Aus dem Institut für Veterinär-Physiologie des Fachbereichs Veterinärmedizin der Freien Universität Berlin

Effects of the Flavonol Quercetin on Glucose Metabolism and Health Status in Neonatal Calves

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Jeannine Gruse

Tierärztin

aus Räckelwitz

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Dekan: Univ.-Prof. Dr. Jürgen Zentek **Erster Gutachter:** Univ.-Prof. Dr. Jörg Aschenbach

Zweiter Gutachter: PD Dr. Harald M. Hammon **Dritter Gutachter:** Univ.-Prof. Dr. Jürgen Zentek

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CONTENT

List	of table	s	III
List	of figure	es	V
Abl	reviatio	าร	. VIII
1.	General	Introduction	1
2.	Literaur	e overview	4
2	.1 Ger	neral aspects on nutritional and immune status in neonatal calves	4
	2.1.1	Postnatal adaptation of glucose metabolism in calves	4
	2.1.2	Maturation of the digestive system	6
	2.1.3	Immune system and antioxidative status	7
2	.2 Imp	ortance of colostrum	9
	2.2.1	General aspects	9
	2.2.2	Effects on GIT and glucose metabolism	10
	2.2.3	Effects on immune system and antioxidative status	11
2	.3 Cha	racteristics of flavonoids	13
	2.3.1	General aspects	13
	2.3.2	Absorption and metabolism of flavonols	14
	2.3.3	Interaction with glucose metabolism	15
	2.3.4	Effects on performance and intestinal health	16
	2.3.5	Effects on antioxidative and anti-inflammatory status	17
3. I	Effects o	f oral quercetin supplementation on splanchnic glucose metabolisi	m in
1-w	eek-old d	calves depend on diet after birth	31
3	.1 Abstra	ct	32
3	.2 Introdu	ction	33
3	.3 Materia	als and Methods	34
	3.3.1 An	imals, husbandry, and feeding	34
	3.3.2 An	alyses of milk	35
	3.3.3 Xy	lose absorption studies	35
	3.3.4 Pla	sma flavonols, metabolites, and hormones	35
	3.3.5 Tra	acer studies	36
	3.3.6 Liv	er sampling and analyses	37
	3.3.7 Sta	atistical analyses	37
3	4 Results	3	38
	3.4.1 Fe	ed intake and growth performance	38
	3.4.2 Xy	lose absorption	38

	3.4.3 Plasma flavonols, metabolites, and hormones	39
	3.4.4 Tracer studies	44
	3.4.5 Analyses of liver tissue	48
3.	5 Discussion	48
3.	.6 Acknowledgments	52
3.	.7 Author Contributions	52
3.	.8 References	53
3.	.9 Supporting Information	57
4. (Quercetin Feeding in Newborn Dairy Calves Cannot Compensate Colostrum	De-
priv	ration: Study on Metabolic, Antioxidative and Inflammatory Traits	60
4.	.1 Abstract	61
4.	2 Introduction	62
4.	.3 Materials and Methods	63
	4.3.1 Animals, Husbandry and Feeding	63
	4.3.2 Treatment	64
	4.3.3 Blood Analyses	64
	4.3.4 Liver Tissue Analyses	66
	4.3.5 Statistical Analyses	68
4.	.4 Results	68
	4.4.1 Feeding, Growth Performance and Health Status	68
	4.4.2 Flavonoid Content	71
	4.4.3 Plasma Metabolites and Hormones	72
	4.4.4 Immunoglobulins and Acute-Phase Proteins	76
	4.4.5 Antioxidative Status	76
	4.4.6 Hepatic Gene Expression	78
4.	5 Discussion	79
4.	.6 Acknowledgments	82
4.	.7 Author Contributions	82
4.	.8 References	83
4.	.9 Supporting Information	88
5. G	eneral discussion	100
	nmary	
	ammenfassung	
App	pendix	XI
	olication list	
	ıksagung	
		VVI

LIST OF TABLES

Table 3.1: Plasma concentrations of urea, NEFA, glucagon and insulin:glucagon ratio in 7-d-old calves fed C or F on d 1–2 and supplemented (Q+) or not supplemented (Q-) with quercetin on d 2–8
Table 3.2: Rates of glucose appearance, first pass glucose uptake, and AUC for $[^{13}C_6]$ -glucose enrichment and blood $^{13}CO_2$ enrichment (derived from $[^{13}C_6]$ -glucose) of 7-d-old calves fed either C or F on d 1 and 2 and supplemented (Q+) or not supplemented (Q-) with quercetin on d 2–8 ¹
Table 3.3: Glycogen concentration and relative mRNA expression in liver biopsy samples 2 h after feeding in 8-d-old calves fed colostrum (C) or formula (F) on d 1 and 2, and supplemented (Q+) or not supplemented (Q-) with quercetin from d 2–8
Supplemental Table 3.1: Composition of colostrum, formula and milk replacer fed to neonatal calves during the first 8 d of life
Supplemental Table 3.2: Characteristics of primers and real-time RT-PCR conditions 58
Table 4.1: Characteristics of Primers and Real-Time RT-PCR Conditions67
Table 4.2: Metabolites and Hormones of Calves during the First Week of Life73
Table 4.3: Hepatic mRNA Expression of Inflammatory and Antioxidative Traits in Newborn Calves
Supplemental Table 4.1: Complete data set of parameters regarding daily milk intake and health status as shown in Figure 4.1
Supplemental Table 4.2: Complete data set of flavonoid measurements in blood plasma as shown in Figure 4.2
Supplemental Table 4.3: Complete data set of plasma concentrations of metabolites and hormones as shown in Figure 4.3 and Table 4.2
Supplemental Table 4.4: Complete data set of immune and inflammatory status in blood plasma as shown in Figure 4.396

Supplemental Table 4.5: Complete data set of antioxidative status in blood plasma as shown in Figure 4.4
Supplemental Table 4.6: Complete data set of hepatic mRNA expression of inflammatory and antioxidative traits as shown in Table 4.399
Appendix
Table A1: Correlation (Pearson coefficients) between flavonol concentration and selected
parameters in plasma of calves fed quercetin twice daily from d 2-8 of lifeXI
Table A2: Correlation (Pearson coefficients) between flavonol concentration and selected
metabolites and hormones on d 7 in plasma in calves fed quercetin twice daily from d
2-8 of lifeXIII
Table A3: Correlations (Pearson correlation coefficients) between plasma flavonol
concentrations (2 h after feeding on d 7) and mRNA abundances of hepatic genes
involved in antioxidative status and inflammation (2 h after feeding on d 8) in calves
orally supplemented with quercetin aglycone (50 mg/(kg BW × d)) for 7 daysXVI

LIST OF FIGURES

Figure	2.1: Percentage composition of macronutrient intake in calves (minor nutrients are disregarded). Composition of colostrum and mature milk was obtained from pluriparous Holstein cows
Figure	2.2: Simplified scheme of the hepatic glucose metabolism. Rate-limiting enzymes of endogenous glucose production (gluconeogenesis, glycogenolysis) are bold black, enzymes of glucose consuming pathways (glycolysis, glycogen synthesis) are italic grey
Figure	2.3: Endogenous formation and neutralization of ROS
Figure	2.4: Plasma xylose concentrations following intestinal absorption in 5d old calves after different initial colostrum supply, modified from Hammon and Blum
Figure	2.5: Antioxidative status of colostrum and plasma of calves fed with respective colostrum, modified from Albera and Kankofer
Figure	2.6: Chemical structure of the most common flavonoid subclasses and names of some representative compounds as well as their dietary sources
Figure	3.1: Plasma concentrations of glucose (A) and xylose (B) before and after feeding milk replacer and 0.5 g/kg BW xylose in 3-d-old calves that were fed either Colostrum or Formula on d 1 and 2 and that were or were not supplemented with quercetin from d 2 onward (50 mg/kg BW daily)
Figure	3.2: Plasma concentrations of total flavonols in 7-d-old calves before and after feeding milk replacer supplemented with quercetin (50 mg/kg BW). Calves received colostrum or formula on d 1 and 2 of life and milk replacer from d 3 onward
Figure	: 3.3: Plasma concentrations of glucose (A), lactate (B), insulin (C), adrenaline (D), and noradrenaline (E) before and after feeding in 7-d-old calves that were fed either colostrum or formula on d 1 and 2 and milk replacer from d 3 onward. In each dietary group, half of the calves received quercetin (50 mg/kg BW daily; CQ+, FQ+; n =6) and half did not receive quercetin (CQ-, FQ-; n =7)

Figure 3.4: Enrichment of [13C ₆]-glucose in plasma (A) and 13CO ₂ in the blood (B) of 7-d-old calves that were fed either C or F on d 1 and 2 and milk replacer from d 3 onward. In each dietary group, half of the calves received quercetin (50 mg/kg daily; CQ+, FQ+ n=6) and half did not (CQ-, FQ-; n=7)
Supplemental Figure 3.1: Concentrations of total flavonols in plasma of individual calves or d 7 before and after tube-feeding milk replacer containing the daily quercetin dose of 50 mg/kg BW. From d 2–6, the daily quercetin dose was divided into equal portions and fed twice daily with colostrum (CQ+, <i>n</i> =5, and calf 19) or formula (FQ+, <i>n</i> =6) respectively (d 1 and 2), or milk replacer (d 3–6)
Figure 4.1: Feed Intake and Health Parameters. Neonatal calves were fed either colostrum (black circles) or formula (gray triangles) on days 1 and 2 and were supplemented with (filled symbols, solid lines) or without (open symbols, dashed lines) querceting aglycone from day 2 to day 8 (50 mg/(kg body weight × day)). (A) Milk intake, (B) percent milk intake by tube feeding, (C) heart rate, (D) respiratory rate, (E) body temperature and (F) fecal score (according to Larson et al. [25]; 1 = normal, 2 = soft 3 = runny, 4 = watery) were observed daily
Figure 4.2: Flavonol Concentrations in Plasma. Stacked concentrations (A) and percentage composition (B) of flavonols in basal plasma samples of calves fed either colostrum (dark grey) or formula (light grey) on days 1 and 2 and supplemented with quercetin aglycone from day 2 to day 8 (50 mg/(kg body weight × day)). Flavonol metabolites quercetin (without fill pattern), isorhamnetin (horizontal lines), tamarixetin (crossline pattern) and kaempferol (horizontal bold lines)
Figure 4.3: Total Protein, Albumin and Immune and Inflammatory Status in Blood Plasma Basal concentrations of total protein (A) and albumin (B), immunoglobulins G1, G2 and M (C, D, E) and acute-phase proteins haptoglobin, serum amyloid A and fibrinogen (F, G, H) in the plasma of calves fed either colostrum (black circles) of formula (gray triangles) on days 1 and 2 and supplemented with (filled symbols, solid lines) or without (open symbols, dashed lines) quercetin aglycone from day 2 to day 8 (50 mg/(kg body weight × day))
Figure 4.4: Antioxidative Status in Plasma. Plasma parameters for antioxidative capacity (AB) and oxidative stress (C, D) in calves fed either colostrum (dark grey) or formula (light grey) on days 1 and 2 and supplemented with (crossline pattern) or without (without fill pattern) quercetin on days 2-8 (50 mg/(kg body weight × day)). TEAC

trolox equivalent antioxidative capacity (in trolox equivalents, TE); FRAP, ferric reducing ability of plasma (in ascorbic acid equivalents, ASCE); TBARS, thiobarbituric acid reactive species (in malondialdehyde equivalents, MDAE).......77

ABBREVIATIONS

8-iso-PGF_{2 α} 8-epimer of Prostaglandin F_{2 α}

ANOVA Analysis of variance

ASCE ascorbic acid equivalents

AUC area under the curve

BW body weight C/Col colostrum

CAT catalase (gene)

cDNA complementary DNA

CRP C-reactive protein (gene)

CV coefficient of variation

DM dry matter

ELISA enzyme-linked immunosorbent assay

Emax maximal enrichment

F/For formula

FBPase frucose-1,6-bisphosphatase (protein)

FGA fibrinogen alpha chain (gene)

FPU first pass uptake

FRAP ferric reducing ability of plasma
G6Pase glucose-6-phosphatase (protein)
G6PC glucose-6-phosphatase (gene)

GCK/GCK glucokinase

GIP glucose-dependent insulinotropic polypeptide

GIT gastrointestinal tract

GLUT faciliated glucose transporter

GNG gluconeogenesis

GPX1 glutathione peroxidase 1 (gene)

H₂O₂ hydrogen peroxide HP haptoglobin (gene)

HPCAL1 hippocalcin-like 1 (gene)

HPLC high-performance liquid chromatography

i.m. intramusculari.v. intravenouslg immunoglobulin

IGF insulin-like growth factor IL1A interleukin-1α (gene) *IL1B* interleukin-1β (gene)

LRP10 low-density lipoprotein receptor-related protein 10 (gene)

LSM least squares mean

MDAE malondialdehyde equivalents

NEFA non-esterified fatty acids

PC pyruvate carboxylase (protein)

PCK1 phosphoenolpyruvate carboxykinase, cytosolic (gene)

PCK2 phosphoenolpyruvate carboxykinase, mitochondrial (gene)

PCR polymerase chain reaction

PEPCK phosphoenolpyruvate carboxykinase (protein)

PGF prostaglandin F

POLR2A RNA polymerase II (gene)

PYG glycogen phosphorylase (protein)

PYGL glycogen phosphorylase (gene)

Q- not supplemented with quercetin

Q+ supplemented with quercetin

Ra rate of appearance RIA radioimmunoassay

ROS reactive oxygen species

s.c. subcutaneous

SAA serum-amyloid A (protein)

SAA2 serum-amyloid A2 (gene)

SE(M) standard error (of the mean)

SGLT sodium-dependent glucose transporter

SOD superoxide dismutase (gene)

TBARS thiobarbituric acid reactive species

TE trolox equivalents

TEAC trolox equivalent antioxidative capacity

Tmax time at maximal enrichment

TNF tumor necrosis factor (gene)

TNFα tumor necrosis factor α (protein)

1. GENERAL INTRODUCTION

Calf losses are still a major problem in the dairy industry, seeing that the overall preweaning calf mortality on U.S. dairy operations is approximately 15% [1]. Newborn calves are particularly prone to sickness, with gastrointestinal imbalances, especially neonatal diarrhea, being responsible for more than 50% of the mortality rate in neonates [2]. The reasons for this are manifold, ranging from infectious to noninfectious causes. After birth, the calf loses the relative protection by the uterus, which entails three fundamental consequences: first, its energy supply shifts from a continuous flow of mainly glucose and amino acids to a discontinuous provision with fat and lactose, the latter being insufficient to meet the neonate's glucose demands [3]. Thus, endogenous glucose production needs to be established fast. Second, the newborn is for the first time exposed to a germ-rich environment, and the acquisition of passive immunity is essential to support the calf's naïve immune system during the first weeks of life [4]. Third, the process of birth itself causes the excessive production of reactive oxygen species, which promote the susceptibility to disease if they are not properly counterbalanced by antioxidative defenses [5]. With the ingestion of first colostrum, the calf is not only provided with macronutrients but also with bioactive factors, e.g. hormones and growth factors supporting the maturation of its gastrointestinal tract and the adaptation of its glucose metabolism [6]. Furthermore, colostrum contains antioxidants and immunoglobulins to support neonatal health; unfortunately, the colostrum supply is often insufficient [7, 8]. The prophylactic application of antibiotic performance enhancers was banned by the European Union in 2006; since then, research has been intensified to identify natural alternatives for enhancing the productivity in farm animal rearing. Special focus has been directed on flavonoids for their attributed health-promoting properties [9]. Flavonoids are polyphenolic plant metabolites with strong antioxidative potential and are involved in the actions of plant enzymes, growth hormones and antimicrobial defense [10]. Because of their antioxidative, antiinflammatory, immunomodulatory and antimicrobial activities, flavonoids have long been used by traditional medicine [11]. In humans, the incidence of some chronic diseases, e.g. obesity, mental and neurological disorders, seems to be lower with increasing dietary flavonoid intake [12, 13]. Furthermore, flavonoids are known to interact with glucose metabolism and to normalize the blood glucose level in diabetic individuals, e.g. by suppressing hepatic glucose release and reducing intestinal carbohydrate digestion, thus they ameliorate metabolic conditions in case of diabetes type II [14]. In farm animals, the usage of flavonoid-rich extracts decreases the number of deaths in calves [15] and exerts immunostimulatory effects in sows and their offspring [16]. Other feed additives with antioxidative potential, e.g. tocopherol, have been shown to improve the performance of farm animals [9]. However, to date, there are no studies on the impact of quercetin on the glucose metabolism and immune

system of newborn farm animals.

The aim of the present work was to characterize the health promoting properties of an oral quercetin supplementation in neonatal calves with regard to its impact on the neonate's glucose metabolism, because any disadvantageous effects of quercetin supplementation on the same would forbid the usage of flavonoids in the upbringing of newborn calves. Another aspect was to determine whether quercetin could be useful in stabilizing the calves' health status by compensating for an insufficient initial colostrum supply.

In the present thesis, chapter 2 introduces major aspects of the glucose metabolism in neonatal calves as well as their immune and antioxidative status, and outlines the importance of colostrum feeding for the maturation of aforesaid aspects. Furthermore, an overview is given on the current state of knowledge concerning flavonoids and their impact on carbohydrate metabolism as well as their health promoting properties. Chapter 3 deals with the effects of an oral quercetin supplementation on the glucose metabolism of neonatal calves according to the initial colostrum supply. Chapter 4 focuses on selected metabolic, antioxidative and inflammatory parameters in newborn calves and how these are affected by initial colostrum deprivation and a seven-day quercetin supplementation. In a final general discussion, the main findings of chapters 3 and 4 are critically reviewed and put into context to the present literature.

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2. LITERATURE OVERVIEW

2.1 GENERAL ASPECTS ON NUTRITIONAL AND IMMUNE STATUS IN NEONATAL CALVES

2.1.1 Postnatal adaptation of glucose metabolism in calves

With birth, energy supply of the calf switches from continuous transplacental provision of mainly carbohydrates (glucose) and amino acids to discontinuous oral intake of colostrum and milk [1, 2]. Colostrum is rich in fat but relatively low in carbohydrates when compared to the fetal diet, thus in most mammalian species, the glucose intake with food is insufficient to meet the glucose requirements of the newborn. Figure 2.1 illustrates the altering percentage composition of the bovine perinatal diet.

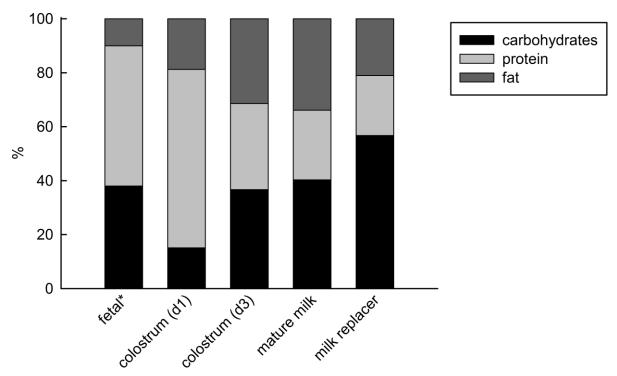


Figure 2.1: Percentage composition of macronutrient intake in calves (minor nutrients are disregarded) [3-6]. Composition of colostrum and mature milk was obtained from pluriparous Holstein cows. *The fat fraction in the fetal diet refers to the short-chain fatty acid acetate.

In neonatal ruminants, milk lactose accounts for only 25% of total glucose demand [7]. To ensure glucose homeostasis, it is crucial for newborn calves to produce glucose endogenously, that is by glycogenolysis and by gluconeogenesis. Towards term, the rise in the

fetal glucocorticoid concentration causes the functional and morphological maturation of several tissues and promotes hepatic glycogen deposition as well as the induction of gluconeogenic enzymes [8-10].

Immediately after birth, glycogenolysis, i.e. the degradation of (hepatic) glycogen stores, is the first action to overcome postnatal starvation and to maintain euglycemia for at least one day of food deprivation [10, 11]. However, within the first days of life, these glycogen stores are depleted [12-14], thus calves have to establish gluconeogenesis (GNG), i.e. the *de-novo* synthesis of glucose from different precursors, as the main glucose-producing metabolic pathway to further keep glucose homeostasis in circulation [15].

Both glucose-producing pathways are controlled by rate-limiting enzymes such as glycogen phosphorylase (PYG), pyruvate carboxylase (PC), phosphoenolpyruvate carboxykinase (PEPCK), and glucose-6-phosphatase (G6Pase), whose activities are regulated by different hormonal and metabolic triggers. Figure 2.2 gives an overview of the hepatic glucose production including key enzymes and different precursors.

In neonates, the gluconeogenic precursors are mainly lactate, amino acids (alanine, glycine, glutamine) and, to a small extent, glycerol [2, 7]. Lactate and glucogenic amino acids are converted to pyruvate, the first substrate of the gluconeogenic pathway. Inside mitochondria, PC catalyzes the reaction of pyruvate to oxaloacetate, which is then converted to phosphoenolpyruvate by PEPCK [7], both being rate-limiting steps of GNG. Depending on whether lactate or other substrates serve as gluconeogenic precursor, the latter conversion occurs either inside mitochondria or in the cytoplasm. Phosphoenolpyruvate is then transformed by five different enzymes to fructose-1,6-bisphosphate. Another rate-limiting step is the conversion of fructose-1,6-bisphosphate to fructose-6-phosphate by fructose-1,6-bisphosphatase (FBPase). The isomerization of fructose-6-phosphate results in glucose-6-phosphate. Glucose-6-phosphate is also generated during glycogenolysis from glucose-1-phosphate, which is sequentially removed from glycogen by PYG. Finally, G6Pase controls the rate of endogenous glucose production as it catalyzes the final step to free glucose, which can then be released into the circulation.

According to Steinhoff-Wagner et al. [15], the fraction of gluconeogenesis on total endogenous glucose production increases during the first week of life, while Haga et al. [12] showed that the activities of PEPCK and PC are lower in weaned than in 1-week-old calves. Furthermore, gluconeogenesis from lactate is eightfold greater in hepatocytes from 1-week-old than from ruminating calves [16]. In neonatal lambs, lactate accounts for 17-31% of total glucose synthesis [17], whereas propionate becomes the preferred gluconeogenic substrate with development of a functional rumen [7].

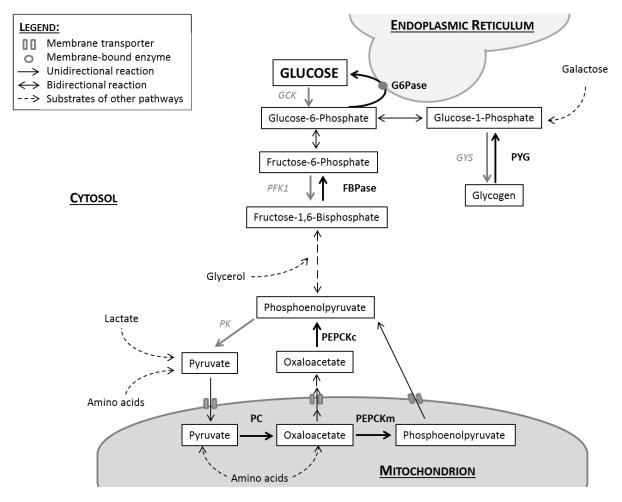


Figure 2.2: Simplified scheme of the hepatic glucose metabolism. Rate-limiting enzymes of endogenous glucose production (gluconeogenesis, glycogenolysis) are bold black, enzymes of glucose consuming pathways (glycolysis, glycogen synthesis) are italic grey. Abbreviations: FBPase, fructose-1,6-bisphosphatase (EC 3.1.3.11); G6Pase, glucose 6-phosphatase (EC 3.1.3.9); GCK, glucokinase (EC 2.7.1.2); GYS, glycogen synthase (EC 2.4.1.11); PC, pyruvate carboxylase (EC 6.4.1.1); PEPCK, phosphoenolpyruvate carboxykinase, (c, soluble; m, mitochondrial; EC 4.1.1.32); PFK, phosphofructokinase (EC 2.7.1.11); PK, pyruvate kinase (EC 2.7.1.40); PYG, glycogen phosphorylase (EC 2.4.1.1).

2.1.2 Maturation of the digestive system

Though calves are born relatively mature, their gastrointestinal tract (GIT) is not fully developed at birth. Therefore, the timing and rate of gastrointestinal maturation are considered as intermediate, meaning that major developmental changes occur both pre- and postnatally [18, 19]. Neonatal calves are regarded as pseudomonogastrics because they are preruminants without considerable forestomach activity [20] making them more dependent on abomasal and intestinal digestive processes than adults to cover their energy needs.

To ensure energy provision *ex utero*, maturation of the GIT is, next to the establishment of hepatic glucose metabolism, a key element of adaptation to postnatal life. Indeed, the most severe changes in the gastrointestinal development of ruminants happen during the first days

of life: Gut regulatory peptides (cholecystokinin, gastrin) rise radically in concentration and promote the development of digestive organs [20, 21]. Functional maturation of the digestive system is evident in elevated activities of brush border membrane enzymes [22] as well as enhanced synthesis and secretion of pancreatic juice within the first week of life [23]. In neonatal calves and lambs, pancreas mass and weight of abomasal mucosa increase by 20-30% within the first week [21]. Morphologic changes of the intestine include accelerated growth rate as well as increase of crypt depth and villus height in calves and piglets, respectively, all contributing to the enlargement of absorptive surface area [22, 24, 25]. Vacuolated fetal-type enterocytes that are able to transport macromolecules disappear resulting in intestinal barrier closure [26, 27].

Microbial colonization of the GIT begins with birth and physiological flora needs to establish during the first week of life by ingestion of first food [28]. Finally, food intake triggers the production of endogenous substances that further enhance maturational processes inside and outside the GIT [18, 24, 29].

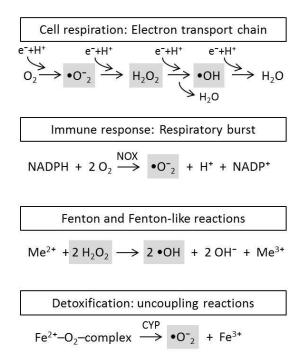
2.1.3 Immune system and antioxidative status

Neonatal calves are prone to diseases because they are considered immunologically naïve at birth [30, 31]. On the one hand, the bovine placenta epitheliochorialis prohibits intrauterine transfer of immunoglobulins hence the calf is born hypogammaglobulinemic [32]. Although endogenous production of immunoglobulins could be demonstrated in colostrum-deprived calves during the first days of life, IgA and IgG do not reach appreciable levels until 16-32 days of life [33, 34]. On the other hand, the immune system is immature at birth and maturation continues until the 6th month of age [31]. Complement activity within the first month of life reaches only 50% of the activity in adults [35] and functional capacity of phagocytes (neutrophils, macrophages) as well as numbers of circulating B- and T-cells are reduced at birth [30, 36]. Due to high concentrations of corticosteroids and cytokines (IL-4, IL-10), the cell-mediated immune response around birth is suppressed [37]. Although present in neonates, many immune components are not (fully) functional until 2-4 weeks of age [31] so the health status of calves mainly depends on the acquisition of passive immunity with colostrum.

Irrespective of age, immune cells are particularly susceptible to oxidative damage because of their phospholipid-rich membrane structure [38-40]. After birth, neonates are for the first time exposed to an oxygen-rich environment. The initiation of aerobic metabolism and reperfusion after hypoxic ischemia during birth imply an increased generation of reactive oxygen species (ROS) [41, 42]. ROS are highly reactive molecules that originate from several physiologic/endogenous processes and are liberated either as mediators or as byproducts [43]. Those processes include immune response, inflammatory reactions, oxidative metabolism or detoxification as well as ischemic processes [44]. During immune response, immune cells,

e.g. neutrophils or macrophages, produce superoxide during respiratory burst to degrade internalized pathogens [45]. However, ROS can also arise from exogenous sources, e.g. radiation, toxins, chemicals or drugs [42, 46, 47]. In a balanced system, ROS are inactivated rapidly by either antioxidative enzymes or antioxidants, which, already at low concentrations, significantly delay or inhibit the oxidation of susceptible substrates and therefore prevent free radical injury. Figure 2.3 opposes the endogenous production of ROS and their inactivation by antioxidants. An imbalance between production of and protection against ROS results in oxidative stress, thus increased ROS are able to react with biomolecules including DNA, RNA, proteins, carbohydrates and lipids, impairing their physiological functions [43, 47].

ENDOGENOUS PRODUCTION



NEUTRALIZATION

Enzymatic inactivation

$$2 \text{ H}^+ + 2 \bullet O^-_2 \xrightarrow{SOD} 2 \text{ H}_2O_2 + O_2$$
 $2 \text{ H}_2O_2 \xrightarrow{CAT} 2 \text{ H}_2O + O_2$
 $GSH + H_2O_2 \xrightarrow{GPx} GSSG + 2 \text{ H}_2O$

Non-enzymatic inactivation

Scavenging Aox + ROS \longrightarrow Stable radicals e.g. albumin, haptoglobin, tocopherols

Quenching Aox + $\bullet O^-_2 \longrightarrow Oxidized$ quencher e.g. β -carotene, ascorbic acid, glutathione

Chelating Aox + $Me^{2+} \longrightarrow Chelate$ complex e.g. transferrin, phospholipids, dietary polyphenols

Figure 2.3: Endogenous formation and neutralization of ROS [44, 48, 49]. Abbreviations: Aox, antioxidant; CAT, catalase (EC 1.11.1.6); CYP, cytochrome P450 (EC 1.14); GPx, glutathione peroxidase (EC 1.11.1.9); GSH, glutathione; GSSG, glutathione disulfide; Me, transition metal ion (e.g. Fe²⁺, Cu²⁺, Zn²⁺, Co²⁺, Mn²⁺); NADP+/NADPH, nicotinamide adenine dinucleotide phosphate; NOX, NADPH oxidase (EC 1.6.3.1); SOD, superoxide dismutase (EC 1.15.1.1).

ROS are involved in the pathogenesis of inflammatory and other diseases [46, 50, 51]. Furthermore, oxidative stress is also considered as a catalyzer for neonatal disease [52, 53]. Gastrointestinal anatomical and functional alterations, reduced intestinal absorption as well as failure of nutrient and immunoglobulin transfer can be consequences of ROS-induced damages in neonates of different species [52]. Conversely, growing calves, puppies and piglets with enteric disease have been shown to have significantly elevated markers for oxidative stress and decreased antioxidant capacity in plasma [54-56]. Supplementing calves and lambs with antioxidants, e.g. vitamin E, reduces morbidity and mortality after exposure to

stress, enhances the immune response and protects against tissue damage [57, 58]. In comparison to adult cattle, concentrations of free radicals in the blood of newborn calves are high while concentrations of antioxidative vitamins are low, leading to an impaired antioxidative status [41, 59-61]. At the same time, the activities of antioxidative enzymes increase. However, data on the antioxidative status are inconsistent and strongly dependent on the regarded parameter and the method of determination [62], thus, comparison of different studies is intricate [63].

2.2 IMPORTANCE OF COLOSTRUM

2.2.1 General aspects

According to the European Commission, colostrum is defined as "the fluid secreted by the mammary glands of milk-producing animals up to three to five days post parturition that is rich in antibodies and minerals, [...]" [European Commission Regulation (EC) No. 1662/2006, Annex II, point 3]. A compositional analysis of 55 Holstein dairy herds revealed macronutrient contents of $6.7 \pm 4.2\%$ fat, $14.9 \pm 3.3\%$ protein and $2.5 \pm 0.7\%$ lactose in colostrum 4 h after calving [64]. While the macronutrients' major purpose is to provide the initial substrate supply for the neonate's energy metabolism, colostrum also contains large amounts of non-nutrient bioactive factors such as immunoglobulins, hormones, enzymes, growth factors and antioxidants [65-67].

The main site of action for milk-borne nonnutritive compounds is the GIT whose growth and development in farm animals is dramatic during the neonatal period [68, 69]. However, some bioactive factors also exert systemic effects: for instance, the absorption of essential fatty acids and vitamins (β-carotene, vitamins A and E) affects the status of plasma of respective compounds [24]. Although some proteins and peptides, e.g. immunoglobulins and lactoferrin, are also intestinally absorbed and appear in the circulation, this does not apply for insulin and IGF-I. However, colostral insulin and IGF-I indirectly modulate the somatotropic axis by enhancing the intestinal nutrient uptake and stimulating endogenous IGF-I production [24, 70]. Thus, the indirect anabolic effect of human IGF-I added to milk replacer becomes apparent by an enhanced lipid status in calves with increased concentrations of essential fatty acids and fat-soluble vitamins in their plasma [65].

Delayed intake of first colostrum alters plasma concentrations of several metabolites and hormones, but is rapidly compensated and does not indicate permanent imprinting effects on hematological, metabolic or endocrine traits [71, 72]. However, absorption of IgG is reduced thus performance is indirectly impaired [11, 73, 74]. Up to now, there is no clear evidence

whether colostral bioactive factors may program the animal and its later productivity [75].

2.2.2 Effects on GIT and glucose metabolism

The development and maturation of the GIT are essential for an efficient uptake of ingested nutrients, which in turn is necessary for proper growth and development of the young.

On the one hand, colostrum modulates the intestinal microbial population as it contains not only antimicrobial substances (e.g. lactoferrin, lactoperoxidase, lysozyme, oligosaccharides) to prevent growth of pathogens, but also pre- and probiotics to establish the physiological intestinal microflora [24, 76, 77].

On the other hand, colostrum is proven to modulate the GIT morphology [10, 69]. Feeding colostrum or colostrum-extracts resulted in greater villus circumferences, areas and heights [29, 78, 79] and increased crypt cell proliferation rates in the small intestine [78] when compared to colostrum-free alimentation. According to this, the intestinal absorption of monosaccharides depends on initial colostrum supply and is reduced when calves are colostrum-deprived (Figure 2.4) [5, 29, 80, 81]. However, the gene expression and protein concentrations of glucose transporters GLUT2 and sodium-dependent glucose transporter-1 (SGLT1) seem not to be affected by colostrum deprivation [25, 79].

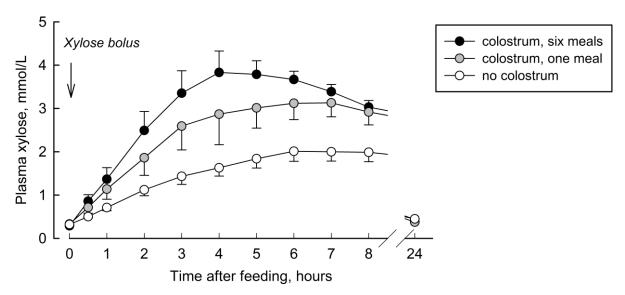


Figure 2.4: Plasma xylose concentrations following intestinal absorption in 5d old calves after different initial colostrum supply, modified from Hammon and Blum [81].

Moreover, the digestive capacity of the GIT is also affected by colostrum intake. Plasma concentrations of gut regulatory peptides, e.g. gastrin, cholecystokinin, and secretin increase after colostrum feeding thus exert favorable effects on gastrointestinal growth and digestive functions by enhancing gastric and pancreatic secretions [20]. In neonatal calves and piglets, activities of intestinal lactase as well as other brush border membrane enzymes seem to be only marginally affected by the initial feeding of formula versus colostrum [25, 78, 79], but

they are positively correlated with the intestinal villus size [29]. However, prolonged colostrum-feeding enhances activities of pancreatic lipase, elastase II and trypsin [78].

Even under conditions of similar lactose intake, colostrum-deprived calves have lower preand postprandial plasma glucose concentrations [5, 82, 83]. In newborn piglets, capacity for GNG from lactate is reduced when colostrum intake was restricted [84]. In contrast to this, GNG and total endogenous glucose production do not differ between colostrum-fed and colostrum-deprived calves [5]. However, dietary effects on mRNA abundance or activities of rate-limiting enzymes of GNG in newborn calves are inconsistent [5, 82, 85].

2.2.3 Effects on immune system and antioxidative status

Colostrum is rich in immunoglobulins, thus it is essential for neonatal calves to acquire passive immunity. As IgG accounts for 85-90% of total immunoglobulins in colostrum [86], its concentration in newborns' plasma is used to assess the success of passive transfer. According to this, a serum IgG concentration of 10 g/L (equivalent to serum protein concentration of 52 g/L) at the age of 24-48 h is considered as threshold for an adequate passive transfer [30, 87, 88]. Closure of the intestinal barrier coincides with the disappearance of vacuolated enterocytes and marks the cessation of macromolecule absorption [89]. Although delayed colostrum feeding can slow down this process, the efficiency of absorption impairs with increasing age at first colostrum ingestion hence incidence for failure of passive transfer rises [90, 91]. Absorbed colostral antibodies activate and regulate innate immune responses in calves [31] therefore health status and productivity are enhanced [92-94]. In addition to immunoglobulins, maternal leukocytes ingested with colostrum also enhance the neonatal immune system by increasing phagocytosis, stimulating cellular immune response and increasing the development of antigen-presenting cells [31, 95, 96]. Despite of immunity-promoting properties, colostrum is also a source of ROS. Those are produced by viable maternal leukocytes as well as enzymes like xanthine oxidase and lactoperoxidase in colostrum [67]. Opposing to this, there is also a variety of antioxidants present in colostrum. While catalase (CAT), superoxide dismutase (SOD) and glutathione peroxidase (GPx) inactivate ROS through enzymatic reactions, vitamins (A, C, E), minerals (Se, Cu, Zn) as well as lactoferrin, caseins and whey proteins are non-enzymatic colostral compounds exerting antioxidative properties [67, 97]. In porcine and bovine colostrum, activities of antioxidative enzymes and lactoferrin increase with time after parturition [98-100] and an assessment of the antioxidative status revealed that the total antioxidant capacity in colostrum and milk of cows rises during the first week after parturition [99, 101]. Abuelo and collegues [61] could show that the levels of lipoperoxides in colostrum are positively correlated with ROS in the serum of newborn calves fed respective colostrum, while colostral antioxidative capacity and serum ROS were negatively correlated. Figure 2.5 opposes the intensity of protein peroxidation and the total antioxidant capacity of colostrum and newborn

calves' plasma of Holstein-Friesian primiparous cows.

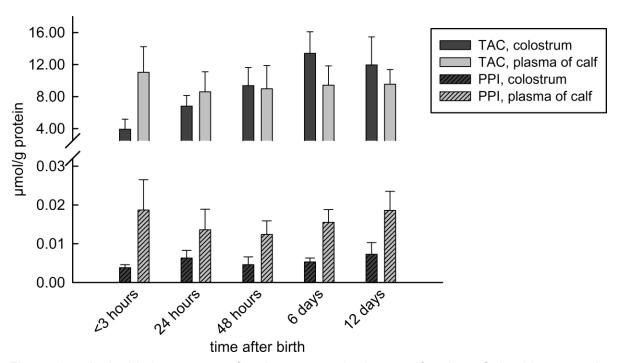


Figure 2.5: Antioxidative status of colostrum and plasma of calves fed with respective colostrum, modified from Albera and Kankofer [99]. Abbreviations: PPI, protein peroxidation intensity; TAC, total antioxidant capacity.

According to Retskii et al. [102], correcting the calf's antioxidant status before the first feeding improves the acquisition of passive immunity. Indeed, serum IgG concentrations have been shown to be positively correlated with the colostral antioxidative capacity [61]. Nonetheless, research on the connection of antioxidative status with IgG acquisition in calves is scarce as studies mostly focus on the effect of single colostral compounds rather than antioxidative capacity in general. Another obstacle is that middle- or long-term effects have been investigated in species whose offspring usually suckle the dam for a longer period of time, therefore the explicit influence of colostral antioxidative capacity on acquisition of passive immunity and health status is ambiguous and results can hardly be transferred to dairy calves.

Finally, the early change from colostrum to artificial milk replacers in calf rearing might attribute to a prolonged imbalance in antioxidant status as milk replacers often contain high levels of pro-oxidative unsaturated fatty acids while their antioxidative capacity is low [103].

2.3 CHARACTERISTICS OF FLAVONOIDS

2.3.1 General aspects

Flavonoids are secondary plant metabolites ubiquitously present in higher plants, where they are involved in photosynthesis, growth, development and pigmentation as well as defense against pathogens and other biotic and abiotic stresses [45, 104, 105]. They are polyphenols with a three-ring structure and are divided into subclasses according to the degree of saturation and oxidation as well as the position of the B-ring (Figure 2.6). Up to now, more than 8,000 different flavonoid compounds have been identified [105]. As animals are not able to synthesize flavonoids, they can only ingest them with vegetarian food components. However, the flavonoid content of green fodder is not fixed, but it depends on many factors, e.g. ripeness, harvesting time, degree of dryness, processing of foodstuff, storage etc. [104, 106]. Flavonols are the most ubiquitous flavonoids with quercetin being the most abundant dietary flavonol [104, 107]. As with other flavonoids except for flavanols, quercetin naturally rarely occurs in its free form (aglycone) but is bound to a sugar moiety by beta-glycosidic linkage, which can be glucose (glucoside) or other mono-, as well as di-, tri- or tetrasaccharides (glycoside) [106].

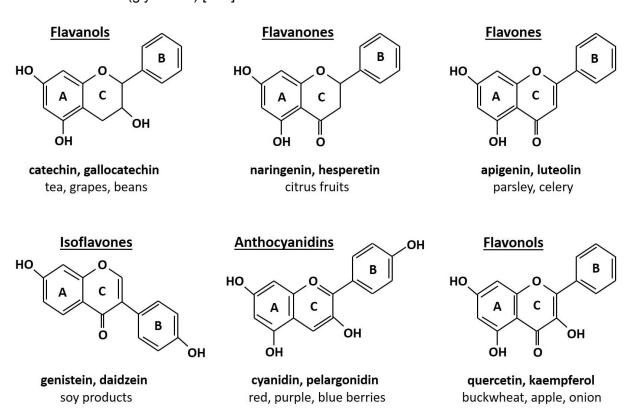


Figure 2.6: Chemical structure of the most common flavonoid subclasses and names of some representative compounds as well as their dietary sources [45, 104, 105].

Flavonoid-rich plants or plant extracts have been used for centuries in traditional medicine [108, 109], taking advantage of their antioxidative, anti-inflammatory, antiproliferative, and antimicrobial properties [110]: In humans, flavonoid-rich diets are related to reduced incidence of diseases associated with oxidative stress, e.g. cancer, cardiovascular, neurodegenerative and inflammatory diseases as well as metabolic disorders [105, 107, 109, 111, 112]. However, the excessive intake of flavonoids may not promote health, but exert adverse effects [49, 113]; thus supplementing flavonoids to the natural diet should be diligently supervised.

2.3.2 Absorption and metabolism of flavonols

Although the oral bioavailability of quercetin is relatively high [107], it depends on many factors: 1) the nature of the compound (hydrophilic glucoside resp. glycoside versus lipophilic aglycone) [114-116], 2) the dietary matrix with which it is ingested [111, 117, 118], 3) availability and activity of intestinal enzymes [107, 119, 120], 4) the gut microflora [107, 121], and 5) the duration of the flavonol administration (single dose vs. long-term intake/application) [111, 122]. For these reasons, interindividual variability of flavonol metabolites in plasma after intestinal absorption is high [123-125].

While quercetin aglycones are passively absorbed from the small intestine by glucose transporter 2 (GLUT2), its glucosides can be taken up actively by sodium-dependent glucose transporter-1 (SGLT1) [126, 127]. More complex glycosides are mainly microbially degraded before absorbed in distal parts of the intestine [104]. If not immediately excreted back into the intestinal lumen by multidrug-resistance-associated proteins [110, 115], absorbed flavonols are metabolized by various organs, e.g. the small intestine, liver and kidneys, where they are mainly glucuronidated, sulphated (intestine, liver, kidney) and methylated (liver, kidney) via phase II metabolism [43, 107, 118]. Thus, after oral administration, there are several glucuronide and sulfate conjugates but almost no aglycones detectable in the circulation of humans [125, 128], rats and pigs [110, 122, 129, 130]. Metabolic detoxification increases the hydrophilicity thus facilitates biliary and urinary excretion [104]. However, the slow elimination from blood, their presence in the bile as well as re-entry peaks in plasma indicate an enterohepatic cycling of flavonols [45, 114, 115, 125].

In adult cattle, intraruminal application of quercetin glycoside (rutin) but not its aglycone increases plasma flavonol concentrations, indicating that glycosylation protects from ruminal degradation [131]. In newborn calves, oral bioavailability changes with increasing age: on d 2 and d 29 of life, feeding of quercetin aglycone results in higher plasma flavonol concentrations than rutin feeding (quercetin-3-O-rutinoside), which is comparable to data obtained in monogastrics or in cows after intraduodenal application [4, 114, 124]. However, the plasma flavonol concentration on d 29 is significantly reduced when compared to d 2 and the proportion of single flavonol metabolites corresponds to patterns found in adult cattle,

indicating maturation of detoxification and elimination processes.

2.3.3 Interaction with glucose metabolism

Quercetin is well-known for its antihyperglycemic properties, which is why it is used as a natural anti-diabetic agent. After oral administration, the first side of action for quercetin and other flavonols is the GIT, where they interact with digestive enzymes and nutrient transporters. Quercetin and kaempferol were shown to inhibit α -glucosidase as well as α -amylase *in vitro* [132, 133] thus reducing polysaccharide digestion. Similar effects on activities of intestinal glucosidases maltase and sucrase were also observed *in vivo* in diabetic quercetinfed rats and mice; however, lactase activity was unaltered [134, 135]. Several flavonols are transported across the intestinal barrier via similar transporters as monosaccharides; hence intestinal transport of the latter is impaired. Quercetin glucosides competitively inhibit intestinal glucose uptake via SGLT1 [114, 129, 136, 137], whereas quercetin aglycone is a non-competitive inhibitor of GLUT2 [126, 137, 138]. GLUT2 is not only located at the basolateral side of enterocytes but it can also be translocated to the apical side if the luminal glucose concentration is high [139]. Furthermore, GLUT2 is also the major glucose transporter in pancreatic β -cells and hepatocytes [137].

Hepatic glucose metabolism is also influenced by flavonols. Kato et al. [140] demonstrated in isolated rat hepatocytes that quercetin inhibits PYG and therefore concluded the inhibition of hepatic glucose production by quercetin. In contrast to this, Gasparin et al. [141] could show in isolated perfused rat liver that quercetin aglycone stimulates glycogenolysis and glucose release in a concentration-dependent manner, which indicates that effects in situ might differ from effects observed in cell culture. Same authors also demonstrated that quercetin in doses above 50 µM inhibited glycolysis from endogenous as well as exogenous glucose [141, 142] by decreasing the activity of glucokinase (GCK), thus glucose could not be phosphorylated for subsequent reaction steps. In diabetic mice, the hepatic expression of *GCK* was unaltered by quercetin-feeding [143]. Yet another enzyme, namely G6Pase, which catalyzes the dephosphorylation of glucose during gluconeogenesis, has been shown to be inhibited by quercetin (50 µM) and its dehydroxylated metabolite kaempferol [141, 144].

Quercetin interacts not only with hepatocytes but also with pancreatic β -cells, hence glucose metabolism might also be indirectly affected by alteration of its hormonal regulation. Anti-inflammatory properties account for the protection against oxidative damage as well as cytokine-induced cell death in the pancreas [145, 146], and it was shown that insulin release increases in diabetic mice and rats treated with quercetin [143, 147]. Insulin is a key regulator of glucose homeostasis as it affects not only hepatic metabolism but is also crucial for glucose uptake in muscle and adipose tissue by GLUT4. However, direct effects of quercetin on insulin-dependent glucose uptake are controversial: while some authors identified quercetin as competitive inhibitor of GLUT4 [148], others postulate that reduction of

insulin-dependent glucose uptake occurs via inhibition of GLUT4 translocation [149]. In contrast to this, Fang et al. [150] postulated that kaempferol and quercetin improve glucose-uptake in the presence of insulin. However, basal, insulin-independent glucose uptake in adipocytes seems to be unaffected by quercetin [148, 150].

In animal models of diabetes mellitus, quercetin treatment was shown to reduce postprandial hyperglycemia and thus ameliorate metabolic conditions [126, 147, 151, 152]. Anyhow, to what extent quercetin might influence overall glucose production *in vivo* depends on its metabolism as well as on biological activity and concentration of flavonols and respective derivatives at the side of action after application of the compound. Until now, the biological relevance of quercetin in healthy, normoglycemic individuals is not clear [153].

2.3.4 Effects on performance and intestinal health

Several flavonoids and flavonoid extracts that have been used by traditional medicine for centuries have attracted special notice during the last decades and thus became subject to intensive scientific research for their health promoting properties.

Diarrhea is a major problem in calves during the first weeks of life and causes substantial economic losses. In 1991, Capasso et al. [154] showed that quercetin inhibited prostaglandin- and leukotriene-induced contractions in isolated guinea pig ileum and concluded that the flavonol might be useful in relieving gastrointestinal colic, diarrhea and other gastrointestinal disorders. The same group also demonstrated that an intraperitoneal injection of quercetin reduced castor oil induced diarrhea in rats and mice [155, 156]. In humans, the spasmolytic effect of a quercetin-rich quava extract reduced abdominal pain, however, the consistency or frequency of diarrhea were not affected [157]. A comparable extract was shown by Chinese scientists to have a good curative effect on infantile rotaviral enteritis, and rotavirus antigen in stool was significantly reduced after oral application of the extract [158]. The antiviral property of quercetin was also evaluated in vitro: although several flavonols showed a favorable anti-rotavirus activity, quercetin was ineffective to inhibit the virus' cytopathic effect [159, 160]. However, quercetin and its 7-rhamnoside are able to reduce the infectivity of several strains of cornonaviruses, which are relevant enteropathogens in neonatal farm animals [161]. Intestinal bacteria have also been investigated for their interference with quercetin: Pakar et al. [162] demonstrated the inhibitory action on growth and cell-adhesion of selected enteropathogenic bacteria in vitro, while at the same time, probiotics were relatively unaffected, concluding that quercetin may help to stabilize the intestinal flora. In laying hens, dietary guercetin improved the performance by beneficially modulating the intestinal microflora populations [163]. In contrast to this, the intraruminal application of quercetin aglycone or its glycorhamnosid rutin did not alter ruminal fermentation processes in dairy cows, thus the ruminal microflora seemed not to be substantially influenced [131].

2.3.5 Effects on antioxidative and anti-inflammatory status

Quercetin possesses strong antioxidant and anti-inflammatory properties. The catechol structure in the B-ring accounts for the metal-chelating properties of quercetin, hence cations are unavailable for the generation of ROS by Fenton-type reactions [118, 164]. As quercetin inhibits myeloperoxidase and NADPH oxidase, the superoxide production during respiratory burst is reduced [45].

The phenolic hydroxyl groups are able to act as electron donors, thus quercetin scavenges ROS [118, 164]: Due to its hydroxyl groups, quercetin also exerts chain-breaking antioxidant action during lipid-peroxidation and is able to inhibit the autoxidation of fatty acids [165, 166]. As mentioned earlier, biological effects always depend on the chemical structure of circulating metabolites: in accordance to this, quercetin was found to act more as an antioxidant than its monoglucosides [165]. Furthermore, Shirai and coworkers [167] could show that although co-incubation with H_2O_2 and quercetin aglycone inhibited H_2O_2 -induced ROS production in mouse fibroblasts, the pre-incubation with quercetin aglycone prevented its antioxidative activity, while quercetin-3-glucuronide retained its effectiveness.

Flavonols do not only exert antioxidative activity by directly scavenging ROS or inhibiting their production but they can also protect the antioxidant ascorbic acid from oxidative degradation [45], conserve tocopherols in biological membranes [168] and reduce α -tocopherol radicals [164]. Further indirect impact on the antioxidative system occurs by the interaction with antioxidative enzymes, e.g. SOD, CAT and GPx [164] but data on potential effects are inconsistent as both stimulating and inhibiting effects have been described *in vivo* and *in vitro* [169-172].

The antioxidative activity of flavonols also affects the immune system: ROS are produced by phagocytes during respiratory burst as part of the innate immune response to fight invading pathogens. Flavonoids are not only able to scavenge ROS, but they can also inhibit their production. In case of immune cells, e.g. macrophages and neutrophils, ROS release is impaired by quercetin as it inhibits relevant enzymes, namely myeloperoxidase and NADPH oxidase [45]. Quercetin was also shown to inhibit the gene expression and enzyme activity of cyclooxygenase-2 and inducible nitric oxide synthetase, which results in a reduced release of pro-inflammatory mediators (nitric oxide, prostaglandin E_2) and thus contributes to the anti-inflammatory properties of quercetin [173]. In isolated macrophages, incubation with quercetin was shown to reduce mRNA levels of proinflammatory cytokines including TNF α and IL-1 β [174, 175]. In non-immune cells, e.g. pancreatic β -cells or adipocytes, quercetin treatment attenuated the TNF α -mediated inflammation by interfering with signal transduction [145, 175, 176]. According to these findings, quercetin has also been shown to ameliorate the metabolic alterations evoking from chronic inflammation in animal models of diabetes mellitus type 2, obesity or atherosclerosis [147, 175, 177].

In rat and mouse models for inflammatory bowel disease, oral treatment with quercetin

aglycone or its O-methylated metabolite isorhamnetin attenuated chemically-induced symptoms of inflammation [178, 179]. Similar intestinal anti-inflammatory effects could be shown in mice fed the quercetin-glucorhamnosid rutin [180]. On a whole body level, however, only the acute administration of quercetin aglycone was useful to prevent endotoxemia and death in mice intraperitoneally injected with lipopolysaccharide, while the anti-inflammatory effect was missing after chronic administration [181]. When administered 12 h after lipopolysaccharide treatment, only quercetin aglycone but not its glucuronide affected the cytokine secretions by peritoneal macrophages [182].

Although numerous experiments have been undertaken *in vitro* or in laboratory rodents to emphasize the anti-inflammatory properties of quercetin, studies in farm animals are scarce and mostly focus on the effects of plant extracts. For example, in piglets of Echinacea-supplemented sows, immunostimulatory effects were evident from increased Ig concentrations when compared to unsupplemented controls [183]. Similar effects could be shown in growing pigs, whose specific immune response to vaccination was higher when animals received Echinacea [184]. Elevated immunoglobulin titers were also observed in buffalo calves fed propolis, a flavonol-rich natural compound produced by honey bees [185]. Furthermore, addition of propolis to an inactivated vaccine against bovine herpesvirus-5 also resulted in significantly higher antibody titers in vaccinated Hereford cattle when compared to animals receiving a propolis-free vaccine [186].

However, as extracts of natural compounds contain mixtures of multiple components, it is not possible to distinguish whether observed immune-stimulatory effects are caused by single substances or by their synergism.

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3. EFFECTS OF ORAL QUERCETIN SUPPLEMENTATION ON SPLANCHNIC GLUCOSE METABOLISM IN 1-WEEK-OLD CALVES DEPEND ON DIET AFTER BIRTH

Jeannine Gruse¹, Solvig Görs¹, Armin Tuchscherer², Winfried Otten³, Joachim M Weitzel⁴, Cornelia C Metges¹, Siegfried Wolffram⁵, and Harald M Hammon^{1*}

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¹ Institute of Nutritional Physiology "Oskar Kellner",

² Institute of Genetics and Biometry,

³ Institute of Behavioural Physiology, and

⁴ Institute of Reproductive Biology, Leibniz Institute for Farm Animal Biology (FBN), Dummerstorf, Germany; and

⁵ Institute of Animal Nutrition and Physiology, Faculty of Agricultural and Nutritional Sciences, Christian-Albrechts-University of Kiel, Kiel, Germany

^{*}Corresponding author: hammon@fbn-dummerstorf.de

4. QUERCETIN FEEDING IN NEWBORN DAIRY CALVES CANNOT COMPENSATE COLOSTRUM DEPRIVATION: STUDY ON METABOLIC, ANTIOXIDATIVE AND INFLAMMATORY TRAITS

Jeannine Gruse¹, Ellen Kanitz², Joachim M. Weitzel³, Armin Tuchscherer⁴, Tadeusz Stefaniak⁵, Paulina Jawor⁵, Siegfried Wolffram⁶ and Harald M. Hammon^{1*}

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¹ Institute of Nutritional Physiology "Oskar Kellner",

² Institute of Behavioural Physiology,

³ Institute of Reproductive Biology, and

⁴ Institute of Genetics and Biometry, Leibniz Institute for Farm Animal Biology (FBN), Dummerstorf, Germany

⁵ Department of Immunology, Pathophysiology and Veterinary Preventive Medicine, Faculty of Veterinary Medicine, Wroclaw University of Environmental and Life Sciences, Wroclaw, Poland

⁶ Institute of Animal Nutrition and Physiology, Faculty of Agricultural and Nutritional Sciences, Christian-Albrechts-University of Kiel, Kiel, Germany

^{*}Corresponding author: hammon@fbn-dummerstorf.de

4.1 ABSTRACT

Immaturity of the neonatal immune system is causative for high morbidity in calves and colostrum intake is crucial for acquiring passive immunity. Pathogenesis is promoted by reactive oxygen species accumulating at birth if counter-regulation is inadequate. The flavonol quercetin exerts antioxidative and anti-inflammatory effects that may enhance neonatal health. The aim of this work was to study effects of quercetin feeding on metabolic, antioxidative and inflammatory parameters in neonatal calves to investigate whether quercetin could compensate for insufficient colostrum supply. Twenty-eight newborn calves were assigned to two dietary groups fed colostrum or milk-based formula on day 1 and 2 and milk replacer thereafter. From day 2 onward, 7 calves per dietary group were additionally fed quercetin aglycone (50 mg/(kg body weight × day)). Blood samples were taken repeatedly to measure plasma concentrations of flavonols, glucose, lactate, total protein, albumin, urea, non-esterified fatty acids, triglycerides, cholesterol, insulin, glucagon, cortisol, immunoglobulins, fibrinogen, haptoglobin and serum amyloid A. Trolox equivalent antioxidative capacity, ferric reducing ability of plasma, thiobarbituric acid reactive species and F2isoprostanes were analyzed to evaluate plasma antioxidative status. Expression of tumor necrosis factor, interleukin-1α, interleukin-1β, serum amyloid A, haptoglobin, fibrinogen, Creactive protein, catalase, glutathione peroxidase and superoxide dismutase mRNA were measured in liver tissue on day 8. Plasma flavonol concentrations were detectable only after quercetin-feeding without differences between colostrum and formula feeding. Plasma glucose, lactate, total protein, immunoglobulins, triglycerides, cholesterol, trolox equivalent antioxidative capacity and thiobarbituric acid reactive species were higher after colostrum feeding. Body temperature, fecal fluidity and plasma concentrations of cortisol and haptoglobin were higher in formula- than in colostrum-fed groups. Hepatic mRNA expression of tumor necrosis factor was higher after quercetin feeding and expression of C-reactive protein was higher after formula feeding. Data confirm that colostrum improves neonatal health and indicate that quercetin feeding cannot compensate for insufficient colostrum supply.

4.2 INTRODUCTION

Calfhood diseases play a key role in the economy of dairy farms because they increase operating costs and reduce long-term productivity of the animal. Incidence of disease is associated with increased mortality rates [1], and enteritis is the most common diagnosis in young calves [2], which, according to Svensson, Linder and Olsson [3], contributes to 23% of calf losses during the first 14 days of life. Neonatal calves are prone to sickness because their immune system is immature. Furthermore, the process of birth itself causes an elevated stress level for the newborn and exposure to an oxygen-rich environment leads to an increased generation of reactive oxygen species [4, 5]. Reactive oxygen species induce peroxidation of lipids and other macromolecules, leading to alteration of cellular components, interaction with signaling cascades and modification of physiological cell functions [6]. If not properly counterbalanced by antioxidative defenses, excessive production of reactive oxygen species results in oxidative stress, which is a cofactor of disease in humans and farm animals [5, 7, 8]. Adequate colostrum supply is vital to calves because colostrum ensures ingestion of nutrients and contains immunoglobulins (Ig), peptides, antioxidants and other bioactive factors supporting maturation, antioxidative and immune defense as well as local intestinal immunity [9].

The ban on antibiotic performance promoters by the European Union in 2006 increased efforts to establish natural alternatives to enhance health and productivity in breeding. Special focus has been directed to phytochemicals because their use can be manifold according to the respective compound [10]. Flavonoids are secondary plant metabolites that are widely distributed in the plant kingdom and are able to modulate inflammation and immune function and exert antioxidative activity [11-13].

Quercetin, which belongs to the subclass of flavonols, is ubiquitous in most plants and is of interest for scientists for its beneficial use in humans and farm animals. Its antioxidative capacity can ameliorate the acquisition of passive immunity in neonates, based on the finding that feeding antioxidant-enriched colostrum enhanced IgG absorption and antioxidative status in newborn calves and piglets [14, 15]. Similarly, Retskii et al. [16] showed that correcting the antioxidative balance in newborn calves prior to first colostrum ingestion increases the acquisition of colostral immunity and reduces the incidence of enteric colibacillosis. Another beneficial effect of quercetin is its local action in the gastrointestinal tract. *In vitro* studies on intestinal epithelium demonstrated that quercetin down-regulates the expression of genes related to inflammation in inflamed epithelium [17], and Lozoya et al. [18] showed in a clinical study that oral quercetin administration reduced abdominal pain in acute diarrheic disease in humans. In guinea pigs, mice and rats, the inhibitory action of quercetin on prostaglandin E₂-induced ileal contractions and on castor-oil-induced diarrhea has been demonstrated [19, 20]. Furthermore, quercetin acts as a prebiotic, thus inhibiting

adhesion of enteropathogens to Caco-2 cells without affecting the viability of probiotics [21], and improves performance in hens by modulating cecal microflora populations [22].

Although a multitude of research on quercetin has been performed *in vitro* or in animal models for medical conditions, studies of the effects in neonatal farm animals are scarce. The aim of the present work was to investigate the potential health-promoting effects of feeding quercetin to newborn calves during the first week of life and to evaluate whether the health-promoting effects of quercetin compensate for initial colostrum deprivation in calves. We hypothesized that quercetin improves antioxidative balance and immune function and that local antibacterial and anti-inflammatory effects reduce the incidence of diarrhea and gastrointestinal dysfunctions.

4.3 MATERIALS AND METHODS

4.3.1 Animals, Husbandry and Feeding

Experimental procedures were conducted in compliance with the German Animal Protection regulations with approval of the authorities of the state Mecklenburg-Western Pomerania, Germany (Landesamt für Landwirtschaft, Lebensmittelsicherheit und Fischerei Mecklenburg-Vorpommern; LALLF M-V/TSD/7221.3-1.1-044/12). Liver biopsies were performed under local lidocaine anesthesia and all calves received metamizole post-operatively for pain relief. Twenty-eight male Holstein Friesian calves were separated from their dams immediately after birth and were housed in single boxes during their first 8 days of life. Before the trial started, separate colostrum pools were prepared from the first and third milkings after parturition. According to the colostrums' macronutrient compositions, milk-based formulas with comparable amounts of macronutrients [23] but without bioactive factors were provided (Bergin MAT-Formula; Bergophor Futtermittelfabrik, Kulmbach, Germany). Calves were randomly assigned to two dietary groups and were bucket-fed twice daily, receiving either colostrum (Col, n = 14) or corresponding formula (For, n = 14) on days 1 and 2 of life (10%) and 12% of body weight/day, respectively). If appetite was reduced, calves were tube fed to ensure complete ingestion of colostrum or formula. From day 3 until day 8, all calves were fed commercial milk replacer (12% of body weight/day; 150 g/L; SALVAlac MIRApro 45; Salvana Tiernahrung, Klein-Offenseth Sparrieshoop, Germany).

On day 2, the dietary groups were subdivided into control (ColQ- and ForQ-; n = 7 per group) and treatment groups (ColQ+ and ForQ+; n = 7 per group), the latter receiving quercetin aglycone twice daily with feeding (50 mg/(kg body weight × day); quercetin aglycone dihydrate \geq 98%, Carl Roth, Karlsruhe, Germany). The control groups received no quercetin aglycone.

4.3.2 Treatment

Navels were disinfected with 10% povidone iodine solution (Vet-Sept; aniMedica, Senden-Bösensell, Germany). Neonatal calves received an oral dose of 1 g iron dextran with their first meal on day 1 (Ursoferran; Serumwerk Bernburg, Bernburg, Germany). To support immunological defense during the first 5 days of life, all calves received chicken egg-derived immunoglobulins with the morning feeding (0.25 g/kg body weight; Globigen Life Start 25%, EW Nutrition, Visbek, Germany) [24]. To prevent cryptosporidiosis, calves were treated with halofuginone (0.1 mg/kg body weight per *os*; Halocur, Intervet, Igoville, France) after the evening feeding from day 1 to day 7.

Colostrum-deprived calves (ForQ+, ForQ-) additionally received B-vitamins (100 mg nicotinamide/calf, 40 mg thiamin chloride hydrochloride/calf, s.c.; Vitamin-B-Komplex, Serumwerk Bernburg, Germany) and bovine colostral immunoglobulins on days 1 (s.c.), 3 and 5 (per os) (2 g gammaglobulins/calf with antibodies against *Escherichia coli*, rotavirus, coronavirus; Aniserin orinject; aniMedica, Seden-Börsensell, Germany). Furthermore, formula-fed calves were treated metaphylactically with colistin sulfate from day 2 to day 8 (3 mg/kg body weight, i.m.; Belacol; BelaPharm, Vechta, Germany).

All calves were weighed immediately after birth and before evening meals on days 2 and 6. Every morning, health status was examined and appetite, general condition, heart rate, respiratory rate, rectal temperature and gut motility were assessed. Fecal fluidity was scored according to Larson et al. [25]. Calves with reduced vitality after the first 2 days of life were allowed one recovery day before further sample taking. In these cases, the times referred to as days 3, 4, 7 and 8 in the results section are days 4, 5, 8 and 9 after birth, respectively. Due to gastrointestinal imbalances, two calves (one calf of group ColQ+ and one calf of group ForQ+) had to be excluded from the study.

4.3.3 Blood Analyses

4.3.3.1 Sample Taking

Basal blood samples were taken before the morning feeding on days 1, 2, 4 and 7 from the jugular vein using evacuated tubes containing either potassium-EDTA (1.2-2 mg/mL EDTA) for analyses of plasma metabolites, insulin, glucagon, immunoglobulins and acute-phase proteins or Li-heparin (12-30 IU heparin) for the determination of the cortisol and flavonol concentrations and the antioxidative status in the plasma. For flavonol analysis, additional blood samples were taken before the morning feeding on days 3 and 8. After centrifugation $(1,500 \times g, 4^{\circ}\text{C}, 20 \text{ min})$, plasma aliquots were stored at -20°C until analyses (-80°C for analyses of flavonol concentrations and antioxidative status, respectively).

4.3.3.2 Plasma Flavonols, Metabolites and Hormones

Plasma concentrations of flavonols (quercetin, isorhamnetin, tamarixetin and kaempferol) were measured via HPLC as previously described [26] with a detection limit of 2 nmol/L and a recovery rate of $92 \pm 2\%$. The intra- and inter-assay coefficients were 0.5 and 7.2%, respectively.

Plasma metabolites were analyzed using an automatic spectrophotometer (ABX Pentra 400; Horiba ABX, Montpellier, France) and respective kits: glucose (#A11A01667), lactate (#A11A01721), albumin (#A11A01664) and triacylglycerides (#A11A01640) from HORIBA ABX, Montpellier, France; total protein (#553-412) and cholesterol (#553-127) from mti-diagnostics, Idstein, Germany; urea (#LT-UR 0010) from Labor+Technik, E. Lehmann, Berlin, Germany; and non-esterified fatty acids (#434-91795, #436-91995) from WAKO Chemicals, Neuss, Germany.

Plasma concentrations of insulin (#RIA-1257) and glucagon (#RIA-1258) were determined by RIA using kits from DRG Instruments, Marburg, Germany, which were adapted to bovines [27]. Intra- and inter-assay coefficients of variation were 3.7% and 5.5% for insulin, and 3.4% and 22.5% for glucagon, respectively. Plasma cortisol concentrations were analyzed in duplicate after extraction with diethylether using a commercially available ELISA kit (#EIA1887; DRG Instruments GmbH, Marburg, Germany) according to the instructions of the manufacturer. Cross reactivities of the antibody to corticosterone and progesterone were 45% and 9%, respectively, and <2% to any further competing plasma steroids. The assay was validated for use with bovine plasma. The test sensitivity was 3.4 μ g/L, and intra- and inter-assay coefficients of variation were 5.3% and 12.1%.

4.3.3.3 Immunoglobulins (Ig) and Acute-Phase Proteins

The concentrations of IgG1, IgG2 and IgM, as well as acute-phase proteins (haptoglobin, serum amyloid A and fibrinogen), were measured in the EDTA plasma samples taken on days 1, 2, 4 and 7. IgG1 was analyzed by radial immunodiffusion [28] (modified by Gasowska and Stefaniak [29]) using bovine reference serum (RS10-103; Bethyl Laboratories Inc., Montgomery, USA) as standard. IgG2 (#E10-117) and IgM (#E10-101) were determined by ELISA using kits from Bethyl Laboratories Inc., Montgomery, USA. Intra-assay coefficients of variation were 10.9% and 4.0% for IgG2 and IgM, respectively. The detection limit of IgG2 was 7.8 μ g/L and that of IgM was 15.6 μ g/L. For detection of serum amyloid A (SAA; #TP-802), we used a multispecies ELISA kit from Tridelta Development, Maynooth, Ireland. The detection limit of SAA was 9.4 mg/L. The intra-assay coefficient of variation was 12.0%. The haptoglobin concentration was analyzed using the guaiacol method developed by Jones and Mould [30] with human haptoglobin Hp 2-2 (Sigma #H9762) as a standard. The detection limit of haptoglobin was 0.01 g/L. Plasma fibrinogen was determined by rapid heat

precipitation according to Millar, Simpson and Stalker [31].

4.3.3.4 Antioxidative Status

Li-heparinized plasma samples taken on days 1, 4 and 7 were used to analyze the Trolox-Equivalent Antioxidative Capacity (TEAC, as trolox equivalents (TE) in mmol/L) and Ferric Reducing Ability of Plasma (FRAP, as Ascorbic Acid Equivalents, (ASCE) in μ mol/L) as parameters of antioxidative capacity as well as Thio-Barbituric Acid Reactive Species (TBARS, as Malondialdehyde Equivalents (MDAE) in μ mol/L) and 8-iso-PGF_{2 α} (F2-isoprostanes, in ng/L) as markers for oxidative stress.

TEAC was analyzed as described by Miller et al. [32] and modified by Re et al. [33]. FRAP and TBARS were determined according to Luehring et al. [34]. For determination of F2-isoprostanes, a commercial ELISA kit (#ADI-900-091; Enzo Life Sciences, Lause, Switzerland) was used. Cross-reactivities of the assay to $PGF_{1\alpha}$ and $PGF_{2\alpha}$ were 4.6% and 1.85%, respectively, and <1% to any further eicosanoids.

4.3.4 Liver Tissue Analyses

On day 8, a liver biopsy was conducted 2 h after the morning meal using a custom-made biopsy trocar [23]. Biopsy tissue was immediately frozen in liquid nitrogen and stored at -80°C until further analysis. Hippocalcin-like 1 (HPCAL1; NM 001098964), low-density lipoprotein receptor-related protein 10 (LRP10; BC149232) and RNA polymerase II (POLR2A; NM_001206313.1) were used as reference genes (given accession numbers related to NIH GenBank). Primer sequences, accession numbers and PCR conditions for target genes related to antioxidative status (catalase (CAT); glutathione peroxidase 1 (GPX1); superoxide dismutase (SOD)) and inflammation (tumor necrosis factor (TNF); interleukin-1α and -1β (*IL1A*, *IL1B*); haptoglobin (*HP*); fibrinogen (*FGA*); serum amyloid A2 (SAA2); and C-reactive protein (CRP)) are listed in Table 4.1. As recently described [23], primer products were verified by sequencing using the BigDye Terminator version 1.1 Cycle Sequencing kit and an ABI 3130 Genetic Analyzer (Life Technologies, Carlsbad, USA). Realtime PCR was performed using a LightCycler (Roche Molecular Biochemicals, Mannheim, Germany); SYBR Green I was used as the fluorescent dye. Melting curve analysis and agarose gel electrophoresis were used to confirm the specificity of the PCR products. Quantification cycle values and amplification efficiencies obtained using LinRegPCR version 2013.0 [35] were imported into qBASE+ version 2.6.1 (Biogazelle, Gent, Belgium) for all subsequent calculations and quality controls. The geometric mean of the reference gene abundances was used for normalization. Data are presented as the ratio of the copy numbers of genes of interest and the geometric mean of the reference genes' abundances.

Table 4.1: Characteristics of Primers and Real-Time RT-PCR Conditions ¹										
Gene	Forward (5'→ 3')	Reverse (5'→ 3')	NIH GenBank	Amplicon	Mean	Efficiency				
			accession number	size (bp)	Cq²					
Proinflan	nmatory Cytokines									
TNF	AGAGGGAAGAGCAGTCCCCAG	TTCACACCGTTGGCCATGAG	NM_173966.3	181	27.30	1.90				
IL1A	TGAACGACGCCCTCAATCAA	GGTGTCTCAGGCATCTCCTTT	NM_174092.1	226	27.98	1.90				
IL1B	AACGTCCTCCGACGAGTTTC	GCTCATGCAGAACACCACTTC	NM_174093.1	163	25.70	1.89				
Acute-Ph	ase Proteins									
HP	GGCCCCGAGATTGCTAATA	CTCTGGGCAGCTGTCATCTT	NM_001040470.2	172	15.53	1.88				
SAA2	CCACTGGGGATCAGCACAAT	CCTCTTTGGGCAGCGTCATA	NM_001075260.2	212	16.25	1.82				
FGA	CGCGATTGAAAGCAAGCACT	GAAGTGTGGATACCTCTGGCA	NM_001033626.1	129	14.71	1.87				
CRP	CAGGCCAGACAGACTTGCATA	TGCTGCTTGGTGGCATAA	NM_001144097.1	181	18.52	1.90				
Antioxida	ative Enzymes									
CAT	TCACTCAGGTGCGGACTTTC	GGATGCGGGAGCCATATTCA	NM_001035386.2	162	18.40	1.87				
GPX1	CTTCCCCTGCAACCAGTT	GGCAATTCAGGATCTCCTCGTT	X13684.1	62	20.97	1.87				
SOD	AAGGCCGTGTGCGTGCTGAA	CAGGTCTCCAACATGCCTCT	M81129.1	246	20.44	1.87				

¹ Initial denaturation = 10 min at 95°C; denaturation =15 s at 95°C; annealing = 10 s at 60°C; extension = 30 s at 72°C.

² Quantification cycle.

4.3.5 Statistical Analyses

Statistical analyses were conducted using SAS software, version 9.3 for Windows, SAS Institute Inc., Cary, USA. Descriptive statistics and tests for normality were calculated using the UNIVARIATE procedure of Base SAS software. Body weight, average daily weight gain and data for hepatic gene expression were analyzed by ANOVA with the MIXED procedure of SAS/STAT taking a model with the fixed factors diet (levels: Col vs. For), quercetin (levels: Q+ vs. Q-) and the interaction diet×quercetin. Feed intake, body temperature, heart and respiratory rate and plasma concentrations of metabolites, hormones, flavonols and markers of antioxidative status were analyzed by repeated measurement ANOVA using the MIXED procedure of SAS/STAT software and a model with the fixed factors diet, quercetin and day of life (repeated variable) and all interactions between the fixed factors. Repeated measures on the same calf were taken into account using the REPEATED statement of the MIXED procedure and a type for the block diagonal residual covariance matrix chosen in dependence on the levels of day of life. For concentrations of plasma metabolites, hormones and data on antioxidative status in plasma, an unstructured type was used. For the plasma concentrations of flavonols, acute-phase proteins and Ig, a compound symmetry type was used. For data on heart and respiratory rate, as well as body temperature and feed intake, an autoregressive (1) type was applied. Least-squares means (LSM) and their standard errors (SE) were computed for each fixed effect in the models and all pairwise differences of LSM were tested by the Tukey-Kramer procedure. The SLICE statement of the MIXED procedure was used to conduct partitioned analyses of the LSM for interactions. Fecal score was analyzed by a generalized linear mixed model using the GLIMMIX procedure of SAS/STAT and a Poisson model with the fixed factors diet, guercetin and day of life (repeated variable) and all interactions between these fixed factors. Repeated measurements on the same animal were taken into account by the RESIDUAL option of the RANDOM statement of the GLIMMIX procedure using a compound symmetry structure for the block diagonal residual covariance matrix. Sick frequencies of calves with respect to diet and quercetin were analyzed with the FREQ procedure of SAS/STAT software using two-way tables of diet by sick and quercetin by sick and the exact Pearson chi-square test. Effects and differences were considered significant at P < 0.05.

4.4 RESULTS

4.4.1 Feeding, Growth Performance and Health Status

Mean body weight at birth was $45.5 \pm 1.9 \,\text{kg}$ and increased with age (P < 0.01) by $368 \pm 140 \,\text{g/d}$, without group differences, respectively. All calves received their first meal $2.2 \pm 0.1 \,\text{h}$ after birth. Milk intake related to body weight did not differ among groups on

days 1 and 2 but was lower on days 3 and 4 in the colostrum-deprived calves (P < 0.01; **Figure 4.1 A**). On day 2, appetite was reduced in the formula-fed calves and the amount of tube-fed milk was higher than in colostrum-fed calves (P < 0.01; **Figure 4.1 B**). Heart rate decreased with age (P < 0.01) and respiratory rate tended to decrease (P = 0.08; **Figures 4.1 C and D**). Rectal temperature was highest on days 3 and 4 and subsequently decreased (P < 0.01). Rectal temperature and fecal score were higher in formula-fed than in colostrum-fed calves (P < 0.01; **Figures 4.1 E and F**). The number of calves with an allowed recovery day was similar among groups. With the exception of ColQ+, we treated one calf per group medically for navel infection. Four calves in group ForQ+ needed antispasmodic/analgesic treatment during a recovery day because of abdominal pain. Thus, Col-fed calves tended to be less susceptible to illness (P = 0.08), whereas quercetin treatment did not affect well-being (P = 0.38). Due to severe disease, we had to remove two calves from the study on day 4 (ForQ+) and day 7 (ColQ+).

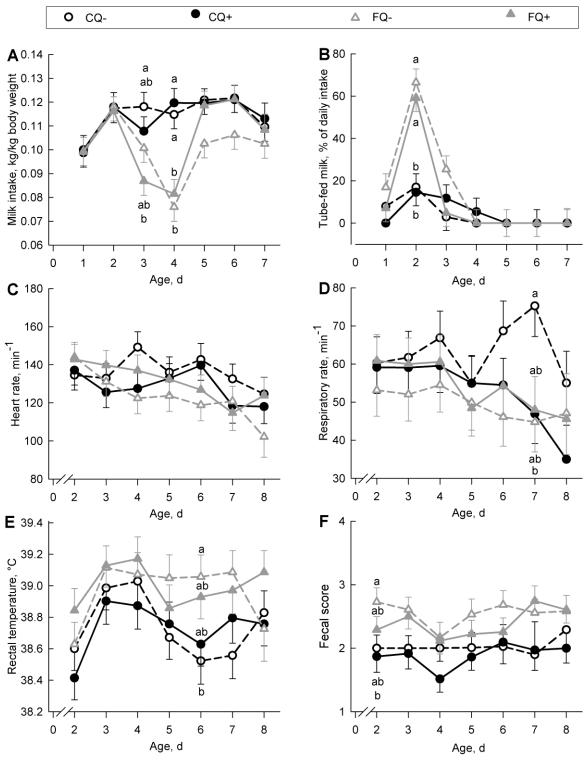


Figure 4.1: Feed Intake and Health Parameters. Neonatal calves were fed either colostrum (black circles) or formula (gray triangles) on days 1 and 2 and were supplemented with (filled symbols, solid lines) or without (open symbols, dashed lines) quercetin aglycone from day 2 to day 8 (50 mg/(kg body weight × day)). (A) Milk intake, (B) percent milk intake by tube feeding, (C) heart rate, (D) respiratory rate, (E) body temperature and (F) fecal score (according to Larson et al. [25]; 1 = normal, 2 = soft, 3 = runny, 4 = watery) were observed daily. Data are presented as the least squares means \pm standard errors. Least squares means with different lowercase letters (a, b) differ among groups within the same day (P < 0.05).

4.4.2 Flavonoid Content

Preprandial concentrations of total flavonols in plasma on days 1 and 2 (before first quercetin supplementation) did not reveal significant differences among groups, but very low concentrations were detectable in five colostrum-fed calves. In the control groups ForQ- and ColQ-, the plasma concentrations did not change with age. In ColQ+ and ForQ+, the plasma flavonol concentrations changed with age (P < 0.01); they reached the maximum on day 3 and decreased subsequently but tended to differ among groups only on day 4 (P = 0.08; **Figure 4.2 A**). Comparing the flavonol fractions in the plasma, the quercetin fraction (60% of total flavonoids) was highest in both dietary groups. The concentrations of isorhamnetin and tamarixetin on day 3 were higher in ColQ+ than in ForQ+ (**Figure 4.2 B**).

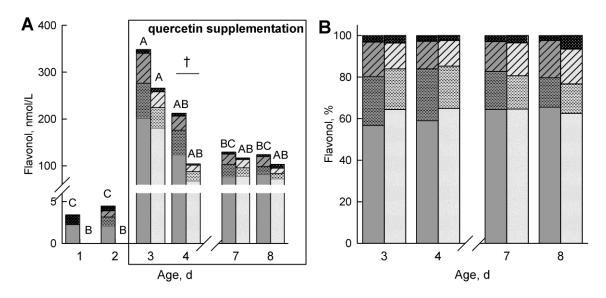


Figure 4.2: Flavonol Concentrations in Plasma. Stacked concentrations (A) and percentage composition (B) of flavonols in basal plasma samples of calves fed either colostrum (dark grey) or formula (light grey) on days 1 and 2 and supplemented with quercetin aglycone from day 2 to day 8 (50 mg/(kg body weight × day)). Flavonol metabolites: quercetin (without fill pattern), isorhamnetin (horizontal lines), tamarixetin (crossline pattern) and kaempferol (horizontal bold lines). Data are presented as the least squares means. Different uppercase letters (A, B, C) symbolize differences in the total flavonol concentration within the same group on different days (P < 0.05). † tend to differ among groups on the same day (P < 0.10)

4.4.3 Plasma Metabolites and Hormones

Plasma glucose concentrations were lowest on day 1, increased on day 2 in ColQ-, ForQ+ and ForQ- (P < 0.05) and decreased on day 4 only in ForQ+ and ForQ- (P < 0.05). The mean plasma concentration of glucose was higher in the colostrum-fed than in the formula-fed calves (P < 0.03; **Table 4.2**). The plasma lactate and non-esterified fatty acid concentrations were highest on day 1 and decreased with age (P < 0.01) without any group differences (Table 4.2). The concentrations of total protein in the plasma increased only in the colostrumfed calves (P < 0.01) and were higher in the colostrum-fed than in the formula-fed calves on day 2, day 4 and day 7 (P < 0.01; Figure 4.3 A). Plasma albumin decreased with age (P < 0.01) in all groups (**Figure 4.3 B**). For both total protein and albumin, the diet×age interaction was significant (P < 0.01). The plasma urea concentrations increased on day 2 in the formula-fed and on day 7 in the colostrum-fed calves (P < 0.01; Table 4.2). The plasma triglyceride concentrations increased until day 4 only in the colostrum-fed calves (P < 0.05) and continuously decreased from day 2 to day 7 in the formula-fed calves (P < 0.01). The plasma cholesterol concentration increased with age (P < 0.01). The plasma triglyceride (day 7) and cholesterol concentrations (day 4 and day 7) were higher in the colostrum-fed than in the formula-fed calves (P < 0.01; Table 4.2).

The plasma insulin concentrations decreased with age (P < 0.01; Table 4.2). Glucagon increased to a maximum plasma concentration on day 2 and subsequently decreased in all calves (P < 0.01). The glucagon concentration on day 4 and day 7 was higher in the colostrum-fed than in the formula-fed calves (P < 0.01; Table 4.2). The cortisol concentrations decreased with age in all groups (P < 0.01) but decreased earlier in the colostrum-fed calves. The mean cortisol concentrations in plasma during the first week of life were lower in the colostrum-fed than in the formula-fed calves (P = 0.02; Table 4.2). Quercetin treatment did not affect the concentrations of metabolites nor hormones in the plasma.

Table 4.2: Metabolites and Hormones of Calves during the First Week of Life.

	Age		Group (Die	t, Quercetin)	ANOVA <i>P</i> -values ¹				
		Colostrum		Forn	nula				
Item	(d)	Q-	Q+	Q-	Q+	Diet	Quercetin	Age	Diet × Age
Glucose (mmol/L)	1	4.3 ± 0.6	4.7 ± 0.6	4.1 ± 0.6	4.1 ± 0.6	0.03	0.89	<0.01	0.06
	2	6.0 ± 0.4	5.5 ± 0.4	5.8 ± 0.4	5.8 ± 0.4				
	4	5.5 ± 0.2ª	5.6 ± 0.2^a	4.4 ± 0.2^{b}	4.4 ± 0.2^{b}				
	7	5.2 ± 0.2ª	5.2 ± 0.2^a	4.4 ± 0.2^{b}	4.6 ± 0.2^{ab}				
Lactate (mmol/L)	1	4.0 ± 0.7	2.8 ± 0.7	3.7 ± 0.7	3.1 ± 0.7	0.88	0.78	<0.01	0.32
	2	2.4 ± 0.3	3.0 ± 0.3	2.7 ± 0.3	2.9 ± 0.3				
	4	1.0 ± 0.2	1.0 ± 0.2	1.0 ± 0.2	1.4 ± 0.2				
	7	0.5 ± 0.1	0.8 ± 0.1	0.6 ± 0.1	0.4 ± 0.1				
Urea (mmol/L)	1	3.3 ± 0.5	3.6 ± 0.5	3.1 ± 0.5	3.3 ± 0.5	0.99	0.55	<0.01	0.02
	2	3.5 ± 0.6	4.3 ± 0.6	5.0 ± 0.6	4.4 ± 0.6				
	4	3.7 ± 0.5	3.9 ± 0.5	3.8 ± 0.5	4.6 ± 0.6				
	7	5.0 ± 0.4	5.4 ± 0.4	4.5 ± 0.4	4.0 ± 0.4				
NEFA (µmol/L)	1	830 ± 156	709 ± 156	767 ± 156	733 ± 156	0.99	0.70	<0.01	0.95
	2	359 ± 43	368 ± 43	387 ± 43	372 ± 43				
	4	290 ± 71	269 ± 71	226 ± 71	285 ± 75				
	7	152 ± 38	157 ± 40	198 ± 38	164 ± 40				
Triglycerides (mmol/L)	1	0.24 ± 0.04	0.25 ± 0.04	0.26 ± 0.04	0.23 ± 0.04	<0.01	0.63	<0.01	<0.01
	2	0.31 ± 0.04	0.26 ± 0.04	0.30 ± 0.04	0.26 ± 0.04				
	4	0.40 ± 0.06	0.43 ± 0.06	0.21 ± 0.06	0.19 ± 0.07				
	7	0.21 ± 0.02ª	0.23 ± 0.02ª	0.10 ± 0.02 ^b	0.09 ± 0.02 ^b				

Table 4.2 Continuation.

	Age		Group (Die		ANOVA <i>P</i> -values ¹				
		Colostrum		Formula					
Item	(d)	Q-	Q+	Q-	Q+	Diet	Quercetin	Age	Diet × Age
Cholesterol (mmol/L)	1	0.62 ± 0.08	0.59 ± 0.08	0.63 ± 0.08	0.60 ± 0.08	0.01	0.90	<0.01	<0.01
	2	0.78 ± 0.07	0.76 ± 0.07	0.92 ± 0.07	0.76 ± 0.07				
	4	1.34 ± 0.09 ^a	1.51 ± 0.09 ^a	1.00 ± 0.09 ^b	0.85 ± 0.09^{b}				
	7	1.49 ± 0.12a	1.71 ± 0.12 ^a	1.03 ± 0.12 ^b	0.96 ± 0.12 ^b				
Insulin (µg/L)	1	0.59 ± 0.43	0.56 ± 0.45	1.59 ± 0.43	0.96 ± 0.43	0.10	0.21	<0.01	0.43
	2	0.77 ± 0.19	0.89 ± 0.19	1.14 ± 0.19	0.85 ± 0.19				
	4	0.34 ± 0.06	0.29 ± 0.06	0.35 ± 0.06	0.26 ± 0.07				
	7	0.36 ± 0.15	0.39 ± 0.16	0.51 ± 0.15	0.18 ± 0.16				
Glucagon (ng/L)	1	82 ± 17	93 ± 17	120 ± 17	118 ± 17	0.01	1.00	<0.01	<0.01
	2	331 ± 27	324 ± 27	288 ± 27	303 ± 27				
	4	194 ± 15ª	181 ± 15 ^{ab}	130 ± 15⁵	138 ± 15 ^{ab}				
	7	163 ± 10ª	159 ± 10ª	82 ± 10 ^b	73 ± 10 ^b				
Cortisol (µg/L)	1	36.8 ± 3.7	39.9 ± 3.7	34.3 ± 3.7	42.8 ± 3.7	0.02	0.30	<0.01	0.01
	2	15.6 ± 4.1 ^{ab}	14.3 ± 4.1 ^b	28.5 ± 4.1 ^{ab}	31.2 ± 4.1 ^a				
	4	13.2 ± 2.7	9.6 ± 2.7	11.5 ± 2.7	15.7 ± 2.8				
	7	9.9 ± 1.3	8.3 ± 1.4	8.1 ± 1.3	11.9 ± 1.4				

Data are given as the least squares means \pm standard errors. Least squares means within a row with different lowercase letters (a, b) differ (P < 0.05). Q+, quercetin-supplemented; Q-, control (no quercetin); n = 7 per group. NEFA, non-esterified fatty acids.

¹Effects: For interactions that are not listed, the probability was P > 0.10.

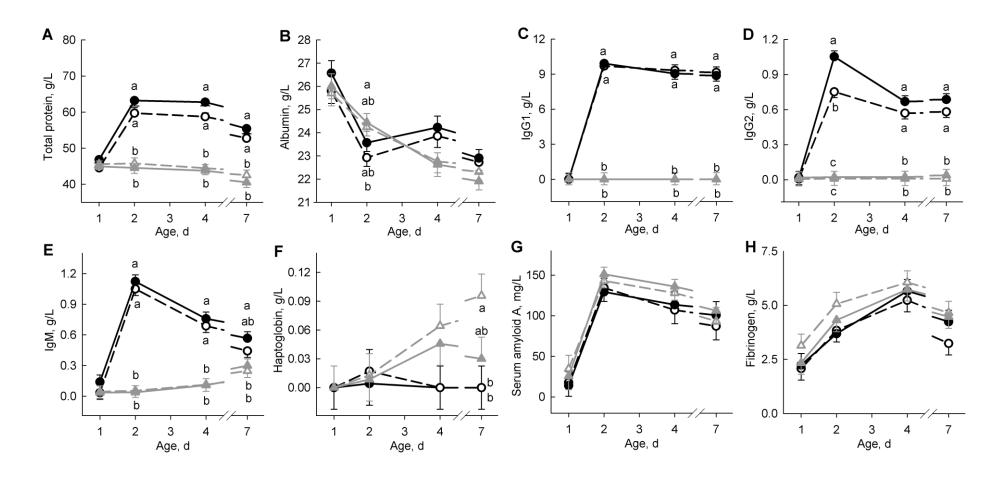


Figure 4.3: Total Protein, Albumin and Immune and Inflammatory Status in Blood Plasma. Basal concentrations of total protein (A) and albumin (B), immunoglobulins G1, G2 and M (C, D, E) and acute-phase proteins haptoglobin, serum amyloid A and fibrinogen (F, G, H) in the plasma of calves fed either colostrum (black circles) or formula (gray triangles) on days 1 and 2 and supplemented with (filled symbols, solid lines) or without (open symbols, dashed lines) quercetin aglycone from day 2 to day 8 (50 mg/(kg body weight × day)). Data are presented as the least squares means ± standard errors. Least squares means with different lowercase letters (a, b, c) differ among groups within the same day (P < 0.05).

4.4.4 Immunoglobulins and Acute-Phase Proteins

The plasma concentrations of IgG1 and IgG2 increased on day 2 only in colostrum-fed calves (P < 0.01; **Figures 4.3 C and D**). The plasma concentrations of IgM increased sharply until day 2 in colostrum-fed calves and then slowly decreased (P < 0.01). In the formula-fed groups, the IgM concentration increased until day 7 (P < 0.01) but was still lower than in the colostrum-fed groups (P < 0.01) at the end of the trial (**Figure 4.3 E**). The concentrations of haptoglobin in blood plasma were below the detection limit on day 1 in all calves and increased only in the formula-fed calves (P = 0.01) and were highest in ForQ- on day 7 (**Figure 4.3 F**). The mean haptoglobin concentration was higher in formula-fed than in the colostrum-fed calves (P = 0.03). The concentrations of SAA increased until day 2 and slowly decreased afterwards in all calves (P < 0.01; **Figure 4.3 G**). The concentrations of fibrinogen increased until day 4 and then decreased (P < 0.01) without differences among groups (**Figure 4.3 H**). However, the mean fibrinogen concentration tended to be higher after formula feeding (P = 0.10).

4.4.5 Antioxidative Status

Plasma TEAC increased in all groups until day 4 of life (P < 0.01) but was higher in the colostrum-fed than in the formula-fed calves (P < 0.01; **Figure 4.4 A**). FRAP decreased from day 1 to day 7 only in the formula-fed groups ($P \le 0.01$) and was higher (P < 0.05) on day 7 in the colostrum-fed groups (**Figure 4.4 B**). Although not statistically significant, the FRAP decrease was delayed in the quercetin-fed calves compared with the control groups. The mean concentrations of TBARS in plasma were higher in the colostrum-fed than in the formula-fed calves (P = 0.01) and revealed a diet×age interaction (P = 0.01) with higher concentrations in the colostrum-fed than in the formula-fed calves on day 4 (**Figure 4.4 C**). The mean concentrations of F2-isoprostanes decreased from day 1 to day 4 (P < 0.01) in all calves and tended to increase on day 7 only in ForQ- (P = 0.08; **Figure 4.4 D**).

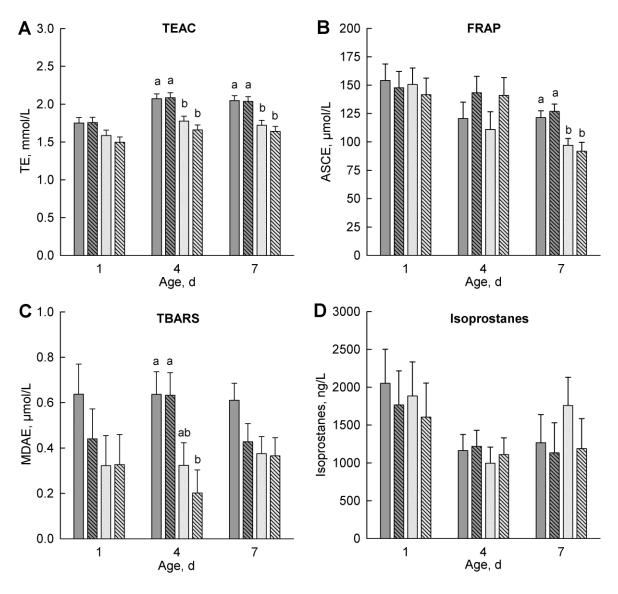


Figure 4.4: Antioxidative Status in Plasma. Plasma parameters for antioxidative capacity (A, B) and oxidative stress (C, D) in calves fed either colostrum (dark grey) or formula (light grey) on days 1 and 2 and supplemented with (crossline pattern) or without (without fill pattern) quercetin on days 2-8 (50 mg/(kg body weight × day)). Data are presented as the least squares means \pm standard errors. Bars with different lowercase letters (a, b) differ among groups within the same day (P < 0.05). TEAC, trolox equivalent antioxidative capacity (in trolox equivalents, TE); FRAP, ferric reducing ability of plasma (in ascorbic acid equivalents, ASCE); TBARS, thiobarbituric acid reactive species (in malondialdehyde equivalents, MDAE)

4.4.6 Hepatic Gene Expression

Regarding inflammation markers, the relative mRNA abundance of TNF was higher after quercetin feeding (P = 0.04; **Table 4.3**). The relative mRNA abundance of CRP was higher (P = 0.03) and that of SAA2 tended to be higher (P = 0.09) in the formula-fed than in the colostrum-fed groups (Table 4.3). For the antioxidative enzymes, the mRNA abundances of GPX1 and SOD were not different among groups, whereas the relative mRNA abundance of CAT tended to be higher in ColQ- than in ColQ+ (P = 0.06) but did not differ between ForQ- and ForQ+, revealing a diet×quercetin interaction (P = 0.04; Table 4.3).

Table 4.3: Hepatic mRNA Expression of Inflammatory and Antioxidative Traits in Newborn Calves.

	Gro	up (Diet	, Querc	etin)			ıes		
	Colostrum		Formula						
	Q-	Q+	Q- Q+		SE	Diet	Quercetin	Interaction	
Proinflam	matory	Cytokine	es						
TNF	0.98	1.15	0.89	1.48	0.10	0.41	0.04	0.39	
IL1A	1.17	0.90	1.05	1.18	0.08	0.62	0.66	0.22	
IL1B	3.00	3.68	3.11	2.69	0.30	0.49	0.83	0.38	
Acute-Pha	ase Prot	eins							
HP	14.74	29.14	39.78	23.09	4.61	0.30	0.90	0.10	
FGA	2.73	3.28	3.73	2.18	0.35	0.95	0.48	0.15	
SAA2	0.91	0.88	1.20	1.12	0.07	0.09	0.72	0.87	
CRP	0.90	0.96	1.11	1.06	0.04	0.03	0.99	0.42	
Antioxidative Enzymes									
CAT	1.22	0.77	1.01	1.08	0.06	0.68	0.11	0.04	
GPX1	1.08	0.95	0.98	1.05	0.03	0.99	0.65	0.13	
SOD	1.03	1.08	1.09	0.90	0.04	0.42	0.36	0.11	

Relative mRNA expression of genes related to inflammation and antioxidative status in the liver of 8-d-old calves; data are given in arbitrary units and are presented as the means \pm standard errors (SE). Q+, quercetin-supplemented; Q-, control (no quercetin); n = 6 per group.

CAT, catalase; GPX1, glutathione peroxidase; SOD, superoxide dismutase; TNF, tumor necrosis factor; IL1A, interleukin-1 α ; IL1B, interleukin-1 β ; HP, haptoglobin; FGA, fibrinogen; SAA2, serum amyloid A2; CRP, C-reactive protein.

4.5 DISCUSSION

In this study, we were able to confirm the oral bioavailability of quercetin in colostrum-fed calves and further showed that after colostrum deprivation, quercetin is absorbed in equal amounts. As indicated earlier, oral bioavailability decreases with age due to intestinal maturation and therefore reduced permeability or more effective elimination processes [24]. The pattern of flavonol fractions in the plasma in this study was different from the patterns in 29-day-old calves or adult cattle but comparable to observations in 2-day-old calves [24, 36]. This result can be explained by the maturation of intestinal metabolism because it has been shown for rats and pigs [37, 38] that orally administered flavonols undergo complete metabolism inside the intestinal mucosa, where glucuronidation is of major importance [39]. Although flavonols were present in the plasma of guercetin-fed calves, we failed to detect any effect on the metabolic parameters or hormone concentrations, which was also the case in previous studies in cattle [24, 36, 40]. However, we could confirm a variety of effects caused by colostrum-deprivation during the first 2 days of life. Cortisol is crucial for initiation of parturition and catabolic activity, especially during hypoxia at birth. However, a decrease of the plasma cortisol concentration is delayed in formula-fed calves, as seen in previous studies [41, 42]. We assume that the delay was possibly caused by either abdominal pain or by reduced utilization of nutrients, which is proven to increase stress parameters in sheep [43, 44].

Higher plasma glucose in colostrum-fed groups was most likely caused by enhanced intestinal maturation and therefore enlarged absorptive surface [45, 46]. Additionally, colostrum feeding seems to accelerate maturation of the pancreas [47] because glucagon concentrations were also higher in colostrum-fed calves from day 4 onwards. Although the basal insulin concentrations were similar among dietary groups, we showed in a companion paper that the postprandial insulin response is more pronounced in colostrum-fed groups [23].

Higher triglyceride and cholesterol concentrations in colostrum-fed calves are in accordance with observations in other studies [41, 42] and are due to the enhanced stimulation of intestinal differentiation by colostral growth factors and hormones. However, it must be considered that diarrhea in formula-fed calves accelerated intestinal transit and therefore reduced absorption time, which might also account for differences in the plasma triglyceride and cholesterol concentrations between dietary groups. Similar concentrations of non-esterified fatty acids indicate that the energy balance in the dietary groups is equal and that lipid mobilization seems to be of minor importance to maintain energy balance; previous studies also failed to demonstrate consistent effects in neonatal calves [41, 42, 48].

As we expected, the concentration of total protein in the plasma increased only in the colostrum-fed groups, although the protein content of the formula and colostrum was similar.

The decrease of the albumin concentration from day 1 to day 2 was probably caused by hemodilution after first feed intake and coincides with previous findings [49, 50]. Because only Iq, but not the albumin concentrations in plasma were affected by colostrum feeding, differences in total protein are most likely caused by absorption of colostral immunoglobulins after colostrum intake [42]. Colostrum intake is important not only for maturational processes but also to acquire passive immunity because the bovine placenta is impermeable to antibodies. As reviewed by Weaver et al. [51], calves with serum IgG concentrations below 10 g/L (total protein < 52 g/L) 24 h after birth suffer from failure of passive transfer, which is associated with increased morbidity and mortality as well as reduced performance. In our study, colostrum-fed calves exceeded this threshold; however, failure of passive transfer was a relevant problem in formula-fed calves, although they were parenterally supplemented with bovine immunoglobulins on day 1. Only formula-fed calves showed slight hyperthermia on the first days of life and a higher incidence of diarrhea, probably caused by gastrointestinal inflammation due to missing local immunity [52], accompanied by abdominal pain and diminished appetite. Obviously, parenteral and oral treatments with immunoglobulins could not prevent local or systemic infections in formula-fed calves, probably because of missing herd-specific immunoglobulins in the formula and the administered drugs. However, we did not perform microbiological analyses in the feces of the calves to determine pathogens that might have caused loose feces. The time course of plasma IgM concentration in formula-fed calves indicates that the indigenous production of IgM is evident as early as day 4 of life, but the concentrations are too low to effectively protect against infections [9].

We assume that inflammatory processes also activated the synthesis of acute phase proteins. Although the increase of the plasma acute phase proteins in all calves emphasized the immunological burden of the new environment, higher haptoglobin concentrations in formula-fed calves suggest more severe inflammatory processes than in colostrum-fed calves [53, 54], which was underlined by the greater incidence of gastrointestinal infections. Because serum amyloid A is more susceptible to stress [53, 55], including physical stress, we suppose that the plasma concentrations were equally high in all groups due to the experimental procedures (e.g., continued venipuncture, restricted feeding, single penning, temporal fixation and dietary changes).

On the mRNA level, we did not find differences in the acute phase proteins between dietary groups because hepatic transcription precedes translation and protein release into circulation. Thus, we obviously missed the time point of elevated mRNA abundances of *HP*. However, mRNA abundance of *CRP*, a moderate acute phase protein, was elevated in the formula-fed groups on day 8, which could indicate the importance of C-reactive protein as a major component of the bovine innate immune system [56]. Hence, increased C-reactive protein production is compensative for the absence of immunoglobulins in colostrum-deprived calves.

Regarding pro-inflammatory cytokines, we found elevated hepatic mRNA abundance of TNF in quercetin-supplemented calves but no differences between dietary groups. Under immunocompetent conditions, the impact of various noxae leads to local production of proinflammatory cytokines, e.g., TNF or interleukin-1, which in turn induce the systemic acute phase response, including synthesis of acute phase proteins [57]. Within signal transduction, reactive oxygen species act as second messengers, thus enhancing TNF-induced gene expression, and oxidative conditions potentiate the activation of respective pathways [57, 58]. Although high doses of quercetin were repeatedly shown to reduce mRNA expression of TNF in vitro [59, 60] application of prophylactic doses increased the ratio of pro- to antiinflammatory cytokines in murine macrophages [60]. We assume that the hepatic guercetin concentration in our experiment did not exceed the prophylactic dose; thus, quercetin might have increased the expression of TNF. Unfortunately, we did not measure TNF protein concentration, but we suppose that posttranscriptional processes anticipated TNF signal transduction [17, 61], because quercetin scavenged reactive oxygen species necessary for signal transduction. Therefore, the observed quercetin effects on *TNF* gene expression could not have been forwarded on the expression of target genes, e.g., IL1B or acute phase proteins, as would have been expected otherwise.

Concerning the antioxidative status in the plasma, previous findings in neonatal calves are inconsistent. Inanami et al. [62] and Stohrer, Lutz and Stangassinger [63] concluded from comparisons between calves and dams that the former are highly susceptible to oxidative stress due to immature defense systems, whereas Gaál et al. [64] deduced from high FRAP values, despite the high reactive oxygen species, that calves are well-prepared to address oxidative stress. Although the increase of TEAC in this and a previous study of our group [65] indicates rising antioxidative capacity during the first week of life, this result was not supported by determination of FRAP, another marker of antioxidative capacity. Furthermore, the values of TBARS, which serve as a proxy to measure the products of lipid peroxidation, were unaffected by age in this study, which contradicts earlier studies [62, 64, 66]. However, the reliability of TBARS is criticized for low sensitivity and specificity, and the use of F2isoprostanes is recommended as the most reliable approach to assess oxidative stress in vivo [67, 68]. The time courses of F2-isoprostanes were significantly decreased in all calves; thus, we support the theory that exposure to an oxygen-rich environment following hypoxia during birth results in severe oxidative stress in the newborn [5]. Although quercetin is known to exert antioxidative effects, we did not find improved antioxidative status in the plasma of calves nor increased hepatic expression of antioxidative enzymes, which is in line with previous findings of our group in research conducted on neonatal calves and lactating dairy cows [65, 69]. Orally administered quercetin is completely metabolized inside the intestinal mucosa; thus, the gastrointestinal tract is the first site of action for quercetin [39], as it is for pathogens and gastrointestinal disorders commonly occurring during the first weeks of life.

Because we did not examine the intestinal tissue, we cannot exclude local effects of quercetin on antioxidative status. Except for F2-isoprostanes, the parameters of the antioxidative system in the plasma were higher after colostrum feeding. Colostrum is a source of reactive oxygen species, but it also contains a variety of antioxidative factors [66, 70], the latter increasing with time after parturition [71]. The absorption of pro- and antioxidants present in colostrum [72] might have contributed to the rise of the respective parameters in the plasma of colostrum-fed calves and triggered immunological processes, which in turn caused elevated plasma levels. However, the hepatic mRNA abundances of antioxidative enzymes seemed not to be affected by diet, but all groups were fed milk replacer from day 3 onwards and potential dietary effects on hepatic antioxidative status might not be long-lasting.

In conclusion, oral administration of quercetin aglycone at a daily dose of 50 mg/kg body weight to newborn calves during the first week of life is unable to compensate for inadequate colostrum supply. Quercetin did not show any positive effects on neonatal antioxidative or anti-inflammatory status, whereas colostrum feeding improves neonatal health status by supporting passive immunity and by promoting antioxidative/oxidative status.

4.6 ACKNOWLEDGMENTS

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4.7 AUTHOR CONTRIBUTIONS

Conceived and designed the experiments: HMH JG SW. Performed the experiments: JG HMH. Analyzed the data: EK JW AT TS SW. Contributed reagents/materials/analysis tools: EK JW TS PJ SW. Wrote the paper: JG HMH.

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4.9 SUPPORTING INFORMATION

Supplemental Table 4.1: Complete data set of parameters regarding daily milk intake and health status as shown in Figure 4.1.

unu	Tiouiti		2 40 0110	*****	9						
Calf	Group	Feeding	Quercetin	Age,	BW,	Milk intake,	Force-fed milk,	Fecal	Rectal	Heart rate,	Resp. rate,
				d	kg	kg/kg BW	% of intake/day	score	temp, °C	per min	per min
1	CoIQ-	COL	Q-	1	38.5	0.0961	27.027				
4	CoIQ-	COL	Q-	1	46	0.09565	0				
5	CoIQ-	COL	Q-	1	54.5	0.09725	28.302	_			
10	ColQ-	COL	Q-	1	46	0.1	0	-	-	•	•
12	ColQ-	COL	Q-	1	44	0.1	Ö	•	•	•	•
	ColQ-			1			0	•	•	•	•
23		COL	Q-		48.5	0.10103		•	•	•	•
28	CoIQ-	COL	Q-	1	39.5	0.10127	0	•	•	•	
2	CoIQ+	COL	Q+	1	44	0.1	0				
3	CoIQ+	COL	Q+	1	52.5	0.09333	0				
7	CoIQ+	COL	Q+	1	42.5	0.09647	0				
8	CoIQ+	COL	Q+	1	50	0.1	0				
11	ColQ+	COL	Q+	1	43	0.10465	Ö	-	-	•	•
								•	•	•	•
19	ColQ+	COL	Q+	1	51	0.10588	0	•	•	•	•
26	CoIQ+	COL	Q+	1	46	0.1	0			•	
9	ForQ-	FOR	Q-	1	51	0.09804	48				
15	ForQ-	FOR	Q-	1	43	0.1	0				
18	ForQ-	FOR	Q-	1	44.5	0.09888	13.636				
21	ForQ-	FOR	Q-	1	45	0.1	37.778				
22	ForQ-	FOR	Q-	1	37	0.1	18.919	-	-	•	•
24	ForQ-	FOR	Q-	1	48	0.1	0	•	•	•	•
								•	•	•	•
29	ForQ-	FOR	Q-	1	47.5	0.09895	0	•	•	•	
13	ForQ+	FOR	Q+	1	44.5	0.09888	31.818				
14	ForQ+	FOR	Q+	1	52	0.1	0				
17	ForQ+	FOR	Q+	1	43	0.09302	0				
20	ForQ+	FOR	Q+	1	33	0.1	0				
25	ForQ+	FOR	Q+	1	46.5	0.10108	Ö	-	-	•	•
27				1			17.021	•	•	•	•
	ForQ+	FOR	Q+		46.5	0.10108		•	•	•	•
30	ForQ+	FOR	Q+	1	47	0.1	0		_:_		·-
1	CoIQ-	COL	Q-	2	38.5	0.1039	82.5		38.7	120	78
4	CoIQ-	COL	Q-	2	46	0.12174	0	2	38.4	150	66
5	CoIQ-	COL	Q-	2	54.5	0.1156	0	2	38.2	138	56
10	ColQ-	COL	Q-	2	46	0.12174	0	3	38.9	138	66
12	ColQ-	COL	Q-	2	44	0.12045	Ö	2	38.4	126	48
23	ColQ-	COL	Q-	2	48.5	0.11959	36.207	2	38.9	132	60
28	CoIQ-	COL	Q-	2	39.5	0.11899	0	1	38.7	138	48
2	CoIQ+	COL	Q+	2	44	0.12273	0		38.3	126	48
3	CoIQ+	COL	Q+	2	52.5	0.1181	85.484	1	38.5	138	96
7	CoIQ+	COL	Q+	2	42.5	0.11765	0	2	38.3	180	48
8	CoIQ+	COL	Q+	2	50	0.12	0	2	38.2	150	54
11	CoIQ+	COL	Q+	2	43	0.11395	16.327		38.6	132	54
19	ColQ+	COL	Q+	2	51	0.11176	0	2	38.5	126	60
				2			0			108	
26	ColQ+	COL	Q+		46	0.12174			38.5		54
9	ForQ-	FOR	Q-	2	51	0.11765	95	-	38.9	162	48
15	ForQ-	FOR	Q-	2	43	0.11628	42		38.2	168	60
18	ForQ-	FOR	Q-	2	44.5	0.11685	80.769	3	38.6	144	60
21	ForQ-	FOR	Q-	2	45	0.11778	100	2	38.9	132	66
22	ForQ-	FOR	Q-	2	37	0.12162	100	2.75	38.6	144	54
24	ForQ-	FOR	Q-	2	48	0.12083	29.31	3	38.9	138	48
29	ForQ-	FOR	Q-	2	47.5	0.11579	18.182	3	38.3	120	36
				2				3			
13	ForQ+	FOR	Q+		44.5	0.11011	0		39.5	166	48
14	ForQ+	FOR	Q+	2	52	0.10962	26.316	2	39.4	126	66
17	ForQ+	FOR	Q+	2	43	0.11395	0	2.5	38.9	138	48
20	ForQ+	FOR	Q+	2	33	0.12424	95.122	2	38.1	156	72
25	ForQ+	FOR	Q+	2	46.5	0.11613	88.889	2.5	38.7	132	72
27	ForQ+	FOR	Q+	2	46.5	0.12043	100	3	38.9	168	60
30	ForQ+	FOR	Q+	2	47	0.11915	103.571	2	38.4	114	60
1	ColQ-	COL	Q-	3	40	0.115	0	2	38.8	108	66
4	ColQ-	COL	Q-	3	46	0.12174	0	2	38.7	132	60
5	ColQ-	COL	Q-	3	56	0.11786	0	1	38.8	126	54
10	CoIQ-	COL	Q-	3	48.5	0.11753	0	3	39.4	150	90
12	CoIQ-	COL	Q-	3	46	0.12391	0	2	39.1	126	66
23	CoIQ-	COL	Q-	3	50.5	0.10693	20.37	2	39	108	48
28	CoIQ-	COL	Q-	3	39.5	0.12405	0	2	39.1	180	48
2	ColQ+	COL	Q+	3	46.5	0.11613	Ö	2	38.8	72	-
3	ColQ+	COL	Q+	3	55	0.09273	19.608	1	38.6	126	90
								'	30.0	120	90
7	ColQ+	COL	Q+	3	44	0.11818	0				<u>.</u> .
8	ColQ+	COL	Q+	3	52	0.11923	0		38.7	132	54
11	CoIQ+	COL	Q+	3	45	0.10889	38.776	3	39.2	120	42
19	CoIQ+	COL	Q+	3	52.5	0.08	23.81	2	38.6	138	42
26	CoIQ+	COL	Q+	3	47	0.11915	0		39.6	132	66
9	ForQ-	FOR	Q-	3	51	0.11765	91.667	3	39.4	144	48
15	ForQ-	FOR	Q-	3	45.5	0.11868	0	2	38.8	132	36
18	ForQ-	FOR	Q-	3	45.5	0.1011	0	3	38.7	174	72
		FOR	Q-	3	45.5	0.12083	0	2		120	
21	ForQ-	1 OK	Q-	3	40	0.12003	U	2	39.5	120	54

Supplemental Table 4.1 Continuation.

Supp	Olemer Group	Tal lar	OIE 4.1 Co	ONTIN Age,	uatior вw,). Milk intake,	Force-fed milk,	Fecal	Rectal	Heart rate,	Resp. rate,
				d	kg	kg/kg BW	% of intake/day	score	temp, °C	per min	per min
22 24	ForQ- ForQ-	FOR FOR	Q- Q-	3 3	38 48	0.12105 0.06042	0 86.207	2.75 2.5	38.8 39.4	114 138	66
29	ForQ-	FOR	Q-	3	50.5	0.06535	0	3	39.2	96	39
13	ForQ+	FOR	Q+	3	47.5	0.08421	17.5	2	39.3	168	48
14	ForQ+	FOR	Q+	3	53	0.11698	0	2	39.2	126	54
17	ForQ+	FOR	Q+	3	44.5	0.11685	0	2	39.7	138	42
20	ForQ+	FOR	Q+	3	34.5	0.08986	16.129	2	38.6	162	90
25	ForQ+	FOR	Q+	3	48	0.12083	0	2.5	39.4	126	72
27	ForQ+	FOR	Q+	3	44	0.01364	0	4	38.6	144	48
30 1	ForQ+ ColQ-	FOR COL	Q+ Q-	3 4	45.5 39.25	0.06593 0.11849	0 0	3 2	39.1 39.2	114 120	66 102
4	ColQ-	COL	Q-	4	46	0.11649	0		38.4	138	54
5	ColQ-	COL	Q-	4	56	0.11786	Ö	2	39.2		
10	CoIQ-	COL	Q-	4	48.5	0.11959	0	2	39.4	168	102
12	CoIQ-	COL	Q-	4	46	0.11957	0	2	39.4	162	72
23	CoIQ-	COL	Q-	4	49.5	0.08233	0	2	38.5	132	48
28	ColQ-	COL	Q-	4	39.5	0.12405	0	2	39.1	180	33
2 3	ColQ+ ColQ+	COL COL	Q+ Q+	4 4	46.5 53.75	0.12043	0 37.903	2 1	39.3 39	102	66 84
3 7	ColQ+	COL	Q+	4	44	0.11996 0.11818	0	1	38.7	156 162	78
8	ColQ+	COL	Q+	4	51	0.12162	Ö		00.7	102	70
11	ColQ+	COL	Q+	4	45	0.12	Ŏ	2	38.9	114	42
19	CoIQ+	COL	Q+	4	52.5	0.1181	0	2	38.5	108	30
26	CoIQ+	COL	Q+	4	46.5	0.11936	0	1	39.2	126	66
9	ForQ-	FOR	Q-	4	51	0.04706	0	3	39.4	120	48
15	ForQ-	FOR	Q-	4	44.25	0.06837	0	2	38.9	150	42
18	ForQ-	FOR	Q-	4 4	45 46 F	0.04948	0 0		38.9		60
21 22	ForQ- ForQ-	FOR FOR	Q- Q-	4	46.5 37.5	0.08153 0.07945	0	2 2	39 39.5	114 108	60
24	ForQ-	FOR	Q-	4	48	0.12083	Ö		39.2	114	54
29	ForQ-	FOR	Q-	4	50.5	0.08515	Ö		38.6	108	48
13	ForQ+	FOR	Q+	4	47.5	0.12	0	2	39.4	126	30
14	ForQ+	FOR	Q+	4	52.5	0.09022	0	2	39.1	138	66
17	ForQ+	FOR	Q+	4	44.5	0.11685	0	2	39.4	150	54
20	ForQ+	FOR	Q+	4	34.5	0.13333	0	2	39.4	168	84
25 27	ForQ+ ForQ+	FOR FOR	Q+ Q+	4 4	47.25 44	0.02718 0.02045	0 0	2	39 38.7	132	78
30	ForQ+	FOR	Q+	4	46.25	0.02043	0	2	39.2	114	66
1	ColQ-	COL	Q-	5	40.23	0.12	0		38.9	116	72
4	CoIQ-	COL	Q-	5	46	0.12174	0		37.6	138	42
5	CoIQ-	COL	Q-	5	56	0.11786	0	2	39.1	132	42
10	CoIQ-	COL	Q-	5	48.5	0.11959	0	2	39.1	156	54
12	CoIQ-	COL	Q-	5	46	0.12174	0	:	38.2	.:.	-:-
23	ColQ-	COL	Q-	5	50.5	0.11881	0	2	38.9	120	60
28 2	CoIQ- CoIQ+	COL COL	Q- Q+	5 5	39.5 46.5	0.12658 0.12043	0 0	2 2	38.9 39.1	156 102	54
3	ColQ+	COL	Q+	5	55	0.12043	0	2	38.9	138	84
7	ColQ+	COL	Q+	5	44	0.12045	Ö	2	38.8	162	
8	CoIQ+	COL	Q+	5	52	0.11923	0	2	37.7	120	42
11	CoIQ+	COL	Q+	5	45	0.12	0		39	138	54
19	ColQ+	COL	Q+	5	52.5	0.1181	0	2	38.8	132	30
26	ColQ+	COL	Q+	5	47 51	0.11915	0	1	39	138	66
9 15	ForQ- ForQ-	FOR FOR	Q- Q-	5 5	51 45.5	0.04706 0.12088	0 0	4 2	39 39.1	114 162	42 42
18	ForQ-	FOR	Q-	5	45.5	0.07253	0	2	39.1	102	42
21	ForQ-	FOR	Q-	5	48	0.12083	0	2	39.4	120	48
22	ForQ-	FOR	Q-	5	38	0.12105	0	2	38.8	132	60
24	ForQ-	FOR	Q-	5	48	0.12083	0		39.3	108	60
29	ForQ-	FOR	Q-	5	50.5	0.11485	0		38.7	102	36
13	ForQ+	FOR	Q+	5	47.5	0.12	0		38.8	126	30
14 17	ForQ+ ForQ+	FOR FOR	Q+ Q+	5 5	53 44.5	0.12075 0.12135	0 0	2 2	39.7 38.7	132 144	54
20	ForQ+	FOR	Q+	5	34.5	0.12133	0	2	38.6	148	76
25	ForQ+	FOR	Q+	5	48	0.12083	Ö	2	38.8	138	60
27	ForQ+	FOR	Q+	5	44	0.14091	0	3	38.4	114	30
30	ForQ+	FOR	Q+	5	45.5	0.06154	0				
1	CoIQ-	COL	Q-	6	40	0.12	0		38.1	126	84
4	ColQ-	COL	Q-	6	46	0.12174	0	•	38.1		<u>.</u> .
5 10	ColQ-	COL	Q-	6 6	56 48.5	0.11964	0 0	2	38.3	168	54
10 12	CoIQ- CoIQ-	COL COL	Q- Q-	6	48.5 46	0.12371 0.12174	0	2	39.1	126	126
23	ColQ-	COL	Q-	6	50.5	0.12174	0		38.7	108	60
28	CoIQ-	COL	Q-	6	39.5	0.12658	ő	2	38.7	186	
2	CoIQ+	COL	Q+	6	46.5	0.12043	0		38.9	126	
3	ColQ+	COL	Q+	6	55	0.12364	0	2	38.5	126	78
7	ColQ+	COL	Q+	6	44	0.12045	0		38.3	180	36
8 11	ColQ+ ColQ+	COL	Q+ Q+	6	52 45	0.12115 0.12222	0 0		38.3	150 144	42 48
11 19	ColQ+	COL COL	Q+ Q+	6 6	45 52.5	0.12222	0	2	38.1 38.9	120	48 48
26	ColQ+	COL	Q+	6	47	0.11915	0	2	39.4	132	72
-		-	•	-			-			-	

Supplemental Table 4.1 Continuation.

Calf	Group	Feeding	Quercetin	Age,	BW,	Milk intake,	Force-fed milk,	Fecal	Rectal	Heart rate,	Resp. rate,
				d	kg	kg/kg BW	% of daily intake	score	temp, °C	per min	per min
9	ForQ-	FOR	Q-	6	51 45 5	0.06078	0	4	38.7	90	30
15 10	ForQ-	FOR	Q-	6 6	45.5	0.11868 0.1033	0 0	3	38.8 39.2	108	
18 21	ForQ- ForQ-	FOR FOR	Q- Q-	6	45.5 48	0.12083	0	2	39.2 38.9	132	54
22	ForQ-	FOR	Q-	6	38	0.12003	0	2	39.1	126	48
24	ForQ-	FOR	Q-	6	48	0.12083	0	_	39.1	126	54
29	ForQ-	FOR	Q-	6	50.5	0.11881	Ö	3	39.6	96	42
13	ForQ+	FOR	Q+	6	47.5	0.12	0	3	38.9	108	
14	ForQ+	FOR	Q+	6	53	0.12075	0		39.2	132	60
17	ForQ+	FOR	Q+	6	44.5	0.12135	0	2	38.7	132	36
20	ForQ+	FOR	Q+	6	34.5	0.12464	0	2	39.1	126	84
25	ForQ+	FOR	Q+	6	48	0.12083	0	2	38.5	156	72
27	ForQ+	FOR	Q+	6	45	0.12	0		38.7	102	30
30	ForQ+	FOR	Q+	6	45.5	0.12088	0	2	39.4	132	
1	ColQ-	COL	Q-	7 7	40 46 F	0.1125	0 0	2	38.9	116	78
4 5	ColQ-	COL	Q-	7	46.5	0.11613	0	2	38.7	116 114	•
10	CoIQ- CoIQ-	COL COL	Q- Q-	7	57.5 51	0.10087 0.10784	0	2	37.6 38.6	126	42
12	ColQ-	COL	Q-	7	48	0.11042	0	•	38.3	138	138
23	ColQ-	COL	Q-	7	52	0.10962	0	•	39.2	114	84
28	ColQ-	COL	Q-	7	41	0.10976	0	2	00.2	204	0.
2	ColQ+	COL	Q+	7	47.5	0.12	0	-	38.3	96	54
3	ColQ+	COL	Q+	7	56.5	0.12035	Ō		38.9	126	66
7	CoIQ+	COL	Q+	7	45.5	0.10989	0				
8	CoIQ+	COL	Q+	7	52.5						
11	CoIQ+	COL	Q+	7	46.5	0.11183	0		38.7	126	36
19	CoIQ+	COL	Q+	7	53.5	0.10654	0	2	39.1		
26	CoIQ+	COL	Q+	7	49	0.1102	0		39.2	120	60
9	ForQ-	FOR	Q-	7	50	0.1	0	4	39.1	96	36
15	ForQ-	FOR	Q-	7	46.5	0.10968	0		38.5	162	•
18	ForQ-	FOR	Q-	7	43.5	0.11264	0	3	39.2	108	<u>-</u> :.
21	ForQ-	FOR	Q-	7	51	0.10588	0	2	38.6	138	54
22	ForQ-	FOR	Q-	7	38	0.11316	0	2.5	38.9	126	42
24 29	ForQ- ForQ-	FOR FOR	Q- Q-	7 7	48 52	0.11042 0.06538	0 0	2 2	39.3 40	120 96	48 42
13	ForQ+	FOR	Q+	7	52 51	0.10784	0	2	39.1	90	42
14	ForQ+	FOR	Q+	7	55	0.10764	0	3	39.5	132	60
17	ForQ+	FOR	Q+	7	47	0.10851	0	3	38.6	102	00
20	ForQ+	FOR	Q+	7	36	0.10833	0	3	38.9	114	60
25	ForQ+	FOR	Q+	7	47.5	0.10947	0				
27	ForQ+	FOR	Q+	7	45	0.11111	0		38.6	108	36
30	ForQ+	FOR	Q+	7	42.5			2	39.3	102	
1	CoIQ-	COL	Q-	8					38.6	104	60
4	CoIQ-	COL	Q-	8					38.9	120	36
5	CoIQ-	COL	Q-	8				3	38.8	-	
10	CoIQ-	COL	Q-	8				2	38.9		
12	CoIQ-	COL	Q-	8		•	•	2	39	126	126
23	ColQ-	COL	Q-	8	•	•	•	2	38.8	126	42
28	ColQ-	COL	Q-	8 8	•	•	•	2	38.8	162	•
2 3	ColQ+ ColQ+	COL COL	Q+ Q+	8		•	•	2	38.7 38.8	114	•
7	ColQ+	COL	Q+	8	•	•	•	2	38.5	96	24
8	ColQ+	COL	Q+	8	•	•	•	2	38.9	30	27
11	ColQ+	COL	Q+	8	•	•	•	•	38.6	120	30
19	ColQ+	COL	Q+	8				2	39.1	120	
26	ColQ+	COL	Q+	8				2	38.7	150	42
9	ForQ-	FOR	Q-	8				3			
15	ForQ-	FOR	Q-	8						•	•
18	ForQ-	FOR	Q-	8				3	39	102	
21	ForQ-	FOR	Q-	8				2	38.7	102	51
22	ForQ-	FOR	Q-	8				3	38.3	108	48
24	ForQ-	FOR	Q-	8		•		•	-	-	•
29	ForQ-	FOR	Q-	8		•				-	•
13	ForQ+	FOR	Q+	8		•		3	39.1		
14 17	ForQ+	FOR	Q+	8		•	•	2	39.2	132	54
17 20	ForQ+ ForQ+	FOR FOR	Q+ Q+	8 8		•	•	3	38.8 38.9	132	
25	ForQ+	FOR	Q+	8	•	•	•	2	38.9	114	48
27	ForQ+	FOR	Q+	8	•	•	•	2	39.1	132	42
30	ForQ+	FOR	Q+	8	•		•	3.5	39.6		
	1 0.00	1 011	٧'	J	<u> </u>	· · · · · · · · · · · · · · · · · · ·	•	0.0	55.0		<u> </u>

doi:10.1371/journal.pone.0146932.s001

Supplemental Table 4.2: Complete data set of flavonoid measurements in blood plasma as shown in Figure 4.2.

Calf	Group	Feeding	Quercetin	Age,	Quercetin,	Isorhamnetin,	Tamarixetin,		Total flavonols,	Quercetin,	Isorhamnetin,	Tamarixetin,	Kaempferol,	Ratio
				d	nmol/L	nmol/L	nmol/L	nmol/L	nmol/L	% of total flav.	KIT:Q			
2	CQ+	COL	Q+	1	7.85	0	0	3.9759	11.826	66.381	0	0	33.6187	0.50645
3	CQ+	COL	Q+	1	7.969	0	0	4.0806	12.05	66.136	0	0	33.8638	0.51203
7	CQ+	COL	Q+	1	0	0	0	0	0	0	0	0	0	0
8	CQ+	COL	Q+	1	0	0	0	0	0	0	0	0	0	0
11	CQ+	COL	Q+	1	0	0	0	0	0	0	0	0	0	0
13	FQ+	FOR	Q+	1	0	0	0	0	0	0	0	0	0	0
14	FQ+	FOR	Q+	1	0	0	0	0	0	0	0	0	0	0
17	FQ+	FOR	Q+	1	0	0	0	0	0	0	0	0	0	0
19	CQ+	COL	Q+	1	0	0	0	0	0	0	0	0	0	0
20	FQ+	FOR	Q+	1	0	0	0	0	0	0	0	0	0	0
25	FQ+	FOR	Q+	1	0	0	0	0	0	0	0	0	0	0
26	CQ+	COL	Q+	1	0	0	0	0	0	0	0	0	0	0
27	FQ+	FOR	Q+	1	0	0	Ō	Ō	0	0	0	0	0	0
30	FQ+	FOR	Q+	1	0	0	0	0	0	0	0	0	0	0
2	CQ+	COL	Q+	2	6.606	2.098	0	4.0518	12.756	51.789	16.4465	0	31.7649	0.93092
3	CQ+	COL	Q+	2	8.006	5.358	5.213	0	18.577	43.098	28.8405	28.0613	0	1.32028
7	CQ+	COL	Q+	2	0	0	0	0	0	0	0	0	0	0
8	CQ+	COL	Q+	2	0	0	0	0	0	0	0	0	0	0
11	CQ+	COL	Q+	2	0	0	0	0	0	0	0	0	0	0
13	FQ+	FOR	Q+	2	0	0	0	0	0	0	0	0	0	0
14	FQ+	FOR	Q+	2	0	0	0	0	0	0	0	0	0	0
17	FQ+	FOR	Q+	2	0	0	0	0	0	0	0	0	0	0
19	CQ+	COL	Q+	2	0	0	0	0	0	0	0	0	0	0
20	FQ+	FOR	Q+	2	0	0	0	0	0	0	0	0	0	0
25	FQ+	FOR	Q+	2	0	0	0	0	0	0	0	0	0	0
26	CQ+	COL	Q+	2	0	0	0	0	0	0	0	0	0	0
27	FQ+	FOR	Q+	2	0	0	0	0	0	0	0	0	0	0
30	FQ+	FOR	Q+	2	0	U	0	0	0	0	0	0	0	0
2	CQ+	COL	Q+	3				:					:	
3	CQ+	COL	Q+	3	62.32	59.12	30	0	151.44	41.152	39.0386	19.8098	0	1.43004
(CQ+	COL	Q+	3	442.07	164.64	173.79	15.75	796.25	55.519	20.6769	21.8261	1.978	0.80119
8	CQ+	COL	Q+	3	146.25	56.897	54.945	8.7674	266.859	54.804	21.321	20.5896	3.2854	0.82469
11	CQ+	COL	Q+	3	251.482	50.674	43.272	12.1299	357.559	70.333	14.1723	12.1022	3.3924	0.42181
13	FQ+	FOR	Q+	3	338.903	85.44	88.522	12.8381	525.703	64.467	16.2525	16.8389	2.4421	0.55119
14	FQ+	FOR	Q+	3	287.612	57.044	33.413	14.0687	392.138	73.345	14.547	8.5207	3.5877	0.36343
17	FQ+	FOR	Q+	3	•	•	•	•	•	•	•	•	•	•
19	CQ+	COL	Q+	3		•	•	•	•	•	•	•	•	•
20	FQ+	FOR	Q+	3	04.400			0.7040			05 5074			
25	FQ+	FOR	Q+	3	24.192	19.666	8.65	2.7849	55.293	43.753	35.5671	15.6434	5.0367	1.28557
26	CQ+	COL	Q+	3 3	96.844	34.327	14.712	2.9816	148.865	65.055	23.0594	9.8831	2.0029	0.53717
27	FQ+	FOR	Q+	-	167.896	32.723	18.219	5.8081	224.646	74.738	14.5666	8.11	2.5854	0.33801
30	FQ+	FOR	Q+	3	96.733	23.013	21.926	3.5314	145.204	66.619	15.849	15.1	2.4321	0.50108
2	CQ+	COL	Q+	4	248.16	94.744	56.899	13.4129	413.215	60.056	22.9284	13.7697	3.246	0.66512
3	CQ+	COL	Q+	4	58.998	30.973	24.604	7.8548	122.43	48.189	25.2984	20.0967	6.4158	1.07516
/	CQ+	COL	Q+	4	142.42	69.31	29.24	0	240.97	59.103	28.7629	12.1343	0	0.69197
8	CQ+	COL	Q+	4	143.141	64.269	51.01	7.3421	265.762	53.861	24.183	19.1938	2.7627	0.85665
11	CQ+	COL	Q+	4	130.985	38.251	20.963	7.3405	197.539	66.308	19.3636	10.6121	3.716	0.50811
13	FQ+	FOR	Q+	4	97 001	. 21 646	12 200	4 0170	127.052	68.822	16 0306	10.4704	2 7602	0.45202
14 17	FQ+	FOR	Q+ Q+	4	87.991 35.007	21.646	13.398	4.8179	127.853	59.759	16.9306	10.4794	3.7683	0.45303 0.67339
17	FQ+	FOR	Q+	4	35.987	12.32	8.637	3.2762	60.221	59./59	20.4581	14.3427	5.4404	0.67339

Supplemental Table 4.2 Continuation.

Calf	Group	Feeding	Quercetin	Age,	Quercetin,	Isorhamnetin,	Tamarixetin,	Kaempferol,	Total flavonols,	Quercetin,	Isorhamnetin,	Tamarixetin,	Kaempferol,	Ratio
				d	nmol/L	nmol/L	nmol/L	nmol/L	nmol/L	% of total flav.	KIT:Q			
19	CQ+	COL	Q+	4	93.096	47.86	30.393	5.6078	176.956	52.609	27.0463	17.1753	3.169	0.9008
20	FQ+	FOR	Q+	4	37.224	15.659	2.205	0	55.088	67.572	28.4259	4.002	0	0.4799
25	FQ+	FOR	Q+	4	69.207	22.767	9.919	2.0801	103.973	66.563	21.8965	9.5401	2.0007	0.50234
26	CQ+	COL	Q+	4	50.863	18.959	0	0	69.822	72.846	27.1538	0	0	0.37276
30	FQ+	FOR	Q+	4	91.096	20.46	31.805	3.105	146.466	62.196	13.9689	21.7152	2.1199	0.60782
2	CQ+	COL	Q+	7	49.054	20.021	9.63	6.1586	84.864	57.803	23.5922	11.3476	7.2571	0.73001
3	CQ+	COL	Q+	7	37.108	21.098	14.833	2.1688	75.209	49.341	28.0527	19.7229	2.8837	1.02672
7	CQ+	COL	Q+	7	114.75	30.36	22.76	3.45	171.32	66.98	17.7212	13.2851	2.0138	0.49298
11	CQ+	COL	Q+	7	82.213	31.798	30.46	6.4607	150.931	54.47	21.0676	20.1815	4.2806	0.83586
13	FQ+	FOR	Q+	7	133.499	29.247	33.533	4.4738	200.752	66.499	14.5685	16.7037	2.2285	0.50377
14	FQ+	FOR	Q+	7	103.231	14.519	12.442	5.0035	135.196	76.357	10.7396	9.2027	3.7009	0.30964
17	FQ+	FOR	Q+	7	16.76	5.331	4.908	2.072	29.07	57.651	18.3365	16.8846	7.1274	0.73456
19	CQ+	COL	Q+	7	151.937	54.495	52.604	7.8448	266.88	56.931	20.4192	19.7106	2.9395	0.75652
20	FQ+	FOR	Q+	7	107.243	22.783	15.851	2.6877	148.564	72.186	15.3352	10.6695	1.8091	0.38531
25	FQ+	FOR	Q+	7	50.308	23.449	20.763	3.2455	97.765	51.458	23.9847	21.238	3.3197	0.94335
26	CQ+	COL	Q+	7	30.657	0	0	0	30.657	100	0	0	0	0
27	FQ+	FOR	Q+	7	62.657	14.244	18.096	3.1707	98.167	63.826	14.5099	18.4337	3.2299	0.56675
2	CQ+	COL	Q+	8	63.343	19.154	9.372	4.6322	96.501	65.639	19.8488	9.7117	4.8001	0.52348
3	CQ+	COL	Q+	8										
7	CQ+	COL	Q+	8										•
11	CQ+	COL	Q+	8				•				•		•
13	FQ+	FOR	Q+	8	161.704	19.12	16.33	5.0544	202.208	79.969	9.4555	8.0757	2.4996	0.25048
14	FQ+	FOR	Q+	8	89.841	10.458	6.72	3.9153	110.935	80.986	9.4272	6.0578	3.5294	0.23479
17	FQ+	FOR	Q+	8				•				•		•
19	CQ+	COL	Q+	8	74.96	16.3	35.961	5.1305	132.351	56.637	12.316	27.1706	3.8764	0.76563
20	FQ+	FOR	Q+	8	76.688	14.205	11.083	29.195	131.17	58.464	10.8292	8.4493	22.2573	0.71045
25	FQ+	FOR	Q+	8	43.227	9.308	13.496	2.47	68.5	63.105	13.5877	19.7017	3.6058	0.58466
26	CQ+	COL	Q+	8	100.191	11.644	15.68	2.0032	129.518	77.357	8.99	12.1067	1.5467	0.29271
27	FQ+	FOR	Q+	8	7.801	6.937	9.782	•	24.52	31.816	28.2917	39.8922	•	

doi:10.1371/journal.pone.0146932.s002

Supplemental Table 4.3: Complete data set of plasma concentrations of metabolites and hormones as shown in Figure 4.3 and Table 4.2.

Calf	Group	Feeding	Quercetin	Age, d	NEFA, μmol/L	Urea, mmol/L	Triglycerides, mmol/L	Lactate, mmol/L	Glucose, mmol/L	Cholesterol, mmol/L	Albumin, g/L	Protein, g/L	Cortisol, μg/L	Insulin, μg/L	Glucagon, ng/L
1	CoIQ-	COL	Q-	1	895	4.27	0.23	2.57	2.62	0.39	26.1	45.3	29.9	0.42164	74.45
4	CoIQ-	COL	Q-	1	1130	3.24	0.23	1.78	3.62	0.41	24.7	42.4	28.8	0.27239	89.945
5	CoIQ-	COL	Q-	1	1156	3	0.24	1.92	4.27	0.82	24	41.6	39.6	0.52239	71.655
10	CoIQ-	COL	Q-	1	223	4.25	0.25	7.68	6.02	0.96	24.9	44	32	1.47948	83.6
12	CoIQ-	COL	Q-	1	1393	3.14	0.36	2.96	6.07	0.59	28.4	47.3	38.4	0.92351	78.35
23	CoIQ-	COL	Q-	1	368	2.47	0.16	6.06	3.39	0.69	26.4	45.4	57.6	0.24254	97.3
28	CoIQ-	COL	Q-	1	646	2.79	0.22	5.29	3.81	0.45	26.1	45.8	31.6	0.23507	80
2	CoIQ+	COL	Q+	1	566	3.39	0.18	2.28	4.35	0.73	25.9	46.7	39.4	0.48694	57.25
3	ColQ+	COL	Q+	1	1478	2.89	0.33	2.2	2.97	0.53	25.8	44.5	44.1	0.375	66.65
7	ColQ+	COL	Q+	1	626	3.28	0.14	4.27	6.89	0.52	26.1	46.9	46.9	0.21269	104.3
8	CoIQ+	COL	Q+	1	46	6.99	0.49		7.67	0.42	27.5	47.9	23.8		187
11	CoIQ+	COL	Q+	1	500	2.6	0.16	3.02	5.08	0.36	27.8	48.8	40.2	1.73507	62.35
19	CoIQ+	COL	Q+	1	1070	3.1	0.32	2.91	2.72	0.64	26.7	47.9	43.2	0.54664	95
26	CoIQ+	COL	Q+	1	679	3.13	0.15	2.28	3.48	0.94	26.2	45	41.8	0.13993	80.85
9	ForQ-	FOR	Q-	1	321	5.96	0.45	8.11	2.63	0.59	23.7	49.3	44.7	0.52425	260.9
15	ForQ-	FOR	Q-	1	1597	3.21	0.42	1.67	5.13	0.85	27.2	50	19.8	0.41978	57.9
18	ForQ-	FOR	Q-	1	308	2.65	0.22	6.31	5	0.48	24.2	42.3	34.9	3.35261	97.45
21	ForQ-	FOR	ã-	i	382	2.84	0.14	2.9	2.72	0.84	27.3	45.9	38.9	0.32649	93.95
22	ForQ-	FOR	ã-	i	950	1.34	0.19	2.555	2.885	0.66	24.4	41.35	32.2	0.54851	127.4
24	ForQ-	FOR	ã-	i .	966	2.72	0.19	2.9	6.28	0.68	27.8	46	29.8	5.45709	127.75
29	ForQ-	FOR	ã-	i	844	3.27	0.21	1.75	3.83	0.29	25.3	44.2	39.5	0.49627	74.25
13	ForQ+	FOR	Q+	i	565	2.93	0.27	2.72	5.16	0.39	26.3	43.5	27.6	0.31343	63.3
14	ForQ+	FOR	Q+	i	448	3.85	0.18	3.25	3.54	0.58	24.2	42.3	44.6	1.13806	85.25
17	ForQ+	FOR	Q+	i	659	2.65	0.29	3.38	3.57	0.54	26.9	47.3	38.5	1.13806	121.2
20	ForQ+	FOR	Q+	i	1160	3.31	0.32	4.01	6.51	0.78	26.5	47.3	46.9	1.59328	146.65
25	ForQ+	FOR	Q+	i	514	2.57	0.11	3.35	3.85	0.85	24.6	43.4	67.6	0.20149	136.95
27	ForQ+	FOR	Q+ Q+	i	833	5.29	0.11	2.84	3.63	0.83	24.8	43.5	38.9	0.66045	88.55
30	ForQ+	FOR	Q+ Q+	i	952	2.57	0.13	2.45	2.52	0.67	28.7	47.4	35.7	1.65485	181.6
1	ColQ-	COL	Q-	2	233	4.11	0.23	1.43	5.8	0.84	22.7	65.7	11.2	0.94963	400
4	ColQ-	COL	Q- Q-	2	233 389	3.64	0.26	2.33	6.13	0.52	23.6	56.3	10	0.94963	398.8
5	ColQ-	COL	Q-	2	199		0.27		5.35		21.2	58.8	15.6	0.93843	241.35
10	ColQ-	COL	Q-	2	399	3.41 4.4	0.36	2.8 1.94	6.31	0.96	22.2	63.2	8.1	0.93643	227.6
12	ColQ-	COL	Q-	2	402	3.3	0.36	2.85	7	1.11 0.61	23.9	55	23.4	1.47575	308.5
	ColQ-		Q-	2	609	3.51			5.61	0.82	23.5		27.3	0.63806	341.5
23		COL			279		0.4	2.69				63 55.9		0.83955	341.5 400
28	ColQ-	COL	Q-	2		2.21	0.47	2.9	5.46	0.57	23.4		13.7		
2 3	ColQ+	COL	Q+	2	348	5.08	0.24	4.13	5.63	0.87	23.3	70.4	10.1	0.375	313.4
	ColQ+	COL	Q+		410	3.82	0.47	2.85	5.08	0.65	22.1	71.2	9.6	1.38246	399.85
7	ColQ+	COL	Q+	2	390	3.71	0.17	3.35	5.87	0.63	23.2	62.8	30.9	0.54291	333.15
8	ColQ+	COL	Q+	2	554	6.49	0.32	2.47	5.27	0.87	24.6	60.7	6.5	1.33396	400
11	ColQ+	COL	Q+	2	293	3.83	0.24	3.75	6.57	0.68	24.7	62.3	16.6	1.5	346.15
19	ColQ+	COL	Q+	2	174	4.05	0.23	2.37	5.43	0.73	23.7	58	14.2	0.93843	265
26	ColQ+	COL	Q+	2	410	3.09	0.16	1.96	4.82	0.87	23.4	57.1	12	0.18097	208.95
9	ForQ-	FOR	Q-	2	314	10.71	0.57	4	7.01	1.21	23	51.4	46.5	1.54104	363.65
15	ForQ-	FOR	Q-	2	535	4.37	0.2	2.43	6.85	0.92	25.2	48.2	19	1.79851	317.2
18	ForQ-	FOR	Q-	2	366	3.57	0.29	2.89	3.59	1.05	23.7	44.3	18	0.86567	323
21	ForQ-	FOR	Q-	2	370	6	0.4	3.22	4.96	1.08	26.3	47.4	32.2	0.59888	251.6
22	ForQ-	FOR	Q-	2	566	2.43	0.21	1.43	4.93	0.86	23.5	40.8	45.4	0.26679	266.25

Supplemental Table 4.3 Continuation.

Calf	Group	Feeding	Quercetin	Age,	NEFA,	Urea,	Triglycerides,	Lactate,	Glucose,	Cholesterol,	Albumin,	Protein,	Cortisol,	Insulin,	Glucagon,
	F0	FOR		d	µmol/L	mmol/L	mmol/L	mmol/L	mmol/L	mmol/L	g/L	g/L	µg/L	μg/L	ng/L
24	ForQ-	FOR	Q-	2	295	3.49	0.14	3.09	7.45	0.74	25	45.9	16.5	1.07463	304.95
29	ForQ-	FOR FOR	Q-	2 2	262 368	4.32 6.26	0.28 0.24	1.67	5.94	0.55 0.53	23 25.3	42.4	21.6	1.84515 0.58582	187.25
13	ForQ+		Q+				0.24	1.49	4.86			45.6	29.6		249.75
14 17	ForQ+ ForQ+	FOR FOR	Q+ Q+	2 2	309 345	4.77 5.29	0.∠5 0.41	2.06 2.89	7.12 5.7	0.76 0.89	23.1 25.6	41.8 47.1	27.6 14.7	1.42537 1.03545	175.9 302.2
20	ForQ+	FOR	Q+ Q+	2	345 387	3.02	0.41	5.23	6.17	0.89	23.8	47.1	33.9	1.13246	400
25	ForQ+	FOR	Q+	2	403	4.64	0.2	2.02	7.23	0.87	23.9	42.8	25.7	1.13433	248.95
25 27	ForQ+	FOR	Q+ Q+	2	403 307	3.4	0.34	3.91	7.23 5.31	0.95	23.9	42.8 44.9	25.7 28	0.40299	248.95 370.7
30	ForQ+	FOR	Q+	2	484	3.7	0.24	2.66	4.12	0.66	24.8	46.4	58.6	0.26306	370.7 374.3
1	ColQ-	COL	Q+ Q-	4	464 179	5.72	0.13	2.66 0.97	4.12	1.58	24.8	59.3	5.7	0.50933	220.8
4	ColQ-	COL	Q-	4	179	3.89	0.24	0.88	6.23	0.95	24.3	56.9	18.1	0.16604	231.05
5	ColQ-	COL	Q-	4	371	3.66	0.43	1.58	5.34	1.8	23.4	60.9	7.7	0.39925	167.55
10	ColQ-	COL	Q-	4	345	3.37	0.43	0.68	5.5 5.5	1.47	23.4	61.4	12.9	0.39925	191.55
12	ColQ-	COL	Q-	4	710	2.8	0.52	0.88	5.74	1.47	25.5	59.7	14.6	0.41045	145.95
23	ColQ-	COL	Q-	4	139	3.89	0.77	0.98	5.96	1.16	23.2	60.2	20	0.19210	227.3
23 28	ColQ-	COL	Q-	4	110	2.74	0.23	1.07	5.14	1.18	24.5	52.7	13.2	0.25933	172.7
2	ColQ+	COL	Q- Q+	4	147	4.39	0.62	0.78	5.14 5.45	1.69	23.1	65.6	2.9	0.25933	187.35
3	ColQ+	COL	Q+ Q+	4	346	4.39 7.45	0.42	0.78	5.86	1.34	21.7	64.1	10.1	0.40299	239.55
3 7	ColQ+	COL	Q+ Q+	4	408	7.45 2.84	0.42	1.07	6.26	1.65	24	64.6	10.1	0.32276	189.8
8	ColQ+	COL	Q+ Q+	4	179	4.09	0.12		5.82	1.54	25.3	62.3	5.6	0.38246	160.5
11	ColQ+	COL	Q+ Q+	4	211	2.36	0.74	1.22 1.27	5.32	1.09	24.6	62.7	14.6	0.28358	164.45
19	ColQ+	COL	Q+ Q+	4	417	3.47	0.74	1.08	4.88	1.76	26.3	63.3	14.7	0.26356	157.15
26		COL	Q+ Q+	4	177	2.62	0.23		5.42				8.8	0.18097	165.3
∠6 9	ColQ+ ForQ-	FOR	Q+ Q-	4	227	2.62 4.18	0.23	0.81 0.48	3.55	1.51 1.04	24.6 20.6	56.8 49.5	3.3	0.18097	156.9
15	ForQ-	FOR	Q-	4	130	3.41	0.36	0.48	4.91	1.04	24.9	49.5 47.6	14.4	0.78731	101.85
18	ForQ-	FOR	Q-	4	95	3.41 4.5	0.16				24.9	42.5		0.20709	112.7
21	ForQ-	FOR	Q-	4	113	4.5 3.47	0.16	1.63 0.72	4.98 4.74	0.85 1.07	22.7	42.5 41.9	11.1 14.1	0.20709	106.55
22	ForQ-	FOR	Q-	4	91	3.15	0.07	0.72	4.74	1.07	22.8	40.7	14.1	0.1903	112
24	ForQ-	FOR	Q-	4	569.5	3.15	0.26	1.075	3.63	0.985	23.85	44.9	3.5	0.09328	164
29	ForQ-	FOR	Q-	4	356	4.9	0.23	1.76	3.87	0.965	23.65	44.5	20.1	0.09328	152.6
13	ForQ+	FOR	Q- Q+	4	315	7.76	0.23	2.47	3.81	0.77	23.5	44.1	20.1	0.28358	85.05
14	ForQ+	FOR	Q+ Q+	4	73	7.76 5.9	0.18	1.23	4.66	0.58	20.8	42.3	6.6	0.41791	55.25
17	ForQ+	FOR	Q+	4	73 211	2.7	0.09	0.91	3.86	0.95	23.3	42.3 45.8	8	0.21828	170.9
20	ForQ+	FOR	Q+	4	692	2.6	0.13	1.57	3.80 4	0.83	24.6	45.6	31.1	0.09328	227.25
25	ForQ+	FOR	Q+	4	55	4.26	0.25	1.32	5.86	1.15	22.4	41.3	11.6	0.36567	136
27	ForQ+	FOR	Q+	4	55	4.20	0.25		5.60	1.15	22.4			0.30307	130
30	ForQ+	FOR	Q+	4	325	5.77	0.17	0.72	3.64	0.93	21.8	44	30	0.1847	159.6
1	ColQ-	COL	Q-	7	236	6.18	0.17	0.72	5.07	2.1	22.6	56.2	12.6	0.34515	194.45
4	ColQ-	COL	Q-	7	105	5.65	0.18	0.44	5.73	1.05	23.3	51.2	4.9	0.46642	206.65
5	ColQ-	COL	Q-	7	145	4.58	0.10	0.55	5.37	2.07	21.4	49.9	14.7	0.42724	141.55
10	ColQ-	COL	Q-	7	133	4.69	0.26	0.47	4.88	1.52	22.3	55.5	10.8	0.28545	177.9
12	ColQ-	COL	Q-	7	233	4.15	0.24	0.47	5.26	1.07	23.2	49	8.1	0.20522	124.3
23	ColQ-	COL	Q-	7	233 93	4.13	0.24	0.47	5.26 5.44	1.15	23.2	57.2	8	0.45149	143.1
28	ColQ-	COL	Q-	7	118	5.06	0.21	0.48	4.93	1.15	23.1	50.2	10.2	0.33022	150.45
2	ColQ+	COL	Q- Q+	7	117	6.89	0.23	1.62	5.32	2.26	22.4	50.2 59.9	7.7	0.33022	183.9
3	ColQ+	COL	Q+ Q+	7	162	6.62	0.23	0.32	5.32	1.56	20.6	59.9 57.3	7.7 5.2	0.47201	160
7	ColQ+	COL	Q+	7	134	5.9	0.24	0.52	6.08	1.71	23.5	58	5.2 6.1	0.375	169.2
8	ColQ+	COL	Q+ Q+	7	134	5.5	0.1	0.31	0.08	1.71	23.5	36	0.1	0.375	105.2
11	ColQ+	COL	Q+	7	141	4.31	0.34	1.17	4.99	1.48	23	53.8	7.5	0.32836	124.55
19	ColQ+	COL	Q+	7	153	4.04	0.26	0.8	5.08	1.71	23.5	52	10.9	0.52985	182.3
19	COIQ+	COL	Q+	,	103	4.04	0.∠6	0.8	5.08	1.71	∠3.5	52	10.9	0.5∠385	104.3

Supplemental Table 4.3 Continuation.

Calf	Group	Feeding	Quercetin	Age,	NEFA,	Urea,	Triglycerides,	Lactate,	Glucose,	Cholesterol,	Albumin,	Protein,	Cortisol,	Insulin,	Glucagon,
				d	μmol/L	mmol/L	mmol/L	mmol/L	mmol/L	mmol/L	g/L	g/L	μg/L	μg/L	ng/L
26	CoIQ+	COL	Q+	7	234	4.38	0.16	0.53	4.61	1.43	23.6	52.8	11	0.125	154.9
9	ForQ-	FOR	Q-	7	316	5.15	0.18	0.5	4.52	1	21	53.2	1.3	0.18284	77.95
15	ForQ-	FOR	Q-	7	179	3.32	0.15	0.53	4.75	1.45	23.1	44.4	9.6	2.21455	55.6
18	ForQ-	FOR	Q-	7	511	6.34	0.07	0.4	3.47	0.5	22.7	40.2	9.1	0.09328	89.45
21	ForQ-	FOR	Q-	7	100	2.98	0.07	0.55	4.46	0.92	23.6	40	9.3	0.15299	75.15
22	ForQ-	FOR	Q-	7	87	5.28	0.08	0.04	5.01	1.07	21.5	38.4	7.2	0.24254	81.05
24	ForQ-	FOR	Q-	7	100	3.54	0.11	0.74	4.92	1.2	23.6	42.8	10	0.33209	80.45
29	ForQ-	FOR	Q-	7	95	4.58	0.06	1.35	3.86	1.06	20.7	38.7	10.4	0.36381	111.65
13	ForQ+	FOR	Q+	7	239	3.8	0.11	0.39	4.06	0.69	21.8	41.4	11.7	0.13993	46.4
14	ForQ+	FOR	Q+	7	153	5.2	0.11	0.28	5.05	0.82	21.8	42.3	8.7	0.35634	47.6
17	ForQ+	FOR	Q+	7	290	3.3	0.1	0.45	3.95	0.98	23	43.7	12	0.09328	95.5
20	ForQ+	FOR	Q+	7	90	3.36	0.13	0.46	5.28	0.89	22	38.4	20.5	0.28731	120.4
25	ForQ+	FOR	Q+	7	80	4.14	0.03	0.5	4.82	1.2	21.6	37.3	5.4	0.22575	69.05
27	ForQ+	FOR	Q+	7	159	3.65	0.07	0.63	5.25	1.08	21.5	38.4	11.2	0.20336	54.05
30	ForQ+	FOR	Q+	7											

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Supplemental Table 4.4: Complete data set of immune and inflammatory status in blood plasma as shown in Figure 4.3.

Calf	Group	Feeding	Quercetin	Age,	lgG1,	lgG2,	IgM,	Fibrinogen,	Haptoglobin,	Serum amyloid A,
	0.10	201		d	g/L	g/L	g/L	g/L	g/L	mg/L
1	ColQ-	COL	Q-	1	0	0	37.29	3.81	0	14.888
4	CoIQ-	COL	Q-	1	0	0	30.82	0	0	13.438
5	CoIQ-	COL	Q-	1	0	8.65	27.15	0	0	1.208
10	CoIQ-	COL	Q-	1	0	11.15	25.41	2.17	0	26.579
12	CoIQ-	COL	Q-	1	0	3.04	20.08	3.5	0	33.281
23	CoIQ-	COL	Q-	1	0	9.81	48.12	2.86	0	14.13
28	CoIQ-	COL	Q-	1	0	6.53	52.32	2.22	0	20.482
2	CoIQ+	COL	Q+	1	0	4.53	13.46	2.71	0	1.499
3	CoIQ+	COL	Q+	1	0			0.81	0	16.725
7	CoIQ+	COL	Q+	1	0	8.1	72.09	0	0	0
8	CoIQ+	COL	Q+	1	0	23.71	57.07	4.52	0	29.727
11	CoIQ+	COL	Q+	1	0	0	20.25	3.75	0	9.764
19	CoIQ+	COL	Q+	1	0	19.59	342.08	2.22	0	27.851
26	ColQ+	COL	Q+	1	0	5.35	20.82	1.67	0	13.159
9	ForQ-	FOR	Q-	1			129.64	5.24	0	107.84
15	ForQ-	FOR	Q- Q-	1			27.75	3.49	0	18.943
18	ForQ-	FOR	Q-	1	0	5.32	10.54	1.95	0	36.547
21	ForQ-	FOR	Q-	1	Ö	4.19	44.36	4.35	Ö	22.804
22	ForQ-	FOR	Q-	1	0	5.01	18.69	1.46	0	17.747
24	ForQ-	FOR	Q-	1	Ö	5.41	39.22	2.86	Ö	14.659
29	ForQ-	FOR	Q-	1	Ö	5.07	42.59	2.6	Ö	23.306
13	ForQ+	FOR	Q+	i	0	4.37	16.99	2.86	ő	26.197
14	ForQ+	FOR	Q+	1	0	2.97	26.66	3.25	Ö	40.492
17	ForQ+	FOR	Q+	1	0	91.95	48.33	2.1	0	32.56
20	ForQ+	FOR	Q+	1	0	5.32	22.17	1.96	0	
		FOR		1				2.04	0	23.143
25	ForQ+		Q+	1	0	3.2 7	15.97	2.04	0	17.394
27	ForQ+	FOR	Q+				15.31	2.34		21.011
30	ForQ+	FOR	Q+	1	0.01	7.81	105.56	1.84	0	18.453
1	ColQ-	COL	Q-	2	10.46	986.05	1386.49	4.6	0.07	139.162
4	ColQ-	COL	Q-	2 2 2	7.58	1021.1	726.9	3.42	0	126.353
5	ColQ-	COL	Q- Q-	2	10.7	605.89	1151.22	2.86	0.05	200.51
10	ColQ-	COL	Q-	2	10.17	762.41	1234.9	4.88	0	168.81
12	ColQ-	COL	Q-	2 2 2	8.9	652.35	852.23	3.9	0	94.11
23	ColQ-	COL	Q-	2	10.46	573.12	1246.63	4.17	0	121.418
28	CoIQ-	COL	Q-	2	9.6	665.97	752.27	3.04	0	90.493
2	CoIQ+	COL	Q+	2 2	12.54	1381.67	1512.83	4.21	0	116.879
3	CoIQ+	COL	Q+	2	13.43	1270.24	1661.41	2.86	0	119.036
7	CoIQ+	COL	Q+	2	10.13	1316.29	1065.97	2.63	0	95.949
8	CoIQ+	COL	Q+	2	9.47	1368.68	773.73	5.53	0.03	130.268
11	CoIQ+	COL	Q+	2	10.13	906.28	1087.25	4.69	0	183.342
19	CoIQ+	COL	Q+	2	6.4	589.25	865.14	3.02	0	128.851
26	CoIQ+	COL	Q+	2	7.28	533.12	893.9	2.88	0	131.3
9	ForQ-	FOR	Q-	2			110.07	7.67	0	239.091
15	ForQ-	FOR	Q- Q-	2			39.03	4.65	0	136.486
18	ForQ-	FOR	Q-	2	0	7.59	27.21	5.12	0	146.964
21	ForQ-	FOR	Q-	2	0	8.36	59.27	6.52	0	170.803
22	ForQ-	FOR	Q-	2	0	9.52	31.99	3.04	0	93.978
24	ForQ-	FOR	Q-	2	0	8.97	49.69	4.22	0	99.581
29	ForQ-	FOR	Q-	2	0	11.88	53.25	4.22	0.09	113.654
13	ForQ+	FOR	Q+	2	0	3.9	12.46	5.53	0.01	203.424
14	ForQ+	FOR	Q+	2	0	3.41	25.95	5.5	0.03	113.708
17	ForQ+	FOR	Q+	2	0	103.22	53.13	4.05	0.02	158.502
20	ForQ+	FOR	Q+	2	0	8.95	31.21	2.32	0	152.436
25	ForQ+	FOR	Q+	2	0	5.96	26.23	4.08	0	132.006
27	ForQ+	FOR	Q+	2	0	19.38	27	5.32	0	124.418
30	ForQ+	FOR	Q+	2	0.01	9.05	89.02	3.41	0	173.474
1	CoIQ-	COL	Q-	4	8.51	638.67	922.58	5	0	91.405
4	CoIQ-	COL	Q-	4	6.98	532.44	432.24	5.13	0	56.168
5	ColQ-	COL	Q-	4	11.86	620.97	881.6	7.32	0	222.568
10	ColQ-	COL	Q-	4	9.25	540.47	789.27	5.13	ŏ	153.284
12	ColQ-	COL	Q-	4	8.6	424.71	549.25	5.23	Ö	98.394
23	ColQ-	COL	Q-	4	10.8	554.4	801.91	5.12	ŏ	44.04
28	ColQ-	COL	Q-	4	9.25	674.06	440.74	3.69	Ö	83.346
2	ColQ+	COL	Q+	4	9.47	758.35	1108.26	7.43	ő	125.193
3	ColQ+	COL	Q+	4	11.14	673	1032.83	4.42	Ö	51.286
7	ColQ+	COL	Q+	4	9.47	659.7	805.85	6.67	0	113.302
8	ColQ+	COL	Q+	4	9.47	916.82	478.51	5.58	0	61.533
	ColQ+			4		664.11	645.61	7.07	0	192.768
11		COL	Q+		9.47					
19	ColQ+	COL	Q+	4	6.98	511.52	765.53	4.65	0	112.308
26	ColQ+	COL	Q+	4	6.98	497.49	476.63	3.78	0	138.403
9	ForQ-	FOR	Q-	4			90.94	7.5	0.12	251.532
15	ForQ-	FOR	Q-	4			103.3	7.14	0	153.666
18	ForQ-	FOR	Q-	4	0	6.66	72.26	6.82	0.04	102.339
21	ForQ-	FOR	Q-	4	0	7.45	166.01	6.52	0	96.74
22	ForQ-	FOR	Q-	4	0	7.93	169.98	0.53	0	65.347
24	ForQ-	FOR	Q-	4	0	7.7	85.58	7.02	0	102.713
29	ForQ-	FOR	Q-	4	0.01	11.22	102.12	6.87	0.29	123.58
13	ForQ+	FOR	Q+	4	0	3.31	57.21	7.27	0.08	249.745

Supplemental Table 4.4 Continuation.

Calf	Group	Feeding	Quercetin	Age,	lgG1,	IgG2,	lgM,	Fibrinogen,	Haptoglobin,	Serum amyloid A,
				d	g/L	g/L	g/L	g/L	g/L	mg/L
14	ForQ+	FOR	Q+	4	0	2.19	112.34	6.67	0.07	64.798
17	ForQ+	FOR	Q+	4	0	128.55	64.52	6.34	0.03	162.574
20	ForQ+	FOR	Q+	4	0	9.02	72.88	4.54	0	68.362
25	ForQ+	FOR	Q+	4	0	5.87	96.97	5.1	0	111.713
27	ForQ+	FOR	Q+	4	0	8.09	85.65	3.12	0	134.697
30	ForQ+	FOR	Q+	4	0.01	7.87	263.18	7.08	0.14	160.24
1	CoIQ-	COL	Q-	7	7.89	630.33	555.9	3.9	0	106.728
4	CoIQ-	COL	Q-	7	7.28	550.14	297.2	3.95	0	56.409
5	CoIQ-	COL	Q-	7	9.8	841.64	426.18	0	0	104.311
10	CoIQ-	COL	Q-	7	12.93	503.05	475.9	4.28	0	85.881
12	CoIQ-	COL	Q-	7	8.8	534.17	424.38	3.02	0	105.054
23	CoIQ-	COL	Q-	7	9.47	516.66	532.58	3.96	0	82.552
28	CoIQ-	COL	Q-	7	7.84	491.83	388.82	3.57	0	69.185
2	CoIQ+	COL	Q+	7	9.47	894.82	759.6	4.62	0	130.413
3	CoIQ+	COL	Q+	7	13.98	694.06	955.99	3.25	0	17.208
7	CoIQ+	COL	Q+	7	9.15	726.69	586.46	4.05	0	40.651
8	CoIQ+	COL	Q+	7	9.47	849.34	335.79	4.5	0	166.134
11	CoIQ+	COL	Q+	7	6.69	615.4	419.04	7.5	0	98.994
19	CoIQ+	COL	Q+	7	6.4	501.25	518.34	2.13	0	115.404
26	CoIQ+	COL	Q+	7	6.98	524.29	395.64	3.62	0	135.623
9	ForQ-	FOR	Q-	7			595.76	7.32	0.51	234.857
15	ForQ-	FOR	Q-	7			129.49	4.77	0	95.298
18	ForQ-	FOR	Q-	7	0	6.7	150.86	4.39	0.15	74.682
21	ForQ-	FOR	Q-	7	0	9.24	208.17	4.13	0	84.651
22	ForQ-	FOR	Q-	7	0	7.31	137.5	3.62	0	38.79
24	ForQ-	FOR	Q-	7	0	7.38	171.53	4.03	0	40.025
29	ForQ-	FOR	Q-	7	0.01	11.33	340.5	4.31	0.01	84.758
13	ForQ+	FOR	Q+	7	0	3.07	333.9	5.36	0.05	198.715
14	ForQ+	FOR	Q+	7	0	5.13	314.56	4.39	0	89.401
17	ForQ+	FOR	Q+	7	0	219.75	150.65	4.88	0	102.721
20	ForQ+	FOR	Q+	7	Ö	8.63	170.05	2.55	0	115.235
25	ForQ+	FOR	Q+	7	Ö	7.73	186.71	3.67	Ö	56.304
27	ForQ+	FOR	Q+	7	Ö	10.59	165.77	4.89	0	33.937
30	ForQ+	FOR	Q+	7	0.01	6.89	774.94	5.53	0.16	147.578

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Supplemental Table 4.5: Complete data set of antioxidative status in blood plasma as shown in Figure 4.4.

Calf	Group	Feeding	Quercetin	Age,	FRAP,	TEAC,	TBARS,	Isoprostanes,
				ď	ASCE (µmol/L)	TE (mmol/L)	MDAE (µmol/L)	ng/L
1	CoIQ-	COL	Q-	1	134.6153846	1.681916039	0.233250176	872.8109473
4	CoIQ-	COL	Q-	1	165.3846154	1.766782194	0.421821589	3323.513168
5	CoIQ-	COL	Q-	1	113.0769231	1.761031366	0.202773465	1845.892853
10	CoIQ-	COL	Q-	1	191.547619	1.995746654	1.693583504	3432.077597
12	CoIQ-	COL	Q-	1	155.952381	2.003353121	1.096887807	1497.934204
23	CoIQ-	COL	Q-	1	153.7777778	1.567096855	0.314685315	1316.667206
28	CoIQ-	COL	Q-	1	164.6666667	1.482471879	0.498251748	2079.781082
2	CoIQ+	COL	Q+	1	106.9230769	1.870161257	0.299751167	2065.667459
3	CoIQ+	COL	Q+	1	133.8461538	1.787868564	0.393335698	2070.592066
7	CoIQ+	COL	Q+	1	151.1904762	1.658526638	0.165721649	1115.406874
8	CoIQ+	COL	Q+	1	221.9047619	1.855027023	0.453178694	1855.912236
11	CoIQ+	COL	Q+	1	126.4285714	2.074663744	0.890731728	3667.339625
19	CoIQ+	COL	Q+	1	144.444444	1.508613992	0.613879004	769.8431956
26	CoIQ+	COL	Q+	1	148.6666667	1.547190497	0.262237762	818.9533998
9	ForQ-	FOR	Q-	1	267.1428571	2.102237185	0.07109375	2352.334682
15	ForQ-	FOR	Q-	1	143.3333333	1.55579468	0.613879004	1097.712239
18	ForQ-	FOR	Q-	1	102.6666667	1.469952654	0.720640569	917.4523105
21	ForQ-	FOR	Q-	1	149.555556	1.498516495	0.5625	2462.551238
22	ForQ-	FOR	Q-	1	136.8888889	1.387163857	0.104895105	391.0270488
24	ForQ-	FOR	Q-	1	121.7777778	1.499549741	0.078671329	704.2283563
29	ForQ-	FOR	Q-	1	132.8888889	1.586838851	0.104895105	5267.394514
13	ForQ+	FOR	Q+	1	166.444444	1.549688221	0.640569395	573.8641745
14	ForQ+	FOR	Q+	1	136.444444	1.465232911	0.427046263	812.8168926
17	ForQ+	FOR	Q+	1	181.7777778	1.572874309	0.266903915	1320.77096
20	ForQ+	FOR	Q+	1	168.5	1.465563684	0.427046263	1118.56926
25	ForQ+	FOR	Q+	1	127.1111111	1.424822272	0.236013986	1937.449516
27	ForQ+	FOR	Q+	1	125.1111111	1.468917919	0.20979021	3083.518788
30	ForQ+	FOR	Q+	1	86.2222222	1.529527086	0.078671329	2385.765122

Supplemental Table 4.5 Continuation.

Calf	Group	Feeding	Quercetin	Age,	FRAP,	TEAC,	TBARS,	Isoprostanes,
				d	ASCE (µmol/L)	TE (mmol/L)	MDAE (µmol/L)	ng/L
1	CoIQ-	COL	Q-	4	154.6153846	1.992423615	0.420020275	681.7809869
4	CoIQ-	COL	Q-	4	108.7179487	2.165506277	0.391534384	1110.758263
5	CoIQ-	COL	Q-	4	102.0512821	2.196177361	0.426345819	866.6391585
10	ColQ-	COL	Q-	4	101.9047619	2.248661666	1.384797643	2145.207851
12	ColQ-	COL	Q-	4	132.6190476	2.217601928	0.756937329	1343.370792
23	ColQ-	COL	Q- Q-	4				
					118.222222	1.89554019	0.576923077	637.0641473
28	ColQ-	COL	Q-	4	126.222222	1.790207411	0.498251748	1350.548847
2	ColQ+	COL	Q+	4	137.3076923	2.324278587	0.61104331	1351.585625
3	ColQ+	COL	Q+	4	278.4615385	2.198094303	0.493036317	808.6143913
7	ColQ+	COL	Q+	4	102.1428571	2.043920942	0.595317869	607.5221159
8	CoIQ+	COL	Q+	4	133.8095238	2.094630719	0.306271478	1079.303575
11	CoIQ+	COL	Q+	4	89.28571429	2.273382682	1.36020748	1860.142827
19	CoIQ+	COL	Q+	4	135.555556	1.851817733	0.774021352	1287.525435
26	CoIQ+	COL	Q+	4	126.222222	1.815953359	0.288461538	1532.568858
9	ForQ-	FOR	Q-	4	107.6190476	2.203022867	0.354947917	1702.926659
15	ForQ-	FOR	Q-	4	96	1.862789135	0.507117438	706.0255733
18	ForQ-	FOR	Q-	4	124.222222	1.681541604	0.346975089	717.9125686
21	ForQ-	FOR	Q-	4	114	1.67239252	0.348214286	533.4365606
22	ForQ-	FOR	Q-	4	115.3333333	1.685813929	0.183566434	529.8769064
24	ForQ-	FOR	Q-	4	106	1.626439486	0.20979021	482.1558393
29	ForQ-	FOR	Q-	4	100	1.702090346	0.314685315	2301.740353
13		FOR	Q- Q+	4	144.444444			
	ForQ+					1.605298125	0.226868327	1248.038414
14	ForQ+	FOR	Q+	4	140.8888889	1.64281134	0.18683274	1011.200921
17	ForQ+	FOR	Q+	4	127.555556	1.710609697	0.18683274	730.7388027
20	ForQ+	FOR	Q+	4	204	1.614742913	0.320284698	942.8627723
25	ForQ+	FOR	Q+	4	97.33333333	1.723530667	0.20979021	321.7424991
27	ForQ+	FOR	Q+	4				
30	ForQ+	FOR	Q+	4	133.5555556	1.684214853	0.157342657	1802.367305
1	CoIQ-	COL	Q-	7	131.5384615	1.998245632	0.621432282	742.713916
4	CoIQ-	COL	Q-	7	136.9230769	2.144419908	0.597931422	1135.866266
5	CoIQ-	COL	Q-	7	98.46153846	2.144419908	0.462749112	703.2490998
10	CoIQ-	COL	Q-	7	123.8095238	2.221088225	0.854614925	1904.189486
12	CoIQ-	COL	Q-	7	124.5238095	2.173230873	0.815552425	2309.539948
23	CoIQ-	COL	Q-	7	102.8888889	1.838914714	0.340909091	598.8128756
28	CoIQ-	COL	Q-	7	131.7777778	1.814688312	0.576923077	1463.093822
2	ColQ+	COL	Q+	7	103.4615385	2.242770349	0.406468531	1155.626195
3	ColQ+	COL	Q+	7	130	2.14633685	0.616258741	703.7163919
7	ColQ+	COL	Q+	7	162.8571429	2.090827486	0.311039519	985.0262514
8	ColQ+	COL	Q+	7	102.037 1429	2.090027400	0.511059519	903.0202314
				7	400 0400470	0.407040050	0.044407047	4000 000000
11	ColQ+	COL	Q+		102.6190476	2.167842959	0.811197917	1920.390602
19	ColQ+	COL	Q+	7	119.7777778	1.800379136	0.346975089	565.4206329
26	ColQ+	COL	Q+	7	141.5555556	1.73951696	0.236013986	1562.882356
9	ForQ-	FOR	Q-	7	76.66666667	2.176400234	0.662239583	2730.653029
15	ForQ-	FOR	Q-	7	99.11111111	1.790522278	0.453736655	872.2010944
18	ForQ-	FOR	Q-	7	94.2222222	1.55631307	0.346975089	4620.767468
21	ForQ-	FOR	Q-	7	93.5555556	1.580970984	0.348214286	556.8655332
22	ForQ-	FOR	Q-	7	99.7777778	1.571634283	0.262237762	607.706398
24	ForQ-	FOR	Q-	7	102.8888889	1.720128713	0.20979021	508.053522
29	ForQ-	FOR	Q-	7	112.6666667	1.642747715	0.340909091	2413.288349
13	ForQ+	FOR	Q+	7			0.18683274	420.791816
14	ForQ+	FOR	Q+	7	91.5555556	1.740558764	0.160142349	527.6681207
17	ForQ+	FOR	Q+	7	98.88888889	1.707662508	0.320284698	713.2756069
				7	90.0000009			
20	ForQ+	FOR	Q+			1.539562447	0.37366548	900.2003889
25	ForQ+	FOR	Q+	7	94.4444444	1.604049263	0.839160839	1540.91929
27	ForQ+	FOR	Q+	7	80.88888889	1.612179518	0.314685315	2460.959659
30	ForQ+	FOR	Q+	7				

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Supplemental Table 4.6: Complete data set of hepatic mRNA expression of inflammatory and antioxidative traits as shown in Table 4.3.

								Targets of	nterest, AU						Reference	ce genes, AU	
Calf	Group	Feeding	Quercetin	CAT	GPX	SOD	CRP	IL1A	SAA2	TNF	FGA	Нр	IL1B	LRP10	POL2A	beta_actin	HPCAL1
1	ColQ-	COL	Q-	1.17143	1.06366	1.15192	0.7976	1.23514	0.62615	0.39087	3.82877	14.275	3.3734	1.11062	1.32371	1.78423	2.98448
4	CoIQ-	COL	Q-	1.09433	1.00378	0.90348	0.718	1.1661	0.53025	1.14482	1.44347	6.0121	4.12951	1.30542	1.56374	1.28497	1.50192
5	CoIQ-	COL	Q-	1.26334	1.38293	1.19406	0.89045	1.3085	0.73128	0.53663	4.34262	22.7932	4.76277	1.19393	1.85821	1.18233	3.27089
10	CoIQ-	COL	Q-	1.68193	1.07982	1.16457	0.87412	0.84428	1.06089	0.99429	1.47065	4.1538	1.53206	1.62734	1.50812	1.0688	1.56687
12	CoIQ-	COL	Q-	0.98027	0.93044	0.8111	1.04122	1.27395	1.4042	1.35465	1.71678	21.3998	2.12321	2.44541	1.07265	1.00	1.62079
23	ColQ-	COL	Q-	1.13387	1.0099	0.93922	1.09176	1.18664	1.12075	0.94523	3.57125	19.7832	2.06005	1.20449	1.28786	1.69099	3.03024
28	CoIQ-	COL	Q-														
2	ColQ+	COL	Q+	0.73922	0.64333	1.02115	0.63041	1.29465	0.86595	1.69928	1.25948	19.6812	2.20152	1.80704	1.36213	1.06567	1.01059
3	ColQ+	COL	Q+	0.44354	1.25023	1.08499	1.02144	0.45686	0.8613	0.74644	3.56168	48.5291	2.98425	1.01548	1.62021	1.59429	1.49852
7	CoIQ+	COL	Q+	0.94772	0.81753	0.92369	0.95628	0.99084	0.43953	0.74649	5.33002	24.5246	2.57569	1.60655	1.48464	1.09975	2.40735
11	CoIQ+	COL	Q+	0.76525	0.92833	1.04868	1.09882	0.80116	1.29573	1.34743	2.70274	39.7927	3.44802	1.14434	1.45122	1.57952	1.99151
19	CoIQ+	COL	Q+	0.54456	0.95776	1.12808	1.13751	0.56584	1.03359	1.47307	3.84848	36.3452	7.37029	1.04446	1.66645	1.50704	2.24306
26	CoIQ+	COL	Q+	1.20786	1.0994	1.27642	0.89764	1.27159	0.80903	0.87578	2.99053	5.9909	3.52465	1	1.33272	1.9682	2.53918
9	ForQ-	FOR	Q-	0.88953	0.93322	1.30995	1.39457	0.93068	1.51382	0.76048	6.07127	97.2782	3.19267	1.03933	1.27779	1.97514	2.13348
15	ForQ-	FOR	Q-	1.27149	0.97648	1.3739	0.90225	1.33036	1.01482	0.58024	7.42837	57.9337	3.63976	1.27616	1.2705	1.61782	3.82103
18	ForQ-	FOR	Q-	1.53709	0.97762	1.26895	1.02068	0.347	1.67499	1.43352	1.43329	13.5158	2.81554	1.33933	1.07143	1.82794	1.50244
21	ForQ-	FOR	Q-	1.152	0.97053	0.83668	1.1456	1.2084	1.00005	0.79529	4.03869	22.7161	5.04311	1.30082	1.38654	1.45432	2.3124
22	ForQ-	FOR	Q-	0.7968	0.91963	0.80882	1.1498	1.00232	0.7198	1.22524	1.72817	9.4138	2.239	1.17379	1.41489	1.57941	1.52159
24	ForQ-	FOR	Q-														
29	ForQ-	FOR	Q-	0.44006	1.0961	0.93117	1.05969	1.47852	1.28236	0.52395	1.70676	37.8081	1.72565	1.19509	1.27504	1.72141	1.14704
13	ForQ+	FOR	Q+	0.92814	1.11748	0.90468	0.94202	0.96915	1.82901	1.39742	4.00911	67.3544	5.47918	1.06309	1.67006	1.47743	2.43242
14	ForQ+	FOR	Q+	1.07636	1.2076	0.71457	0.92361	0.73109	1.15721	1.5481	1.34891	14.6176	2.11813	1.38901	1.11465	1.69421	1.40983
17	ForQ+	FOR	Q+	1.11213	0.8946	1.09038	0.93096	1.16911	1.03578	0.60566	3.4702	26.4486	2.42419	1.08941	1.4429	1.66873	2.19312
20	ForQ+	FOR	Q+	0.99311	0.98436	1.02362	0.9965	2.08857	1.37232	1.62735	1.80067	17.9123	3.68457	1.2915	1.27819	1.58899	1.68524
25	ForQ+	FOR	Q+	0.92992	0.88046	0.90038	1.3269	1.49731	0.6904	1.06856	1.46064	11.2364	1.4396	1.42924	1.27971	1.43415	1.75185
27	ForQ+	FOR	Q+	1.42541	1.21503	0.75697	1.21643	0.61317	0.65656	2.66255	1.00	1.00	1.00	1.43461	1.59914	1.14338	1.98326

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5. GENERAL DISCUSSION

For newborn farm animals, the energy supply during the first stage of life significantly determines developmental and maturational processes and thus, affects health and productivity of the animal. In dairy calves, the external energy supply with colostrum and milk needs to be complemented by endogenous glucose production to fully meet the demands of the organism [1], and colostrum feeding is essential to promote gastrointestinal development and advance nutrient uptake [2]. By adding flavonoids to the neonate's diet, productivity may be improved as flavonoids possess antioxidative, anti-inflammatory, and antimicrobial properties that may promote health [3, 4]. However, as flavonoids also interact with glucose metabolism [5], it is of major importance to exclude potential inhibitory effects on the development of the neonate's glucose metabolism before considering the use of flavonoids as health-promoting feed additives in calf rearing.

During the last decade, various studies have been conducted to evaluate the health-promoting properties of quercetin, one of the main flavonoids [6]. Most of these studies focused on quercetin effects either *in vitro* or in experiments with laboratory rodents, where the great antioxidative potential of the compound was highlighted [7, 8]. This antioxidative capacity indeed was identified to be causative for the health-promoting properties in model animals of metabolic disorders, e.g. diabetic mice or obese rats, where quercetin treatment lowers elevated plasma glucose concentrations to normal levels, thus enhancing the metabolic status of the animal [9, 10]. While this effect is intended to ameliorate chronic or degenerative disorders, it would be highly undesirable in newborn calves: here, a quercetin-induced decrease of carbohydrate digestion and inhibition of hepatic glucose production would cause a detrimental energy deficiency, as glucose is a main energy source during the first days of life. As a consequence, not only growth in general, but also immune function and overall health would be diminished [11].

For the above-mentioned reasons, the main objective of this thesis was to investigate the effects of an oral quercetin supplementation on the glucose metabolism in newborn calves during the first week of life, according to the initial colostrum supply (**Chapter 3**). Calves were fed quercetin aglycone from d 2 until d 8 of life, the daily dose of 50 mg/kg BW resulting from data obtained in companion studies in calves [12, 13]. After an initial phase of colostrum or non-colostrum feeding, intestinal monosaccharide absorption and splanchnic glucose extraction, as well as the hepatic expression of genes related with glucose metabolism were investigated. As, based on the data obtained in this study, quercetin-feeding does not affect the neonates' glucose metabolism, the health-promoting potential of quercetin in newborn calves was further explored. For this purpose, blood plasma and liver biopsy samples obtained during the trial were analyzed for parameters of oxidative stress and immune

function, and performance and health parameters of the calves were evaluated in the second study of this thesis (**Chapter 4**).

In both studies, although the impact of an initial colostrum deprivation was significant and persistent during the first week of life, the little effects caused by quercetin treatment on some of the parameters regarding glucose metabolism and calf health seemed to be random and inconsistent. As the quercetin supplementation was increased five-fold compared to previous studies [12, 13] and quercetin-supplemented calves showed increased plasma flavonol concentrations, the absence of quercetin effects was unlikely to be caused by inaccurate application or insufficient absorption of the compound. In humans, about 99% of plasma quercetin is bound to proteins, mainly albumin [14], and *in vitro* experiments revealed that the covalent attachment of quercetin to bovine serum albumin reduces its total antioxidant activity [15]. Even after proteolysis of the quercetin-albumin complex, the free flavonol does not fully regain its antioxidative activity [15, 16]. If excessive binding between quercetin and albumin also occurred in the plasma of the calves in the present study, quercetin might have lost its biological activity, which would explain the lack of effects on plasma antioxidative status.

As mentioned in Chapter 2, flavonols rarely occur as aglycones, but mainly in conjugated form in plasma of pigs and humans [17, 18]. In the present studies, plasma flavonol concentrations were determined after enzymatic treatment of the plasma [19] hence the fraction of glucuronidated and sulfated conjugates to total flavonols in plasma is unknown. This is an important aspect, given that the biological effects of quercetin vary with its derivatives [20, 21]. *In vitro* experiments reveal that the antioxidative and anti-inflammatory properties of conjugated quercetin are reduced compared to the non-conjugated parent compound [18, 22]. In adult cattle, approximately 15% of total flavonols in plasma are unconjugated, whereas the predominant fraction appears as glucuronidated or sulfated derivatives [19]. If quercetin metabolism in the newborn calf is comparable to that in cows, this might be an explanation for the lack of quercetin effects on plasma metabolites and hormones analyzed herein.

Referring to the main objective of the present work, it was shown here that quercetin does not impair glucose metabolism when fed to newborn calves during the first week of life in a daily dosage of 50 mg/kg BW (Chapter 3). Contrariwise, data indicate that quercetin feeding might accelerate intestinal glucose absorption on d 7 of life, which contradicts previous studies where quercetin was shown to inhibit carbohydrate digestion and intestinal glucose transport [23, 24]. In milk-fed calves, the most important intestinal disaccharidase is lactase, whose activity seems to be unaffected by quercetin [25]. Additionally, the extensive flavonol metabolism inside the enterocytes and the subsequent luminal excretion of flavonol derivatives [26] is not considered in studies on quercetin's inhibitory potential on specific glucose transporters; thus, it is possible that intestinal glucose transport inhibition is of minor

importance after quercetin feeding *in vivo*. The results of the present study could be insofar beneficial to newborn calves as the improved intestinal monosaccharide absorption might enhance the overall energy supply. Thus, the glucose excess could be used as a fuel to cover the elevated energy demand in terms of infection and accruing immune defense [27]. Unfortunately, it could not be shown in the second study (Chapter 4) that quercetin supplementation alone is effectual to improve the newborn's health status. Although a specific immune challenge was not conducted, it is assumed from general health and performance parameters that quercetin feeding did not favor supplemented calves to better resist or more effectively fight infections. This conclusion is further supported by plasma concentrations of immunoglobulins and acute phase proteins, which were similar between quercetin-treated and untreated animals. Rather, it was shown that an adequate colostrum provision is essential for the newborn to stand against the new environment and support the immune system. Data obtained here indicate that negligence concerning the initial colostrum supply cannot be compensated by adding plant bioactives such as quercetin to the ensuing diet.

In growing pigs, it has been shown that after feeding a single quercetin dose, the flavonol concentration in the liver exceeds the plasma flavonol concentration, although this difference diminishes when quercetin is fed for a long-term period [28]. Further, it has been pointed out that in contrast to flavonols in plasma, quercetin and its metabolites isorhamnetin and tamarixetin appear to approximately 90% as unconjugated aglycones in the liver. If this is transferable to newborn calves, it would be expected that, first, effects in the liver might emerge earlier than in plasma, and second, that findings gained in cell culture experiments with aglycones could be confirmed in liver tissue of calves fed quercetin. To follow up this theory, PCR analyses were performed to assess the expression of different genes involved in glucose metabolism, immune function or antioxidative defense in the course of the studies that underlie the present thesis. In Chapter 3, the focus was put on key enzymes of glucose metabolism. As it has been shown in rat liver, it was anticipated that guercetin may stimulate glycogenolysis on the one hand [29], but on the other hand, inhibition of GNG was apprehended [30]. Thus it was tested to what extent key enzymes of both pathways are influenced by quercetin treatment. Interpreting the mRNA abundances of PYGL, coding for a key enzyme of glycogen breakdown, combined with hepatic glycogen concentrations that were also measured in the first study of this work, quercetin feeding seems to not affect hepatic glycogen metabolism in newborn calves. It is assumed that, because the neonate's energy supply is already delicate, glycogenolysis is physiologically exceedingly active. Therefore, glycogen stores are depleted shortly after birth [1, 31] which was confirmed by very low hepatic glycogen concentrations in all calves involved in this study. At the same time, GNG is upregulated, which is crucial to cover the energy demand during the first stage of life [32]. Because of the high priority of endogenous glucose production, it would be

reasonable if hepatic glucose metabolism in the newborn is autonomous and insensible to foreign impacts such as quercetin feeding. This conclusion is supported by another study in newborn calves which shows that endogenous glucose production is even independent of colostrum feeding [33], which is yet crucial for other maturational processes.

As mentioned above, quercetin not only interacts with glucose metabolism but also possesses antiinflammatory and antioxidative properties. For that reason, gene expression of proinflammatory cytokines, acute phase proteins and antioxidative enzymes were analyzed in the second study (Chapter 4) but again, effects evoking from quercetin treatment were scarce. Solely the mRNA abundance of TNF was increased in quercetin-fed calves, which contradicts studies *in vitro* and *in vivo* demonstrating that quercetin downregulates this cytokine. However, those experiments were conducted after inducing a high TNF expression, and quercetin treatment approximated $TNF\alpha$ levels back to the range of unstimulated, non-inflamed cells [34-36]. Therefore, it is assumed that the calves investigated in this study did not fight a general infection, thus hepatic TNF was not elevated and quercetin was needless to antagonize inflammatory gene expression.

A general influence of quercetin on the hepatic targets investigated in both studies, however, cannot be excluded because due to the limited amount of liver tissue, translation products and enzyme activities were disregarded. Furthermore, the flavonol concentration and the pattern of conjugates and derivatives in the liver was not analyzed, thus it could not be proven whether flavonol distribution patterns and subsequent effects obtained in growing pigs are actually deducible to newborn calves.

Effects resulting from treatment with flavonoids are generally difficult to summarize, as they are very variable, depending on the bioavailability of the compound. In this context, different doses of the same substance can cause adverse effects, and flavonoids known for their antioxidative properties can also act as prooxidants when supplemented in doses that exceed the physiologically achievable range [37]. The duration of flavonoid treatment influences its biological properties, too: Reviewing results from different studies, it can be concluded that acute and chronic treatment with polyphenols do not produce parallel results and that different compounds belonging to the same subclass can have opposite effects [38]. Finally, the physiological condition of the biological system seems to play an important role, because the actions of polyphenols on intestinal epithelium largely differ between inflamed and non-inflamed cells [39]. Concerning the glucose metabolism, Shao and colleagues have shown that the oral treatment with quercetin impairs glucose tolerance and attenuates insulin-stimulated glucose uptake in healthy mice. In mice with inflammation-induced insulin resistance, however, quercetin reversed glucose intolerance and improved insulin sensitivity, hence beneficially modulated glucose homeostasis [40]. There are several studies demonstrating that although quercetin has ameliorative effects on metabolic conditions in animal models of chronic inflammation, e.g. diabetes type II, it is of little value in healthy animals or humans [41-43]. A possible explanation for this is that the generation of the biologically active aglycone from its inactive glucuronidated/sulfated derivatives exceedingly happens at sites of inflammation [44]. Hence, it is assumed that the metabolic stress in newborn calves that occurs with the adaptation to extrauterine life is a necessary and physiological process incomparable to deranged metabolic conditions evoking from chronic inflammation; therefore, data obtained in both studies underlying this thesis indicate that the metabolism of newborn calves cannot be altered substantially by quercetin feeding.

In contrast to quercetin treatment, the systemic effects of colostrum deprivation were distinct and consistent throughout the first week of life and agree with the results described in the literature [31]. Whether quercetin might improve the local defense against pathogens in the gastrointestinal tract cannot be concluded from the data gained herein. Although quercetin did not affect the fecal score or general health status of the animals, it must be kept in mind that the calves were probably stressed because of intensive sample taking. Stress is known to accelerate gastrointestinal motility [45] and reduce immune function, which could mask local effects of quercetin. To make a reliable statement on quercetin effects inside the gastrointestinal tract, the characterization of its microbiota under field conditions could expose whether quercetin is able to stabilize the calf's microflora, which has already been demonstrated for flavonoids of other subclasses in calves [46] and for quercetin in laying hens [47]. Furthermore, it would be of value to analyze tissue samples with particular focus on the gut associated lymphoid tissue, and to conduct special tests to evaluate its immune function.

In conclusion, the studies underlying this thesis demonstrated that quercetin feeding does not impair the glucose metabolism of newborn calves, but that it enhances the gastrointestinal absorption of labelled glucose. However, this effect alone seems to be insufficient to improve the overall immune function. Whether adding quercetin to the diet would be more advantageous in weakish calves if at the same time the glucose (as lactose) content of the meal is increased, needs to be probed in further studies.

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SUMMARY

Effects of the Flavonol Quercetin on Glucose Metabolism and Health Status in Neonatal Calves

With birth, calves are confronted with a new, extrauterine environment and have to meet their energy demand autonomously. For this reason, it is essential that not only overall energy intake, but also endogenous glucose production work effectively, because glucose is one of the main energy sources during the first stage of life. With the intake of first colostrum calves ingest not only nutrients, but also a plethora of bioactive factors which support the maturation and development of metabolic processes. Colostrum is further essential to support the naïve immune system of newborn calves during the first weeks of life and to mediate passive immunity. Unfortunately, colostrum supply is often inadequate. To support the health status, flavonoids could play an important role in the upbringing of newborn calves. These are secondary plant metabolites with numerous attributed health-promoting properties. One of the most important flavonoids is quercetin, which is proven to possess antioxidative, antiinflammatory and antimicrobial capacity and to modulate the intestinal microflora. However, quercetin also interacts with glucose metabolism by inhibiting intestinal carbohydrate absorption and reducing plasma glucose concentrations. For newborn calves, such an effect would be crucial. Therefore, it was the main objective of the present work to investigate the effects of an oral quercetin supplementation on the glucose metabolism of newborn calves, because this knowledge would be the prerequisite to consider the usage of quercetin as a health promoting feed additive in calf rearing.

In the first part of the study, 28 newborn male Holstein Friesian calves were assigned to two dietary groups and fed colostrum or a milk-based formula with same macronutrient composition, but without bioactive factors, during the first two days of life. On d 2 of life, groups were subdivided into control and treatment groups, the latter receiving quercetin aglycone with meals during the first week of life. On d 3, intestinal xylose absorption was probed, on d 7, a tracer study was conducted to investigate the first pass uptake of glucose, and on d 8, a liver biopsy sample was taken and analyzed via PCR. The postabsorptive recovery rate of orally administered xylose and ¹³C₆-labelled glucose in plasma was higher in calves that initially received colostrum, indicating a better intestinal absorption capacity and a lower splanchnic glucose extraction when compared to colostrum-deprived calves. Irrespective of quercetin supplementation, the mRNA abundance of hepatic mitochondrial phosphoenolpyruvate carboxykinase and several plasma metabolites were also reduced in formula-fed calves, pointing to a delayed maturation of metabolic pathways after colostrum deprivation. Although quercetin-fed calves initially showed higher peak concentrations of ¹³C₆-glucose in plasma on d 7, this effect did not last for the following hours.

Based on the finding that an oral quercetin supplementation does not seem to disadvantage

the glucose metabolism in newborn calves, the health parameters recorded during the trial were evaluated in the second part of this work. Additionally, plasma samples of the calves were analyzed for concentrations of immunoglobulins, acute-phase proteins, as well as parameters of the antioxidative system. Furthermore, the expression of genes for some proinflammatory cytokines, antioxidative enzymes and acute-phase proteins was analyzed in cDNA samples generated from the liver biopsy tissue. Data indicate that an adequate initial colostrum supply supports neonatal health and prepares the calves very well to cope with the new environment. Colostrum-fed calves showed reduced signs of inflammation and were more vital than calves initially fed formula, further, the metabolic status was improved. In contrast to this, the quercetin supplementation did not induce any detectable health-promoting effects, a finding that underlines that results proven *in vitro* are not always transferable to the organism as a whole.

In summary, it can be concluded from the present work that an oral quercetin supplementation does not impair the glucose metabolism of newborn dairy calves. However, health-promoting properties cannot be reproduced *in vivo*, hence quercetin feeding cannot compensate for an inadequate initial colostrum supply.

ZUSAMMENFASSUNG

Einfluss des Flavonols Quercetin auf Glucosestoffwechsel und Gesundheitsstatus beim neugeborenen Kalb

Mit der Geburt werden Kälber mit einer neuen, extrauterinen Umgebung konfrontiert und müssen ihren Energiebedarf selbstständig decken. Dafür ist es essentiell, dass nicht nur ausreichend Nährstoffe aufgenommen werden, sondern auch die endogene Glucoseproduktion effizient funktioniert, da Glucose in der ersten Lebensphase als eine Hauptenergiequelle dient. Mit der Aufnahme von Kolostrum werden neben Nährstoffen auch zahlreiche bioaktive Substanzen zugeführt, die die Reifung und Entwicklung von Stoffwechselprozessen fördern. Kolostrum ist außerdem essentiell, um das naive Immunsystem des Kalbes während der ersten Lebenswochen zu unterstützen und passive Immunität zu vermitteln. Leider ist die Kolostrumversorgung oft suboptimal. Um die Gesundheit zu fördern, könnte dem Einsatz von Flavonoiden hier eine wichtige Rolle zukommen. Es handelt sich dabei um sekundäre Pflanzenstoffe, denen zahlreiche gesundheitsfördernde Eigenschaften zugeschrieben werden. Einer der Hauptvertreter ist Quercetin, welches nachweislich antioxidativ, antiinflammatorisch und antimikrobiell wirken und die Darmflora modulieren kann. Allerdings interagiert es auch mit dem Glucosestoffwechsel, wodurch die intestinale Aufnahme sowie der Blutglucosespiegel gesenkt werden. Für das Kalb wäre dies fatal. Somit war es das Hauptziel dieser Arbeit, die Wirkung einer oralen Quercetinsupplementierung auf den Glucosestoffwechsel neugeborener Kälber zu untersuchen, da die Kenntnis darüber die Voraussetzung ist, um den Einsatz von Quercetin als gesundheitsfördernden Futterzusatzstoff in der Kälberaufzucht überhaupt in Betracht ziehen zu können.

Zu diesem Zweck wurden 28 neugeborene männliche Kälber der Rasse Deutsche Holstein in zwei Fütterungsgruppen unterteilt und während der ersten beiden Lebenstage entweder mit Kolostrum oder einer Formula mit gleicher Makronährstoffzusammensetzung, jedoch ohne bioaktive Substanzen, gefüttert. Am 2. Lebenstag erfolgte eine Unterteilung in Kontrollund Behandlungsgruppen, wobei letztere während der ersten Lebenswoche Quercetin mit der Fütterung erhielten. Am 3. Lebenstag wurde eine Xyloseabsorptionsstudie durchgeführt, am 7. Lebenstag erfolgte eine Tracerstudie zum first pass uptake von Glucose, und am 8. Tag wurde Lebergewebe mittels Biopsie entnommen und mit Hilfe von PCR analysiert. Die postabsorptive Wiederfindung von oral verabreichter Xylose und $^{13}C_6$ -markierter Glucose im Plasma war höher in initial Kolostrum-gefütterten Kälbern, was im Vergleich zu Formulagefütterten Tieren auf eine bessere Absorptionskapazität sowie eine geringere Glucoseextraktion im Splanchnikusgewebe schließen lässt. Ferner waren die mRNA Abundanz der hepatischen mitochondrialen Phosphoenolpyruvatcarboxykinase sowie verschiedene Plasmametaboliten in den Formulagruppen unabhängig von der Quercetin-

supplementierung reduziert, was auf eine verzögerte Reifung des Stoffwechsels nach Kolostrumentzug hindeutet. Obwohl die Quercetinsupplementierung am 7. Lebenstag den intestinalen Glucoseabsorptionspeak steigern konnte, verlor sich dieser Effekt im Verlauf der darauffolgenden Stunden.

Basierend darauf, dass eine orale Quercetinsupplementierung den Glucosestoffwechsel bei neugeborenen Kälbern nicht nachteilig beeinflusst, wurden im zweiten Teil der Arbeit die während der Studie aufgenommenen Gesundheitsparameter genauer ausgewertet. Weiterhin wurden Plasmaproben der Tiere auf Konzentrationen von Immunglobulinen, Akute-Phase-Proteinen sowie auf Parameter des antioxidativen Systems untersucht und die Genexpression verschiedener proinflammatorischer Zytokine, antioxidativer Enzyme und akuter-Phase-Proteine in den zuvor angelegten cDNA Umschreibungen analysiert. Die Ergebnisse zeigten, dass die adäquate Versorgung mit Kolostrum während der ersten Lebenstage die Kälber gut für die erste Lebensphase ausstattet und die Kälbergesundheit fördert. Die Tiere zeigten verminderte Anzeichen von Inflammation und waren in den ersten Lebenstagen vitaler als Kälber der Formulagruppe, ferner war der metabolische Status verbessert. Die Quercetinsupplementierung hingegen brachte keine gesundheitsfördernden Effekte, was beweist, dass *in vitro* nachgewiesene Effekte nicht auf den Gesamtorganismus übertragbar sind.

Zusammenfassend lässt sich aus dieser Arbeit ableiten, dass die Fütterung von Quercetin Aglykon beim neugeborenen Kalb zwar keine nachteiligen Effekte auf den Glucosestoffwechsel hat. Dennoch sind die gesundheitsfördernden Eigenschaften nicht *in vivo* reproduzierbar, sodass der Einsatz von Quercetin als Futterzusatz keinesfalls eine unzureichende Kolostrumversorgung kompensieren kann.

APPENDIX

Table A1: Correlation (Pearson coefficients) between flavonol concentration and selected parameters in plasma of calves fed quercetin twice daily from d 2-8 of life.

-		Colostrum (CQ+)		Formula	Formula (FQ+)		es (Q+)
Parameter	Day	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value
Metabolites							
Glucose	4	0.11	0.83	0.12	0.87	0.44	0.16
	7	0.34	0.54	0.14	0.81	0.25	0.44
	∑ (4,7)	0.33	0.28	0.16	0.64	0.39	0.06
Lactate	4	0.10	0.84	-0.48	0.45	-0.25	0.54
	7	0.06	0.92	-0.36	0.51	0.04	0.90
	∑ (4,7)	0.16	0.62	-0.36	0.29	0.07	0.76
Albumin	4	-0.08	0.87	-0.88	0.05	-0.14	0.68
	7	0.39	0.48	-0.58	0.25	0.17	0.60
	∑ (4,7)	0.26	0.39	-0.64	0.03	0.23	0.29
Total protein	4	0.69	0.09	-0.63	0.29	0.64	0.02
	7	-0.33	0.56	-0.18	0.75	-0.01	0.99
	∑ (4,7)	0.47	0.11	-0.37	0.27	0.45	0.02
NEFA	4	-0.30	0.53	-0.49	0.45	-0.22	0.51
	7	-0.46	0.39	-0.24	0.68	-0.29	0.36
	∑ (4,7)	-0.01	0.97	-0.35	0.30	-0.11	0.61
Urea	4	-0.02	0.97	0.98	<0.01	0.06	0.85
	7	-0.41	0.45	0.25	0.66	-0.13	0.70
	∑ (4,7)	-0.30	0.32	0.51	0.11	-0.05	0.81
TAG	4	0.36	0.45	-0.61	0.32	0.52	0.08
	7	0.21	0.71	0.38	0.49	0.23	0.48
	∑ (4,7)	0.47	0.11	-0.23	0.50	0.50	0.01
Cholesterol	4	0.35	0.46	-0.10	0.89	0.60	0.04
	7	0.05	0.93	-0.68	0.16	-0.01	0.98
	∑ (4,7)	0.04	0.91	-0.43	0.19	0.28	0.18
Hormones							
Cortisol	4	-0.59	0.17	0.01	0.99	-0.42	0.18
	7	0.17	0.76	0.23	0.69	0.12	0.72
	∑ (4,7)	-0.26	0.41	0.01	0.98	-0.22	0.30
Insulin	4	0.35	0.46	0.48	0.46	0.37	0.35
	7	0.72	0.12	0.33	0.55	0.56	0.06
	∑ (4,7)	0.34	0.27	0.33	0.34	0.41	0.05
Glucagon	4	-0.08	0.88	-0.65	0.28	0.04	0.90
	7	0.23	0.69	-0.32	0.57	0.06	0.85
	∑ (4,7)	0.16	0.60	-0.45	0.17	0.15	0.48

Table A1 Continuation.

		Colostrum (CQ+)		Formula	Formula (FQ+)		All calves (Q+)	
Parameter	Day	Pearson r	P-value	Pearson r	P-value	Pearson r	<i>P</i> -value	
Antioxidativ	e status							
FRAP	4	-0.28	0.57	-0.45	0.50	-0.23	0.49	
	7	-0.06	0.92	-0.50	0.58	0.14	0.71	
	∑ (4,7)	-0.13	0.68	-0.20	0.63	-0.04	0.88	
TEAC	4	0.62	0.15	0.14	0.85	0.74	<0.01	
	7	-0.15	0.80	-0.40	0.55	0.06	0.86	
	∑ (4,7)	0.33	0.27	-0.22	0.56	0.49	0.02	
TBARS	4	0.09	0.85	-0.70	0.22	0.39	0.21	
	7	0.02	0.97	-0.32	0.57	-0.10	0.76	
	∑ (4,7)	0.19	0.55	-0.22	0.52	0.25	0.23	
Isoprostanes	4	-0.06	0.89	0.55	0.38	0.21	0.52	
	7	-0.40	0.46	-0.33	0.56	-0.33	0.30	
	∑ (4,7)	-0.15	0.62	-0.06	0.87	-0.04	0.85	
Immunoglol	oulins							
lgG1	4	0.31	0.53	0.66	0.26	0.61	0.04	
	7	-0.41	0.46			-0.05	0.89	
	∑ (4,7)	-0.04	0.90	0.25	0.47	0.34	0.10	
lgG2	4	0.63	0.13	-0.55	0.38	0.68	0.01	
	7	-0.31	0.58	-0.77	0.08	-0.06	0.85	
	∑ (4,7)	0.25	0.41	-0.68	0.02	0.38	0.07	
lgM	4	0.44	0.34	0.81	0.11	0.68	0.01	
	7	-0.24	0.67	0.74	0.10	0.05	0.89	
	∑ (4,7)	0.31	0.31	0.73	0.01	0.50	0.01	
Acute phase	protein	s						
SAA	4	0.02	0.98	0.12	0.87	0.03	0.94	
	7	-0.04	0.95	0.63	0.20	0.24	0.47	
	∑ (4,7)	0.10	0.75	0.42	0.21	0.16	0.47	
Haptoglobin	4	·	·	0.79	0.13	-0.17	0.61	
	7	·	•	0.70	0.14	0.35	0.47	
	∑ (4,7)			0.43	0.19	-0.01	0.97	
Fibrinogen	4	0.81	0.02	0.67	0.25	0.56	0.06	
	7	-0.17	0.77	-0.05	0.93	-0.14	0.68	
	∑ (4,7)	0.48	0.10	0.02	0.95	0.33	0.12	

Table A2: Correlation (Pearson coefficients) between flavonol concentration and selected metabolites and hormones on d7 in plasma in calves fed quercetin twice daily from d 2-8 of life.

		Colostrum (CQ+)		Formula	Formula (FQ+)		All calves (Q+)	
Parameter	Time	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value	
Metabolites								
Glucose	0 h	0.34	0.54	0.14	0.81	0.25	0.44	
	1 h	-0.37	0.50	0.72	0.11	0.06	0.85	
	2 h	-0.68	0.24	-0.80	0.06	-0.42	0.21	
	3 h	-0.15	0.79	-0.25	0.66	-0.11	0.74	
	4 h	-0.94	0.01	-0.39	0.47	-0.71	0.01	
	5 h	-0.31	0.65	-0.09	0.88	-0.53	0.09	
	6 h	0.73	0.19	0.53	0.30	-0.14	0.68	
	8 h	-0.01	0.98	0.35	0.53	-0.09	0.79	
	10 h	0.90	0.04	0.56	0.27	-0.02	0.95	
	All	-0.04	0.78	0.08	0.58	-0.07	0.46	
Lactate	0 h	0.06	0.92	-0.36	0.51	0.04	0.90	
	1 h	0.25	0.66	0.69	0.14	0.46	0.13	
	2 h	-0.63	0.30	-0.68	0.16	-0.41	0.21	
	3 h	0.33	0.55	-0.9	0.01	0.23	0.49	
	4 h	0.30	0.66	-0.66	0.17	-0.15	0.67	
	5 h	0.57	0.52	-0.34	0.54	-0.11	0.78	
	6 h	-0.62	0.47	-0.37	0.50	-0.30	0.42	
	8 h	-0.08	0.89	-0.33	0.56	-0.10	0.76	
	10 h	0.94	0.01	-0.43	0.43	-0.01	0.98	
	All	0.07	0.63	-0.13	0.34	-0.04	0.67	
Urea	0 h	-0.41	0.45	0.25	0.66	-0.13	0.70	
	1 h	-0.66	0.17	-0.14	0.81	-0.19	0.56	
	2 h	-0.78	0.13	0.34	0.54	-0.07	0.85	
	3 h	-0.69	0.14	0.84	0.04	-0.11	0.75	
	4 h	-0.59	0.33	0.33	0.55	-0.43	0.19	
	5 h	-0.59	0.34	0.15	0.79	-0.49	0.13	
	6 h	0.51	0.43	0.35	0.52	-0.28	0.41	
	8 h	-0.39	0.47	-0.44	0.41	-0.42	0.19	
	10 h	0.50	0.43	0.19	0.74	-0.31	0.37	
	All	-0.39	0.01	0.29	0.03	-0.20	0.04	

Table A2 Continuation.

		Colostrum (CQ+)		Formula	Formula (FQ+)		All calves (Q+)	
Parameter	Time	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value	
NEFA	0 h	-0.46	0.39	-0.24	0.68	-0.29	0.36	
	1 h	-0.39	0.48	-0.73	0.11	-0.53	0.08	
	2 h	0.21	0.77	0.71	0.13	0.47	0.15	
	3 h	0.64	0.19	0.16	0.78	0.43	0.17	
	4 h	0.30	0.66	0.14	0.81	0.17	0.63	
	5 h	0.80	0.12	0.40	0.46	0.47	0.15	
	6 h	0.45	0.49	0.14	0.81	0.11	0.76	
	8 h	0.38	0.49	0.35	0.52	0.29	0.38	
	10 h	-0.21	0.76	-0.68	0.15	-0.55	0.08	
	All	-0.07	0.63	-0.28	0.04	-0.17	0.08	
Hormones								
Insulin	0 h	0.72	0.12	0.33	0.55	0.56	0.06	
	1 h	-0.29	0.61	0.77	0.08	0.03	0.92	
	2 h	-0.18	0.79	0.34	0.54	0.04	0.92	
	3 h	0.46	0.39	0.76	0.08	0.53	0.08	
	4 h	<0.01	1.00	0.05	0.93	0.15	0.67	
	5 h	-0.43	0.51	-0.45	0.41	-0.56	0.08	
	6 h	0.11	0.88	0.74	0.10	-0.13	0.72	
	8 h	-0.18	0.75	-0.21	0.72	-0.25	0.44	
	10 h	0.84	0.09	0.18	0.76	0.18	0.60	
	All	0.10	0.50	0.13	0.34	0.07	0.51	
Glucagon	0 h	0.23	0.69	-0.32	0.57	0.06	0.85	
	1 h	0.29	0.61	0.25	0.66	0.38	0.23	
	2 h	-0.92	0.02	-0.15	0.79	-0.17	0.63	
	3 h	0.89	0.02	-0.27	0.63	0.31	0.33	
	4 h	-0.44	0.51	-0.16	0.78	-0.57	0.07	
	5 h	-0.32	0.64	0.26	0.64	-0.40	0.23	
	6 h	0.68	0.24	0.05	0.93	-0.41	0.21	
	8 h	-0.64	0.18	0.59	0.25	-0.23	0.49	
	10 h	0.11	0.88	0.84	0.04	-0.37	0.28	
	All	0.05	0.71	0.07	0.63	-0.10	0.32	

Table A2 Continuation.

		Colostrum (CQ+)		Formula	Formula (FQ+)		All calves (Q+)	
Parameter	Time	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value	Pearson r	<i>P</i> -value	
Noradrenaline	0 h	-0.01	0.99	-0.12	0.83	<0.01	0.99	
	1 h	-0.10	0.87	-0.63	0.19	0.08	0.81	
	2 h	-0.21	0.77	-0.77	0.08	-0.20	0.57	
	3 h	-0.17	0.76	-0.02	0.97	<0.01	0.99	
	4 h	0.88	0.05	-0.12	0.84	0.13	0.71	
	5 h	0.28	0.68	-0.32	0.56	-0.27	0.43	
	6 h	0.44	0.51	-0.23	0.68	-0.10	0.78	
	8 h	-0.12	0.83	-0.77	0.08	-0.25	0.44	
	10 h	0.85	0.08	-0.39	0.47	0.04	0.91	
	All	-0.11	0.47	-0.23	0.09	-0.19	0.05	
Adrenaline	0 h	-0.64	0.19	-0.23	0.69	-0.30	0.35	
	1 h	-0.25	0.65	-0.58	0.25	-0.15	0.65	
	2 h	-0.03	0.97	-0.56	0.28	-0.26	0.45	
	3 h	-0.63	0.20	-0.09	0.88	-0.31	0.33	
	4 h	0.56	0.37	-0.26	0.65	-0.17	0.62	
	5 h	0.66	0.26	-0.86	0.02	-0.50	0.12	
	6 h	0.50	0.44	-0.47	0.37	-0.14	0.69	
	8 h	-0.34	0.55	-0.68	0.15	-0.34	0.30	
	10 h	0.42	0.53	-0.57	0.26	-0.29	0.40	
	All	-0.28	0.05	-0.21	0.13	-0.26	0.01	

Table A3: Correlations (Pearson correlation coefficients) between plasma flavonol concentrations (2 h after feeding on d 7) and mRNA abundances of hepatic genes involved in antioxidative status and inflammation (2 h after feeding on d 8) in calves orally supplemented with quercetin aglycone (50 mg/(kg BW × d)) for 7 days.

	Colostrum (CQ+)		Formula	a (FQ+)	All calves (Q+)				
Gene	Pearson r	P-value	Pearson r	P-value	Pearson r	<i>P</i> -value			
Cytokines									
TNF	-0.50	0.44	-0.15	0.80	-0.22	0.52			
IL1A	-0.35	0.60	-0.13	0.82	-0.18	0.60			
IL1B	0.94	0.02	0.82	0.05	0.82	<0.01			
Antioxidativ	e status								
CAT	0.28	0.68	-0.59	0.24	-0.20	0.58			
GPX	0.63	0.30	0.29	0.61	0.36	0.29			
SOD	0.41	0.53	-0.14	0.81	0.03	0.93			
Acute phase	Acute phase proteins								
SAA2	0.35	0.61	0.85	0.03	0.67	0.03			
HP	0.16	0.82	0.86	0.03	0.68	0.02			
CRP	0.87	0.06	-0.46	0.39	-0.04	0.91			
FGA	0.39	0.57	0.56	0.28	0.45	0.18			

CAT, catalase; *GPX1*, glutathione peroxidase; *SOD*, superoxide dismutase; *TNF*, tumor necrosis factor; IL1A, interleukin-1 α ; IL1B, interleukin-1 β ; HP, haptoglobin; FGA, fibrinogen; SAA2, serum amyloid A2; CRP, C-reactive protein.

PUBLICATION LIST

Publications (peer reviewed)

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DECLARATION

med. vet. Jeannine Gruse

Statutory Declaration:

Herewith I declare on oath that the submitted dissertation under the title

"Effects of the Flavonol Quercetin on Glucose Metabolism and Health Status in Neonatal Calves"

has been authored independently and without illegitimate external help and that it has not been formerly submitted to another university department.

Jeannine Gruse Berlin, 24th October 2016

Herewith I declare that I am not subject to any pending case of public prosecution.

Jeannine Gruse Berlin, 24th October 2016