

1 **Increased level of DNA damage in some organs of obese Zucker rats by γ -H2AX analysis.**

2 Alessia Azzarà¹, Anna Chiaramonte¹, Erika Filomeni¹, Barbara Pinto², Stefano Mazzoni², Simona
3 Piaggi², Maria Angela Guzzardi³, Fabrizio Bruschi², Patricia Iozzo³ and Roberto Scarpato¹

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5 ¹Unità di Genetica, Dipartimento di Biologia, University of Pisa, Via Derna 1, 56126 Pisa, Italy

6 ²Dipartimento di Ricerca Traslazionale e delle Nuove Tecnologie in Medicina e Chirurgia, Via Savi
7 10, 56126 Pisa, Italy

8 ³CNR Institute of Clinical Physiology, Via Giuseppe Moruzzi, 1, 56124 Pisa, Italy

9

10 Corresponding author: Roberto Scarpato, Unità di Genetica, Dipartimento di Biologia, University of
11 Pisa, Via Derna 1, 56126 Pisa, Italy

12 Phone number: +390502211509

13 Fax number: +390502211527

14 e-mail: roberto.scarpato@unipi.it

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17

1 **Abstract**

2

3 In a recent study, we showed that lymphocytes of obese Italian children/adolescents displayed
4 levels of double strand breaks (DSB), assayed as serine 139-phosphorylated histone H2AX (γ -
5 H2AX), about eight-fold higher than normal weight controls, and that 30% of this damage
6 generated micronuclei. These findings suggested that obese children could be at increased risk of
7 obesity-mediated cancer later in life. We therefore aimed to assess the level of γ -H2AX in a genetic
8 animal model of obesity (Zucker rat) to identify a genotoxic/carcinogenic risk in some organs. The
9 DSB marker was studied in 3-4-week-old rats and in 9-13-week old rats. Paraffin-embedded
10 sections of heart, thyroid, liver, pancreas, lung, kidney, oesophagus and gut from the fa-/fa- (obese)
11 and the fa+/fa- (lean) control animals were processed for immunohistochemistry detection of γ -
12 H2AX. Pancreas (0.0624 ± 0.0195), lung (0.1197 ± 0.0217), oesophagus (0.1230 ± 0.0351), kidney
13 (0.1546 ± 0.0149) and gut (0.1724 ± 0.0352) of 9-13-week-old obese rats showed a higher proportion
14 of γ -H2AX positive nuclei, than their lean counterparts (0.0092 ± 0.0033 , 0.0416 ± 0.0185 ,
15 0.0368 ± 0.0088 , 0.0686 ± 0.0318 and 0.0703 ± 0.0239 , respectively). No difference was seen in the 3-
16 4-week-old age group with regard to obesity, indicating that the DNA damage increased with older
17 age of the rats. We hypothesize that the organs of the obese animals showing high levels of DSB,
18 could represent target tissues for the development of obesity-related cancers.

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1 **1. Introduction**

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3 Obesity is a complex disease characterised by an excess of fat in the body in which genetic,
4 physiological, environmental and socio-economic factors, interacting with each other, predispose
5 and **concur** **contribute** to its pathogenesis as well as to the appearance of adverse effects later in life
6 [Wright et al., 2012; Trésallet et al., 2014; Marcello et al., 2014; Brenner, 2014; Drew, 2012]. This
7 pathological condition, in fact, can have severe outcomes for human health, causing chronic
8 illnesses such as type 2 diabetes mellitus, asthma, cardiovascular diseases [Mitchell et al., 2013;
9 Rosenkranz et al., 2005; Weiss et al., 2004; DeVallance et al., 2015] and also some types of cancers
10 [Bardou et al., 2013; Calle et al., 2003; Karagozian et al., 2014; Renehan et al., 2008]. In this
11 context, formation of double strand breaks (DSB) within the genome, if unrepaired, lead to
12 mutations that, in turn, can initiate the carcinogenesis process. However, cells activate the DNA
13 damage response (DDR) to reconstitute the integrity of DNA, in which phosphorylation at serine
14 139 of several histone H2AX molecules surrounding the site of DSB (γ -H2AX foci) represents an
15 early and key event. Hence, it has been assumed that the number of γ -H2AX foci can reflect
16 approximately the number of nuclear DSB formed, thus making γ -H2AX an excellent marker of
17 early DNA damage [Rogakou et al., 1998; Watters et al., 2009; Scarpato et al., 2011; Redon et al
18 2011]. As a consequence, it was also possible to relate the presence of γ -H2AX in the nuclei to how
19 and how much a given cell population responds to carcinogenesis stimuli [Bartkova et al., 2005;
20 Nuciforo et al., 2007]. Recently, we have provided evidence that a group of obese Italian
21 children/adolescents had about eight-fold higher levels of DSB, expressed as γ -H2AX nuclear foci,
22 in their peripheral lymphocytes than normal weight subjects, and that 30% of this damage generated
23 micronuclei [Scarpato et al., 2011; Azzarà et al., 2016]. These findings suggested that obese
24 children could be at increased risk for developing obesity-mediated cancer later in life.
25 Genetic animal models have been developed in rodents to efficiently study obesity and/or the
26 metabolic syndrome, and among these, the Zucker fatty rat offers a reliable *in vivo* genetic system.

1 The genome of this animal model carries the homozygous missense mutation that causes loss of
2 function of the leptin receptor gene [Takaya et al., 1996]. As a consequence, the rats show
3 hyperphagia, hypercholesterolaemia, hyperglycaemia, and hyperinsulinaemia [Cozzi et al., 2009;
4 Geurts et al., 2009]. Leptin is encoded by the *lep* gene and produced by adipocytes [Zhang et al.,
5 1994], regulating food intake in mammals via a cross-talk between the adipose tissue and central
6 nervous system that maintains energy balance and normal body weight [Schwartz et al., 2000]. In
7 addition, as described in the literature, these rats were also used to investigate some mechanisms
8 connected to the obesity- or diabetes-mediated carcinogenesis [Koch et al., 2008; Ishii et al., 2011;
9 Ishizaki et al., 2013; Imai et al., 2013]. Thus, we aimed to gain information on the obesity-mediated
10 genotoxic risk by investigating, at tissue level, the expression and time-course of the γ -H2AX
11 marker in eight organs of the Zucker rat model. We performed the study in the 3-4 week-old rats
12 and in 9-13-week-old rats in order to mimic and cover, as much as possible, the human life period
13 spanning from childhood and adolescence to maturity.

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15 **2. Materials and Methods**

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17 **2.1. Zucker rat models and biochemistry determinations**

18 Male Zucker fatty rats were purchased from Charles River Laboratories International, Inc.,
19 (Wilmington, MA, USA), housed in the animal facility under constant conditions of
20 temperature (24–25 °C) and artificial lighting (12 h light–dark cycle), fed with standard rat
21 chow (calories provided by: 27.0% protein 13.1% fat and 59.9% carbohydrates; Sandown
22 Scientific, Middlesex, England) and water *ad libitum* until used for the study. The Zucker
23 genome carries the homozygous missense mutation (C→A) at nucleotide 806 of the leptin
24 receptor gene, with consequent Gln239Pro amino-acidic substitution in the extra-cellular
25 domain of the receptor (fa^-/fa^- animals), loss of function and acquisition of an obese phenotype.
26 The lean control group was represented by heterozygous counterparts (fa^+/fa^- animals). We

1 studied eight organs (heart, kidney, lung, oesophagus, gut, pancreas, thyroid and liver) in 44
2 rats subdivided into 23 animals aged 9-13 weeks and into 21 animals aged 3-4 weeks.
3 According to their body weight and the phenotypic characteristic previously described, the 9-
4 13-week-old rats were categorized as obese (n=13) and lean (control) rats (n=20). In the 3-4-
5 week-old rats, 19 and 12 animals were classified as obese and normal weight (lean) animals,
6 respectively. Due to specific difficulties in collecting the thyroid and pancreas from animals
7 aged 3-4 weeks, we could not perform DNA damage analysis in these organs.
8 Standard laboratory methods were used for glycaemia, insulin and total cholesterol level
9 determinations. A scil Vet ABC™ Haematology Analyzer (scil animal care company S.r.l.,
10 Milan, Italy) was used to measure blood glucose and total cholesterol, while insulin plasma
11 levels were measured by a commercial immunoassay kit (Crystal Chem Zaandam, Netherlands,
12 sensitivity = 1.5 µIU/mL, CV < 10%). On the day of the study, the animals were sacrificed by
13 inhalation of an overdose of isoflurane after collection of blood samples for clinical
14 biochemistry measurements. In order to minimize experimental variability, groups of four to-
15 five obese and lean animals were processed at once. We followed the recommendations of
16 Italian legislation (DL No. 116, 1992), which implements the EEC directive 609/86 for the care
17 and use of laboratory animals.

18 Body weight, genetic and phenotypic characteristics and clinical biochemistry measurements
19 (glycaemia, insulin and total cholesterol levels) of all the Zucker rats are reported in Table 1.

20

21 **2.2. Tissue histology and immunohistochemistry protocol**

22 After sacrifice, a piece of tissue was randomly collected from each organ of each animal,
23 washed of blood residues, cut into two small pieces and fixed in formalin until processed for γ -
24 H2AX analysis by the immunohistochemistry protocol described elsewhere [Del Ry et al.,
25 2015]. Briefly, the formalin fixed tissues were washed in tap water to remove fixative,
26 dehydrated in an ethanol series, immersed two times in Isoparaffin for 1 h each (Panreac, Nova

1 Chimica, Milan, Italy), and included in paraffin. The paraffin-embedded organs were then
2 sectioned into 5- μ m slices using a Leica RM 2155 microtome (Leica instruments, Wetzlar,
3 Germany). The slices were then placed on Superfrost Ultra Plus® slides (Manzel-Glaser,
4 Thermo-Fisher, Milan, Italy) to guarantee firm electrostatic adhesion of the sections. Two
5 coded slides, containing two sections of each organ, were set up, processed and scored blindly
6 by three experienced persons. Following deparaffination/rehydration done in xylene/ethanol
7 series and heat-mediated antigen unmasking, the sections were incubated in 3% hydrogen
8 peroxide (Sigma-Aldrich, Milan, Italy) for 10 min, and then for 1 h in blocking solution (10%
9 FBS (Life Technologies, Monza, Italy), 0.3% Triton-X, in 1X PBS). Then the slides were
10 incubated overnight at 4°C with an 1:200 diluted primary polyclonal rabbit anti- γ -H2AX
11 antibody (Abcam, Prodotti Gianni, Milan, Italy), followed by incubation for two hours at RT
12 with an 1:100 diluted anti-rabbit horseradish peroxidase-conjugated secondary antibody
13 (Sigma-Aldrich, Milan, Italy) , subsequently revealed by application of the 3,3'-
14 diaminobenzidine tetrahydrochloride substrate (DAB; Sigma-Aldrich, Milan, Italy). Finally,
15 the slides were counterstained in haematoxylin, dehydrated and mounted in glass coverslips for
16 microscopy observation.

17 The presence of γ -H2AX positive nuclei within a haematoxylin-counterstained cell population
18 was recognised by the formation of a homogeneous brown precipitate covering partially or
19 entirely the DSB-damaged nuclei, which were easily distinguished by their undamaged, blue-
20 coloured nuclei. Figures 1-2 show a panel of positive and negative sections for each organ.

21 Slides incubated without the primary or the secondary antibodies, as well as with 1000 U/ml of
22 alkaline phosphatase (Sigma-Aldrich, Milan, Italy), were also made to ensure on the proper
23 recognition of histone H2AX molecules phosphorylated at ser 139.

24 A (semi)-quantitative analysis was made with the aid of Image J software (downloaded at
25 <http://imagej.nih.gov/ij/>). Briefly, four areas in each section showing the signal for γ -H2AX
26 were randomly identified, and the ratio between γ -H2AX positive (brown-stained) to negative

1 (blue-stained) nuclei was calculated. The average value from the four areas, adjusted for the
2 degree of γ -H2AX positivity of the section expresses the DNA DSB marker in each section.
3 This was achieved by scanning the entire section with the 10X objective.

4 To delineate the time-course of nuclear phosphorylation in each organ from 3-4-week-old rats
5 to 9-13-week-old rats, γ -H2AX levels were expressed as the difference or the ratio between the
6 average values obtained in the two age groups. Thus, for the lean or obese rats, two parameters
7 were calculated subtracting or dividing the average values of the 9-13-week-old rats to the
8 corresponding values of the 3-4-week-old group.

9

10 **2.3. Statistical analysis**

11 To detect differences in the level of DNA damage (γ -H2AX values) of the organs or in the
12 values of the metabolic parameters between lean (fa+/fa+) and obese (fa+/fa-) rats, data were
13 statistically analysed, in each age group (3-4-week-old rats and 9-13-week-old rats) by the
14 Student's t-test using the Statgraphics Centurion XV software package (Statistical Graphics
15 Corp., Rockville, MD, USA). *P*-value was considered significant when < 0.05 .

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18 **3. Results**

19

20 The 9-13-week-old obese rats weighted (353.4 ± 19.3 g) approximately 20% more than the lean
21 counterparts (289.3 ± 21.4 g), and compared to this group, they showed significantly higher levels
22 of basal glycaemia (305.2 ± 28.1 g vs. 209.4 ± 35.6 g, $p < 0.05$), insulin (2944 ± 341 pg/ml vs. 750
23 ± 199 pg/ml, $p < 0.001$) and total cholesterol (162.5 ± 12.8 mg/dl vs. 110.4 ± 6.8 mg/dl, $p < 0.05$). As
24 previously mentioned, no difference was observed in the 3-4-week-old rats between obese and lean
25 animals (see Table 1).

1 In the two rat age groups, the results of the DNA damage analysis, expressed as γ -H2AX levels, are
2 shown in Figure 3 or Figure 4 according to the organs in which a genotype effect was or was not
3 observed, respectively. For each sampled organ, we did not detect a significant difference in the
4 basal DNA damage between fa^{+}/fa^{+} and fa^{+}/fa^{-} 3-4-week-old rats. In the adult rats, the levels of γ -
5 H2AX positive cells was quite variable among the analysed tissues, with values ranging from
6 0.0092 ± 0.0032 (pancreas) to 0.1210 ± 0.0323 (thyroid) in the lean group, and from $0.0495 \pm$
7 0.0246 (heart) to 0.1724 ± 0.0352 (gut) in the obese group. Statistically significant increases were
8 observed in five organs of obese as compared to lean rats (Figure 3), with the highest and lowest
9 difference observed in the pancreas (0.0624 ± 0.0195 vs. 0.0092 ± 0.0033 , $p = 0.0447$) and in the
10 kidney (0.1546 ± 0.0149 vs. 0.0686 ± 0.0318 , $p = 0.0186$), respectively. The other values were
11 (obese vs. lean): lung (0.1197 ± 0.0217 vs. 0.0416 ± 0.0185 , $p = 0.0352$), gut (0.1724 ± 0.0352 vs.
12 0.0703 ± 0.0239 , $p = 0.0470$) and oesophagus 0.1230 ± 0.0351 vs. 0.0368 ± 0.0088 , $p = 0.0249$).
13 Also grouping together the γ -H2AX data from all the positive organs, we registered significantly
14 higher values of DNA damage in the obese animals than in the lean rats (0.1264 ± 0.0287 vs. 0.0453
15 ± 0.0139 , $p = 0.0234$). In the heart, liver and thyroid, the baseline frequencies of obese animals
16 were, on average, comparable to those of the lean group (Figure 4).
17 With regard to the time-course of DNA damage, from animals aged 3-4 weeks to animals aged 9-13
18 weeks, the γ -H2AX levels of lean and obese rats showed a general increase in all the analysed
19 organs, suggesting that the basal frequencies of DNA DSB increased with older age of the animal.
20 However, this increase was more pronounced in the obese rats than in the lean ones, with the
21 exception of heart and liver (thyroid was not sampled in 3-4-week-old rats): in these organs, in fact,
22 the elevated increase in the basal DSB levels occurred at a comparable extent in both fa^{+}/fa^{+} and
23 fa^{+}/fa^{-} animals.

24

25 4. Discussion

1
2 The response of animal organs to genotoxic insults has been widely demonstrated using
3 immunohistochemical detection of γ -H2AX levels in lung tissue of rats exposed to fine and
4 ultrafine dusts [Rittinghausen et al., 2013]. Elevated immunostaining for this DSB marker has also
5 been reported in renal tissue of obese mice as well as in the lungs of obese Zucker rats with altered
6 expression of the endothelin axis [Del Ry et al., 2014; Mozaffari et al., 2012]. Despite the
7 physiological presence of individual and organ variability, our results indicate that, irrespectively of
8 their genotype, the levels of the DNA damage marker were significantly lower in all analysed
9 organs of the 3-4-week-old rats as compared to the 9-13-week-old ones. Cell tissue alterations
10 occurred in the 3-4-week-old obese rats were minor and less widespread among body organs than
11 those observed in the corresponding 9-13-week-old age group. In other words, the tissue damage
12 would seem to increase, at DNA level, as the age of the animals increased. This might be explained
13 by the fact that in 3-4-week-old rats the level of inflammation/oxidative stress has not yet reached a
14 sufficient intensity to affect DNA integrity. It is only in the 9-13-week old rats that the frequencies
15 of cells containing DSB increase, in some organs, as a function of the presence of genetic
16 predisposition to obesity. In fact, the levels of DNA damage we detected in the analysed tissues of
17 the obese group delineated the following descending order: gut, kidney, oesophagus, lung, pancreas
18 (organs that resulted significantly different from those of the lean animals), thyroid, liver and heart.
19 As the parameter we used to quantify γ -H2AX (percent of positive nuclei adjusted for the degree of
20 positivity to the γ -H2AX signal of the whole section) gives a rough estimate of the actual
21 spontaneous levels of γ -H2AX in each tissue, any comparison among the various organs should be
22 properly interpreted. Thyroid was found to express γ -H2AX signal to an extent much higher than
23 other tissues such as, for example, the pancreas. However, it is not relevant that thyroid was, in
24 general, the most reactive organ to γ -H2AX immunostaining, probably due to an overproduction of
25 H_2O_2 by thyreocytes [Driessens et al., 2009], it is important that γ -H2AX levels in the

1 corresponding tissue of the obese rats resulted unchanged (thyroid) or significantly increased
2 (pancreas). Moreover, although no information on the extent of DNA damage at single cell level
3 could be drawn, as neither the number of γ -H2AX foci nor any other parameter related to the
4 presence of DSB was assayed, the percent of γ -H2AX positive nuclei is considered a useful marker
5 of DNA damage, and other authors used it to quantify DSB in some rodent tissues. Rittinghouse et
6 al. [2013] found about 150 positive nuclei per mm^2 in the lung of female Wistar rats, while urinary
7 bladder of 5 weeks old F344 rats showed approximately 2 γ -H2AX positive nuclei per 1000 cells
8 counted [Toyoda et al., 2015]. Finally, no cell with the γ -H2AX signal or a very slight amount of
9 nuclei was found in the brain of Wistar rats developing lymphosarcomas [Masutani et al., 2014] or
10 in the duodenum of B6C3F1 mice, respectively [Thompson et al., 2015].

11 The fa^-/fa^- animals are also affected, on average, by the highest glucose levels that, indeed, is an
12 intrinsic characteristic of these rats [Serpillon et al., 2009]. Hyperglycaemia is connected to
13 oxidative stress as it causes, via ROS- and NOX1-mediated glycooxidation, a pro-inflammatory
14 response and oxidative damage in the target tissue [Yoshida et al., 2000; Yang et al., 2011; Vlassara
15 et al., 2011]. We have recently demonstrated that adolescents affected by type 1 diabetes but with
16 BMI in the range of normality, have elevated levels of γ -H2AX foci in their peripheral lymphocytes
17 [Giovannini et al.; 2014]. In this context, Nox 1-deficient hyperglycaemic mice exhibited reduced
18 levels of γ -H2AX in their kidneys [Zhu et al., 2015]. We therefore suggest that the increased
19 frequency of γ -H2AX positive cells observed among our fa^+/fa^- rats was probably due to the
20 obesity status, in which a role might also be played by their constitutive hyperglycaemic condition,
21 which was preferentially expressed in gut, kidney, oesophagus, lung and pancreas rather than in
22 thyroid liver and heart.

23 With regard to a possible oncologic risk for humans from the animal data, we should take into
24 account that DSB are still reparable DNA lesions, and the presence of γ -H2AX positive cells
25 indicates the recruitment of the appropriate cellular response to remove the damage. However,

1 although only eight organs were studied, the obese rats, showing an increased occurrence of this
2 type of DNA lesion, might be considered at increased risk of developing malignancies, especially in
3 the above mentioned tissues. The excess of H2AX phosphorylation we have observed in the fa⁻/fa⁻
4 animals actually reflects the obesity condition that is capable of generating DSB and, in turn, organ
5 damage. However, it cannot be ruled out that these γ -H2AX foci represent unrepaired DSB that
6 persist and accumulate in the tissues, as also suggested by other authors [Siddiqui et al., 2013;
7 Bhogal et al., 2010].

8 The presence of endocrine-metabolic alterations is compatible with an imbalance in plasma and
9 intracellular antioxidant/pro-oxidant activities that produce DNA-reactive molecules through
10 generation of chronic inflammation. In accordance with our data, some studies observed, in
11 humans, a tight association between inflammatory-based lung diseases, colon and pancreatic
12 cancers, and obesity, probably due to the excess of serum levels of leptin that can cause an
13 imbalance of the immune response [Shore, 2010; Schatz et al., 2013; Youssef et al., 2013], or
14 trigger a pro-inflammatory and anti-apoptotic response [Bardou et al., 2013; Huang et al., 2009;
15 Yeo, 2015; Drew, 2012]. In addition, a prolonged obesity condition sustained by oxidative stress
16 can trigger renal damage, as demonstrated in 3-month-old fa⁻/fa⁻ Zucker rats [Poirier et al., 2000].
17 Also oesophageal cancer and Barrett's oesophagus were clearly linked to adult obesity through
18 overproduction of the insulin-like growth factor 1 (IGF-1), together with a reduced presence of
19 IGF-binding proteins, due to hyperinsulinaemia [Calle et al., 2003; Kamat et al., 2009; Greer et al.,
20 2012]. Indeed, the increase in the occurrence of γ -H2AX we observed in the cells of the respective
21 tissues could reflect these conditions. It has been demonstrated that in mice, rats and hamsters the
22 induction and persistence of γ -H2AX by ionizing radiation occurs in a tissue-specific manner, this
23 in turn depends on the different ability of the organ's tissues to divide themselves [Koike et al.,
24 2008; Firsanov et al., 2012]. The lack of an increase in the levels of DSB in the heart, thyroid and
25 liver of obese rats, might, at least partially, rely on the proliferative status of the three organs. In

1 fact, heart is a non-dividing tissue and thyroid and liver are considered quiescent tissues with a slow
2 proliferation ability. These tissues, therefore, would be less responsive to the low-grade oxidative
3 stimuli present in the obesity condition than more proliferating tissues such as gut and lung.
4 Variation among the analysed organs may also depend on a different expression and/or functioning
5 of their antioxidant systems. At the same time, the protective action of some natriuretic peptides
6 produced by the cardiac tissue and of adipokines such as visfatin cannot be ruled out [Rittinghausen
7 et al., 2013; Habbu et al., 2006; Xiao et al., 2013]. Moreover, additional researches to investigate
8 these topics should be performed.

9 The present study conducted in Zucker fatty rats indicates there may be an association between
10 obesity and the occurrence of potentially genotoxic/carcinogenic lesions such as DSB. In addition,
11 although we could not detect these signs of DNA damage earlier in animal life (*i.e.* in the rats of age
12 comparable to that of children/adolescents) as we did in a previous study [Scarpato et al., 2011], the
13 Zucker model has shown the potential to identify organs which could be considered at risk of
14 developing obesity-mediated cancers later in human life.

16 **Author Contributions**

17 RS designed the study and wrote the paper. AA performed all the experimental parts of the study
18 and wrote the paper. AC and EF performed all the experimental parts of the study. PI participated to
19 the design of the study, she and was the responsible for the Zucker rat model. MG performed the
20 collection and processing of rat organs. SM prepared paraffin-embedded tissue sections. FB, BP and
21 SP participated in writing the manuscript. All authors approved the final manuscript.

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Table 1. Genetic, phenotypic characteristics and biochemistry measurements of male Zucker rats grouped according to age. All values are expressed as mean±S.E.

Animal groups	Genotype ^a	Body weight (g)	Hyperphagia	Blood glucose (mg/dl)	Insulin (pg/ml)	Total cholesterol (mg/dl)
9-13 weeks						
Lean	fa+/fa-	289.3±21.4	no	209.4±35.6	750±199	110.4±6.8
Obese	fa-/fa-	353.4±19.3	yes	305.2±28.1 ^b	2944±341 ^c	162.5±12.8 ^b
3-4 weeks						
Lean	fa+/fa-	67.6±4.6	no	159.8±9.4		
Obese	fa-/fa-	64.1±5.3	no	175.0±14.5		

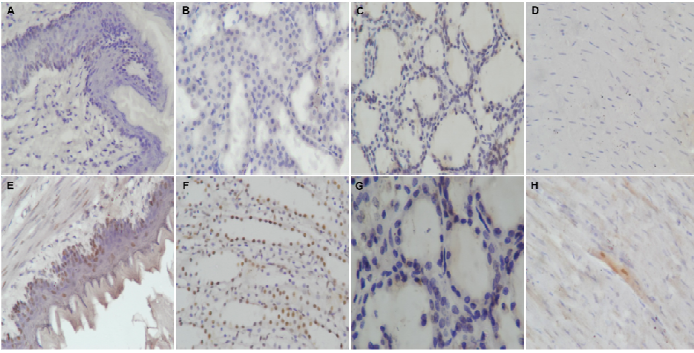
^afa- indicates the presence of the missense mutation (C→A) at nucleotide 806 of the leptin receptor gene.

^bSignificantly different vs. lean rats ($p < 0.05$, Student's t-test).

^cSignificantly different vs. lean rats ($p < 0.001$, Student's t-test).

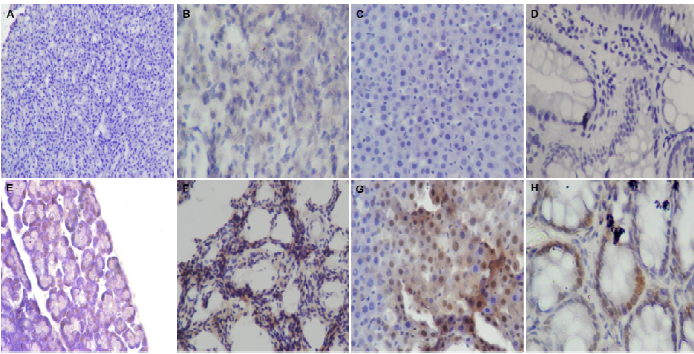
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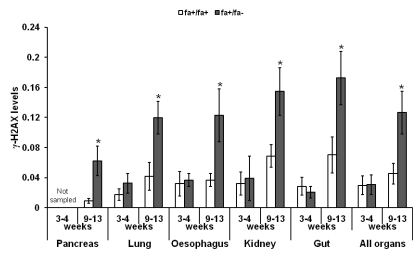
2 **Figure 1**



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4 **Figure 2**

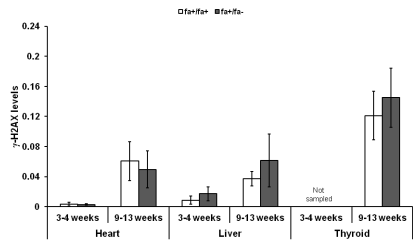
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2 **Figure 3**

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5 **Figure 4**

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Figure legends

Figure 1. Images of organ sections from Zucker rat after immunohistochemistry with γ -H2AX primary antibody detected with an anti-rabbit horseradish peroxidase-conjugated secondary antibody (hematoxylin counterstaining). A-D: areas of oesophagus, kidney, thyroid and heart showing only blue stained γ -H2AX-negative nuclei; E-H, areas of the same organs showing several brown coloured γ -H2AX-positive nuclei. A-F, 200X magnification; G and H, 400X magnification.

Figure 2. Images of organ sections from Zucker rat after immunohistochemistry with γ -H2AX primary antibody detected with an anti-rabbit horseradish peroxidase-conjugated secondary antibody (hematoxylin counterstaining). A-D: areas of pancreas, lung, liver and gut showing only blue stained γ -H2AX-negative nuclei; E-H, areas of the same organs showing several brown coloured γ -H2AX-positive nuclei. A, 100X magnification; C, D, E, F and G, 200X magnification; B and H, 400X magnification.

Figure 3. Levels of the DNA damage marker (γ -H2AX) in 3-4 week-old rats and 9-12 week-old rats. Data refer to those organs in which a genotype effect was observed ($fa+/fa+$ = lean rats, $fa+/fa-$ = obese rats). γ -H2AX levels are expressed as percentage of γ -H2AX positive nuclei adjusted for the degree of positivity to the γ -H2AX signal of the whole section. Bars represents the mean \pm S.E. of each animal group (two replicate sections per animal). Asterisks denote a significant difference ($* p < 0.05$, Student's t-test) vs. the respective $fa+/fa-$ rats.

Figure 4. Levels of the DNA damage marker (γ -H2AX) in 3-4 week-old rats and 9-12 week-old rats. Data refer to those organs in which a genotype effect was not observed ($fa+/fa+$ = lean

1 rats, fa+/fa- = obese rats). γ -H2AX levels are expressed as percentage of γ -H2AX positive
2 nuclei adjusted for the degree of positivity to the γ -H2AX signal of the whole section. Bars
3 represents the mean \pm S.E. of each animal group (two replicate sections per animal).

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