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EFFECTS OF DIETARY GAMMA-CYCLODEXTRIN ON VOLUNTARY ACTIVITY AND MUSCLE STRENGTH IN MICE

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Gamma-cyclodextrin (γ CD) is a cyclic oligosaccharide consisting of eight α -(1,4)-linked glucopyranose subunits, which is often used in the food and pharmaceutical industries. However, little is known regarding the metabolic activity of “empty” γ CD *per se*. Therefore, in the present study young C57BL/6 male mice received a control diet (CON) or an experimental diet that was supplemented with 12.88% γ CD exchanged against corn starch. After 6 weeks of treatment, the voluntary wheel running activity was monitored and the muscle strength of mice was measured by employing Kondziela’s inverted screen test and forelimb grip strength assay. The γ CD-treated mice covered a significantly larger distance per night (CON 8.6 km, γ CD 12.4 km) and were significantly longer active (CON 340 min, γ CD 437 min). Moreover, γ CD-treated mice significantly performed better at the inverted screen test indicated by an enhanced Kondziela score (CON 3.10, γ CD 4.63). These data suggest that dietary γ CD leads to an increased endurance. We also found a slightly anti-glycemic effect of γ CD during oral glucose tolerance test. However, our mice from the γ CD group exhibited no difference in terms of GLUT2 protein level in ileum tissue nor increased muscle glycogen storage. Furthermore, γ CD exhibited no DPP-4 inhibitory activity *in vitro*. By analysing candidate muscle genes and proteins related to endurance and muscle performance we did not observe any differences in terms of Sirt1, Pgc1 α , Cpt1b, Mef2c, Myh1 and Myh2 gene expression levels as well as total oxidative phosphorylation (OXPHOS), mtTFA and GLUT4 protein expression levels in skeletal muscle in response to γ CD. We could not fully establish the exact underlying molecular mechanisms of the fitness improvement by dietary γ CD which warrants further investigations.

Key words: *gamma-cyclodextrin, exercise, skeletal muscle, muscle strength, wheel running, glucose transporter, sirtuin 1, total oxidative phosphorylation, sports nutrition*

INTRODUCTION

Cyclodextrins (CDs) are natural conversion compounds of starch, first discovered in 1891 by Villiers (1). CDs are cyclic oligosaccharides consisting of α -(1,4)-linked glucopyranose subunits. The most prominent members of the CD family are α - (six glucopyranose units), β - (seven glucopyranose units) and γ CD (eight glucopyranose units, *Fig. 1*). CDs are characterized by a toroid molecular structure consisting of a hydrophilic outer wall and a less hydrophilic inner wall (1). Through this steric arrangement of glucose units, CDs are soluble in water and can function as important molecular complexation agents (2, 3). In fact, we have previously shown that various nutritional supplements including R- α lipoic acid (4, 5), tocotrienols (6) and Brazilian green propolis (7) exhibit an improved bioactivity in mice attributable to the encapsulation into γ CD. γ CD differs from α CD and β CD in that it is almost completely degraded by salivary and pancreatic α -amylase, similar to digestion of starch and linear dextrans (8-10). Furthermore, γ CD shows the highest water solubility and bioavailability with the least toxicity among all parental CDs (1, 2, 11). A dietary dose of up to 20% γ CD has been shown to be tolerated without toxic side effects in studies

with Beagle dogs and Wistar rats (9, 12). In humans, a single dose of 8 g γ CD in 100 g yogurt was tolerated equally well as the same amount of maltodextrin (13). Accordingly, γ CD acts as a drug carrier and is often used in the food and pharmaceutical industries (2).

Adequate exercise training is a robust strategy to increase strength, endurance and neuroplasticity (14, 15). Furthermore, it is well established that diet composition can affect exercise capacity by altering the availability and utilization of the major energy source (16). Ingestion of carbohydrates prior to and during exercise can enhance endurance capacity. Expanding and maintaining glycogen stores is thereby an often discussed mechanism behind this performance improvement through carbohydrates (17, 18). Moreover, the beneficial effect may depend on the ingested carbohydrate type, which can differently influence factors like glycemic response, gastric emptying, fluid delivery, absorption rates and gastrointestinal comfort (18). These factors define the rate of carbohydrate availability, thereby determining the pre- or post-exercise glycogen stores (18). Thus, identifying natural conversion compounds of starch which increase exercise capacity and deciphering their underlying molecular mechanisms may lead to the development

of new strategies to further improve the beneficial effects of exercise by dietary means.

Voluntary activity in mice can be determined *via* recording wheel running behaviour. This wheel running behaviour of mice is often used to establish physical performance and endurance (19). Voluntary wheel running occurs within routine diurnal rhythmicity patterns in a non-stressed laboratory environment. Further advantages from voluntary wheel running are similarity to natural running behaviour of mice and the absence of direct interference from the researcher. Besides, it appears to satisfy playing and escaping (20, 21). The voluntary wheel running activity differs by sex, strain, age, design of the running wheel, diet and environment conditions (21). The alternative to voluntary wheel running is forced exercise. Forced exercise models, which use aversive stimuli to induce exercise, have the advantage to ensure reproducible distances and speed. However, one has to be aware that experimental parameters such as performance in daylight hours contrary to the murine diurnal pattern may be confounding factors in terms of an appropriate physiological response (19-21).

In our recently published mouse study, γ CD-fed mice were originally considered as an experimental group receiving the vehicle control diet (22). However, phenotypic data obtained from voluntary wheel running and measuring of muscle strength revealed significant differences between control mice (not receiving the vehicle control) and mice supplemented with γ CD. Mice from the γ CD group showed increased voluntary activity in the running wheel and enhanced muscle strength in exercise assays indicating that γ CD has beneficial effects on exercise capacity. To illuminate the underlying molecular mechanism, gene and protein expression levels of candidate genes of the energy metabolism and myogenesis were examined in skeletal muscle tissue.

MATERIALS AND METHODS

Mice and diet

In the present study, we employed the murine inbred strain C57BL/6NRj that has been shown to be appropriate for voluntary wheel running studies (21). Five-week-old male C57BL/6NRj mice were purchased from Janvier Labs (Saint Berthevin Cedex, France). Mice (initial body weight = 17.5–22.4 g) were housed in groups (5 animals per cage) in Makrolon cages with environmental enrichment under controlled climatic conditions (22 – 24°C, 55% relative humidity, 12 h light/dark cycle) at the Institute of Human Nutrition and Food Science at the University of Kiel. The mice had free access to tap water and the experimental diets throughout the feeding study (22). Mice were fed a purified, semisynthetic, energy-dense, high-fat, high-fructose diet (Ssniff S0065-E230) based on casein, corn starch and pork lard as previously described (22). After a 2-week adaptation phase, the experiment began by dividing the mice randomly into groups with 10 mice each. Next to the control group (CON), there was the supplementation group that received the experimental diet supplemented with 12.88% γ CD (CycloChem Bio Co., Ltd., Kobe, Japan) exchanged against corn starch.

Over the entire study period, the health conditions of mice were controlled daily. Mice were weighed weekly and fed daily. We lost two animals over the entire time period, one before the intervention started and one from the γ CD group. After 6 weeks on the experimental diets, the mice were fasted for 5 hours prior to euthanization with carbon dioxide. Blood and tissue samples were collected as described previously (22).

Animal studies were performed according to German and international regulations of animal welfare. The experimental protocol was approved by the local authority on 31 August 2018 (Ministry of Energy, Agriculture, the Environment, Nature and Digitalization, Schleswig-Holstein, V 242-28307/2018).

Voluntary wheel running

A 11.8 cm diameter upright running wheel as part of the PhenoMaster system (TSE Systems GmbH, Bad Homburg, Germany) was placed in isolated cages to measure voluntary wheel running activity. Mice were placed separately in a cage with running wheel access for 12 hours overnight with free access to tap water and experimental diets. To record total distance, time and speed, the software TSE PhenoMaster V5.1.5 (2014-4301) was used.

Measuring the muscle strength of mice

To determine the muscle strength of animals using all four limbs, the Kondziela's inverted screen test was performed (23). Each mouse was placed in the centre of the screen, which was built according to Deacon (23). The screen was smoothly inverted, with the head of the mouse declining first representing the starting point t_0 . The screen was held steadily 20 – 30 cm above a soft surface covered with litter. The time point when the mouse fell off the screen was recorded to a maximum of 20 min. Depending on the hanging time, score points were awarded from 1 (< 5 min) to 5 (\geq 20 min).

Forelimb grip strength was measured by a weight lifting test employing seven weights, which constitutes the apparatus according to Deacon (23). Each weight consists of a ball of tangled fine gauge stainless steel wire, a scale collector (12 g) attached to a series of steel chain links (12 – 14 g). The mass of the weights were therefore: 12, 26, 40, 54, 66, 80 and 94 g. The number of links ranged from zero to six. Each mouse was held by the middle of the tail and lowered to grasp the first weight. As it grasped the scale collector, it was raised until the link was clear off the bench. A hold for 3 s was the threshold criterion. If the weight was dropped before 3 s, the mouse was given a second and third chance (23). A final total score was calculated as the product of the number of links in the heaviest chain held for the full 3 sec, multiplied by the time (s) it was held. If the heaviest weight was dropped before 3 s an appropriate intermediate value was calculated for the time of chain which could not be lifted for full 3 s. Thus, a mouse holding a 5-link weight for 3 s, but unable to lift a 6-link weight, is assigned a score of 15 (5*3). If it held the 6-link weight 3 times for 1 sec, it scores 16 (5*3+1).

Oral glucose tolerance test in mice

After 6 weeks on the experimental diets, mice from each experimental group were fasted for 5 – 6 hours, only receiving water *ad libitum*, prior to the oral glucose tolerance test (oGTT). For the oGTT, 2 g glucose/kg body weight was administered orally by gavage, and glucose levels were measured in blood taken from the tail tip (glucometer, Abbott Freestyle Lite). The blood glucose level was determined immediately prior glucose administration (at 0 min) as fasting blood glucose level and then after 15, 30 and 60 min of glucose administration.

Glycogen in muscle

The glycogen content of muscle was measured with a commercially available fluorometric assay (MAK016, Sigma-Aldrich Chemie GmbH, Darmstadt, Germany) according to the manufacturer's protocol.

RNA isolation and one-step quantitative reverse transcription real-time polymerase chain reaction

Total RNA was isolated from muscle tissue stored in RNAlater (Qiagen, Hilden, Germany) with peqGOLD TriFast (VWR International, Darmstadt, Germany) following the manufacturer's instructions and as described previously (24). A one-step quantitative reverse transcription real-time polymerase chain reaction (one-step qRT-PCR) was carried out with a SensiFAST SYBR No-ROX One-step Kit (Bioline, Luckenwalde, Germany) and SYBR Green detection according to the manufacturer's protocol using a Rotor-Gene 6000 thermocycler (Corbett Research, Sydney, Australia). Primers were designed with Primer 3 Input software (version 4.1.0) and purchased from Eurofins MWG (Ebersberg, Germany). Corresponding primer sequences and annealing temperatures are given in Table 1. RNA amplification for each sample was conducted in duplicate, and each run included a standard curve and a no-template control. Relative mRNA levels of

target genes were normalized to Ap3d1 gene expression, and fold changes relative to the control group are given.

Western blot analysis

Protein expression was determined in whole cell lysates prepared from frozen muscle and ileum tissue, respectively. Protein concentrations were determined with a Pierce bicinchoninic acid (BCA) Protein Assay Kit (Thermo Fisher Scientific, Bremen, Germany) according to the manufacturer's instructions.

For Western blotting, samples containing 30 µg of protein each for muscle tissue and 35 µg of protein each for ileum tissue were heated at 37°C for 20 min with loading buffer, and a Western blot analysis was performed as previously described in detail (25). Target proteins were identified using the corresponding primary antibodies (glucose transporter 2, GLUT2: Novus Biologicals, Colorado, USA; glucose transporter 4, GLUT4: Sigma-Aldrich; transcription factor A, mitochondrial, mtTFA:

Table 1. Primer sequences and annealing temperatures for one-step quantitative reverse transcription real-time polymerase chain reaction.

| Primer | Gene | Gene ID | Primer sequence | T _a [°C] ¹ |
|--------|-------------------------------------------------------------------------|---------|--------------------------------------------------------------------------------|----------------------------------|
| Ap3d1 | Adaptor-related protein complex 3, delta 1 subunit | 11776 | F ² : AGGAGCTGAAGCAGGACAAC R ³ : CGCTTGAATGTGAACTTGGA | 55 |
| Cpt1b | Carnitine palmitoyltransferase 1b | 12895 | F: CTACCACAAAGGTCGCTTCT R: GGATTCTCTGGAACTGCATC | 57 |
| Mef2c | Myocyte enhancer factor 2C | 17260 | F: CTGTCAGCACACTGGGAAAC R: CCTGTGTTACCTGCACTTGG | 60 |
| Myh1 | <i>Mus musculus</i> myosin, heavy polypeptide 1, skeletal muscle | 17879 | F: GCGAATCGAGGCTCAGAACAA R: GTAGTCCGCTTCGGTCTTG | 61 |
| Myh2 | <i>Mus musculus</i> myosin, heavy polypeptide 2, skeletal muscle | 17882 | F: AAGTGAAGTGTGAAAACAGAAGCA R: GCAGCCATTTGTAAGGGTTGAC | 55 |
| Pgc1α | Peroxisome proliferative activated receptor, gamma, coactivator 1 alpha | 19017 | F: TGCCCAGATCTTCTGAACT R: TCTGTGAGAACCCTAGCAA | 57 |
| Sirt1 | Sirtuin 1 | 93759 | F: GTCTCCTGTGGGATTCTGA R: ACACAGAGACGGCTGGAAC | 61 |

¹T_a, annealing temperature; ²F, forward; ³R, reverse.

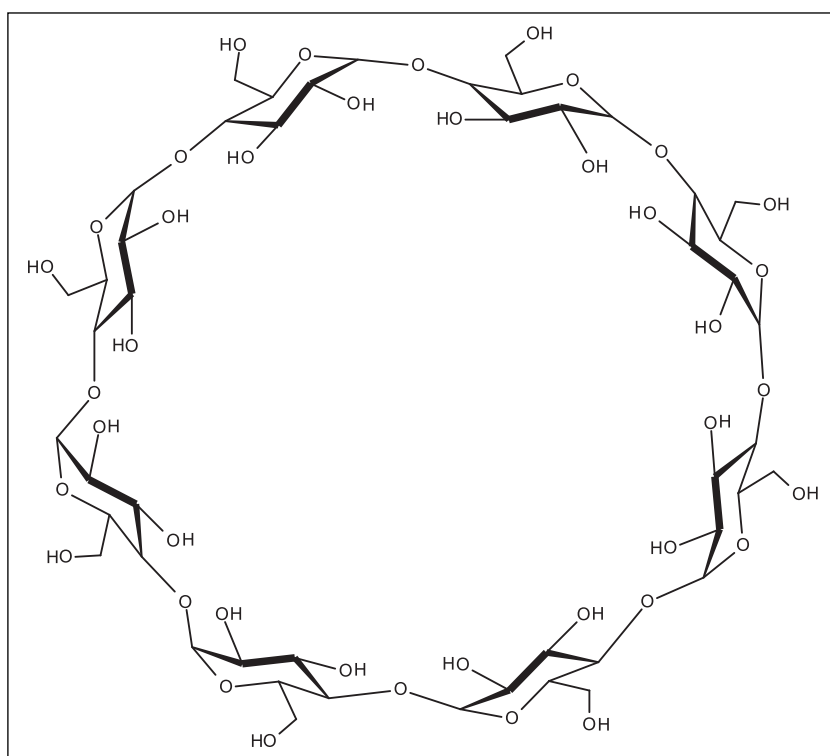


Fig. 1. Chemical structure of γ -cyclodextrin.

Santa Cruz Biotechnologies, Dallas, USA; OXPPOS rodent antibody cocktail: Abcam, Cambridge, United Kingdom) and secondary antibodies. Immunoreactive proteins were detected with enhanced chemiluminescence (ECL) reagents (Thermo Fisher Scientific). Bands were visualized in a ChemiDoc XRS system (Bio-Rad, Munich, Germany). Target protein expression was related to the total protein load per lane.

Dipeptidyl peptidase-4 inhibition assay

Human dipeptidyl peptidase-4 (DPP-4) inhibition assay with γ CD as potential inhibitor was performed *in-vitro* using a human inhibitor screening kit (MAK203, Sigma-Aldrich) according to the manufacturer’s protocol. γ CD was solved in assay buffer and analysed in concentrations of 1, 10, 100 μ g/mL and 1 g/mL.

Statistical analysis

For the statistical analysis, the model of variance analysis was used. The voluntary wheel running performance was

analysed using a mixed-effect model in which mice nested within cages was used as a random variable, whereas diet and run were considered as fixed variables. However, run was not significant and therefore not used for statistical analysis. The same mixed-model was used to analyse the muscle strength data, area under the curve (AUC) from oGTT, fasting blood glucose data and glycogen in muscle. The oGTT was analysed using a mixed-effect model with repeated measures in time. In the model, mice nested within cages was entered as a random variable and diet, run, time and the two-way interaction of diet and time were considered as fixed effects. However, run and the two-way interaction of diet and time were not significant and therefore excluded from the model. Least squares means were separated into significant effects using the Tukey adjustment. Differences among treatments were considered to be significant when $P < 0.05$. Statistical analyses were performed with SAS (release 9.2, SAS Institute). Data from qRT-PCR and Western blot were analysed for normality of distribution (Kolmogorov-Smirnov and Shapiro-Wilk test). In the case of normally distributed data, the *t*-test was applied. In

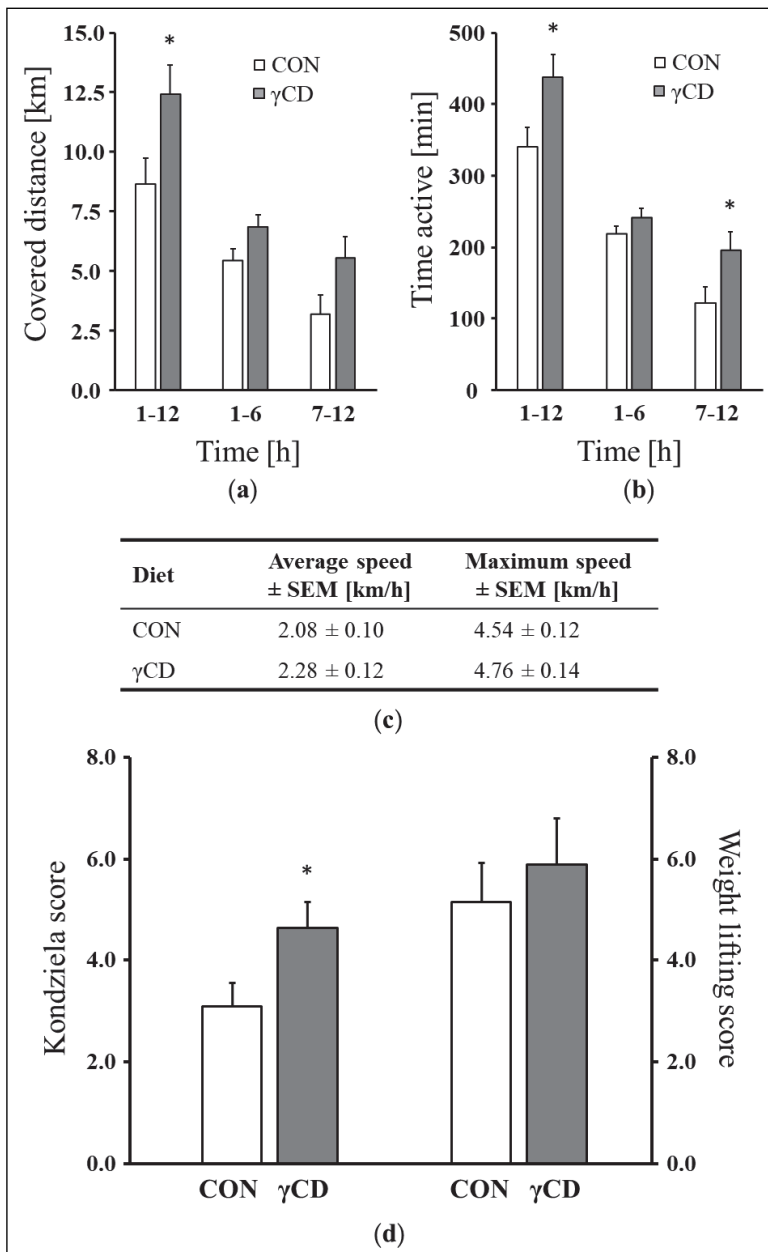


Fig. 2. The mice supplemented with γ -cyclodextrin (γ CD) showed a higher voluntary activity and increased muscle strength compared to the control mice. Mice were fed (*ad libitum*) for 6 weeks with a Western-type control diet (CON) and control diet supplemented with 12.88% γ CD (γ CD) exchanged against corn starch, respectively, before they were placed in a cage with running wheel access for 12 hours overnight.

a) γ CD mice covered a significantly larger distance per night (1 – 12) compared to control mice in the running wheel. When looking at the first and the second half of the night separately, the γ CD mice covered barely significantly larger distances ($P = 0.061$ and $P = 0.057$, respectively). Bars represent the least squares means (LSM) + standard error of the mean (SEM).

b) However, the mice supplemented with γ CD were significantly longer active during the whole night and in the second part of the night compared to CON. Bars represent the LSM + SEM.

c) Average and maximum speed in running wheel were not significantly different between groups. Data are expressed as LSM \pm SEM.

d) Kondziela’s inverted screen test was performed and the maximum time of 20 minutes was the abort criterion. The score from the Kondziela’s inverted screen test was significantly higher in γ CD mice compared to CON mice. The score from the weight lifting test showed no significant differences between the groups. Bars represent the LSM + SEM. All data were analysed with ANCOVA, * $P < 0.05$, $n = 8 - 10$ mice/diet.

the absence of normally distributed data, the Mann-Whitney U test was applied. The results were considered statistically

significant at $P < 0.05$. IBM SPSS statistical software (version 25, IBM: Armonk, NY, USA) was used for the analyses.

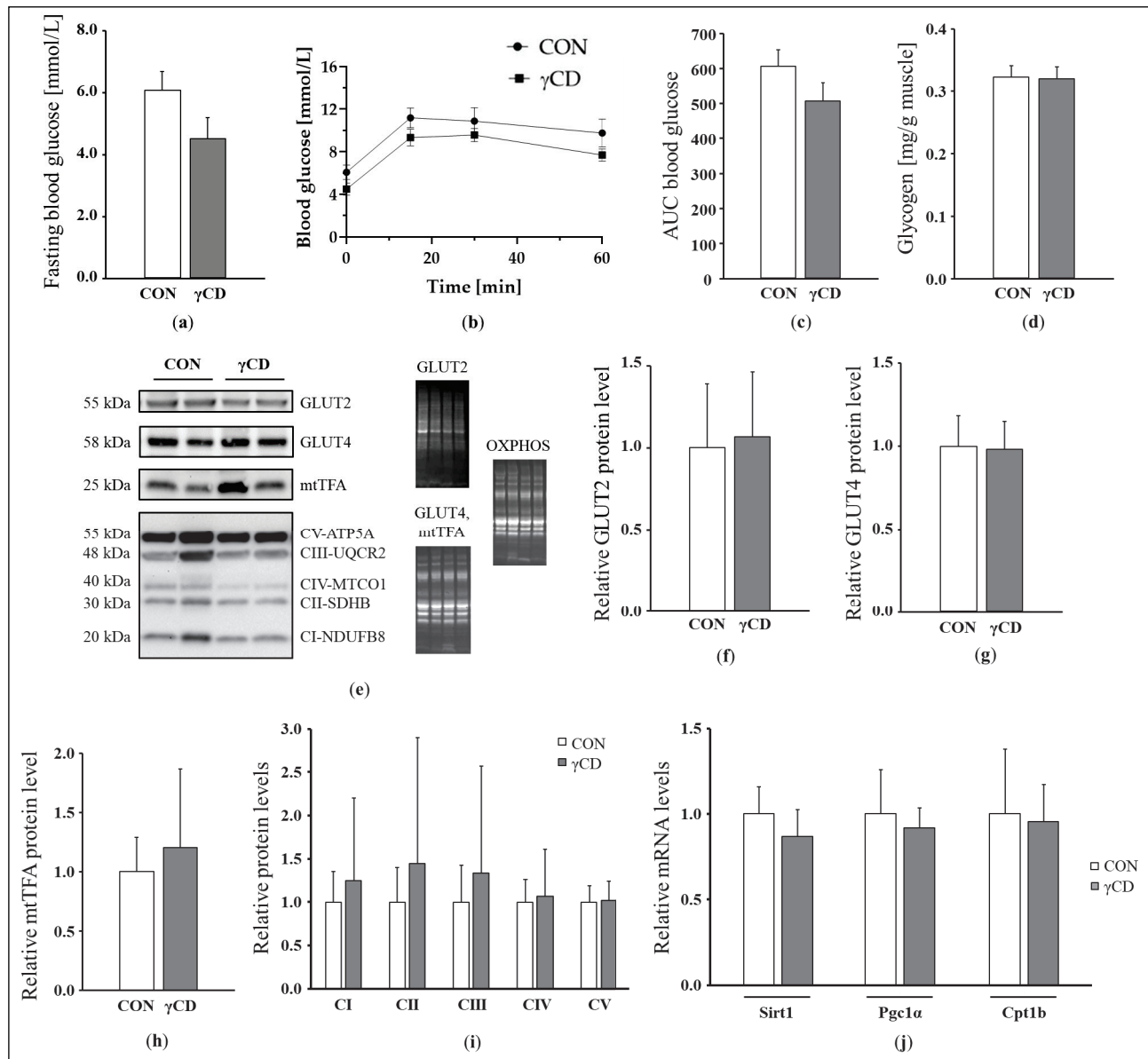


Fig. 3. Effect of γ -cyclodextrin on glucose metabolism and muscle parameters that are known to be related to muscle strength and endurance.

a) Fasting blood glucose levels from oral glucose tolerance test (oGTT) in mice fed (*ad libitum*) a Western-type diet supplemented with 0 (control, CON) or 12.88% γ -cyclodextrin (γ CD) exchanged against corn starch over a 6-week experimental period. Bars represent the least squares means (LSM) + standard error of the mean (SEM).

b) oGTT (LSM \pm SEM) and

c) area under the curve (AUC) of oGTT. Bars represent the LSM + SEM.

d) Glycogen content in muscle was not different between CON and γ CD. Bars represent the LSM + SEM. Data were analysed with ANCOVA, $n = 8 - 10$ mice/diet.

Western blot (e) and densitometric analysis (f, g, h, i) of glucose transporter 2 (GLUT2) in ileum tissue and glucose transporter 4 (GLUT4), transcription factor A, mitochondrial (mtTFA) and mitochondrial respiratory complexes subunits in muscle tissue. Protein levels were determined by Western blotting, and a representative blot with the corresponding stain-free UV image is shown. Protein bands were analysed densitometrically. Values were related to total protein load per lane, and are expressed in relation to the CON group. Bars represent the mean + standard deviation ($n = 8 - 10$ mice/diet), analysed with t-test.

j) No effect of dietary γ CD on mRNA levels of sirtuin 1 (Sirt1), peroxisome proliferative activated receptor, gamma, coactivator 1 alpha (Pgc1a) and carnitine palmitoyltransferase 1b (Cpt1b) in skeletal muscle. Gene expression was analysed *via* a one-step quantitative reverse transcription real-time polymerase chain reaction (one-step qRT-PCR). All qRT-PCR data were normalized to Ap3d1 gene expression and are expressed in relation to the CON group. Data are given as the mean + standard deviation ($n = 8 - 10$ mice/diet) and were analysed with t-test.

RESULTS

Dietary γ CD increased voluntary activity and muscle strength

The mice supplemented with γ CD showed a higher voluntary activity in the running wheel compared to CON mice, especially in the second part of the night. The γ CD mice covered significantly larger distances with an average distance of 12.4 ± 1.2 km (CON 8.6 ± 1.1 km) per night (Fig. 2a) and were significantly longer active during the whole night (CON 340 ± 28 min, γ CD 437 ± 31 min; Fig. 2b). However, average and maximum speed were not significantly different between the groups (Fig. 2c). To get a better understanding of running characteristics, we separated the night period into two halves (first six hours versus last six hours). In the first half of the night, the running distance of the γ CD supplemented mice was barely significantly different from controls (CON 5.4 ± 0.5 km, γ CD 5.6 ± 0.9 km, $P = 0.061$; Fig. 2a), while the active time of both groups was quite similar (CON 218 ± 10 min, γ CD 242 ± 12 min; Fig. 2b). In the second half of the night, γ CD-treated mice were significantly longer active (CON 121 ± 23 min, γ CD 196 ± 25 min; Fig. 2b). However, the covered distance was only barely significantly different (CON 3.2 ± 0.8 km, γ CD 5.6 ± 0.9 km, $P = 0.057$; Fig. 2a).

To test the muscle strength of our mice, the Kondziela's inverted screen test as well as the weight lifting test were performed. From the mice supplemented with γ CD, 5 out of 8 animals (62.5%) reached the abort criterion timepoint of 20 min, whereas in the CON group only 4 out of 10 animals reached the 20 min (40.0%). Therefore, γ CD mice hang significantly longer at the inverted screen, shown as Kondziela score (CON 3.1 ± 0.5 , γ CD 4.6 ± 0.5 ; Fig. 2d). However, the weight lifting score representing forelimb grip strength showed no significant differences between the two groups (CON 5.1 ± 0.8 , γ CD 5.9 ± 0.9 ; Fig. 2d).

 γ CD supplementation did not significantly improve glucose metabolism

The effect of dietary γ CD on glucose metabolism is shown in Fig. 3. Fasting blood glucose levels of γ CD-treated mice were 26% lower than in control mice but did not reach statistical significance ($P = 0.105$; Fig. 3a). Furthermore, administration of dietary γ CD slightly decreased glycemia over time during the oGTT and the corresponding AUC (Fig. 3b and 3c). The peak blood glucose occurred at 15 – 30 min post glucose administration in both groups (Fig. 3b). To examine if γ CD affects glucose transport in the intestine, GLUT2 protein levels in ileum tissue were determined. There was no significant difference between CON and γ CD in terms of GLUT2 protein level (Fig. 3e and 3f). The glycogen content in muscle of γ CD mice was not significantly different between groups (Fig. 3d). GLUT4 protein level in muscle tissue from CON and γ CD mice showed no significant difference (Fig. 3e and 3g). The *in vitro* human DPP-4 inhibition assay exhibited no DPP-4 inhibitory activity of γ CD at 1, 10, 100 μ g/mL or 1 mg/mL, respectively (data not shown).

No effect of γ CD on gene expression levels of regulators of mitochondrial biogenesis and muscle development in skeletal muscle

To examine if γ CD affects the mitochondria of skeletal muscle cells, the steady-state mRNA levels of the genes encoding proteins centrally involved in mitochondrial biogenesis and function were determined *via* qRT-PCR. Dietary γ CD had no effect on gene expression of sirtuin 1 (Sirt1),

peroxisome proliferative activated receptor, gamma, coactivator 1 alpha (Pgc1 α) and carnitine palmitoyltransferase 1b (Cpt1b) in skeletal muscle (Fig. 3j).

Furthermore, myocyte enhancer factor 2C (Mef2c) was examined *via* qRT-PCR to determine skeletal muscle development and function in CON versus γ CD mice. There was no significant difference between the groups in terms of Mef2c mRNA levels (CON 1.00 ± 0.18 , γ CD 0.93 ± 0.08 , data not shown). To determine muscle fibre type composition, the mRNA levels of *Mus musculus* myosin, heavy polypeptide 1, skeletal muscle (Myh1) and *Mus musculus* myosin, heavy polypeptide 2, skeletal muscle (Myh2) were measured *via* qRT-PCR. γ CD supplementation had no significant effect on muscle fibre type composition regarding Myh1 (CON 1.00 ± 0.20 , γ CD 1.16 ± 0.30 , data not shown) and Myh2 (CON 1.00 ± 1.39 , γ CD 0.89 ± 0.80 , data not shown).

No difference in mitochondrial total oxidative phosphorylation protein levels in muscle

The levels of mitochondrial- and nuclear-encoded protein subunits involved in OXPHOS were evaluated using muscle extracts (Fig. 3e and 3i). The densitometric analysis (Fig. 3i) showed no significant differences in protein levels of mitochondrial respiratory complexes subunits between CON and γ CD mice. While complexes IV (CIV) and CV exhibited almost similar protein levels between the two groups, CI, CII and CIII were slightly increased in γ CD-treated mice. Even the protein level of mtTFA in muscle tissue showed no significant differences between the groups (Fig. 3e and 3h).

DISCUSSION

Little is known regarding the metabolic activity of "empty" γ CD *per se*. Therefore, in the present mouse study we supplemented our experimental diet with 12.88% γ CD exchanged against corn starch. Accordingly, the cyclic oligosaccharide γ CD was the main complex carbohydrate source for γ CD-treated mice when compared to the remaining 3.62% corn starch. A similar γ CD concentration has been previously used in a feeding study in dogs without toxic side effects (12).

We observed a significant improvement in nocturnal voluntary wheel running activity by 44% and an enhanced muscle strength in our γ CD-treated mice. The mice from CON group ran on average 8.6 km overnight, which is similar to the running distances reported for untreated male C57BL/6 mice in previous studies (20, 21, 26). In addition, the activity pattern of our untreated mice is consistent with published data. Under normal conditions, mice ran most intensely the first several hours of the active dark cycle (19, 20, 26). Accordingly, our CON mice ran a significantly higher distance during the first half of the night comparing with the second half. The main observed difference between CON and γ CD mice was in the second half of the night regarding covered distance and running time. γ CD mice spent 61% more time running than CON mice during the last six hours of the dark cycle. A detailed analysis of mice running trends revealed that during their actual running activity there was no difference in average and maximum running speed between untreated and treated mice. Our data suggest that dietary γ CD caused an increased endurance in our mice. Phelps *et al.* (27) described an exhaustion phenotype when they compared the running activity of wild-type C57BL/6 female mice and α -klotho-deficient mice. Here, the klotho-deficient mice ran significantly less than the wild-type control. Similar to our results for γ CD supplemented animals, the mutant mice ran at the same speed as the wild-type mice. However, in contrast to

our γ CD mice, α -klotho-deficient mice spent less time running than the control wild-type animals, indicating that the lack of α -klotho protein resulted in a more quickly exhaustion (27). Accordingly, the running distance in combination with running time reflects endurance capacity in an appropriate way.

It has been assumed that voluntary wheel running could serve also as a measure of muscle performance (19). Consistent with that, we observed a significant increase in all four limbs muscle strength in γ CD mice *via* Kondziela's inverted screen test although the underlying molecular mechanisms remain to be further elucidated. So far, administration of γ CD in mice led to a more active and stronger phenotype, probably through increasing endurance.

Previous studies have shown that γ CD is almost fully digested and absorbed within the small intestine (10, 28), whereas absorption of intact γ CD from gastrointestinal tract is extremely low (0.02%) (29). Moreover, according to the findings by Spears *et al.* (28), an impact of γ CD on intestinal microbiota is rather unlikely. Hence, the gastrointestinal tract with its digesting enzymes is most probably the primary site of action of γ CD. Although γ CD is resistant towards degradation by β -amylase, an enzyme that hydrolyzes starch from the non-reducing end, it is a substrate of α -amylase, which hydrolyzes α -bonds from large polysaccharides like starch (9). Accordingly, γ CD is almost completely degraded to malto-triose, maltose and glucose by amylases (8-10). The degradation of γ CD by α -amylase starts with a ring-opening reaction which results in linear malto-octaose that can be further degraded by amylases (10). Owing to its cone-shaped structure and its hydrophobic core, the ring-opening reaction by α -amylase represents the slowest step of the degradation process of γ CD. Accordingly, γ CD is most probably slower digested than the easily digestible linear polysaccharide maltodextrin (10, 30). In this context, it is interesting that depending on its actual complex structure starch is classified as slowly or rapidly digestible starch. Compared to the rapidly digestible form, the amylase-mediated breakdown of slowly digestible starch is hindered, which affects the rate of glucose release and glucose absorption (31). The impact of meals with different glycemic indexes (GI) on exercise was examined by Kirwan *et al.* (32). A pre-exercise meal with a moderate GI significantly increased performance time in young men in contrast to a high GI meal prior to exercise or water as control. In addition, consuming the moderate GI meal maintained euglycemia during exercise over an extended period of time (32). In line with this assumption, it has been reported that γ CD results in a lower postprandial glucose response when compared with maltodextrin (30) and provides glucose over a prolonged period of time during exercise. Asp *et al.* (30) performed a study with 32 healthy adult subjects who received 50 g of carbohydrate from γ CD or maltodextrin *via* beverages. In this study, γ CD reduced the postprandial glycemic and insulinemic response compared with the rapidly digested maltodextrin in humans without carbohydrate malabsorption. Additionally, inhibitory effects of CDs on the activity of amylase have been demonstrated (11, 33, 34). When amylose is used as substrate, α -amylase activity is competitively inhibited by γ CD (34). This can additionally attenuate the postprandial glycemic and insulinemic response and may contribute to a steady and longer lasting supply of glucose through a slowed polysaccharide digestion rate in γ CD supplemented animals.

Remarkably, Takii *et al.* (35) reported an enhancement of swimming endurance in mice by highly branched cyclic dextrin (HBCD). Furuyashiki *et al.* (36) compared the efficacy of HBCD with maltodextrin during endurance exercise on the rating of perceived exertion (RPE) in humans. Interestingly, the increase in RPE was significantly lower after ingesting HBCD than maltodextrin. It has to be noted though that HBCD differs

structurally from γ CD, but this likewise digested cyclic dextrans may affect endurance capacity in a similar way as γ CD.

However, the improvement in voluntary wheel running activity and enhanced muscle strength by γ CD could also be a result of behavioral change in mice. Two-hydroxypropyl- γ CD showed positive effects in Niemann-Pick type C (NPC1) patient-derived fibroblasts through restoring cellular homeostasis (37). Besides, there is evidence that hydroxypropyl- β -CD treatment *via* subcutaneous injection improved learning and memory deficits in a transgenic mouse model of Alzheimer disease (38). An overview of β CDs and its derivatives in the treatment of neurodegenerative diseases is given by Coisne *et al.* (39). Unlike our experimental design, the injection or nasal delivery of CD partly avoid the gastrointestinal tract with its digesting enzymes, thus resulting in primary intact CD molecules in the circulatory system. The extent to which dietary cyclodextrins have an influence on brain function remains open.

We examined several muscle parameters that are known to be related to muscle strength and endurance in order to find out whether γ CD supplementation affected muscle physiology. The glucose transporter GLUT4 mediates insulin-stimulated glucose transport in muscle tissue (40) and muscle glycogen plays an important role in athletic performance (41), since pre-exercise muscle glycogen content has been shown to determine muscle time to fatigue (42). However, there was no effect of dietary γ CD on muscle GLUT4 protein and glycogen content in our mice.

Endurance capacity depends among other things on muscle fibre type proportions (43). So, we took a deeper look in skeletal muscle structure of our mice. Dietary γ CD did not alter gene expression of Mef2c, a key regulator of muscle development. Even the muscle fibre type composition was not shifted in γ CD-treated mice after 6 weeks on experimental diet and one night in the running wheel compared to CON mice.

Pgc1 α is a transcription factor in muscle, adipose tissue and other tissues, controlling oxidative metabolism including muscle fibre type switching (44). Recently published studies with PGC1 α transgenic mice have shown that increased skeletal muscle PGC1 α expression decreased fatigability (45) and increased exercise ability (46, 47). Additionally, enhanced oxidative capacity and endurance were accompanied with enriched muscle expression of genes and proteins involved in mitochondrial biogenesis and function, including Sirt1, Pgc1 α and mtTfa (48) as well as increased mitochondrial OXPHOS efficiency (49). Besides, it has been shown that endurance training increased the expression of oxidative and lipid metabolism markers like CPT1 (50) because mitochondrial fatty acid oxidation with its key enzyme CPT1 is the main muscle energy source during endurance exercise (51, 52). Moreover, Miklosz *et al.* (53) showed that a single exercise bout induced mRNA and protein expression of pro-lipolytic-related proteins in skeletal muscle of male rats, such as adipose triglyceride lipase and hormone sensitive lipase. However, we did not observe differences in terms of Sirt1, Pgc1 α and Cpt1b gene expression levels as well as OXPHOS and mtTFA protein expression levels in skeletal muscle in response to dietary γ CD and exercise.

A strength of this study was that we conducted a controlled feeding trial with comprehensive phenotyping. Furthermore, we determined many biomarkers related to skeletal muscle function and strength in our mice. In this context, it has to be noted that we partly measured potential targets of gCD only on the mRNA or protein levels, respectively. Besides, GLUT4 membrane translocation would have been interesting in a comparison between control and γ CD mice. Another limitation of our study was that we examined the effects of only a single concentration of γ CD over a relatively short experimental period of 6 weeks. Otherwise, it may be possible that the significant improvement in running wheel activity and muscle strength was not mainly

mediated *via* a direct transcriptional response or part of a structural and metabolic adaptation in the skeletal muscle but rather mediated *via* altered glucose availability. The underlying mechanisms by which cyclic dextrans improve endurance are still not fully understood and further studies are needed to elucidate how γ CD can provide a blunted postprandial glucose response and affect endurance. In fact, a higher dietary γ CD concentration may result in a more pronounced attenuation in postprandial glucose response and insulin secretion. In this context, a limitation of the present mouse feeding trial was that we conducted blood glucose measurements without having measured insulin and only at one time point. Corresponding analyses before, during and after exercise would have been interesting (32).

Overall, based on present and literature data it is suggested that γ CD moderately improves endurance and muscle strength, which renders it as a putative carbohydrate source in sports nutrition. Endurance athletes may benefit from the blunted postprandial glucose response after γ CD intake. Therefore, present findings in mice should be verified in further studies in humans including athletes. In particular, since it has been shown that regular physical activity improves nitric oxide-dependent endothelial function in healthy adults (54), it would be of interest to study if γ CD could enhance the cardiovascular effects of aerobic training. Furthermore, endurance and muscle strength decline with age (55, 56). Thus, it is plausible to address the question whether γ CD may partly counteract age-dependent decline in endurance and muscle strength in ageing rodent models as well as in the elderly.

In conclusion, our results suggest that administration of γ CD moderately enhanced endurance capacity, indicated by increased voluntary activity and enhanced muscle strength in mice. Exact underlying molecular mechanisms still need to be elucidated.

List of abbreviations: AUC, area under the curve; BCA, bicinchoninic acid; C, complex; CD, cyclodextrin; CON, control; Cpt1b, carnitine palmitoyltransferase 1b; DPP-4, dipeptidyl peptidase-4; ECL, enhanced chemiluminescence; F, forward; GI, glycemic index; GLUT2, glucose transporter 2; GLUT4, glucose transporter 4; HBCD, highly branched cyclic dextrin; LSM, least squares means; Mef2c, myocyte enhancer factor 2C; mtTFA, transcription factor A, mitochondrial; Myh1, *Mus musculus* myosin, heavy polypeptide 1, skeletal muscle; Myh2, *Mus musculus* myosin, heavy polypeptide 2, skeletal muscle; oGTT, oral glucose tolerance test; one-step qRT-PCR, one-step quantitative reverse transcription real-time polymerase chain reaction; OXPHOS, total oxidative phosphorylation; Pgc1 α , peroxisome proliferative activated receptor, gamma, coactivator 1 alpha; R, reverse; RPE, rating of perceived exertion; SEM, standard error of the mean; Sirt1, sirtuin 1; Ta, annealing temperature; γ CD, γ -cyclodextrin.

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