerbate or even give rise to depressive symptoms
382 or may be associated to other psychiatric conditions (Dantzer et al., 2018, 2008). However, cytokines are also
383 constitutively released in the healthy brain by resident myeloid cells to keep proper synaptic plasticity. As an
384 example, the modulation of both IL-1β and TNF-α plays an important role in the processes of LTP and synaptic
385 scaling (a form of homeostatic plasticity) (Rizzo et al., 2018; Salim et al., 2012). Here we found that lower
386 mRNA levels of *Il-1β* in the **dorsal** hippocampus of PNS rats were associated to increased time spent immobile
387 in the EPM (as previously assessed in (Panetta et al., 2017)). Moreover, and in line with our previous work
388 (Panetta et al., 2017), PNS rats were characterised by decreased levels of the neurotrophin *Bdnf* (in the dorsal

389 hippocampus); this decrease was also associated to an increase in the time spent immobile in the EPM,
390 suggesting that PNS might affect foetal brain programming by reducing neuronal plasticity. Interestingly, the
391 interaction between PNS and adult HFD (PNS-HFD group) resulted in increased levels of *Il-1 β* and *Tnf- α*
392 mRNA. Alboni and colleagues have recently provided evidence for increased rather than decreased levels of
393 pro-inflammatory cytokines in the brain of stressed mice treated with the antidepressant fluoxetine. Such an
394 effect was associated to increased BDNF levels and stronger LTP in the hippocampus (Alboni et al., 2017,
395 2016), suggesting that brain plasticity may also be related to the activation of basal metabolism that in turn is
396 positively associated to the ability to properly mount and control inflammatory responses. To this regard, we
397 cannot exclude that prolonged exposure to the HFD or to a stronger metabolic insult, such as a western pattern
398 diet (rich in fats and sugar), should lead to an overall excessive brain inflammation (not observed in this study)
399 with main consequences on emotional/cognitive behaviour. **Worth to notice, we have previously shown that**
400 **upon PNS, male rats were characterized by increased corticosterone levels under basal conditions, this**
401 **effect was associated with a decrease in reactive oxygen species (ROS) as well as a with decreased NF-**
402 **kB signalling in the hippocampus suggesting a lower set-point under basal conditions in PNS male rats**
403 **(Anacker et al., 2013). Here we observed a similar blunted activation upon PNS with HFD triggering a**
404 **response only in PNS subjects (increased *Il-1 β* and *Tnf- α* in the prefrontal cortex and *Tnf- α* in the dorsal**
405 **and ventral hippocampus). To our knowledge this is one of the few instances in which a reduction in**
406 **inflammatory mediators is described a result of PNS. Because cytokines have been shown to modulate**
407 **hippocampal development and plasticity (Bourgognon and Cavanagh, 2020; Goshen et al., 2007) a**
408 **decreased expression profile as a result of a “first hit” (PNS) might set the stage for an increased**
409 **response to a “second hit” (metabolic challenge).** Obesity is characterised by low-grade systemic and central
410 inflammation and TNF- α and IL-1 β appear to be key players in this condition and in the associated pathologies
411 (Mighiu et al., 2012). Although the hypothalamus is a well-known area playing a main role in the crosstalk
412 between brain and periphery in metabolic pathologies associated to obesity, other brain regions have been
413 shown to be affected by high-fat diet- or obesity-induced inflammation such as the hippocampus, cortex,
414 brainstem, or amygdala (Guillemot-Legris and Muccioli, 2017). There are data to suggest that the effects of a
415 metabolic challenge may be different depending on the region of interest dealing with brain plasticity
416 (hippocampus) or metabolic regulation and homeostasis (hypothalamus) (Rasgon and McEwen, 2016). Thus,
417 our data also indicate that PNS might set the stage for a differential regulation of central inflammatory
418 mediators in specific brain regions, an effect that can be unmasked only upon the occurrence of a second hit,
419 such as the HFD at adult age.

420 Hyperactivity of the HPA axis and increased circulating GC have been implicated in the pathogenesis of mood
421 and anxiety disorders suggesting an impairment in the ability of the HPA axis to self-regulate its function by
422 shutting down the system (Binder, 2009; Holsboer, 2000; Pariante and Miller, 2001). **Glucocorticoids have**
423 **also an important function in metabolic regulations as they mobilize glucose to fuel the energy demands**
424 **of the stress response and furthermore promote energy storage, feeding, and weight gain. Thus, it is**
425 **possible to hypothesize that both psychological as well as metabolic stress might be able to affect HPA**

426 **axis functionality (Balsevich et al., 2019).** A growing body of evidence suggests that levels of *Fkbp5* mRNA
427 GR co-chaperone have been associated with higher levels of circulating cortisol and reduced negative feedback
428 inhibition of the stress response associated with a depressive phenotype (Binder, 2009). We have previously
429 shown that, following PNS, male rats were characterised by elevated GC levels when compared to controls
430 (Anacker et al., 2013); **in this study we show elevated levels of *Fkbp5* upon PNS in the prefrontal cortex**
431 **and upon HFD administration in the dorsal hippocampus, suggesting that both psychological and**
432 **metabolic stressors affect this mediator. When we looked at the hypothalamus, *Fkbp5* expression levels**
433 **were reduced in the HFD group, possibly accounting for a compensatory mechanism. Interestingly, we**
434 **also showed that levels of *Fkbp5* in the dorsal hippocampus were positively related to the time spent in**
435 **the open arm of the EPM and that this positive correlation is lost as a result both of PNS as well as of**
436 **HFD. This result might suggest that under physiological conditions *Fkbp5* is required to engage animals**
437 **in the proper exploration of novel environments through a fine modulation of the GR receptors and that**
438 **this balance can be greatly affected by both early life stressors as well as by metabolic challenges.**

439 The prefrontal cortex is one of the most important cortical area among the network of regions being involved
440 in the pathogenesis of depression that plays a main role in the ability to process positive and negative emotions
441 (Harms et al., 2017; Kaya and McCabe, 2019). Thus, the observed increase in levels of *Fkbp5* in this brain
442 area should not be surprising and may underlie a condition of potential increased vulnerability to stress.

443 **Changes in *Nr3c1* and *Fkbp5* in the hippocampus, a result of HFD, indicate a cross-regulation between**
444 **a metabolic challenge and *Nr3c1/Fkbp5* balance, strengthening the notion that indeed these molecules**
445 **are involved in “sensing” environmental challenges, be these of psychological or metabolic nature.** We
446 also found that HFD increased the levels of *Nr3c1* in the prefrontal cortex; by contrast, changes in *Fkbp5* in
447 this brain area were observed only as a result of PNS. These results possibly suggest a fine and complex
448 multiple level regulation of the GR-negative feedback in this specific brain region highlighting the role of this
449 brain area in mood disorders such as depression. Transcriptional control of GR relies in part upon the DNA
450 methylation status at multiple alternative initiation sites that are tissue specific in fact, *Nr3c1* gene is
451 characterised by an unusually complex promoter structure (Turner et al., 2010). This might possibly explain
452 the unexpected decrease in the dorsal hippocampus observed as a result of HFD or the lack of effects observed
453 in the hypothalamus (see also concluding remarks).

454 When we investigated levels of *Ghs-R* we found a specific decrease upon HFD in the dorsal hippocampus
455 while an increase was observed in both the ventral hippocampus and the prefrontal cortex in CTRL-HFD and
456 PNS-CD animals. It has become increasingly apparent that the dorsal and the ventral portions of the
457 hippocampus are preferentially involved in different physiological functions the first playing a role in the
458 regulation of cognitive processes (spatial memory) the second in the modulation of emotions - such that they
459 have been defined the cold and the hot hippocampus respectively - (Fanselow and Dong, 2010). Chen and co-
460 workers provided evidence that infusion of ghrelin in the dorsal hippocampus enhanced synaptic plasticity and
461 spatial memory (Chen et al., 2020). By contrast, Kanoski and colleagues showed that ghrelin delivery to the
462 ventral but not the dorsal hippocampus increased the ability of environmental food-related cues to stimulate

463 meal initiation and to enhance motivation to obtain it by increasing food intake frequency (Kanoski et al.,
464 2013) confirming that this hippocampal sub-region might be of importance in feeding/appetitive rewarding
465 behaviours. To this regard it is interesting to note that we have previously found an increase in caloric intake
466 upon HFD (Panetta et al., 2017). Recently Guo and colleagues have shown that GHS-R knock-out mice were
467 characterised by improved abilities to cope to social stressors, decreased emotionality and depressive-like
468 behaviours (Guo et al., 2019). Thus, we can hypothesize that both the pre-natal and the postnatal stressors
469 might act by increasing the ghrelin signal in two brain regions of main importance for the ability to process
470 emotions (the ventral hippocampus and the prefrontal cortex) setting the stage for increased vulnerability to
471 mood disorders.

472

473 **CONCLUDING REMARKS AND FUTURE DIRECTIONS**

474 We have here provided evidence that PNS can affect brain development (first hit) setting the stage for increased
475 vulnerability to further insults during life (HFD, second hit). In particular, we observed important changes in
476 the expression levels of GR, their chaperons and *Ghs-R* in response to both PNS and HFD, confirming that the
477 there is an important overlap between pathways and effectors involved in the regulation of emotions and in
478 food intake and metabolic balance. **Further investigations are warranted to assess more in detail the role
479 played by these molecules in setting the stage for co-morbidity between metabolic and psychiatric
480 disorders. In addition, the role of additional co-regulators, such as endocannabinoids should be tested
481 (Balsevich et al., 2019).**

482 There are some limitations of the current study that should be mentioned. First, we examined here the
483 expression level of selected genes of interest but did not complement this with the levels of the corresponding
484 proteins, the functional end products of the genes. Secondly, we have used real-time PCR in combination with
485 biopsy punching as a quantitative approach instead of in situ hybridization. Due to this approach we had to
486 make a selection of brain areas and lose anatomical resolution. **Thirdly, genomic and epigenomic regulation
487 of glucocorticoids in the brain may affect mood and metabolism but is based upon the coordinated
488 activity of many chaperone proteins, in addition to proper circadian and ultradian fluctuation of
489 hormone release (Gray et al., 2017). Future studies should assess the interrelationship between
490 psychological and metabolic stressors. In addition, given the observed changes of *Ghs-R* and *Nr3c1* in
491 the dorsal hippocampus following a HFD, further research should be devoted to explore more in depth
492 the effects of HFD and of PNS, and their interaction, on brain plasticity.**

493 **Conflict of Interest Statement**

494 The authors have nothing to disclose.

495

496 **Author's contribution**

497 **AB and CM analysed, interpreted data and wrote the manuscript; MM and AC collected all gene**
498 **expression data; FC and MR designed the experiment and provided data interpretation.**

499

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512

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For Peer Review

695 **FIGURE LEGENDS**

696 **GRAPHICAL ABSTRACT.** Based on the “two hit model” of the developmental origin of diseases, PNS
697 (first hit) affects foetal brain developmental trajectories; signalling pathways related to brain plasticity
698 and function (*Bdnf*, *Tnf- α* and *Il- β*) are down-set potentially providing increased liability to later-life
699 mood and metabolic disorders. A later-life challenge such as HFD feeding (second hit), experienced at
700 adulthood, might impinge upon the same signalling pathways primed by PNS leading to enhanced
701 responses to the metabolic challenge. The hippocampus and prefrontal cortex, being main targets of
702 glucocorticoids, are finely modulated by PNS and readily respond to the HFD metabolic stress.

703

704 **FIGURE 1.** Overall subjects who experienced prenatal stress (PNS-CD) were characterised by decreased m-
705 RNA levels of both *Il-1 β* (panel A) and *Tnf- α* (panel B) in almost all the brain regions investigated; *Il-6* showed
706 a decrease in its expression levels upon HFD in the dorsal hippocampus (panel C). *Il-1 β* showed statistical
707 significance only in the prefrontal cortex while *Tnf- α* was tightly modulated upon pre- and postnatal stressors.
708 Data are presented graphically as means \pm SEM box plot (observations outside the ranges are represented with
709 dots outside the boxes). Post hoc comparisons: **\$\$\$p<0.01, main effect of diet (dorsal hippocampus, *Il-6*);**
710 **post hoc comparisons: *p<0.05, interaction effect (dorsal hippocampus, *Tnf- α* : PNS-CD vs. CTRL-CD);**
711 ****p<0.01, interaction effect (ventral hippocampus, *Tnf- α* : PNS-CD vs. PNS-HFD; Prefrontal cortex, *Tnf-***
712 ***α* and *Il-1 β* : PNS-CD vs. PNS-HFD). Number of subjects: 8-10 within each experimental group.**

713

714 **FIGURE 2.** Male rats exposed to prenatal stress (PNS) were characterised by an inverse relation between
715 levels of *Il-1 β* in the dorsal hippocampus and immobility duration as assessed in the elevated plus maze test
716 (EPM). *p<0.05. Number of subjects: 6-10 within each experimental group.

717

718 **FIGURE 3.** Prenatal stress decreased *Bdnf* m-RNA levels specifically in the dorsal hippocampus suggesting
719 a reduced neuronal plasticity. This effect is reversed by HFD administration at adulthood. Data are presented
720 graphically as means \pm SEM box plot (observations outside the ranges are represented with dots outside the
721 boxes). Post hoc comparisons *p<0.05, **interaction effect in the dorsal hippocampus, PNS-CD vs. CTR-**
722 **CD and PNS-HFD.** Number of subjects is between 9-11 within each experimental group.

723

724 **FIGURE 4.** Male rats exposed to prenatal stress (PNS) were characterised by an inverse relation between
725 levels of *Bdnf* in the dorsal hippocampus and immobility duration as assessed in the elevated plus maze test
726 (EPM). Post hoc comparisons *p<0.05. Number of subjects: 9-10 within each experimental group.

727

728 **FIGURE 5. Independently from the prenatal condition, post-natal HFD administration increases levels**
729 **of *Nr3c1* in the ventral hippocampus and prefrontal cortex by contrast, in the dorsal hippocampus a**
730 **significant interaction between PNS and HFD results in decreased levels of this gene both as a result of**
731 **pre- and post-natal stressors. No change in *Nr3c1* was observed in the hypothalamus as a result of PNS**
732 **nor of HFD (see panel A). As for *Fkbp5*, HFD independently from the prenatal condition, increased its**
733 **expression levels in the dorsal hippocampus while a decrease was observed in the hypothalamus; no**
734 **change was observed in the ventral hippocampus. PNS overall increased levels of *Fkbp5* specifically in**
735 **the prefrontal cortex. $\$p<0.01$, main effect of diet (*Nr3c1*: ventral hippocampus and prefrontal cortex;**
736 ***Fkbp5*: dorsal hippocampus and hypothalamus; *Fkbp5*: dorsal hippocampus; hypothalamus); $\pounds p<0.05$,**
737 **main effect of PNS (*Fkbp5*: prefrontal cortex); post hoc comparisons: $**p<0.01$, interaction effect**
738 **(*Nr3c1*: CTRL-CD vs. CTRL-HFD, PNS-CD and PNS-HFD); Number of subjects:8-10 within each**
739 **experimental group.**

740

741

742 **FIGURE 6. Control subjects are characterized by a direct relationship between expression levels of**
743 ***Fkbp5* in the dorsal hippocampus and the time spent in the open arms of the Elevated Plus Maze, overall**
744 **suggesting a permissive role for explorative behavior of this GC-related genes under physiological**
745 **conditions. Post hoc comparisons $**p<0.01$. Number of subjects: 6-9 within each experimental group.**

746

747

748 **FIGURE 7. Levels of *Ghs-R* were found to be specifically **decreased** upon HFD in the dorsal hippocampus**
749 **while an increase was observed in both the ventral hippocampus and the prefrontal cortex in CTRL-HFD and**
750 **PNS-CD animals. Main effect of HFD: **main effect of diet**, $\$ p<0.01$; post hoc comparisons **for the**
751 **interaction effects: $*p<0.05$: prefrontal cortex, CTRL-CD vs. CTRL-HFD; $**p<0.01$: ventral**
752 **hippocampus, CTRL-CD vs. CTRL-HFD and PNS-CD, prefrontal cortex, CTRL-CD vs. PNS-CD.**
753 **Number of subjects: 8-10 within each experimental group.****

754

1 **High-fat diet at adulthood interacts with prenatal stress, affecting both brain**
2 **inflammatory and neuroendocrine markers in male rats**

3

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25 **Abstract**

26 Prenatal stress (PNS) affects foetal programming and, through an interaction with subsequent challenges, can
27 increase vulnerability to mood and metabolic disorders. We have previously shown that, following PNS, adult
28 male rats are characterised by increased vulnerability to a metabolic stressor experienced at adulthood (8-
29 week-high-fat diet - HFD). In this study, we specifically assessed whether PNS might interact with an adult
30 metabolic challenge to induce an inflammatory phenotype. Changes in the expression levels of inflammatory
31 (*Il-1 β* , *Tnf- α* , *Il-6*) and of stress response mediators (*Nr3c1*, *Fkbp5*) as well as of mood and metabolic
32 regulators (*Bdnf*, *Ghs-R*) were investigated in the hippocampus, prefrontal cortex and hypothalamus, brain
33 regions involved in the pathogenesis of depression and prone to inflammation in response to stress. Overall,
34 PNS reduced the expression of *Bdnf* and *Tnf- α* , while HFD administered at adulthood counteracted this effect
35 suggesting that PNS impinges upon the same pathways regulating responses to a metabolic challenge at
36 adulthood. Furthermore, HFD and PNS affected the expression of both *Nr3c1* and *Fkbp5*, two neuroendocrine
37 mediators involved in the response to stress to metabolic challenges and in the modulation of the emotional
38 profile (as shown by the correlation between *Fkbp5* and the time spent in the open arms of the elevated plus-
39 maze). Overall, these results indicate that the same metabolic and neuroendocrine effectors engaged by PNS
40 are affected by metabolic challenges at adulthood, providing some mechanistic insight into the well-known
41 comorbidity between mood and metabolic disorders.

42

43 **Keywords:** Animal model, Prenatal stress, High-fat diet, Neuroinflammation, Mood disorders, Co-morbidity.

44

45 **LIST OF ABBREVIATIONS**

- 46 BDNF= Brain-Derived Neurotrophic Factor
- 47 CD= Control Diet
- 48 CRP= C-Reactive Protein
- 49 CTRL= control
- 50 EPM= Elevated Plus-Maze
- 51 EU= European
- 52 GC= Glucocorticoids
- 53 GD= Gestational Day
- 54 GHS-R= Ghrelin Receptor
- 55 GR= Glucocorticoid Receptor
- 56 HFD= High-Fat Diet
- 57 HPA= Hypothalamic–Pituitary–Adrenal
- 58 IL= interleukin
- 59 LTP= Long-Term Potentiation
- 60 NF- κ B= Nuclear Factor Kappa-Light-Chain-Enhancer of Activated B Cells
- 61 PND= Post-Natal Day
- 62 PNS= Prenatal Stress
- 63 ROS= Reactive Oxygen Species
- 64 SEM= Standard Error Mean
- 65

66

67 **1. INTRODUCTION**

68 Mood disorders are common conditions with major public health implications. It is estimated that each year
69 around 40% of the EU population suffers from a mental disorder. Adjusted for age and comorbidity, this
70 corresponds to 164.8 million people affected (Health Organization Regional Office for Europe, 2015). Among
71 the risk factors that might trigger the onset of psychiatric conditions, or an exacerbation of symptoms, stressful
72 life events play a pivotal role. In fact, according to a “two-hit model” of vulnerability to diseases, stress
73 experienced during early life phases, might affect brain development leading to a reduced ability to cope with
74 further stressors during life (Daskalakis et al., 2013). To this regard, a suboptimal intrauterine environment,
75 prompted by maternal stress (ranging from poor socio-economic status and maternal obesity to depression or
76 maltreatments) may predispose the offspring to lifelong negative health outcomes, including neuropsychiatric
77 disorders as well as metabolic and immune dysregulations (Barker, 1995; Berry et al., 2015; Boersma et al.,
78 2013; Cattaneo et al., 2020; Cirulli et al., 2020; Krontira et al., 2020).

79 Consistent evidence suggests that depressed patients are characterised by alterations in the functional activity
80 of the immune system, affecting both peripheral and central tissues and that early life stress represents an
81 important factor influencing the inflammatory status (Cattaneo et al., 2015; Dantzer et al., 2008; Dowlati et
82 al., 2010; Liu et al., 2012; Valkanova et al., 2013). As an example, Danese and colleagues reported increased
83 blood C-reactive protein (CRP) levels in maltreated children, an effect that was larger in those subjects that
84 also developed depression later in life (Danese et al., 2009, 2008). Likewise, Slopen and co-workers reported
85 increased CRP and interleukin-6 (IL-6) in adolescents who experienced early life adversities (Slopen et al.,
86 2014). Offspring of women who experienced stressful life events during pregnancy show higher levels of IL-
87 1 β , IL-4, IL-5, IL-6, and IL-8 in umbilical cord blood at delivery (Andersson et al., 2016). Interestingly, it has
88 been recently proposed that inflammation may act as a key mediator linking exposure to prenatal stress (PNS)
89 and enhanced vulnerability to psychopathology in the offspring (Hantsoo et al., 2019).

90 Preclinical studies in rodent models provide strong support to the above mentioned clinical findings, showing
91 for example that PNS exposure causes extended inflammation in the foetal brain (Ślusarczyk et al., 2015). To
92 this regard, Gur and colleagues found increased levels of IL-1 β in the placenta of mouse dams undergoing
93 stress during pregnancy and in the foetal brain of their offspring; this was also associated with decreased levels
94 of the neurotrophin Brain-Derived Neurotrophic Factor (BDNF) specifically in the amygdala (Chen et al.,
95 2020; Gur et al., 2017) suggesting that increased expression of placental immune responsive genes upon
96 maternal stress might be considered as a potential mechanism (Bronson and Bale, 2014; Mueller and Bale,
97 2008).

98 Despite the large body of clinical and preclinical evidence showing associations among PNS, the
99 hyperactivation of the immune system and an enhanced vulnerability to mood disorders, cause-effect
100 mechanisms are still largely unknown, and the prenatal environment may not act alone to set the stage for long-
101 term onset of mood disorders. Interestingly, psychiatric conditions and metabolic pathologies are often found

102 to co-occur within the same individual showing a feed-forward pattern overall suggesting the activation of
103 shared mechanisms/pathways (Cattane et al., 2020; Cirulli et al., 2020; Milaneschi et al., 2019). Among these,
104 BDNF and glucocorticoids (GC) hormones have been identified as main effectors of brain plasticity and
105 metabolic function in response to stressful events. Thus, it is possible that these actors may represent both
106 effectors as well as targets of stress - during sensitive developmental periods - leading, in the developing
107 organism, to a remodelling of the mechanisms associated with stress responsiveness (Cattaneo et al., 2015;
108 Cirulli and Alleva, 2009; McEwen, 2000). The neurotrophin BDNF plays a pivotal role in brain and
109 behavioural plasticity as well as in the control of energy homeostasis (Cirulli and Alleva, 2009; Marosi and
110 Mattson, 2014); its expression is finely tuned by GC in response to acute or chronic stressful stimuli (Cirulli,
111 2017; Cirulli & Alleva, 2009) and it has been also suggested to be negatively regulated by pro-inflammatory
112 cytokines (Barrientos et al., 2004; Bilbo et al., 2008). Thus, BDNF appears as a sensitive target as well as a
113 biomarker of stress response as it could actively contribute to maintain brain health/homeostasis in different
114 challenging conditions. To this regard, the interplay between the activation of stress-related pathways by
115 adverse experiences during development and adult metabolic challenges might be crucial in shaping individual
116 vulnerability to stress (Nederhof and Schmidt, 2012).

117 A growing body of evidence suggests that the effects of PNS are sex-specific and that males' foetal brain might
118 be more vulnerable to changes in the inflammatory mediators overall showing learning deficits and decreased
119 brain plasticity particularly with regard to the hippocampus prefrontal cortex (McCarthy, 2019; Weinstock,
120 2007). Indeed, in a previous study we have shown that male rats who underwent PNS were more vulnerable
121 than females to the effects of a metabolic challenge experienced at adulthood (high-fat diet feeding - HFD),
122 while females showed an overall greater plasticity, possibly mediated by increased total *Bdnf* mRNA
123 expression levels both in the hippocampus and in the hypothalamus. Based on the "two hit model" of diseases,
124 we hypothesized that PNS might disrupt the intrauterine environment affecting brain developmental
125 trajectories leading, in turn, to increased brain inflammation and reduced plasticity in the offspring (first hit).
126 We also hypothesized that a metabolic challenge such as a HFD, experienced at adulthood, might interact with
127 inflammatory pathways to exacerbate the PNS-induced pro-inflammatory phenotype. Thus, the aim of this
128 study was to investigate the effect of a second hit - represented by an HFD - specifically in male rats. We
129 focused on selected brain regions - characterised by high degree of sexual dimorphism (Handa et al., 1994) -
130 such as the dorsal and the ventral hippocampus, the prefrontal cortex and the hypothalamus known to be
131 involved in the pathogenesis of depression and to be prone to inflammatory status in response to external
132 challenges. We specifically assessed in these brain regions levels of *Bdnf*, *Tnf- α* , *Il-6* and *Il- β* .

133 Stress powerfully affects both mood and energy homeostasis. Such effects are achieved not only through the
134 interaction between GC hormones and their receptors (that regulate HPA axis function in addition to glucose
135 metabolism) but also by triggering a multitude of signalling cascades reciprocally modulating one another in
136 a regional-, temporal, and functional-dependent manner (Balsevich et al., 2019). Among these, ghrelin is a
137 peptide hormone produced in the stomach and involved in the signalling of meal initiation. Its action is
138 mediated through the growth hormone secretagogue receptor (Ghs-R) that is found to be highly expressed in

139 several brain regions including the hypothalamus, pituitary gland and hippocampus (see Zarouna, 2015 and
140 references therein). Both Ghrelin and Ghs-R are modulated by stress (Patterson et al., 2010); there is evidence
141 that ghrelin may decrease anxiety-like and depressive-like behaviours in mice (Lutter et al., 2008) moreover,
142 it appears to be involved in food anticipatory activity and consumption (Blum et al., 2009; Verhagen et al.,
143 2011) as well as in mood disorders (Zarouna, 2015). Interestingly, both genetic variants within ghrelin and GC
144 signalling pathways have been associated with obesity, stress-related mental disorders, or both (see Balsevich
145 et al., 2019 and references therein). To this regard, we also investigated changes in the expression levels of
146 *Ghs-R* as well as of *Nr3c1* (encoding for the glucocorticoid receptors - GR) and the GC co-chaperon *Fkbp5*
147 (encoding for the FK506 binding protein 51) in all the above-mentioned brain regions.

148

149 2. MATERIALS AND METHODS

150 2.1 Animals and experimental design

151 Twelve adult nulliparous female (230–260 g) and 6 male Sprague-Dawley rats (400 g) were purchased from a
152 commercial breeder (Charles River, Calco, Italy). Upon arrival, animals were pair-housed with same sex
153 conspecifics under standard laboratory conditions (see Panetta et al. 2015 for further details). Pellet food
154 (Altromin-R, Rieper, Italy) and tap water were continuously available. Following one week of adaptation, two
155 females and one male were mated for 24 hours. To assess pregnancy, changes in body weight were monitored.
156 Pregnant females were randomly assigned to either the control (CTRL) or prenatal stress (PNS) groups. CTRL
157 rats were left undisturbed throughout gestation, while PNS females underwent a repeated restraint stress
158 procedure during the last third of gestation until the day delivery day (postnatal day 0 - PND-0). On PND-1 all
159 pups were weighted and litters culled to an average of 5 male and 5 female pups. On PND-21 pups were
160 weaned and housed in groups of 2 or 3 same-sex littermates until 2 months of age. By this time, n=20 CTRL
161 and n=21 PNS males were assigned either to high-fat diet (HFD) or control diet (CD) regimen for 8 weeks.
162 The final number of animals per group was: n=10 CTRL-CD, n=10 CTRL -HFD, n=10 PNS-CD and n=11
163 PNS-HFD. Groups were composed by 1 pup/litter (CTRL-CD; CTRL-HFD; PNS-CD; PNS-HFD. After 4
164 weeks on the respective diets, all animals underwent a number of metabolic and behavioural assessments; they
165 were successively sacrificed and brains dissected out to investigate the expression levels of *Il-6*, *Il-1 β* , *Tnf- α* ,
166 *Nr3c1*, *Fkbp5*, *Ghs-R*, in the dorsal hippocampus, ventral hippocampus, hypothalamus and prefrontal cortex;
167 total *Bdnf* was investigated only in the dorsal hippocampus and prefrontal cortex (since it was already assessed
168 in the other two areas, (Panetta et al., 2017).

169 To better characterise the effects of possible changes in gene expression levels, regression analyses were
170 carried out by taking into account specific outcomes deriving from the Elevated Plus-Maze (EPM), as
171 previously performed in (Panetta et al., 2017). We focused on immobility time since this is a well-known proxy
172 for emotionality and reactivity to novel environment (Fernandes and File, 1996) and correlated it with *Bdnf*
173 and *Il-1 β* since both these mediators are involved in behavioural plasticity (Bourgognon and Cavanagh, 2020;

174 Cirulli and Alleva, 2009; Goshen et al., 2007). Moreover, we also assessed the ability of *Fkbp5* to affect
175 emotionality in the EPM by correlating expression levels of this gene in the dorsal hippocampus to the time
176 spent in the open arms of the maze. As for the EPM protocol used, briefly, the apparatus was made of Plexiglas
177 and consisted of two opposite open arms and two arms closed by transparent walls (50×10×40 cm). Each rat
178 was placed in the center of the maze and video-recorded for 5 minutes under dim light conditions. Each session
179 was recorded and behavioral analysis was carried out using a commercial software (“The Observer 3.0”,
180 Noldus, The Netherlands) (see (Panetta et al., 2017) for further details).

181 All experimental procedures were reviewed by the ethical body of the Istituto Superiore di Sanità for animal
182 welfare and conducted in conformity with the European Directive 2010/63/EU and the Italian legislation on
183 animal experimentation, D.Lgs. 26/2014. They were authorized by the Italian Ministry of Health.

184

185 **2.2 Dams’ stress procedure**

186 Pregnant females (at gestational day - GD - 14) were restrained in a transparent Plexiglas cylinder (7.5×19 cm)
187 under a bright light (6.500 lux) for 45 minutes three times daily at random times (between 9:00 a.m.-5:00 p.m.)
188 during the dark phase until the expected delivery day (GD-21, see Maccari et al., 1995 for further details). This
189 stress has been selected since it is one of the most well characterised prenatal stressors with well-known effects
190 on developmental trajectories of the offspring.

191

192 **2.3 High-fat diet administration**

193 CTRL and PNS offspring were fed *ad libitum* either with HFD (energy: 5.24 kcal/g; composition: fat 60%,
194 carbohydrate 20% and protein 20%) or CD (energy: 3.3 kcal/g; composition: fat 17%, carbohydrate 60% and
195 protein 23%) starting from two months of age. The HFD diet (D12492) was purchased from Research Diets,
196 Inc., New Brunswick, NJ, USA; the CD was purchased from Altromin-R, Rieper, Italy.

197

198 **2.4 Tissue collection**

199 At 4 months of age offspring was weighted and sacrificed. Brains were removed and the dorsal and ventral
200 hippocampus, hypothalamus and prefrontal cortex were dissected out and immediately frozen at -80°C (Panetta
201 et al., 2017).

202

203

204

205 2.7 Molecular analysis

206 2.7.1 RNA isolation and Real Time PCR Analyses

207 Total RNA was isolated using PureZol RNA isolation reagent (Bio-Rad Laboratories, Italy), treated with
208 DNase to avoid DNA contamination and quantified by spectrophotometric analysis. The quantified RNA was
209 analyzed by TaqMan q-RT PCR Instrument (CFX384 real time system, Bio-Rad Laboratories) using the
210 iScript™ one-step RT-PCR kit for probes (Bio-Rad Laboratories) and Applied BioSystem Assays (Gene
211 Expression Assays: *Il6*, *Il-β*, *Tnf-α*, *Bdnf*, *Nr3c1*, *Fkbp5*, and *Ghs-R*). Samples were run in triplicate and each
212 target gene analyzed has been normalized to the expression of the housekeeping (HK) gene β-actin (ActB).
213 The expression of target genes was calculated by using to the Ct method (-ΔΔCt method) (Schmittgen and
214 Livak, 2008), where CTRL-CD rats have been used as a reference group.

215

216 2.8 Statistical analysis

217 Data were evaluated by a two-way ANOVA with Diet (HFD vs CD) and Prenatal condition (PNS vs CTRL)
218 as between-subject factors. *Post-hoc* comparisons between groups were performed using the Tukey's test. A
219 linear regression model was used to assess the ability of *Bdnf* levels and of *Il-1β* in the dorsal hippocampus to
220 affect immobility duration (a parameter indicative of increased emotionality) in the EPM test. A level of
221 probability set at $p < 0.05$ was used as statistically significant. Statistical tendency was set at $p < 0.1$. For
222 interactions at $p < 0.1$, we also examined lower order effects. Data are presented graphically as means ± SEM
223 box plot (observations outside the ranges are represented with dots outside the boxes). The raw data supporting
224 the conclusions of this article will be made available by the authors upon request without undue reservation.

225

226 3. RESULTS

227

228 3.1 Inflammatory mediators *Il-1β*, *Tnf-α* and *Il-6* mRNA expression

229 *Il-1β* - We first analyzed the expression of *Il-1β*, a prototype pro-inflammatory cytokine. We observed that
230 PNS resulted in a nearly significant decrease in the dorsal hippocampus while neither a main effect of HFD
231 nor a PNS x HFD interaction effect were found ($F(1,35)=3.818, 0.004, 2.184; p=0.0587, 0.9518, 0.1484$).
232 No main effects of PNS nor of HFD were observed in the ventral hippocampus ($F(1,34)=0.152, 0.335;$
233 $p=0.6995, 0.5665$). A strong PNS x HFD interaction was found ($F(1,34)=4.655; p=0.0381$) PNS exposed
234 animals being characterised by decreased levels of this cytokine, while the administration of postnatal HFD
235 counteracted the effects of the PNS (post-hoc comparisons just missed statistical significance).

236 In the prefrontal cortex while no main effects of PNS was observed ($F(1,33)=0.969; p=0.3321$) postnatal HFD
237 overall increased levels of *Il-1β* (main effect of HFD: $F(1,33)=5.112, p=0.0305$). Similarly to what was
238 observed in the ventral hippocampus, a nearly significant PNS x HFD interaction was found with decreased

239 levels of *Il-1 β* in PNS animals, this effect being reverted by postnatal administration of HFD ($F(1,33)=3.858$;
240 $p=0.0580$, post hoc comparisons: $p<0.01$, PNS vs. PNS-HFD); (see Figure 1A).

241 In the hypothalamus, levels of *Il-1 β* were not affected by PNS, HFD nor an interaction was found
242 ($F(1,36)=0.042, 0.0001633, 0.070$; $p=0.8386, 0.9899, 0.7934$ respectively for PNS, HFD and PNS x HFD, see
243 Figure 1A).

244

245 *Tnf- α* - In the dorsal hippocampus no main effects of PNS nor of HFD were observed ($F(1,37)=1.288, 0.209$;
246 $p=0.2638, 0.6506$). By contrast, a significant interaction was found between PNS and HFD ($F(1,37)=5.772$;
247 $p=0.0214$). In particular, PNS animals were characterised by decreased levels of *Tnf- α* , although this effect
248 was no longer observed in PNS-HFD animals that showed levels of this cytokine comparable to those observed
249 in CTRL animals (post hoc comparisons: $p<0.05$, PNS vs. CTRL).

250 No main effects of PNS nor of HFD were found in the ventral hippocampus ($F(1,34)=0.033, 2.941$; $p=0.8573,$
251 0.0955 , respectively for PNS and HFD), while a PNS x HFD interaction was found showing that administration
252 of postnatal HFD in the PNS group selectively increased levels of this cytokine ($F(1,34)=6.747$; $p=0.0138$);
253 post hoc comparisons: PNS vs. HFD-PNS, $p<0.01$).

254 No main effects of PNS nor of HFD were observed in the prefrontal cortex ($F(1,33)=0.079, 1.648$; $p=0.7809,$
255 0.2082 respectively for PNS and HFD) however, as also observed for the ventral hippocampus a PNS x HFD
256 interaction effect was found showing that the administration of postnatal HFD in the PNS group selectively
257 increased levels of this cytokine ($F(1,33)=7.946$; $p=0.0081$); post hoc comparisons: PNS vs. HFD-PNS,
258 $p<0.01$); (see Figure 1B).

259 Levels of *Tnf- α* in the hypothalamus (see Figure 1B) were not affected neither by PNS nor by postnatal HFD
260 ($F(1, 35)=0.286, 0.702, 2.111$; $p=0.5964; 0.4079; 0.1551$ respectively for PNS, HFD and PNS x HFD).

261

262 *Il-6* - Postnatal HFD decreased levels of *Il-6* in the dorsal hippocampus (main effect of HFD: $F(1,37)=0.814$;
263 $p=0.0066$), whereas no effect of the PNS nor of the PNS x HFD interaction effect were found (PNS main
264 effect: $F(1,37)=1.387$; $p=0.2465$; PNS x HFD interaction: $F(1,37)=0.164$; $p=0.6881$).

265 Levels of *Il-6* did not change as a result of PNS, HFD or their interaction neither in the ventral hippocampus
266 ($F(1,36)=0.759, 2.631, 0.029$; $p=0.3893, 0.1135, 0.8647$ respectively for PNS, HFD and PNS x HFD) nor in
267 the prefrontal cortex ($F(1,35)=0.144, 0.053, 0.202$ respectively for PNS, HFD and PNS x HFD); (see Figure
268 1C). PNS decreased nearly significantly the levels of *Il-6* also in the hypothalamus ($F(1,36)=3.889$; $p=0.0563$),
269 but no effect of HFD nor an interaction PNS x HFD were found to be significant ($F(1,36)=0.903, 0.757$;
270 $p=0.3482, 0.3899$, respectively for HFD and PNS x HFD).

271

272 -----FIGURE 1 about here-----

273

274 Interestingly, when expression levels of *Il-1 β* in the dorsal hippocampus were related to immobility duration
275 in the EPM (a test run in a companion paper on the same experimental animals, see (Panetta et al., 2017)), we

276 observed a significant strong negative association linking these two parameters only in the PNS-CD animals,
 277 overall suggesting that reduced levels of this cytokine can be related with reduced behavioral plasticity (see
 278 discussion; CTRL-CD: $F(1,8)=0.007$; $p=0.9367$; $R^2=0.01$; CTRL-HFD: $F(1,8)=3.461$; $p=0.1051$; $R^2=33.1$;
 279 PNS-CD: $F(1,8)=7.538$; $p=0.0287$; $R^2=51.8$; PNS-HFD: $F(1,9)=1.739$; $p=0.2237$, $R^2=17.9$; see Figure 2).

280

281 -----FIGURE 2 about here-----

282

283 3.2 Neuronal plasticity marker - *Bdnf* mRNA expression

284 We first investigated the expression of *Bdnf*, a neurotrophin that is considered a prototype marker of plasticity.

285 Within the dorsal hippocampus, overall PNS and postnatal HFD did not affect levels of *Bdnf* ($F(1, 36)=3.678$;

286 1.964 ; $p=0.0631$; 0.1696), main effect of PNS and HFD respectively), although we observed a significant

287 interaction between PNS and postnatal HFD ($F(1, 36)=5.606$; $p=0.0234$). More in detail, PNS decreased *Bdnf*

288 expression, while the postnatal administration of HFD was able to counteract this effect (PNS vs. CTRL,

289 $p<0.05$; PNS vs. PNS-HFD, $p<0.05$). In the prefrontal cortex no effect of PNS, HFD or their interaction was

290 observed ($F(1, 36)=0.200$, 1.377 , 0.174 ; $p=0.6572$; 0.2482 ; 0.6794 , respectively for PNS, HFD and PNS x

291 HFD); (see Figure 3).

292

293 -----FIGURE 3 about here-----

294

295 Interestingly, levels of *Bdnf* in the dorsal hippocampus were negatively associated with the time spent

296 immobile in the EPM (a test run in a companion paper on same experimental animals, see (Panetta et al.,

297 2017)) only in the PNS-CD group (CTRL-CD: $F(1,9)=0.730$; $p=0.4261$; $R^2=0.081$; CTRL-HFD: $F(1,9)=$

298 0.090 ; $p=0.7713$; $R^2=0.011$; PNS-CD: $F(1,8)=11.084$; $p=0.0127$; $R^2=0.612$; PNS-HFD: $F(1,9)=0.716$;

299 $p=0.4221$; $R^2=0.082$), supporting a role of this neurotrophin in the emotional phenotype (see Figure 4).

300

301 -----FIGURE 4 about here-----

302

303 3.3 Stress response mediators - *Nr3c1*, *Fkbp5* mRNA expression

304 *Nr3c1* - In the dorsal hippocampus the postnatal HFD decreased the levels of *Nr3c1* when compared to controls

305 and a similar nearly significant effect was observed for PNS (main effect: $F(1,36)=3.827$, 7.063 , $p=0.0582$;

306 0.0117 respectively for PNS, HFD). Moreover, a significant PNS x HFD interaction was found showing that

307 both PNS-CD and CTRL-HFD as well as the combination PNS-HFD were characterised by a decrease of GR

308 mRNA levels ($F(1,36)=6.228$; $p=0.0173$ and PNS x HFD; post hoc comparisons $p<0.05$).

309 In the ventral hippocampus levels of *Nr3c1* were increased a result of postnatal HFD (main effect:

310 $F(1,36)=10.405$; $p=0.0027$), while no effects of PNS nor of the PNS x HFD interaction were found ($F(1,$

311 $36)=0.011$, 0.258 ; $p=0.9167$; 6146). In the prefrontal cortex the postnatal HFD overall increased the levels of

312 *Nr3c1* (main effect of HFD: $F(1, 37)=12.453$; $p=0.0011$); no effects of PNS nor of the PNS x HFD interaction

313 were observed ($F(1, 37)=1.001, 0.702; p=0.3236; 0.4076$). No effects of PNS, HFD nor of their interaction
 314 were found in the hypothalamus ($F(1, 37)=0.157, 0.009, 1.067; p= 0.6943; 0.9257; 0.3084$); (see Figure 5).

315

316 *Fkbp5* - In the dorsal hippocampus the postnatal HFD increased the levels of *Fkbp5*
 317 ($F(1,35)=10.331;p=0.0028$), while no effect of PNS nor of the PNS x HFD interaction was observed
 318 ($F(1,35)=1.502, 0.958; p= 0.2285; 0.3343$). No effect of PNS, HFD or their interaction was found in the ventral
 319 hippocampus ($F(1,36)=0.781, 3.222, 0.007; p=0.3828; 0.0811; 0.9359$). In the prefrontal cortex, PNS increased
 320 the levels of *Fkbp5* ($F(1,37)=5.244; p=0.0278$), while no effect of HFD nor of the PNS x HFD interaction was
 321 observed ($F(1,37)=1.219, 2.019; p=0.2767; 0.1637$). A main effect of the postnatal HFD was also observed In
 322 the hypothalamus, the Postnatal HFD decreased levels of *Fkbp5* ($F(1,37)=9.156; p=0.0045$); no effect of PNS
 323 nor of the PNS x HFD interaction was observed ($F(1,37)=0.094, 3.353; p= 0.7608; 0.0751$); (see Figure 5).

324

325 -----**FIGURES 5 about here**-----

326

327 Interestingly, levels of *Fkbp5* in the dorsal hippocampus were positively related with increased time spent in
 328 the open arm of the EPM (a test run in a companion paper on same experimental animals, see (Panetta et al.,
 329 2017)) only in the CTRL-CD group (CTRL-CD: $F(1,8)=26.617; p=0.0013; R^2= 0.792$; CTRL-HFD: $F(1,9)=$
 330 $0.104; p=0.7713; R^2=0.013$; PNS-CD: $F(1,8)=0.253; p=0.6307; R^2=0.035$; PNS-HFD: $F(1,9)=0.503;$
 331 $p=0.6307; R^2= 0.4982$). This result might suggest that *Fkbp5* is physiologically required to engage animals in
 332 the proper exploration of novel environments through a fine modulation of GR receptors (see Figure 6).

333

334 -----**FIGURES 6 about here**-----

335

336 **3.4 Metabolic/mood regulator marker - *Ghs-R* mRNA expression**

337 *Ghs-R* - In the dorsal hippocampus the postnatal HFD overall decreased levels of *Ghs-R* ($F(1,33)=1,948;$
 338 $p=0.0007$). By contrast, PNS increased *Ghs-R* levels ($F(1,33)=5.192; p=0.0293$), but no interaction between
 339 PNS x HFD was observed ($F(1,33)=0.290; p= 0.5941$). In the ventral hippocampus HFD overall increased the
 340 levels of *Ghs-R* ($F(1,32)=6.278; p=0.0175$), while no main effect of PNS was observed ($F(1,32)=0.593;$
 341 $p=0.4468$). However, *Ghs-R* was increased in the CTRL-HFD and in the PNS-CD groups (interaction between
 342 PNS and HFD: $F(1,32)=14.778; p=0.0005$, post hoc: CTRL vs. CTRL-HFD, $p<0.01$; CTRL vs. PNS, $p<0.01$).
 343 In the prefrontal cortex, PNS increased the levels of *Ghs-R* ($F(1,29)=5.865; p=0.0219$) and no effect of HFD
 344 was observed ($F(1,29)=2.191; p=9.1496$). However, in this brain area, *Ghs-R* was increased in the CTRL-HFD
 345 and in the PNS-CD groups (interaction between PNS and HFD: $F(1,29)=6.870; p= 0.0138$, post hoc: CTRL
 346 vs. CTRL-HFD, $p<0.05$; CTRL vs. PNS, $p<0.01$); (see Figure 7). As for the hypothalamus, neither PNS nor
 347 HFD affected levels of *Ghs-R* ($F(1,34)=3.155, 3.033, 0.907; p=0.0846; 0.0906; 0.3477$).

348

349 -----FIGURE 7 about here-----

350

351 4. DISCUSSION

352 In the present study we assessed the long-term effects of exposure to PNS (a first hit) on foetal brain
353 programming with regard to neuronal plasticity and brain inflammation. Moreover, we used a metabolic
354 challenge (HFD) at adulthood (second hit) to assess potential interaction effects with the PNS-induced
355 phenotype. We have previously shown (Panetta et al., 2017) that although PNS resulted in a strong metabolic
356 liability in male rats fed with HFD at adulthood it also dampened the negative effects driven by such a
357 metabolic challenge on the emotional phenotype as assessed in the EPM. Here we extended such body of
358 evidence by providing information on the possible mechanisms underlying this interaction. PNS-CD rats were
359 characterised by reduced brain plasticity showing decreased expression levels of the neurotrophin *Bdnf* and of
360 pro-inflammatory cytokines in a number of brain regions; moreover, HFD feeding, experienced at adulthood,
361 induced an increase in pro-inflammatory cytokines in those animals that had experienced PNS. This suggests
362 that PNS might impinge upon the same mechanisms underlying vulnerability to metabolic challenges at
363 adulthood. Furthermore, HFD greatly affected the expression levels of the main effectors of HPA axis function
364 (the glucocorticoid receptor - GR - gene *Nr3c1* and of the GR's co-chaperone *Fkbp5*) an effect possibly
365 underlying the metabolic vulnerability previously observed in male rats.

366 In our study, we found that PNS animals were characterised by differential expression of mRNA levels of *Il-*
367 *1β* and *Tnf-α* in almost all the brain regions investigated except for the hypothalamus, an area that, in our
368 experimental condition, appeared to be resilient to inflammatory changes driven by both pre- and postnatal
369 stressors. While *Tnf-α* was tightly modulated upon pre- and postnatal stressors, changes in *Il-1β* were observed
370 only when considering the interaction with the second hit (HFD), suggesting that the former cytokine (*Tnf-α*)
371 might be considered as a reliable marker of stress adaptation in response to a prenatal challenge. As for *Il-6*,
372 no effect was observed upon PNS in neither of the brain areas considered, though HFD overall decreased its
373 expression levels in the dorsal hippocampus. TNF-α, IL-1β and IL-6 are the main activators of the HPA and
374 are in turn modulated (inhibited) by GC hormones. As described by O'Connors and colleagues these cytokines
375 hold differential sensitivity for adrenal steroids with TNF-α being the most sensitive to such inhibition in the
376 range of physiological levels, IL-1β being second and IL-6 being the most resistant (O'Connor et al., 2000).
377 There is general consensus that elevated inflammation can exacerbate or even give rise to depressive symptoms
378 or may be associated to other psychiatric conditions (Dantzer et al., 2018, 2008). However, cytokines are also
379 constitutively released in the healthy brain by resident myeloid cells to keep proper synaptic plasticity. As an
380 example, the modulation of both IL-1β and TNF-α plays an important role in the processes of LTP and synaptic
381 scaling (a form of homeostatic plasticity) (Rizzo et al., 2018; Salim et al., 2012). Here we found that lower
382 mRNA levels of *Il-1β* in the dorsal hippocampus of PNS rats were associated to increased time spent immobile
383 in the EPM (as previously assessed in (Panetta et al., 2017)). Moreover, and in line with our previous work
384 (Panetta et al., 2017), PNS rats were characterised by decreased levels of the neurotrophin *Bdnf* (in the dorsal

385 hippocampus); this decrease was also associated to an increase in the time spent immobile in the EPM,
386 suggesting that PNS might affect foetal brain programming by reducing neuronal plasticity. Interestingly, the
387 interaction between PNS and adult HFD (PNS-HFD group) resulted in increased levels of *Il-1 β* and *Tnf- α*
388 mRNA. Alboni and colleagues have recently provided evidence for increased rather than decreased levels of
389 pro-inflammatory cytokines in the brain of stressed mice treated with the antidepressant fluoxetine. Such an
390 effect was associated to increased BDNF levels and stronger LTP in the hippocampus (Alboni et al., 2017,
391 2016), suggesting that brain plasticity may also be related to the activation of basal metabolism that in turn is
392 positively associated to the ability to properly mount and control inflammatory responses. To this regard, we
393 cannot exclude that prolonged exposure to the HFD or to a stronger metabolic insult, such as a western pattern
394 diet (rich in fats and sugar), should lead to an overall excessive brain inflammation (not observed in this study)
395 with main consequences on emotional/cognitive behaviour. Worth to notice, we have previously shown that
396 upon PNS, male rats were characterized by increased corticosterone levels under basal conditions, this effect
397 was associated with a decrease in reactive oxygen species (ROS) as well as a with decreased NF- κ B signalling
398 in the hippocampus suggesting a lower set-point under basal conditions in PNS male rats (Anacker et al.,
399 2013). Here we observed a similar blunted activation upon PNS with HFD triggering a response only in PNS
400 subjects (increased *Il-1 β* and *Tnf- α* in the prefrontal cortex and *Tnf- α* in the dorsal and ventral hippocampus).
401 To our knowledge this is one of the few instances in which a reduction in inflammatory mediators is described
402 a result of PNS. Because cytokines have been shown to modulate hippocampal development and plasticity
403 (Bourgognon and Cavanagh, 2020; Goshen et al., 2007) a decreased expression profile as a result of a “first
404 hit” (PNS) might set the stage for an increased response to a “second hit” (metabolic challenge). Obesity is
405 characterised by low-grade systemic and central inflammation and TNF- α and IL-1 β appear to be key players
406 in this condition and in the associated pathologies (Mighiu et al., 2012). Although the hypothalamus is a well-
407 known area playing a main role in the crosstalk between brain and periphery in metabolic pathologies
408 associated to obesity, other brain regions have been shown to be affected by high-fat diet- or obesity-induced
409 inflammation such as the hippocampus, cortex, brainstem, or amygdala (Guillemot-Legris and Muccioli,
410 2017). There are data to suggest that the effects of a metabolic challenge may be different depending on the
411 region of interest dealing with brain plasticity (hippocampus) or metabolic regulation and homeostasis
412 (hypothalamus) (Rasgon and McEwen, 2016). Thus, our data also indicate that PNS might set the stage for a
413 differential regulation of central inflammatory mediators in specific brain regions, an effect that can be
414 unmasked only upon the occurrence of a second hit, such as the HFD at adult age.

415 Hyperactivity of the HPA axis and increased circulating GC have been implicated in the pathogenesis of mood
416 and anxiety disorders suggesting an impairment in the ability of the HPA axis to self-regulate its function by
417 shutting down the system (Binder, 2009; Holsboer, 2000; Pariante and Miller, 2001). Glucocorticoids have
418 also an important function in metabolic regulations as they mobilize glucose to fuel the energy demands of the
419 stress response and furthermore promote energy storage, feeding, and weight gain. Thus, it is possible to
420 hypothesize that both psychological as well as metabolic stress might be able to affect HPA axis functionality
421 (Balsevich et al., 2019). A growing body of evidence suggests that levels of *Fkbp5* mRNA GR co-chaperone

422 have been associated with higher levels of circulating cortisol and reduced negative feedback inhibition of the
423 stress response associated with a depressive phenotype (Binder, 2009). We have previously shown that,
424 following PNS, male rats were characterised by elevated GC levels when compared to controls (Anacker et
425 al., 2013); in this study we show elevated levels of *Fkbp5* upon PNS in the prefrontal cortex and upon HFD
426 administration in the dorsal hippocampus, suggesting that both psychological and metabolic stressors affect
427 this mediator. When we looked at the hypothalamus, *Fkbp5* expression levels were reduced in the HFD group,
428 possibly accounting for a compensatory mechanism. Interestingly, we also showed that levels of *Fkbp5* in the
429 dorsal hippocampus were positively related to the time spent in the open arm of the EPM and that this positive
430 correlation is lost as a result both of PNS as well as of HFD. This result might suggest that under physiological
431 conditions *Fkbp5* is required to engage animals in the proper exploration of novel environments through a fine
432 modulation of the GR receptors and that this balance can be greatly affected by both early life stressors as well
433 as by metabolic challenges.

434 The prefrontal cortex is one of the most important cortical area among the network of regions being involved
435 in the pathogenesis of depression that plays a main role in the ability to process positive and negative emotions
436 (Harms et al., 2017; Kaya and McCabe, 2019). Thus, the observed increase in levels of *Fkbp5* in this brain
437 area should not be surprising and may underlie a condition of potential increased vulnerability to stress.

438 Changes in *Nr3c1* and *Fkbp5* in the hippocampus, a result of HFD, indicate a cross-regulation between a
439 metabolic challenge and *Nr3c1/Fkbp5* balance, strengthening the notion that indeed these molecules are
440 involved in “sensing” environmental challenges, be these of psychological or metabolic nature. We also found
441 that HFD increased the levels of *Nr3c1* in the prefrontal cortex; by contrast, changes in *Fkbp5* in this brain
442 area were observed only as a result of PNS. These results possibly suggest a fine and complex multiple level
443 regulation of the GR-negative feedback in this specific brain region highlighting the role of this brain area in
444 mood disorders such as depression. Transcriptional control of GR relies in part upon the DNA methylation
445 status at multiple alternative initiation sites that are tissue specific in fact, *Nr3c1* gene is characterised by an
446 unusually complex promoter structure (Turner et al., 2010). This might possibly explain the unexpected
447 decrease in the dorsal hippocampus observed as a result of HFD or the lack of effects observed in the
448 hypothalamus (see also concluding remarks).

449 When we investigated levels of *Ghs-R* we found a specific decrease upon HFD in the dorsal hippocampus
450 while an increase was observed in both the ventral hippocampus and the prefrontal cortex in CTRL-HFD and
451 PNS-CD animals. It has become increasingly apparent that the dorsal and the ventral portions of the
452 hippocampus are preferentially involved in different physiological functions the first playing a role in the
453 regulation of cognitive processes (spatial memory) the second in the modulation of emotions - such that they
454 have been defined the cold and the hot hippocampus respectively - (Fanselow and Dong, 2010). Chen and co-
455 workers provided evidence that infusion of ghrelin in the dorsal hippocampus enhanced synaptic plasticity and
456 spatial memory (Chen et al., 2020). By contrast, Kanoski and colleagues showed that ghrelin delivery to the
457 ventral but not the dorsal hippocampus increased the ability of environmental food-related cues to stimulate
458 meal initiation and to enhance motivation to obtain it by increasing food intake frequency (Kanoski et al.,

459 2013) confirming that this hippocampal sub-region might be of importance in feeding/appetitive rewarding
460 behaviours. To this regard it is interesting to note that we have previously found an increase in caloric intake
461 upon HFD (Panetta et al., 2017). Recently Guo and colleagues have shown that GHS-R knock-out mice were
462 characterised by improved abilities to cope to social stressors, decreased emotionality and depressive-like
463 behaviours (Guo et al., 2019). Thus, we can hypothesize that both the pre-natal and the postnatal stressors
464 might act by increasing the ghrelin signal in two brain regions of main importance for the ability to process
465 emotions (the ventral hippocampus and the prefrontal cortex) setting the stage for increased vulnerability to
466 mood disorders.

467

468 **CONCLUDING REMARKS AND FUTURE DIRECTIONS**

469 We have here provided evidence that PNS can affect brain development (first hit) setting the stage for increased
470 vulnerability to further insults during life (HFD, second hit). In particular, we observed important changes in
471 the expression levels of GR, their chaperons and *Ghs-R* in response to both PNS and HFD, confirming that the
472 there is an important overlap between pathways and effectors involved in the regulation of emotions and in
473 food intake and metabolic balance. Further investigations are warranted to assess more in detail the role played
474 by these molecules in setting the stage for co-morbidity between metabolic and psychiatric disorders. In
475 addition, the role of additional co-regulators, such as endocannabinoids should be tested (Balsevich et al.,
476 2019).

477 There are some limitations of the current study that should be mentioned. First, we examined here the
478 expression level of selected genes of interest but did not complement this with the levels of the corresponding
479 proteins, the functional end products of the genes. Secondly, we have used real-time PCR in combination with
480 biopsy punching as a quantitative approach instead of in situ hybridization. Due to this approach we had to
481 make a selection of brain areas and lose anatomical resolution. Thirdly, genomic and epigenomic regulation
482 of glucocorticoids in the brain may affect mood and metabolism but is based upon the coordinated activity of
483 many chaperone proteins, in addition to proper circadian and ultradian fluctuation of hormone release (Gray
484 et al., 2017). Future studies should assess the interrelationship between psychological and metabolic stressors.
485 In addition, given the observed changes of *Ghs-R* and *Nr3c1* in the dorsal hippocampus following a HFD,
486 further research should be devoted to explore more in depth the effects of HFD and of PNS, and their
487 interaction, on brain plasticity.

488 **Conflict of Interest Statement**

489 The authors have nothing to disclose.

490

491 **Author's contribution**

492 **AB and CM analysed, interpreted data and wrote the manuscript; MM and AC collected all gene**
493 **expression data; FC and MR designed the experiment and provided data interpretation.**

494

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507

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For Peer Review

690 **FIGURE LEGENDS**

691 **GRAPHICAL ABSTRACT.** Based on the “two hit model” of the developmental origin of diseases, PNS
692 (first hit) affects foetal brain developmental trajectories; signalling pathways related to brain plasticity
693 and function (*Bdnf*, *Tnf- α* and *Il- β*) are down-set potentially providing increased liability to later-life
694 mood and metabolic disorders. A later-life challenge such as HFD feeding (second hit), experienced at
695 adulthood, might impinge upon the same signalling pathways primed by PNS leading to enhanced
696 responses to the metabolic challenge. The hippocampus and prefrontal cortex, being main targets of
697 glucocorticoids, are finely modulated by PNS and readily respond to the HFD metabolic stress.

698

699 **FIGURE 1.** Overall subjects who experienced prenatal stress (PNS-CD) were characterised by decreased m-
700 RNA levels of both *Il-1 β* (panel A) and *Tnf- α* (panel B) in almost all the brain regions investigated; *Il-6* showed
701 a decrease in its expression levels upon HFD in the dorsal hippocampus (panel C). *Il-1 β* showed statistical
702 significance only in the prefrontal cortex while *Tnf- α* was tightly modulated upon pre- and postnatal stressors.
703 Data are presented graphically as means \pm SEM box plot (observations outside the ranges are represented with
704 dots outside the boxes). Post hoc comparisons: **\$\$\$p<0.01, main effect of diet (dorsal hippocampus, *Il-6*);**
705 **post hoc comparisons: *p<0.05, interaction effect (dorsal hippocampus, *Tnf- α* : PNS-CD vs. CTRL-CD);**
706 ****p<0.01, interaction effect (ventral hippocampus, *Tnf- α* : PNS-CD vs. PNS-HFD; Prefrontal cortex, *Tnf-***
707 ***α* and *Il-1 β* : PNS-CD vs. PNS-HFD). Number of subjects: 8-10 within each experimental group.**

708

709 **FIGURE 2.** Male rats exposed to prenatal stress (PNS) were characterised by an inverse relation between
710 levels of *Il-1 β* in the dorsal hippocampus and immobility duration as assessed in the elevated plus maze test
711 (EPM). *p<0.05. Number of subjects: 6-10 within each experimental group.

712

713 **FIGURE 3.** Prenatal stress decreased *Bdnf* m-RNA levels specifically in the dorsal hippocampus suggesting
714 a reduced neuronal plasticity. This effect is reversed by HFD administration at adulthood. Data are presented
715 graphically as means \pm SEM box plot (observations outside the ranges are represented with dots outside the
716 boxes). Post hoc comparisons *p<0.05, **interaction effect in the dorsal hippocampus, PNS-CD vs. CTR-**
717 **CD and PNS-HFD.** Number of subjects is between 9-11 within each experimental group.

718

719 **FIGURE 4.** Male rats exposed to prenatal stress (PNS) were characterised by an inverse relation between
720 levels of *Bdnf* in the dorsal hippocampus and immobility duration as assessed in the elevated plus maze test
721 (EPM). Post hoc comparisons *p<0.05. Number of subjects: 9-10 within each experimental group.

722

723 **FIGURE 5. Independently from the prenatal condition, post-natal HFD administration increases levels**
724 **of *Nr3c1* in the ventral hippocampus and prefrontal cortex by contrast, in the dorsal hippocampus a**
725 **significant interaction between PNS and HFD results in decreased levels of this gene both as a result of**
726 **pre- and post-natal stressors. No change in *Nr3c1* was observed in the hypothalamus as a result of PNS**
727 **nor of HFD (see panel A). As for *Fkbp5*, HFD independently from the prenatal condition, increased its**
728 **expression levels in the dorsal hippocampus while a decrease was observed in the hypothalamus; no**
729 **change was observed in the ventral hippocampus. PNS overall increased levels of *Fkbp5* specifically in**
730 **the prefrontal cortex. $\$p<0.01$, main effect of diet (*Nr3c1*: ventral hippocampus and prefrontal cortex;**
731 ***Fkbp5*: dorsal hippocampus and hypothalamus; *Fkbp5*: dorsal hippocampus; hypothalamus); $\pounds p<0.05$,**
732 **main effect of PNS (*Fkbp5*: prefrontal cortex); post hoc comparisons: $**p<0.01$, interaction effect**
733 **(*Nr3c1*: CTRL-CD vs. CTRL-HFD, PNS-CD and PNS-HFD); Number of subjects:8-10 within each**
734 **experimental group.**

735

736

737 **FIGURE 6. Control subjects are characterized by a direct relationship between expression levels of**
738 ***Fkbp5* in the dorsal hippocampus and the time spent in the open arms of the Elevated Plus Maze, overall**
739 **suggesting a permissive role for explorative behavior of this GC-related genes under physiological**
740 **conditions. Post hoc comparisons $**p<0.01$. Number of subjects: 6-9 within each experimental group.**

741

742

743 **FIGURE 7. Levels of *Ghs-R* were found to be specifically **decreased** upon HFD in the dorsal hippocampus**
744 **while an increase was observed in both the ventral hippocampus and the prefrontal cortex in CTRL-HFD and**
745 **PNS-CD animals. Main effect of HFD: **main effect of diet**, $\$ p<0.01$; post hoc comparisons **for the****

746 **interaction effects: $*p<0.05$: prefrontal cortex, CTRL-CD vs. CTRL-HFD; $**p<0.01$: ventral**
747 **hippocampus, CTRL-CD vs. CTRL-HFD and PNS-CD, prefrontal cortex, CTRL-CD vs. PNS-CD.**
748 **Number of subjects: 8-10 within each experimental group.**

749

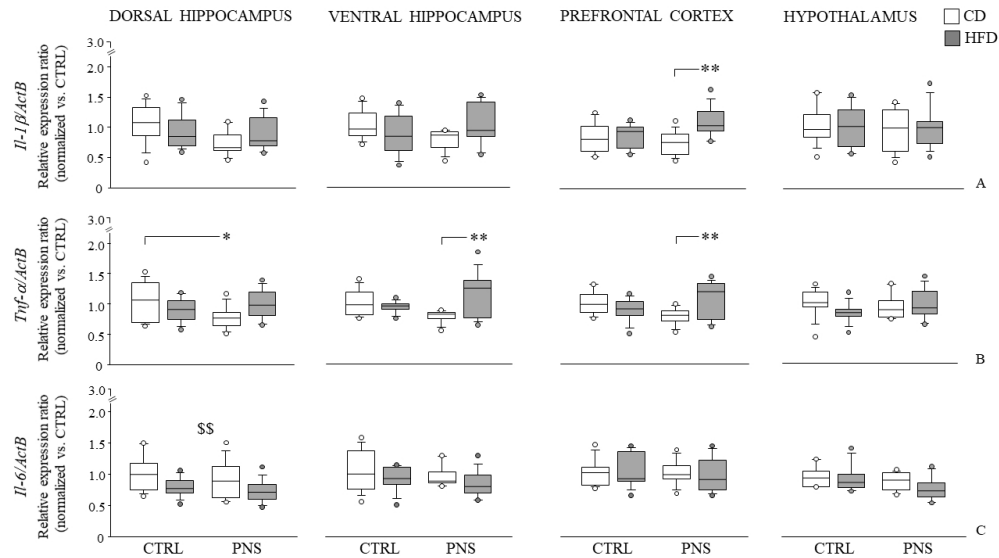


FIGURE 1. Overall subjects who experienced prenatal stress (PNS-CD) were characterised by decreased mRNA levels of both *Il-1beta* (panel A) and *Tnf-alpha* (panel B) in almost all the brain regions investigated; *Il-6* showed a decrease in its expression levels upon HFD in the dorsal hippocampus (panel C). *Il-1beta* showed statistical significance only in the prefrontal cortex while *Tnf-alpha* was tightly modulated upon pre- and postnatal stressors. Data are presented graphically as means \pm SEM box plot (observations outside the ranges are represented with dots outside the boxes). Post hoc comparisons: \$\$ p <0.01, main effect of diet (dorsal hippocampus, *Il-6*); post hoc comparisons: * p <0.05, interaction effect (dorsal hippocampus, *Tnf-alpha*: PNS-CD vs. CTRL-CD); ** p <0.01, interaction effect (ventral hippocampus, *Tnf-alpha*: PNS-CD vs. PNS-HFD; Prefrontal cortex, *Tnf-alpha* and *Il-1beta*: PNS-CD vs. PNS-HFD). Number of subjects: 8-10 within each experimental group.

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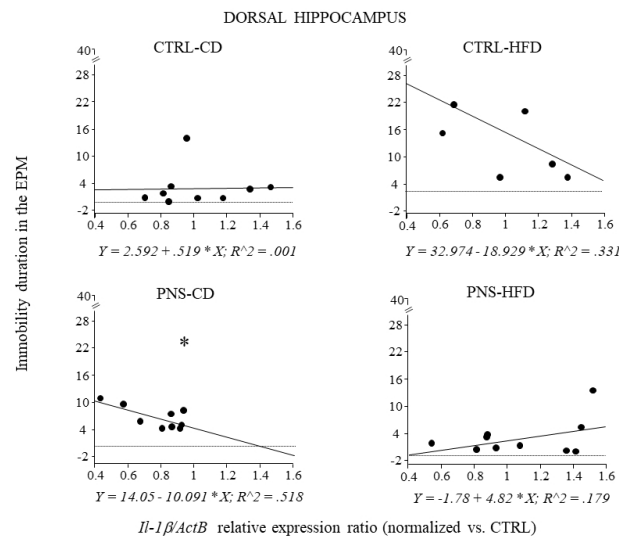


FIGURE 2. Male rats exposed to prenatal stress (PNS) were characterised by an inverse relation between levels of *Il-1beta* in the dorsal hippocampus and immobility duration as assessed in the elevated plus maze test (EPM). * $p < 0.05$. Number of subjects: 6-10 within each experimental group.

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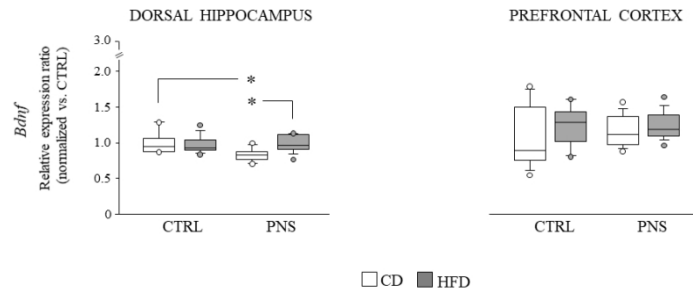


FIGURE 3. Prenatal stress decreased *Bdnf* m-RNA levels specifically in the dorsal hippocampus suggesting a reduced neuronal plasticity. This effect is reversed by HFD administration at adulthood. Data are presented graphically as means \pm SEM box plot (observations outside the ranges are represented with dots outside the boxes). Post hoc comparisons $*p < 0.05$, interaction effect in the dorsal hippocampus, PNS-CD vs. CTRL-CD and PNS-HFD. Number of subjects is between 9-11 within each experimental group.

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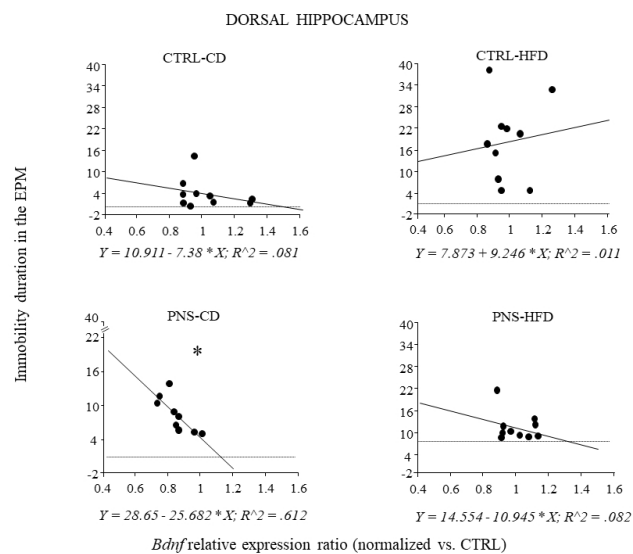


FIGURE 4. Male rats exposed to prenatal stress (PNS) were characterised by an inverse relation between levels of *Bdnf* in the dorsal hippocampus and immobility duration as assessed in the elevated plus maze test (EPM). Post hoc comparisons $*p < 0.05$. Number of subjects: 9-10 within each experimental group.

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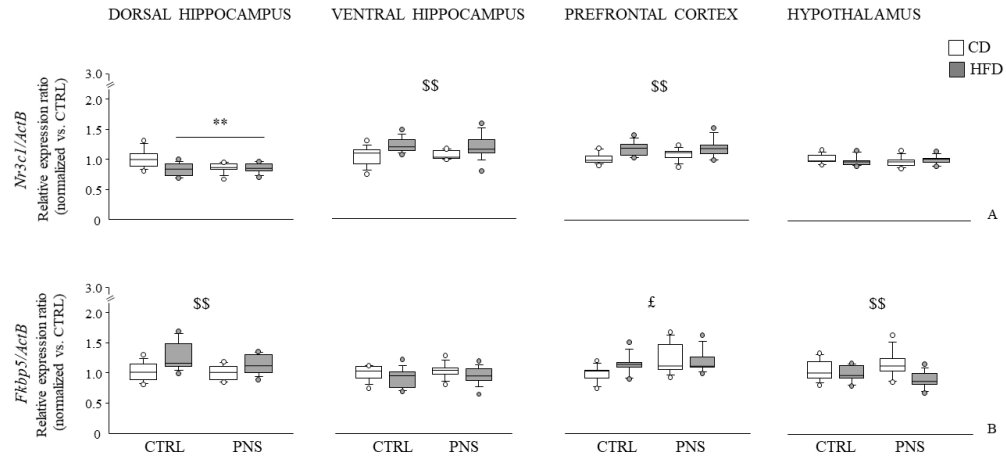


FIGURE 5. Independently from the prenatal condition, post-natal HFD administration increases levels of Nr3c1 in the ventral hippocampus and prefrontal cortex by contrast, in the dorsal hippocampus a significant interaction between PNS and HFD results in decreased levels of this gene both as a result of pre- and post-natal stressors. No change in Nr3c1 was observed in the hypothalamus as a result of PNS nor of HFD (see panel A). As for Fkbp5, HFD independently from the prenatal condition, increased its expression levels in the dorsal hippocampus while a decrease was observed in the hypothalamus; no change was observed in the ventral hippocampus. PNS overall increased levels of Fkbp5 specifically in the prefrontal cortex. SS $p < 0.01$, main effect of diet (Nr3c1: ventral hippocampus and prefrontal cortex; Fkbp5: dorsal hippocampus and hypothalamus; Fkbp5: dorsal hippocampus; hypothalamus); $£$ $p < 0.05$, main effect of PNS (Fkbp5: prefrontal cortex); post hoc comparisons: $**$ $p < 0.01$, interaction effect (Nr3c1: CTRL-CD vs. CTRL-HFD, PNS-CD and PNS-HFD); Number of subjects: 8-10 within each experimental group.

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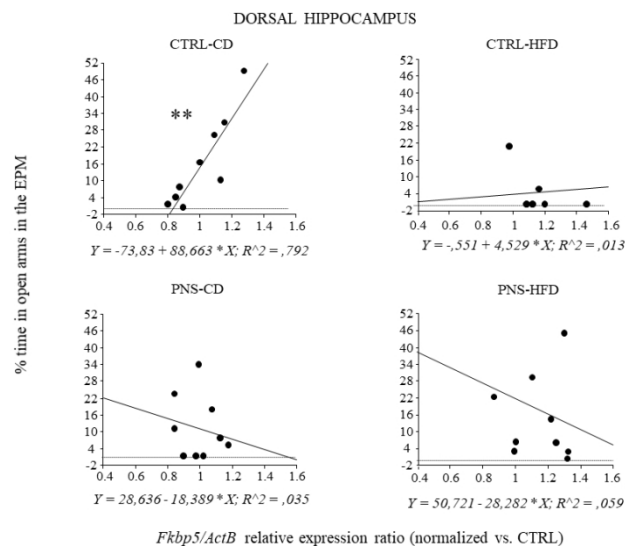


FIGURE 6. Control subjects are characterized by a direct relationship between expression levels of *Fkbp5* in the dorsal hippocampus and the time spent in the open arms of the Elevated Plus Maze, overall suggesting a permissive role for explorative behavior of this GC-related genes under physiological conditions. Post hoc comparisons $**p < 0.01$. Number of subjects: 6-9 within each experimental group.

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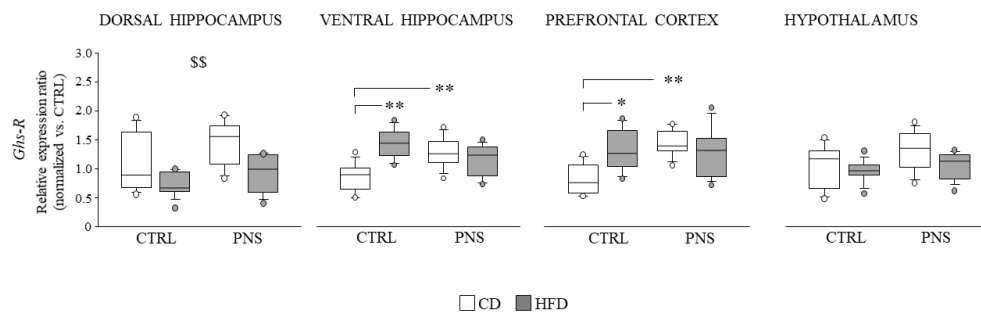
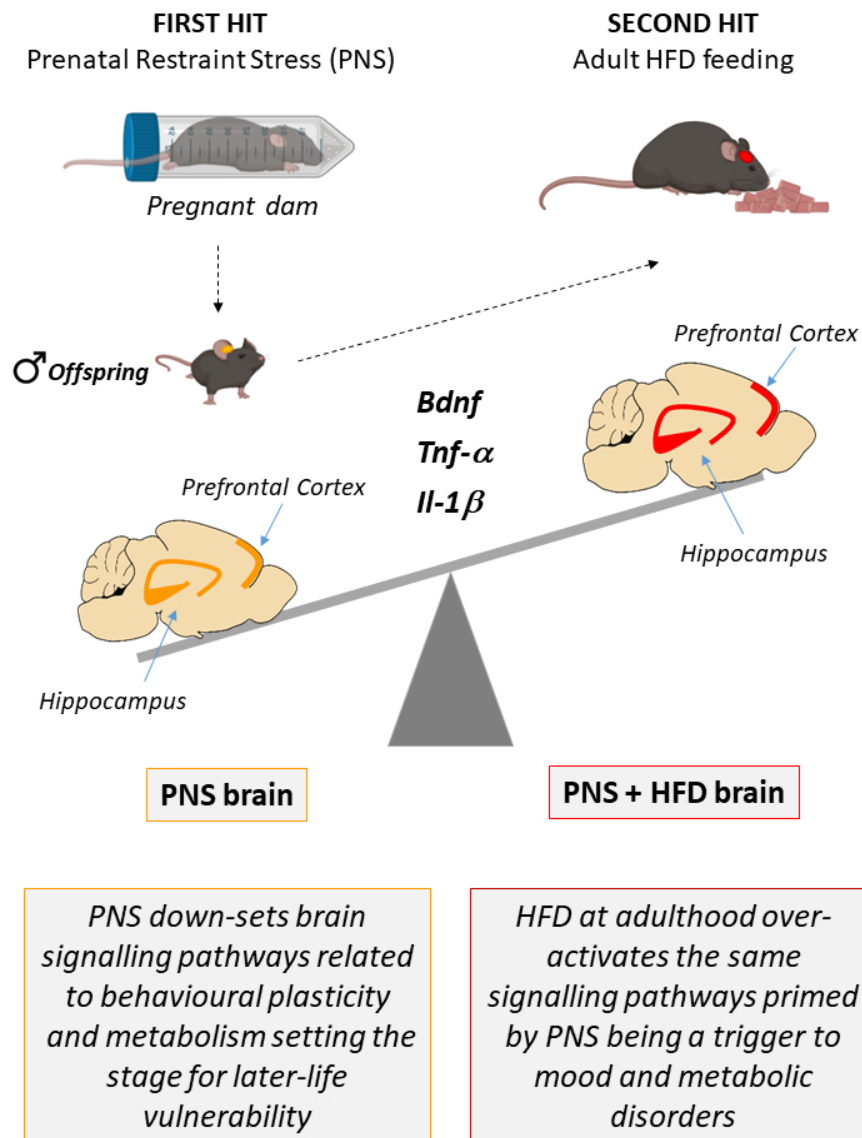


FIGURE 7. Levels of Gh/R were found to be specifically decreased upon HFD in the dorsal hippocampus while an increase was observed in both the ventral hippocampus and the prefrontal cortex in CTRL-HFD and PNS-CD animals. Main effect of HFD: main effect of diet, \$\$ $p < 0.01$; post hoc comparisons for the interaction effects: * $p < 0.05$: prefrontal cortex, CTRL-CD vs. CTRL-HFD; ** $p < 0.01$: ventral hippocampus, CTRL-CD vs. CTRL-HFD and PNS-CD, prefrontal cortex, CTRL-CD vs. PNS-CD. Number of subjects: 8-10 within each experimental group.

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GRAPHICAL ABSTRACT. Based on the “two hit model” of the developmental origin of diseases, PNS (first hit) affects foetal brain developmental trajectories; signalling pathways related to brain plasticity and function (Bdnf, Tnf-alpha and Il-beta) are down-set providing increased liability to later-life mood and metabolic disorders. A later-life challenge such as HFD feeding (second hit), experienced at adulthood, might impinge upon the same signalling pathways primed by PNS leading to enhanced responses to the metabolic challenge. The hippocampus and prefrontal cortex, being main targets of glucocorticoids, are finely modulated by PNS and readily respond to the HFD metabolic stress.

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