RESEARCH ARTICLE

Effects of a wide range of dietary nicotinamide riboside (NR) concentrations on metabolic flexibility and white adipose tissue (WAT) of mice fed a mildly obesogenic diet

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Scope: Metabolic flexibility is the ability to switch metabolism between carbohydrate oxidation (CHO) and fatty acid oxidation (FAO) and is a biomarker for metabolic health. The effect on metabolic health of nicotinamide riboside (NR) as an exclusive source of vitamin B3 is unknown and is examined here for a wide range of NR.

Design and methods: Nine-week-old male C57BL/6Jrc mice received a semi-purified mildly obesogenic (40 en% fat) diet containing 0.14% L-tryptophan and either 5, 15, 30, 180, or 900 mg NR per kg diet for 15 weeks. Body composition and metabolic parameters were analyzed. Metabolic flexibility was measured using indirect calorimetry. Gene expression in epididymal white adipose tissue (eWAT) was measured using qRT-PCR.

Results: The maximum delta respiratory exchange ratio when switching from CHO to FAO (maxΔRERCHO→FAO) and when switching from FAO to CHO (maxΔRERFAO→CHO) were largest in 30 mg NR per kg diet (30NR). In eWAT, the gene expression of Pparα, a master regulator of adipogenesis, and of Sod2 and Prdx3, two antioxidant genes, were significantly upregulated in 30NR compared to 5NR.

Conclusion: 30NR is most beneficial for metabolic health, in terms of metabolic flexibility and eWAT gene expression, of mice on an obesogenic diet.

Keywords: Metabolic flexibility / NAD+ / Niacin / Nicotinamide riboside / Vitamin B3

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1 Introduction

Nicotinamide adenine dinucleotide (NAD+) is essential to maintain cellular redox state and basal energy metabolism. Beyond its role as a reusable coenzyme, NAD+ is also permanently degraded by various signalling enzymes, including sirtuins [1]. Although the degradation product, nicotinamide (Nam), can be recycled for NAD+ generation, maintenance of NAD+ levels is dependent on exogenous supplementation with vitamin B3, from which NAD+ can be synthesized. Indeed plasma NAD+ levels decrease when humans or rodents are exposed to a vitamin B3-deficient diet [2–4]. Severe vitamin B3 deficiency causes pellagra, a disease characterized by skin rash, diarrhoea, and dementia.

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Abbreviations: 2PY, 2-pyridone; BW, Body weight; CHO, Carbohydrate oxidation; CLS, Crown-like structure; EE, Energy expenditure; eWAT, epididymal white adipose tissue; FAO, Fat acid oxidation; HFD, High fat diet; LFD, Low fat diet; MeNam, N-methylnicotinamide; mtDNA, mitochondrial DNA; NA, Nicotinamide acid; NAD+, Nicotinamide adenine dinucleotide; Nam, Nicotinamide; NaMN, Nicotinic acid mononucleotide; nDNA, nuclear DNA; NEFA, Non-esterified fatty acids; NMN, Nicotinamide mononucleotide; NR, Nicotinamide riboside; RER, Respiratory exchange ratio; TG, Triglycerides; Trp, Tryptophan; WAT, white adipose tissue.

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by dermatitis, diarrhea, and dementia, ultimately resulting in death [5]. Even though pellagra has become rare in developed countries, it remains endemic in underdeveloped countries [6].

While severe vitamin B3 deficiency results in disease, hardly any information is available on physiological consequences of marginal vitamin B3 levels. This is particularly relevant because it was shown that obesogenic diets reduce NAD+ levels in several tissues [7–10]. Surprisingly, requirements for dietary vitamin B3 are still not well-established for rodents, especially mice [11, 12]. The National Research Council suggested the vitamin B3 requirement in rats to be 15 mg/kg diet, based on three scientific publications in the 1940s. However, these studies did not use purified diets and only used growth rate as a read-out parameter [13–15]. More importantly, requirements were based on studies with rats as no direct data were available for mice. Estimation of vitamin B3 requirements is not straightforward, since NAD+ can also be synthesized de novo from the essential amino acid tryptophan (Trp) [16]. Dietary Trp may thus rescue vitamin B3 requirement thresholds precludes research into the molecular, metabolic and physiological consequences of marginal status of various forms of vitamin B3.

Nicotinic acid (NA) and Nam are the classic forms of vitamin B3, present in the diet. NA and Nam can be converted to NAD+ via the Preiss-Handler and the salvage pathway, respectively [20]. Nicotinamide riboside (NR) is another source of vitamin B3 that naturally exists in cow’s milk and yeast-containing food products. NR can be metabolized into NAD+ directly via the Nrk pathway [21]. Recent studies have shown that supplementation of NR at pharmacological levels can provide physiological or metabolic benefits by boosting NAD+ levels [8, 22–26]. No side effects of NR have been reported, in contrast to high dose treatment with NA, that may cause skin flushing, or Nam, that may lead to liver damage [27]. Therefore, NR has been proposed as the vitamin B3 of choice. Despite the benefits from pharmacological NR supplementation [8, 22, 24, 26], it would be necessary to re-evaluate the NR dose, as it appears strikingly high (400 mg NR/kg body weight/day) compared to most commercially available supplements (60–500 mg/person/day) [28].

The metabolic health effects of NR as an exclusive source of vitamin B3 at nutritional relevant levels are unknown. Recently, the ability to rapidly switch metabolism between carbohydrate oxidation and fatty acid oxidation, so-called metabolic flexibility, has been recognized as a sensitive biomarker for metabolic health in nutritional interventions [29–32]. Metabolic flexibility can be assessed by analysing the respiratory exchange ratio (RER) during a fast-refeeding challenge in non-invasive indirect calorimetry. To examine the effects of nutritional relevant levels of NR on metabolic health, we performed a dose-response dietary intervention study using a wide range of NR, from 5 to 900 mg NR per kg of a defined semi-purified obesogenic diet, combined with a metabolic flexibility measurement. We used a mildly obesogenic diet (40 en% fat) to mimic a Western diet. To limit NAD+ biosynthesis from Trp, we implemented a low level of Trp (0.14%). This Trp level is still sufficient for normal growth in mice [33]. Since white adipose tissue (WAT) may play a crucial role in metabolic flexibility at the whole body level [34–36], we focused on metabolic flexibility and WAT function.

2 Materials and methods

2.1 Dietary intervention

The animal experiment was approved by the Animal Welfare Committee of Wageningen University, Wageningen, The Netherlands (DEC2014029). Nine-week-old male C57BL/6J Rcc mice (Envigo, Horst, The Netherlands) were individually housed (12 h light–dark cycle, 23 ± 1°C, 55 ± 15% humidity) with ad libitum access to feed and water. During a 4-week adaptation period, mice received a semi-synthetic diet (10% energy from fat), containing 0.14% L-tryptophan and 30 mg NR/kg diet (Research Diet Services, Wijk bij Duurstede, The Netherlands). At 13 weeks of age, mice were stratified into 5 experimental groups on mean body weight (n = 12/group) and received a semi-synthetic obesogenic diet (40% energy from fat) containing 0.14% L-tryptophan and either 5, 15, 30, 180, or 900 mg NR per kg diet (referred to as 5NR, 15NR, 30NR, 180NR, and 900NR, respectively) for 15 weeks (diet composition in Table 1). Feed intake and body weight as well as lean and fat mass (by NMR, EchoMRI, Houston, USA) were measured weekly. In the beginning of week 16 mice were sacrificed by decapitation after 2 h of fasting. Blood was collected for immediate blood glucose measurement using a Freestyle blood glucose meter (Abbott Diabetes Care, Hoofddorp, the Netherlands). The remaining blood was centrifuged at 3000 × g and stored at −80°C. Tissues were rapidly dissected, snap frozen in liquid nitrogen, and stored at −80°C (i.e. left epididymal white adipose tissue (eWAT), liver, soleus muscle, brain, mucosa scraped from small intestine). In addition, right eWAT was weighted, divided in half, and fixed for 24 h at 4°C in PBS with 4.0% formaldehyde (pH = 7.40) as described [37].

2.2 Indirect calorimetry

Indirect calorimetry was performed in week 14, using a PhenoMaster System (TSE Systems, Bad Homburg, Germany) as described [31]. Briefly, mice were individually housed with a steady normal air flow, 12 h light–dark cycle (07:00 h lights...
Table 1. Composition of diets

| Ingredients (g·kg⁻¹ diet) | Run-in | 5NR | 15NR | 30NR | 180NR | 900NR |
|---------------------------|--------|-----|------|------|-------|-------|
| Casein                    | 120.0  | 120.0 | 120.0 | 120.0 | 120.0 | 120.0 |
| Wheat starch              | 385.9  | 215.7 | 215.7 | 215.7 | 215.7 | 215.7 |
| Gelatin (hydrolyzed)      | 100.0  | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 |
| Maltodextrin              | 100.0  | -    | -    | -    | -    | -    |
| Sugar                     | 100.0  | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 |
| Dextrose                  | 50.0   | 50.0  | 50.0  | 50.0  | 50.0  | 50.0  |
| Arbocel B800              | 50.0   | 50.0  | 50.0  | 50.0  | 50.0  | 50.0  |
| Linseed oil               | 5.2    | 4.0   | 4.0   | 4.0   | 4.0   | 4.0   |
| Palm oil                  | -      | 206.0 | 206.0 | 206.0 | 206.0 | 206.0 |
| Coco oil                  | 7.7    | -    | -    | -    | -    | -    |
| Sunflower oil             | 30.1   | -    | -    | -    | -    | -    |
| Mineral mixture AIN-93    | 35.0   | 35.0  | 35.0  | 35.0  | 35.0  | 35.0  |
| Vitamin mixture AIN-93    | 10.0   | 10.0  | 10.0  | 10.0  | 10.0  | 10.0  |
| L-Cystine                 | 3.3    | 3.3   | 3.3   | 3.3   | 3.3   | 3.3   |
| L-Phenylalanine           | 0.4    | 3.5   | 3.5   | 3.5   | 3.5   | 3.5   |
| Choline chloride 50%      | 2.5    | 2.5   | 2.5   | 2.5   | 2.5   | 2.5   |
| Nicotinamide riboside (mg·kg⁻¹ diet) | 30.0 | 5.0   | 15.0  | 30.0  | 180.0 | 900.0 |
| Calculated amount of L-tryptophan | 1.4 | 1.4   | 1.4   | 1.4   | 1.4   | 1.4   |
| Calculated energy (kJ·g⁻¹) | 3825  | 4660  | 4660  | 4660  | 4660  | 4660  |

Energy (% of total energy content)

| Carbohydrate | 66 | 40 | 40 | 40 | 40 | 40 |
|--------------|----|----|----|----|----|----|
| Fat          | 10 | 40 | 40 | 40 | 40 | 40 |
| Protein      | 23 | 20 | 20 | 20 | 20 | 20 |

a) without vitamin B3.

Oxygen consumption, carbon dioxide production, activity (infrared beam breaks) and food and drink intake were automatically recorded. After 20 h of ad libitum feeding, mice were monitored for 24 h starting from 07:00 h. Next, the mice were exposed to a fast and refeeding challenge. For this, the mice were provided with 1.5 g of fresh experimental diet at 16:00 h. The mice fully consumed this, after which they changed to a fully fasted state. The next day at 16:00 h they were provided with 1.8 g of fresh experimental diet (refeeding), which was fully consumed. Respiratory exchange ratio (RER) and energy expenditure (EE) were calculated by TSE software (TSE systems). Metabolic flexibility was assessed as described [36], with the following modifications: the maximum delta RER was calculated when switching from CHO1 (mean of 5 highest values) to FAO (mean of 5 lowest values) during fasting (maxΔRER_{CHO1→FAO}), and when switching from FAO (mean of 5 lowest values) to CHO2 (mean of 5 highest values) during refeeding (maxΔRER_{FAO→CHO2}). How a specific RER value relates to percentage lipid or carbohydrate oxidation can be found in [38].

2.3 Serum parameters

Serum triglycerides (TG) and non-esterified fatty acids (NEFA) were measured as described [39]. Serum insulin, leptin, and adiponectin were measured with a Bio-Plex Pro Mouse Diabetes Assay using a Bio-Plex 200 system according to manufacturer’s instructions (Bio-Rad, Veenendaal, The Netherlands). Samples were diluted 1:12.5 with sample diluent for insulin and leptin measurement and 1:1600 with serum-based diluent for adiponectin measurement. To assess insulin resistance HOMA-IR a was calculated as (glucose (mmol/l) × insulin (µU/ml)/22.5) [40].

2.4 NR metabolites in serum and liver

NR metabolites were extracted from serum as well as liver and subsequently identified and quantified using LC-ESI-MS/MS according to the procedure described in Supporting Methods.

2.5 Gene expression

Tissues were grinded in liquid nitrogen, after which RNA was isolated from liver, brain, and mucosa with a RNeasy Mini kit (Qiagen, Venlo, The Netherlands) and from eWAT with Trizol, as described [41]. RNA purity and integrity was verified using Nanodrop (NanoDrop, Wilmington, USA) and Experion (Bio-Rad, CA, USA), respectively. cDNA synthesis, followed by regular qRT-PCR was performed as described [41]. Low expressed genes were pre-amplified for 12 cycles before qRT-PCR using SsoAdvanced PreAmp Supermix (Bio-Rad, CA, USA). The expression of each gene was normalized.
Figure 1. Effects of dietary NR on whole body physiological parameters. Mice were on obeseogenic diets different levels of NR for 15 weeks. Body weight (A), cumulative feed intake (B), lean mass (C), fat mass (D). 5NR open circle, 15NR open upward triangle, 30NR open square, 180NR closed downward triangle, 900NR closed diamond. NR in mg/kg diet. Data are analyzed using two-way ANOVA and presented as mean ± SEM (n = 11–12 mice per treatment).

by the stably expressed reference genes using CFX Manager software (Bio-Rad, CA, USA). Primer sequences and PCR annealing temperatures for each gene are in Supporting Information Table 1. References genes were selected based on stable expression in the selected tissues.

2.6 Mitochondrial density

EWAT mitochondrial DNA (mtDNA) and nuclear DNA (nDNA) were measured by qRT-PCR as described [31]. mtDNA/nDNA was calculated to assess mitochondrial density.

2.7 EWAT morphology

Adipocyte cell size determination as well as macrophage staining and counting were performed as described previously [37]. Detailed description of these methods can be found in the Supporting methods.

2.8 Statistics

Data are expressed as mean ± SEM for n = 10-12 mice, but n = 6 for IHC. Statistical analyses were performed using GraphPad Prism v5.04 (Graphpad, San Diego, CA, USA). Data were verified for normality using the D’Agostino and Pearson omnibus normality test and log transformed if needed. Body weight, cumulative feed intake, lean mass, fat mass, and mean RER were analyzed using two-way repeated measures ANOVA (factor 1 = NR, factor 2 = WEEK) followed by Bonferroni post-hoc analysis. All other data were analyzed using one-way ANOVA following by Dunnett’s multiple comparison test, with 30NR as control (unless otherwise stated).

Pearson correlation analysis was performed on serum NR and NAD⁺, liver NR and NAD⁺ as well as leptin/adiponectin ratio and ΔRERFAO→CHO2. P-values < 0.05 were considered to be statistically significant.

3 Results

3.1 Body physiological parameters and metabolic flexibility

No differences were found in body weight, cumulative feed intake, lean mass or fat mass, when comparing mice with 5, 15, 30, 180, or 900 mg NR/kg diet, after the 15-week intervention (Fig. 1). Metabolic flexibility was assessed by calculating maxΔRER in response to a fast and refeeding in week 14 (Fig. 2A). Fasting metabolic flexibility, maxΔRERCHO1→FAO, was significantly better in 30NR than in 5NR (Fig. 2B). Refeeding metabolic flexibility, maxΔRERFAO→CHO2, was significantly greater in 30NR than in 5NR, 15NR, or 900NR (Fig. 2C). No differences were found in energy expenditure or physical activity (Supporting Information Fig. 1). Under the non-challenged condition no significant differences in RER or feed intake were found among NR doses (Supporting Information Fig. 2).

3.2 Blood glucose and serum lipids and adipokines

No differences were seen in blood glucose, serum TG or NEFA among NR doses (Fig. 3A–C). Serum insulin, leptin, adiponectin, leptin/adiponectin ratio, and HOMA-IR index exhibited a tendency toward a dose-response curve, without reaching statistical significance (Fig. 3D–H). In all cases, except adiponectin which shows opposite behavior,
the measured value decreased and then increased, with 30NR being the turning point. Of note, 30NR resulted in lower leptin/adiponectin ratio compared to 5NR using t-test ($p = 0.0499$), supporting our observations in metabolic flexibility.

### 3.3 Serum and liver NR metabolites

No differences were found in levels of NR, nicotinic acid mononucleotide (NaMN), Trp or Nam ($p = 0.0546$ when comparing 900NR with 15, 30, or 180NR groups) in serum.
Figure 4. Effects of dietary NR on NR metabolites. Serum (A-D), liver (E-K). The correlations between NAD\(^+\) and NR concentration in liver (L). NR in mg/kg diet. Data are analyzed using one-way ANOVA and presented as mean ± SEM (n = 10–12 mice per treatment, serum NaMN n = 5–10 mice).

3.4 eWAT morphology and gene expression

We focused the remainder of the analyses on the 5NR, 30NR, and 900NR interventions, the three most dose-representative treatments which also showed differences in maxA\(\text{RER}_{\text{FAO}}\rightarrow\text{CHO}_{2}\).

mRNA levels of rate limiting enzymes involved in the NR pathway, Ido1, Ido2, Tdo2, Qprt (de novo pathway), Nrk1, Nrk2 (Nrk pathway) and Nampt, Nmnat1, Nmnat3 (salvage pathway), were determined. No differences were found in the expression of these genes in either liver (Supporting Information Fig. 3A), small intestinal mucosa (Supporting Information Fig. 3B), skeletal muscle (Supporting Information Fig. 3C), or eWAT (Supporting Information Fig. 3D).

EWAT morphology was assessed by adipocyte size and CLS number. 900NR showed a smaller average adipocyte cell surface area (4015 ± 224.3 \(\mu\)m\(^2\)) than 5NR (5012 ± 363.8 \(\mu\)m\(^2\)) or 30NR (5015 ± 873.5 \(\mu\)m\(^2\)) (Fig. 5A–D). Compared to the two other treatments, 30NR showed the largest number of small adipocytes (100–1500 \(\mu\)m\(^2\)), while 900NR showed the smallest number of large adipocytes (>6000 \(\mu\)m\(^2\)) (Fig. 5E–H). 900NR showed the lowest number of CLS, compared to 5NR and 30NR (Fig. 5I–L).

The expression of genes related to mitochondrial function, adipogenesis, antioxidant response, lipid metabolism, glucose metabolism as well as the adipokines leptin and adiponectin were analyzed. \(C\) was not affected by the treatments (Fig. 6A). The expression of \(Ppary\), \(Sod2\) and \(Prdx3\) was highest in 30NR, a difference that was significant compared to 5NR (Fig. 6A and B). The expression of \(C/ebp\), \(C/ebp\beta\) and \(Pgc1\alpha\), three adipogenesis related genes involved in the regulation of \(Ppary\), showed a similar dose-response pattern as \(Ppary\), but differences did not reach statistical significance (Fig. 6A). The expression of the anti-oxidant genes \(Cat\), \(Gpx3\), \(Sod1\), \(Trp53\) showed a dose-response pattern similar to \(Sod2\) and \(Prdx3\), but without reaching significance (Fig. 6B). Although we did not find differences in the expression of lipid metabolism, glucose metabolism or adipokine genes, 30NR tended to show higher levels of \(Acox1\), \(Fasn\), \(Ppar\), \(Glut4\) and \(Adipoq\) (Fig. 6C and D). \(Lept\) was not different, in agreement with no difference in adiposity.

4 Discussion

To assess beneficial and adverse effects of metabolic health, it is important that metabolic health is compromised, but not deteriorated. The mice in our study were metabolically compromised as they displayed a higher weight gain and worsened metabolic parameters compared to the same substrain of mice on an LFD in another study and comparable or slightly worse than the mice on the HFD in that study [42]. On the other hand, these parameters were not fully deteriorated as, for example, in Ob/Ob mice [43]. Metabolic flexibility is a biomarker for metabolic health that can be assessed non-invasively using indirect calorimetry [31, 32]. Using a fast-refeeding challenge, we here found a dose-response effect, showing that mice fed 30 mg NR/kg diet (30NR) were more metabolically flexible than the wide range of other NR concentrations.

By challenging homeostatic systems subtle metabolic effects, difficult to detect using static biomarkers, may be identified [44, 45]. Indeed, despite an absence of changes in whole body physiological parameters and RER under the non-challenged conditions, we did find differences in metabolic flexibility measured during a fast-refeeding challenge. NR influenced metabolic flexibility in a dose-response

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Figure 5. Effects of dietary NR on eWAT morphology. Representative images for 5NR (A), 30NR (B), 900NR (C); NR in mg/kg diet. Bar represents 100 μm and A–C are same magnification (20×). Cell surface area (D). Frequency distribution (E) and area under the curve (AUC) of small (<100–1500 μm², F), medium (<1500–6000 μm², G) and large (>6000 μm², H) fractions. Representative images of crown-like structures (CLS, indicated by stars) for 5NR (I), 30NR (J), 900NR (K). Bar represents 150 μm and I–K are same magnification (20×). Total CLS number per 100 adipocytes (L). Striped 5NR, continuous 30NR, dotted 900NR. NR in mg/kg diet. Data are analyzed using one-way ANOVA followed by Dunnett’s multiple comparison test, with 30NR as control (F–H), or Bonferroni post-hoc test (D and L) and presented as mean ± SEM (n=6 mice per treatment). *p<0.05, **p<0.01, ***p<0.005, ****p<0.001.

manner, with 30NR being most metabolically flexible. This was not caused by differences in activity or energy expenditure as these parameters were not influenced by NR. The dose-dependent changes of dietary NR in metabolic flexibility might be associated with the activity of sirtuins, enzymes that are regulated by intracellular NAD+ levels and are linked to longevity and other health benefits such as improved insulin sensitivity [46, 47]. The HOMA-IR index [48] and the leptin/adiponectin ratio [49] are markers for insulin resistance. Because insulin resistance interacts with glucose and glucose disposal [29, 50], a causal relation exists. Indeed, insulin resistance and metabolic flexibility were negatively correlated in another study [51]. That the HOMA-IR index and the leptin/adiponectin ratio show a dose-response pattern opposite to the metabolic flexibility dose-response pattern therefore enforces our observation. In this study we

Figure 6. Effects of dietary NR on eWAT gene expression. mRNA levels (normalized to mean of indicated reference mRNAs) of adipogenesis and mitochondria (Clebpα, Clebpβ, Ppargα, Pparg1α); A), antioxidant response (Cat, Gpx3, Sod, Sod2, Trp53; B), glucose metabolism and adipokine (Glut4, Pdk4, Pk, Adipoq, Lep; C) genes, lipid metabolism (Acox1, Cd36, Cpt1, Fasn, Hsl, Ppar; D). White 5NR, gray 30NR, black 900NR. # indicates qRT-PCR determination was after 12-cycle pre-amplification. NR in mg/kg diet. Full gene names are in Supporting Information Table 2. Data are analyzed using one-way ANOVA followed by Dunnett’s multiple comparison test, with 30NR as control and presented as mean ± SEM (n=10–11 mice per treatment). *p<0.05.

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used a semi-synthetic obesogenic diet with moderate levels of fat (40% en%) resembling the average fat intake in the Netherlands and identified 30NR as the most optimal dose of NR in terms of metabolic health. Metabolic flexibility was not improved at 900NR (equivalent to 102 mg NR/kg body weight (BW)/day at the start of the experiment to 65 mg NR/kg BW/day in the end); if anything it was decreased. This contrasts with Canto et al. who found that metabolic flexibility, assessed by the difference of RER between dark phase and light phase, was increased by supplementation with 400 mg NR/kg BW/day of a high fat diet that already contained 30 mg/kg diet of vitamin B3 (Nam) [8]. This improvement in metabolic flexibility, on the other hand, was not observed in mice fed a high-fat-high-sucrose (HFFHS) diet [22], nor in mice fed a chow diet [24], even though the same high dose of NR was used. This suggests a modifying effect of the diet on improvement of metabolic flexibility at high NR concentrations. Alternatively, differences may be due to the absence and presence of the nicotinamide nucleotide transhydrogenase (Nnt) gene. The C57BL/6JRcc mice used in our study have, similar to humans, an intact Nnt gene, which is absent in the strain used by Canto et al. [8]. NNT has a key role in redox and peroxide metabolism [52, 53] and is responsible for maintenance of mitochondrial NAD⁺ levels [54]. Its absence impairs regulation of insulin secretion and thus affects glucose homeostasis [55]. Absence or presence of Nnt thus affects disease development as well as NAD⁺ levels which may explain the observed differences in protective effects of high dose NR supplementation, which needs to be tested experimentally.

WAT plays an important role in metabolic flexibility, in addition to skeletal muscle and liver [34]. Although several studies have shown a decrease in fat mass or WAT weight upon high dose NR supplementation, no WAT morphological parameters were investigated [8,22,56]. Since adipocyte morphology and especially adipocyte size mirrors overall WAT biological function [57], we newly investigated this. 900NR decreased the average adipocyte size compared to 5NR or 30NR. However, the calculated total adipocyte number (according to Skurk T et al., 2007) [57], tended to be higher in the 900NR group as there were no differences found in eWAT weight (data not shown). 30NR showed less medium-size adipocytes, compared to 5NR and 900NR, but had higher numbers of small (compared to 5NR and 900NR) as well as large (compared to 900NR) adipocytes. The functional differences between small and large adipocytes have been described in several studies. Smaller adipocytes tend to be more sensitive to insulin stimulation, showing a two times larger translocation of GLUT4 to the plasma membrane and increased adiponectin release [58, 59]. Large adipocytes, on the other hand, have a higher ability to buffer fatty acid flux and promote lipid mobilization [60]. We sacrificed the mice in the fasted state and it may have been necessary to sacrifice them at the peak of RER (3h after refeeding) to detect clear differences in the serum parameters related to adipose tissue function. Nevertheless, the adipocyte size frequency distribution in 30NR agrees with the dose-response profile of HOMA-IR index and serum adipokines, supporting the best metabolic flexibility at 30NR.

We examined the inflammatory state of the adipocytes by Mac-2 staining and measuring CLS in eWAT. CLS per 100 adipocytes were lowest in 900NR compared to 5 NR and 30NR, although also there the numbers of CLS remained low. This supports the association of CLS with (clearance of) large adipocytes. Our data may also indicate that 900NR affects WAT inflammation beneficially. In agreement, treatment with 200 mg NA/kg body weight/day for 5 weeks was shown to attenuate WAT inflammation by reducing the expression of gene MCP-1, IL-1b and the pro-inflammatory M1 macrophage marker CD11c in HFD-fed mice [61]. However, when taking the total adipocyte number into account, there is no statistical difference in total number of CLS in eWAT (data not shown). To definitely conclude on the effect of NR on inflammation in our study, more inflammatory markers, such as Tumor Necrosis Factor α, should be measured.

PPARγ is the master regulator of adipogenesis [62]. Ppary mRNA levels were significantly higher at 30NR compared to 5NR, the same trend, although not significant, was seen for Acox1 and Glut4, key genes involved in fatty acid oxidation and glucose disposal. PPARγ activation in adipocytes has been shown to promote glucose metabolism by facilitating glucose uptake and enhancing glucose oxidation, as well as to increase lipid metabolism by stimulating fat uptake and mobilization and enhancing free fatty acid oxidation [63, 64]. Thus, PPARγ may contribute to the increased metabolic flexibility of 30NR compared to 5NR. Ppary was not further increased at 900NR. A similar pattern is seen for antioxidant defence.

SOD2 and PRDX3 were significantly elevated at the transcriptional level at 30NR compared to 5NR, suggesting a higher mitochondrial antioxidant defence activity. This was not due to a change in mitochondrial density (Supporting figure 3), confirmed by no change in expression of Cs. Since selective downregulation of Sod2 and Prdx3 in eWAT upon high fat diet feeding resulted in oxidative stress [65], their up-regulation suggests increased protection against damage by reactive oxygen species (ROS). As for Ppary, 900NR did not increase the expression of Sod2 and Prdx3 any further. Our data suggest that adipogenesis and antioxidant response in eWAT are sensitive to low dietary NR, but not to supplemental NR.

Vitamin B3 deficiency leads to low tissue or blood NAD⁺ level and growth retardation in rodents [3,4,66]. In our study, no decrease in body weight or body mass were observed in 5NR, neither were any vitamin B3 deficiency symptoms, such as rough skin, diarrhea seen (data are not shown). Most importantly, there were no differences between the treatments in the expression of the genes involving in NAD⁺ biosynthesis in many tissues, e.g. liver, skeletal muscle, WAT and mucosa. In line with this, NAD⁺ levels and the other NR metabolites in liver and serum did not show differ between treatments. It should be noted that the Trp content used in this study
can maintain normal vitamin B3 status in mice fed a vitamin B3-free diet [17, 18]. Furthermore, NR has been shown be converted to Nam before being absorbed or reaching tissues [67, 68], where it is rapidly metabolized [69]. Considering that fasted plasma Nam levels are very low [70], blood and tissue collection at the fasted status may provide another explanation for the small alteration in NR metabolites. Regrettfully, we did not collect whole blood at section for the whole blood NAD measurement, which has been shown as a good biomarker for vitamin B3 status [71]. Nevertheless, metabolic flexibility and gene expression in eWAT suggest that the 5NR mice were are less metabolically healthy. 5 mg NR/kg diet, with low but sufficient Trp, may therefore constitute a potential cut-off for marginal vitamin B3 sufficiency, with metabolic flexibility as a sensitive marker. However, before this can be definitively concluded, dose-response studies with lower vitamin B3 concentrations and other forms of vitamin B3 are needed.

In conclusion, we investigated the effects of a wide range of dietary NR on metabolic health, focusing on metabolic flexibility and WAT function. To the best of our knowledge, this is the first time that NR is used as an exclusive source of vitamin B3, applied in a range between marginal sufficiency and supplemental amounts in the context of an mildly obesogenic diet in a rodent study. Based on metabolic flexibility, the data linked to WAT function and no effect on growth, we conclude that 30 mg NR/kg diet constitutes the optimal concentration to support metabolic health.

W.S., M.A.H., D.A.M.D., and J.K. were responsible for the study concept and design; W.S. and H.S. conducted the animal experiment; W.S., J.T., M.S., L.A., and B.H. performed biochemical, molecular and histological analyses. W.S. performed statistical analysis and made an initial draft of the manuscript. W.S., M.A.H., and J.K. wrote the manuscript. All authors revised and approved the final version of the paper and agree to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. All persons designated as authors qualify for authorship, and all those who qualify for authorship are listed.

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