Effect of citrus polyphenol- and curcumin-supplemented diet on inflammatory state in obese cats

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Abstract
Among obesity-associated disorders, low-grade inflammation has been described. The putative therapeutic properties of citrus and curcumin polyphenols could be associated with their anti-inflammatory properties. Two diets supplemented either with hesperidin (0·05 %) and naringin (0·1 %) from citrus extract or with highly bioavailable curcumin from Curcuma longa extract (0·09 %) were fed to eight obese cats for two 8-week periods (cross-over study design) while maintaining animals in an obese state. Plasma acute-phase protein (APP; α1-acid glycoprotein (AGP), serum amyloid A and haptoglobin) levels were assessed before and at the end of each test period. TNF-α, IL-1β, IL-2, IL-4, IL-5, IL-10, IL-12, IL-18, transforming growth factor-β, interferon (IFN)-γ mRNA levels were determined in peripheral blood mononuclear cells (PBMC) by real-time PCR. Compared with pre-study values, supplementation with citrus polyphenols resulted in lower plasma AGP and haptoglobin concentrations, while that with curcumin resulted in lower plasma AGP concentration. There were no differences between the supplementations. TNF-α, IL-1β, IL-2, IL-4, IL-5, IL-10, IL-12, IL-18, transforming growth factor-β, mRNA levels remained unaffected by either dietary supplementation. In contrast, IFN-γ and IL-2 mRNA levels were lower at the end of the citrus and the curcumin supplementation, respectively. There were no differences between the supplementations. The present study results show a slight effect of citrus and curcumin supplementation on inflammatory markers expressed by PBMC, and a decreased concentration of APP, which are mainly expressed by the liver. This would confirm that hesperidin and naringin or highly bioavailable curcumin extract have beneficial effects, targeted in the liver and could improve the obesity-related inflammatory state.

Key words: obesity; polyphenols; citrus; curcumin; inflammation; cat

The prevalence of obese or overweight cats is about 39 % of the feline population1,2. As obesity can predispose or exacerbate several serious medical conditions, its prevention or treatment is now a crucial issue. Indeed, obesity in cats is associated with hepatic lipodisosis, dermatological disease, urinary tract disease and diabetes2,3. Moreover, chronic inflammation has been described in obese cats3.

In this context, any tool able to correct obesity or its effects would be useful, including a food supplement that could be effective in contributing to the loss of body fat mass and/or improving the inflammatory state of obese cats.

Plants synthesise numerous compounds, i.e. polyphenols, that may explain most of the beneficial health-related effects that have been reported with different plant extracts in rodents or with diets rich in fruits and vegetables in humans. Citrus and Curcuma longa are known sources of polyphenols. In companion animals, citrus extract containing the polyphenols hesperidin and naringin has been shown to decrease plasma lipids and to have anti-inflammatory activity4,5. Curcuma longa (turmeric) extracts, particularly the dietary polyphenol curcumin, have been shown to possess beneficial properties, especially anti-inflammatory properties6. Recent studies indicate that turmeric extracts are poorly absorbed following oral ingestion. However, a modified extraction process is known to enhance bioavailability of natural turmeric extract.

The present study aimed to determine the effects of hesperidin and naringin or highly bioavailable curcumin dietary supplements on plasma acute-phase protein (APP) concentration, and cytokine mRNA expression in isolated peripheral blood mononuclear cells (PBMC) from obese cats. The hypothesis was that such diets could specifically improve the inflammatory state of obese cats.

Abbreviations: AGP, α1-acid glycoprotein; APP, acute-phase protein; PBMC, peripheral blood mononuclear cells; SAA, serum amyloid A

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Cytokine mRNA levels in isolated peripheral blood mononuclear cells

Cells were isolated on Ficoll/Hypaque gradients. Total RNA was extracted using TRIzol reagent according to the manufacturer’s protocol (Gibco BRL, Grand Island, NY, USA). After isolation, RNA pellets were dissolved in water and were then quantified spectrophotometrically. Total RNA (1 μg) was reverse transcribed in a 20-μl reaction volume using random primers (Pharmacia, Saclay, Orsay Cedex, France) and SuperScript III Reverse Transcriptase (Invitrogen, Cergy Pontoise, France). The cDNA solution (2 μl) was subjected to real-time PCR conducted in an iCycler System (Bio-Rad, Hercules, CA, USA) using the MasterMix kit (Eurogentec, Angers, France). PCR consisted of a 10-min denaturation at 95°C followed by 40 cycles (15 s at 95°C and 1 min at 60°C). The sense/anti-sense primer sequences were designed using the web-based Primer3 program (http://frodo.wi.mit.edu/primer3/) and are shown in Table 1. The glyceraldehyde-3-phosphate dehydrogenase mRNA level was used as a reference value. Relative quantitative expression was calculated by the $2^{-\Delta\Delta C_t}$ method. The level of expression before the supplementation was arbitrarily set at 100%.

**Plasma acute-phase protein concentrations**

Cephalic vein blood samples were collected from cats unfed overnight and centrifuged at 3000 g for 10 min at 4°C. Plasma APP (serum amyloid A (SAA); haptoglobin and α1-acid glycoprotein (AGP)) were assayed using commercial ELISA kits (Abcys, Paris, France).

**Statistical procedures**

Data were checked for normality and equality of variance. For plasma peptide concentrations, comparison was performed using paired Student’s t test. For gene expression, comparison among means was performed with Wilcoxon test for non-parametric values. A P-value < 0.05 was considered to be statistically significant. All statistical procedures were performed using StatView 5.0 software (SAS Institute, Cary, NC, USA).

**Results**

**Plasma acute-phase protein concentrations**

Table 1 shows the plasma concentrations of SAA, haptoglobin and AGP before (T0), and at the end of the citrus- (Citrus) or the curcumin-supplementation period (Curcumin). There was no effect of either supplement on plasma SAA concentrations, compared with the beginning of the study (T0). At the end of the citrus-supplementation period, plasma haptoglobin and AGP concentrations were reduced in Curcumin-supplemented cats.

**Table 1. Sense/anti-sense primers used for IL-1α, IL-2, IL-4, IL-10, IL-18, interferon (IFN)-γ, TNF-α, transforming growth factor (TGF)-β and glyceraldehyde-3-phosphate dehydrogenase (GAPdh) relative quantification**

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<td>5′-CACCAAGTAGTACCCGGAGG-3′, 5′-TTTGCGAGGCGAGGATG-3′</td>
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<td>TNF-α</td>
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</tr>
<tr>
<td>TGF-β</td>
<td>5′-CAATCCTGAGGAGGAGCAG-3′, 5′-GGAGGACAAAGCCCTTACTG-3′</td>
</tr>
<tr>
<td>GAPdh</td>
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European domestic shorthair neutered cats (n 8, three males, five females, 6.5 (SEM 0.5)-year old, mean body weight: 6.0 (SEM 0.4) kg) were used in this study. They were classified as obese on the basis of their body condition score (mean 3.8 (SEM 0.3), 1–5 possible) on admission. The cats had been obese for more than 1 year before the beginning of the study. Weight gain had originally been obtained by allowing ad libitum food consumption. All the cats were healthy based on physical examination and routine clinical laboratory data, except regarding obesity and associated insulin resistance. They were fed so as to maintain their obese body weight. The daily food quantity needed to reach this objective had been previously determined during the observation phase (4 weeks before the start of the study). Then, the cats were divided into two groups. Each group was fed a diet supplemented with either citrus extracts (protein 34.3%, fat 15.4%, starch 30.6%, hesperidin (Natural Orange Extract; Exquim SA, Barcelona, Spain) 0.05%, naringin (Citrus Eclectic; Exquim SA, Barcelona, Spain) 0.09% diet), or with highly bioavailable curcumin extract from C. longa (protein 34.2%, fat 16.5%, starch 28.9%, Bio-Curcumin (BCM-95; Frutarom, Londerzel, Belgium) 0.09% diet), for an 8-week period, providing 15.4%, starch 30.6%, hesperidin (Natural Orange Extract; Exquim SA) 0.1% diet. Cells were isolated on Ficoll/Hypaque gradients. Total RNA (1 μg) was extracted using TRIzol reagent according to the manufacturer’s protocol (Gibco BRL, Grand Island, NY, USA). After isolation, RNA pellets were dissolved in water and were then quantified spectrophotometrically. Total RNA (1 μg) was reverse transcribed in a 20-μl reaction volume using random primers (Pharmacia, Saclay, Orsay Cedex, France) and SuperScript III Reverse Transcriptase (Invitrogen, Cergy Pontoise, France). After reverse transcription, 80 μl distilled water was added. cDNA solution (2 μl) was subjected to real-time PCR conducted in an iCycler System (Bio-Rad, Hercules, CA, USA) using the MasterMix kit (Eurogentec, Angers, France). PCR consisted of a 10-min denaturation at 95°C followed by 40 cycles (15 s at 95°C and 1 min at 60°C). The sense/anti-sense primer sequences were designed using the web-based Primer3 program (http://frodo.wi.mit.edu/primer3/) and are shown in Table 1. The glyceraldehyde-3-phosphate dehydrogenase mRNA level was used as a reference value. Relative quantitative expression was calculated by the $2^{-\Delta\Delta C_t}$ method. The level of expression before the supplementation was arbitrarily set at 100%.

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AGP concentrations were lower compared with the initial values. At the end of the curcumin-supplementation period, plasma AGP concentration was lower, compared with T0 value. There were no significant differences in any of the plasma APP concentrations between the two supplementations (Table 2).

**Cytokine mRNA levels in isolated peripheral blood mononuclear cells**

Compared with the pre-study values (T0), cytokine mRNA levels were not different at the end of each period, except the interferon-γ mRNA level that was lower at the end of the citrus-supplementation period (12 (SEM 3) v. 100 (SEM 40) %, \( P<0.05 \)) and the IL-2 mRNA level that was lower at the end of the curcumin-supplementation period (48 (SEM 14) v. 100 (SEM 18) %, \( P<0.05 \)) than at T0. There were no significant differences in any of cytokine mRNA levels in isolated PBMC between the two supplementations.

**Discussion**

A chronic low-grade inflammatory state has been described in obese human subjects\(^{(9)}\) and dogs\(^{(10)}\). Many features associated with obesity could be involved in the development of an inflammatory state, even if the link between them is still not fully elucidated. First of all, because of the expansion of adipose tissue, adipocytes are more distant from blood vessels, and could therefore suffer from hypoxia which is associated with the release of cytokines\(^{(11)}\), and an alteration of the insulin signalling pathways that leads to a lower insulin sensitivity\(^{(12)}\). The attraction of macrophages to adipose tissue and their activation result in an amplification of local inflammation\(^{(13,14)}\). The recruitment of T cells has been described as playing a fundamental role in the initiation and propagation of adipose tissue inflammation\(^{(15)}\), leading to a systemic inflammation that is correlated with insulin resistance.

In the present study, we aimed to assess the specific effects of citrus- or curcumin-supplemented diet on low-grade inflammation in obese cats. For this, obese cats were fed quantities of diet necessary to maintain them in an obese state, and we measured the APP levels in plasma and the expression of cytokine mRNA in isolated PBMC. In these conditions, the results showed a slight effect of citrus- and curcumin-supplemented diets on IL-2 and interferon-γ expression, respectively. There were no significant differences in any cytokine mRNA level between the citrus and the curcumin supplementations. In contrast, these diets seemed to lower APP levels in plasma, especially the citrus-supplemented diet after which haptoglobin and AGP concentrations were lowered. There were no differences in plasma APP concentrations between the citrus and the curcumin supplementation.

APP belongs to a class of plasma proteins, levels of which increase (positive APP) or decrease (negative APP) in response to inflammation. Haptoglobin, SAA and AGP are positive APP, associated with a chronic inflammatory state, they are synthesised mainly by the liver but also by other tissues such as adipose tissue\(^{(16,17)}\), and the plasma concentration that we measured in this study reflected therefore the secretion from both the liver and other tissues.

Hesperidin, naringin and curcumin belong chemically to the polyphenol family, and have been used for centuries for their biological activity. Curcumin is a spice derivative that has been reported to lower cytokine production in diabetic rats\(^{(18)}\), and to improve inflammatory state in a mouse model of obesity\(^{(19)}\). In the present study, we did not confirm these results. This could be explained at least in part by the difference between the doses of curcumin that animals received. In the present study we wanted to test concentrations that would reflect typically recommended consumption (curcumin 0.09% diet), whereas in the previous in vivo studies, animals were fed diets with very high concentrations of curcumin (3%\(^{(19)}\), 100 mg/kg body weight\(^{(18)}\). In the same way, citrus plant extracts have been shown to have anti-inflammatory activities in canine leucocytes\(^{(4)}\). However in that study, leucocytes were from dogs that were not obese, and the genes whose expression was altered by citrus extracts were different from the genes we studied here.

One study\(^{(20)}\) performed in mice has shown a lower plasma haptoglobin concentration in an adjuvant arthritis model pretreated with *Curcuma*. The present study is, at least to the best of our knowledge, the first one concerning the effects of dietary citrus or curcumin on concentration of APP responsible for obesity-associated inflammation in cats.

Two phenomena could explain why citrus and curcumin supplementation had no effect on cytokine expression by PBMC while they decreased APP concentrations in plasma. First, it is possible that the supplements, at least at the dose we chose, could be efficient in their action on the liver after their absorption, and in lowering hepatic synthesis of APP;

<table>
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<th></th>
<th>T0</th>
<th>Citrus</th>
<th>Curcumin</th>
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<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>SEM</td>
<td>Mean</td>
</tr>
<tr>
<td>SAA (ng/ml)</td>
<td>2.36</td>
<td>0.14</td>
<td>3.79</td>
</tr>
<tr>
<td>Haptoglobin (mg/ml)</td>
<td>1.52</td>
<td>0.12</td>
<td>1.13</td>
</tr>
<tr>
<td>AGP (mg/ml)</td>
<td>524</td>
<td>66</td>
<td>251</td>
</tr>
</tbody>
</table>

SAA, serum amyloid A; AGP, α1-acid glycoprotein.

* Comparison was performed between Citrus and T0, and Curcumin and T0 by paired Student’s t test. There were no differences in plasma acute-phase protein concentrations between the citrus and the curcumin supplementation.
Supplementation, especially through an action targeted on cats that could be improved by dietary citrus and curcumin knowledge for the first time, an inflammatory state in obese supplementation lower plasma APP levels. This suggests, to our numerical decrease could become significant by increasing the statistical power of the test.

The present study shows that citrus and curcumin supplementation lower plasma APP levels. This suggests, to our knowledge for the first time, an inflammatory state in obese cats that could be improved by dietary citrus and curcumin supplementation, especially through an action targeted on the liver.

Acknowledgements

V. L., J. J., C. T. and P. N. conceived, conceptualised and supervised the present study. V. L., J. L. B. and B. F. acquired and analysed data. The authors thank S. Ninet and P. Bleis for their technical assistance. All authors have contributed to the preparation of the manuscript and agree with the submitted manuscript content, and have no conflict of interest.

References