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Biomodification of acenocoumarol by bifidobacteria

Melisa Fragomeno¹, Sabrina Assad¹, Pablo Mobili¹, Pablo J. Peruzzo³,
Jessica Minnaard^{1,2,*†} and Pablo Fernando Pérez^{1,2}

¹Centro de Investigación y Desarrollo en Criotecología de Alimentos (CIDCA)-Universidad Nacional de La Plata (UNLP), Consejo Nacional de Investigaciones Científicas y Técnicas (CONICET- CCT La Plata) and Consejo de Investigaciones Científicas de la Provincia de Buenos Aires (CICPBA). Calle 47 y 116, CP 1900, La Plata, Argentina, ²Área Microbiología e Inmunología, Departamento de Ciencias Biológicas, Facultad de Ciencias Exactas, UNLP. Calle 47 y 115, CP 1900, La Plata, 13, Argentina and ³Instituto de Investigaciones Fisicoquímicas Teóricas y Aplicadas - INIFTA (UNLP – CONICET CCT La Plata), Diag. 113 y 64, CC 16 Suc. 4 (B1904DPI) La Plata, Argentina

*Corresponding author: Centro de Investigación y Desarrollo en Criotecología de Alimentos, Calle 47 y 166, CP1900, La Plata, Argentina. Tel: +54 221 4254853; E-mail: jm@biol.unlp.edu.ar

One sentence summary: Bifidobacterial isolates of human origin biomodified in vitro the essential structure of anticoagulant acenocoumarol.

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†Jessica Minnaard, <https://orcid.org/0000-0003-3782-0187>

ABSTRACT

The increased interest of consumers in probiotic foods requires a deeper knowledge on the possible interactions with drugs, because their pharmacological properties could be modified. In this context, these studies are relevant for drugs such as acenocoumarol, whose dosage must be controlled due to, among other factors, food-drug interactions. Acenocoumarol is an oral anticoagulant with a narrow therapeutic range. The aim of the present research is to evaluate, *in vitro*, the effect of bifidobacteria on acenocoumarol. The drug was incubated with *Bifidobacterium bifidum* CIDCA 5310 or *Bifidobacterium adolescentis* CIDCA 5317 in MRS broth at 37°C for 24 h in anaerobic conditions. The effect of incubation with sterilized spent culture supernatants (SSCS) was also evaluated. Analysis by RP-HPLC showed that both bifidobacterial strains reduced the area of the acenocoumarol peak and two new peaks were evidenced. In addition, a decrease in the intensity of the bands at 1650, 1390 and 1110/cm was observed in the FTIR spectroscopic determinations. Moreover, a new band appeared at 1720/cm. No effect on the drug was observed when incubation was performed with SSCS. The present study showed a significant change in the concentration of the anticoagulant after incubation with bifidobacteria and results are compatible with biomodification of the drug due to enzymatic activity of bifidobacteria.

Keywords: bifidobacteria; biomodification; enzymes; acenocoumarol; probiotic; anticoagulants

INTRODUCTION

Probiotic microorganisms have been extensively studied, evidencing many beneficial properties when administrated in appropriate doses to the host. To highlight some of them, antioxidant activity, enhanced immune response and prevention of diseases caused by pathogenic bacteria can be mentioned (Joung

et al. 2021; Kim et al. 2018; Plaza-Diaz et al. 2019; Cruz et al. 2021; Xia et al. 2021).

It is worth noting that probiotics have shown the ability to modify drug pharmacokinetics (Kato et al. 2007; Lee et al. 2012; Matuskova et al. 2014; Stojančević et al. 2014; Kim et al. 2018). In an *in vivo* assay in rats, an increase in amiodarone bioavailability was observed in animals administered with *Escherichia*

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coli strain Nissle 1917 (Matuskova et al. 2014). Similar results were observed when nifedipine was administered concomitantly with *Lactobacillus casei* (Kato et al. 2007). On the other hand, the metabolic activity of probiotics on sulfasalazine was demonstrated, although this activity did not alter the blood concentrations compared to the control group of rats which did not receive the probiotic strains (Lee et al. 2012).

Probiotics could modify pharmacokinetics by affecting either the permeability of the intestinal epithelium or the activity of cytochromes, which are mainly responsible for the metabolism of many drugs (Matusková et al. 2011, 2014; Selwyn et al. 2016). Additionally, indirect effects of probiotic microorganisms, such as modification of intestinal microbiota, also contribute to the metabolization of many xenobiotics (Sousa et al. 2008; Stojančević et al. 2014; Jourova, Anzenbacher and Anzenbacherova 2016). Many studies describe the interaction with drugs of bacteria residing in the human gut, and how they can affect their pharmacodynamics and pharmacokinetics (Sousa et al. 2008; Jourova, Anzenbacher and Anzenbacherova 2016; Wilkinson, Ilhan and Herbst-Kralovetz 2018). It has been described, for example, that the action of bacterial reductases and hydrolases can generate nonpolar compounds (Sousa et al. 2008). The large number and variety of microorganisms in the intestine would place this organ at the same level as the liver in terms of drug metabolism (Stojančević et al. 2014).

Acenocoumarol is an anticoagulant derived from 4-hydroxicoumarin. It is structurally similar to warfarin but it presents a nitro group in the 4' position (Thijssen, Baars and Reijnders 1983; Saraeva et al. 2007). Acenocoumarol is a vitamin K antagonist that inhibits the enzyme vitamin K epoxide reductase, thus interfering in the carboxylation of the coagulation factors (Militaru et al. 2015). It is one of the most widely used oral anticoagulants after warfarin (Gschwind et al. 2013; Militaru et al. 2015) and the commercial presentations are racemic mixtures of S (-) and R (+) enantiomers. The R form elicits most of the therapeutic effects due to its longer half-life, compared to the S form.

Metabolism of acenocoumarol is carried out mainly by cytochrome P4502C9 (CYP2C9) and in a lesser extent by cytochromes CYP1A2 and CYP2C19 (Tassies et al. 2002; Ufer 2005). In the first phase of metabolism, acenocoumarol is biotransformed to 6- and 7-hydroxy-acenocoumarol, amine and acetamide acenocoumarol, and two diastereomeric alcohols. In addition, 8-hydroxy-acenocoumarol has been reported as a metabolite of minor quantitative importance. This hydroxylation is CYP-dependent (Saraeva et al. 2007).

Acenocoumarol presents a narrow therapeutic range and thus, patients under anticoagulant therapy have to be monitored frequently for correct dosage (Trejo 2004; Keeling 2017). Several intra- and inter-individual factors like body weight, sex, age and polymorphisms in genes involved in the drug or vitamin K metabolism affect the response to anticoagulant treatment (Wadelius et al. 2004; Saraeva et al. 2007; Militaru et al. 2015; Culler et al. 2020). The interaction of oral anticoagulants with foods and/or concomitant treatment with other medications has also been described (Vranckx, Valgimigli and Heidebuchel 2018). Nevertheless, there is no scientific evidence to support the interaction between oral anticoagulants and probiotic foods.

Microorganisms belonging to the genus *Bifidobacterium* early colonize the intestine of newborns and remain as part of the commensal protective microbiota in adults. Different species of this genus are frequently found in probiotic food formulations (Bottacini et al. 2014; O'Callaghan and van Sinderen 2016).

Intra- and extra-cellular enzymes such as hydrolases, reductases, epimerases and mutases have been described in *Bifidobacterium* (Nakamura et al. 2002; Manasian et al. 2020). These findings place this genus in a prominent position in the degradation of food ingredients and exogenous substances, thus contributing to the beneficial effects on human hosts (Pokusaeva, Fitzgerald and Van Sinderen 2011; Jung et al. 2020; Manasian et al. 2020; Modrackova et al. 2020; Dias de Queirós et al. 2020). However, there are no reports on the interaction of bifidobacteria with oral anticoagulants.

The aim of this work was to study, *in vitro*, the effect of bifidobacteria on acenocoumarol. Considering that nowadays lots of people are switching their dietary habits by increasing consumption of probiotic foods, the present study is relevant to understand the effect of those foods on consumers that are also under anticoagulant therapy.

MATERIALS AND METHODS

Bacterial strains and growth conditions

A total of two bacterial strains from the CIDCA collection were selected: *Bifidobacterium bifidum* CIDCA 5310 and *Bifidobacterium adolescentis* CIDCA 5317 (Pérez, Minnaard Disalvo and De Antoni 1998).

Bacteria were stored at -80°C with 10% w/v glycerol as cryoprotectant. Frozen bacterial suspensions were thawed, inoculated in MRS broth (Biokar Diagnostics, Beauvais, France), supplemented with 0.05% w/v L-cysteine (final concentration) and incubated at 37°C for 48 h in anaerobiosis (AnaeroPack™ anaerobic system, Mitsubishi Gas Chemical America, Inc. New York, NY). For the experiments, bacteria were inoculated (1% v/v) in the same medium and incubated for 24 h in the conditions specified above.

Acenocoumarol

Stock (100x) solutions of acenocoumarol (Laboratories Bagó, La Plata, Argentina) were prepared by dissolving 16 mg of drug in 1 mL of dimethylformamide (DMF) (Laboratories Anedra, Buenos Aires, Argentina) and filtering the resulting solution through a $0.45\ \mu\text{m}$ pore size membrane.

Effect of acenocoumarol on bacterial growth

Growth curves of both microorganisms were studied in the presence or not of 0.16 mg/mL acenocoumarol. Bacterial counts were assessed by plating appropriate dilutions on MRS agar and incubating for 48 h at 37°C in anaerobic conditions (AnaeroPack™ anaerobic system, Mitsubishi Gas Chemical America, Inc.).

Effect of bacteria on acenocoumarol

Bacteria were grown in the presence of the drug as previously described. After incubation for 0 and 24 h, samples were centrifuged for 5 minutes at 3000 g, filter sterilized ($0.45\ \mu\text{m}$) and frozen at -20°C until analysis.

To assess the effect of spent culture supernatants, 24 h-old cultures of each strain were centrifuged (15000 g for 15 min) and collected supernatants were filter sterilized ($0.45\ \mu\text{m}$). Then, supernatants were incubated with 0.16 mg/mL acenocoumarol for 24 h at 37°C and samples were stored at -20°C until analysis.

Chromatographic analysis-HPLC

Acenocoumarol concentration was determined by using reverse-phase high-performance liquid chromatography (RP-HPLC) by modification of a previously reported method (De Orsi et al. 1998). Hewlett Packard HP 1100 HPLC equipment with UV detection was used. Diode array detector was set at 280 nm. Chromatographic separations and subsequent quantifications were carried out at room temperature using a LiChrospher 100 RP-18 (4×250 mm, 5 μm) column. The mobile phase consisted of acetonitrile (Biopack, Zárate, Buenos Aires, Argentina)/phosphoric acid (Cicarelli, San Lorenzo, Santa Fe, Argentina) 60/40, prepared with MilliQ water, and the flow rate was adjusted to 1.2 mL/min at 25°C. Mobile phase was filtered before use (Nylon membranes, 0.45 μm, 13 mm, Osmonics Inc, Fisher Scientific, Pittsburgh, PA) and samples were previously filtered through 0.22 μm filters (GVS ABLUO, Sanford, FL). The injection volume was 20 μL and each sample was analyzed in triplicate. Peak areas were used for quantitative analyses. Calibration curve of peak areas versus acenocoumarol concentration was made with different dilutions of a stock solution (1.6 mg/mL). Linear relationship was observed in the range of 0.001–0.050 mg/mL. Retention times (t_R) and peak areas were evaluated using PeakFit (Systat Software, Inc, San Jose, CA).

Spectroscopic analysis-FTIR

Approximately 4 μL of each sample were placed on the sample holder of an Attenuated Total Reflectance FTIR (ATR-FTIR) Thermo Nicolet iS10 spectrometer (Thermo Scientific, Waltham, MA). Spectra were registered in the 4000–600/cm range by co-adding 64 scans with 4/cm spectral resolution, using OMNIC software (version 8.3, Thermo Scientific). At least five spectra were recorded for each sample.

Whenever necessary, residual contributions due to atmospheric water vapor and CO₂ were eliminated by subtraction of the corresponding spectra from the registered sample spectra, in order to obtain a flat baseline.

Statistical analysis

Statistical analysis was performed by the Mann–Whitney U Test by using InfoStat v 2020 (Di Rienzo et al. 2020). Differences were considered as statistically significant when $P < 0.05$. Figures were made using Microsoft Office Excel or OMNIC software (v8.3, Thermo Scientific; for FTIR figures).

RESULTS

Effect of acenocoumarol on bacterial growth

Figure 1 represents the growth curves for strains CIDCA 5310 (A) and CIDCA 5317 (B) in the presence or not of acenocoumarol. Both strains were able to grow in the presence of the anticoagulant and no significant differences were found between growth kinetics ($P > 0.05$).

Effect of bacteria on acenocoumarol

When RP-HPLC chromatograms were analyzed, the characteristic peak of acenocoumarol at $t_R = 15.25 \pm 0.15$ min was observed (Figs 2 and 3, peak 1).

RP-HPLC chromatograms corresponding to samples from 0- and 24-hour-old cultures as well as the controls without drug are

shown in Figs 2 and 3, and Figure S2 (Supporting Information). For both strains, at 24 h of incubation, the area of the acenocoumarol peak (peak 1) decreases and two new peaks are evident: peak 2 and peak 3. For both strains under study, these two new peaks are observed at t_R values ranging from 6.25 to 6.30 min for peak 3 and 10.60 to 10.74 min for peak 2 (Fig. 2B, C and 3B, C). It is important to note that the samples from cultures without the anticoagulant did not show peaks in the time range of interest (5–15 min) as shown in Figure S2 (Supporting Information) as an example.

The initial concentrations of acenocoumarol for peak 1 were 0.120 ± 0.017 mg/mL and 0.119 ± 0.016 mg/mL for the samples obtained from the incubation with strain CIDCA 5310 and strain CIDCA 5317, respectively. If these concentration values are compared with those obtained after 24 h of incubation a decrease is evident for both strains. Indeed, drug concentrations lowered down to 0.020 ± 0.008 mg/mL and 0.015 ± 0.001 mg/mL for samples corresponding to the strains CIDCA 5310 and CIDCA 5317, respectively.

Regarding the effect of the spent culture supernatants on acenocoumarol, after 24 h of incubation, there were no changes neither in the t_R nor in the area of the peaks (Figure S2, Supporting Information). No additional peaks were observed, and chromatograms resemble those for controls without bacteria (Figure S2, Supporting Information).

FTIR analysis

To gain further insight on the effect of bifidobacteria on acenocoumarol, we conducted FTIR analysis. As presented in Fig. 4A, the FTIR spectra of acenocoumarol in aqueous solution is characterized by a high intensity band at 1650/cm, corresponding to the stretching of the carbonyl bond in the lactone ring and two other medium intensity bands at 1390/cm and 1110/cm, ascribed to the -NO₂ symmetric stretching and to the stretching of the lactone C–O bond, respectively (Karci and Ertan 2005; Kostova and Nikolova 2006; Hubert Joe et al. 2009). The intensity of these bands correlated well with the concentration of the drug in MRS broth.

In the presence of *B. bifidum* CIDCA5310, the band at 1650/cm diminished its intensity during incubation. This band overlaps that of acenocoumarol at the beginning of the bacterial growth and decreased by around 50% after 24 h incubation (Fig. 4B).

In addition, the intensity of the bands at 1390/cm and in the 1110/cm region also decreased after incubation with bifidobacteria (Fig. 4B). At the same time, a new band, ascribed to the stretching of the C = O bond of carboxylic acid, appeared at 1720/cm (Fig. 4B).

DISCUSSION

As reports on the beneficial effects of probiotic microorganisms on the health of consumers increase, probiotic consumption has grown steadily in the population. However, the interaction of probiotic microorganisms with oral anticoagulants is seldom reported in the scientific literature (Lindh 2010). It is worth noting that people who are prescribed with acenocoumarol require dietary surveillance (e.g. restriction in foods with high vitamin K content; Holmes, Hunt and Shearer 2012). In this context, the presence of microorganisms in foods could modify the effectiveness of the anticoagulation therapy. In this study we assessed the effect on acenocoumarol of bacteria belonging to the genus *Bifidobacterium*. This is relevant since this genus is often included in the formulation of fermented foods and, in addition, is a main

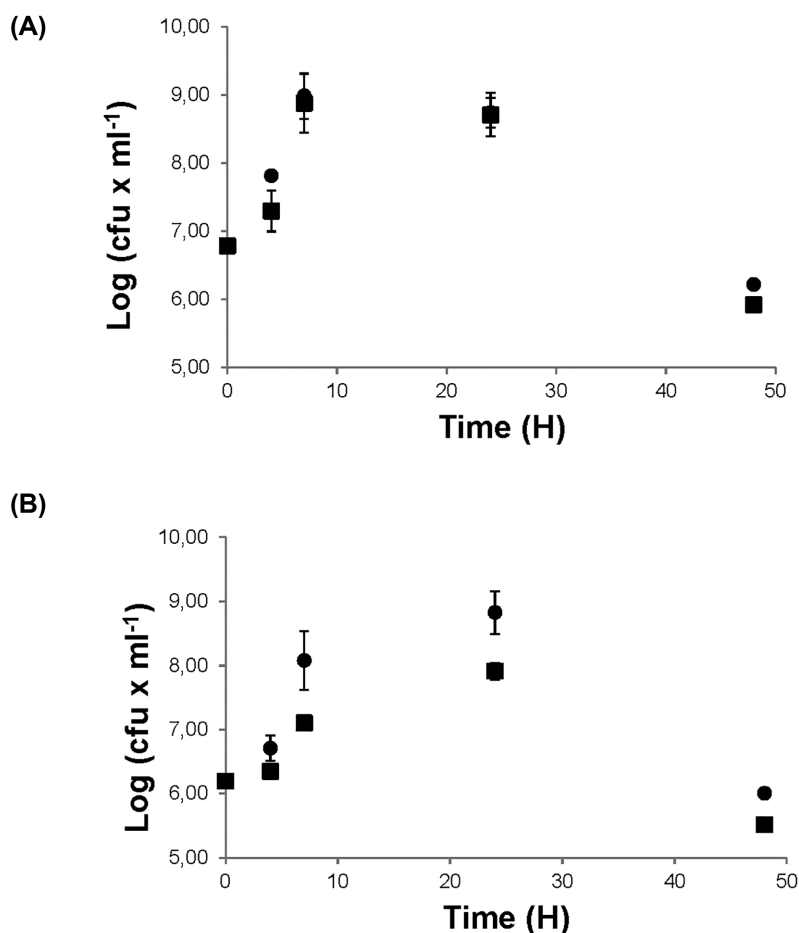


Figure 1. Acenocoumarol did not affect the growth kinetic of bifidobacteria. Growth curves of *B. bifidum* CIDCA 5310 (A) and *B. adolescentis* CIDCA 5317 (B) incubated in MRS broth with acenocoumarol (■) or control medium (●). Results show a representative experiment from three independent experiments. Bars show standard deviation.

member of the normal microbiota of children and adults (Turroni et al. 2009; Bottacini et al. 2014; Sun et al. 2015; Redondo-Useros et al. 2019).

The strains used in this work were isolated from the feces of healthy infants (Pérez et al. 1998) and they have demonstrated their potential as probiotic microorganisms (Trejo et al. 2006; Trejo, Pérez and De Antoni 2010; Trejo, De Antoni and Pérez 2013; Assad et al. 2020). Interestingly, some regulatory agencies consider that the species *B. bifidum* are representative of probiotic bifidobacteria (principle of “shared benefits”) (Marco et al. 2021).

Since in the present study we demonstrated that bifidobacteria decreased the concentration of anticoagulant, HPLC and FTIR were used as complementary analytical approaches.

Analysis by HPLC showed that the area of the characteristic peak of acenocoumarol (peak 1) was decreased in the presence of bifidobacteria and new peaks were detected (Figs 2 and 3). These findings are compatible with a biotransformation of acenocoumarol into other products. Since the column used in the present study separates species according to their polarity (Ceresole et al. 2008) we can hypothesize that bifidobacteria modifies acenocoumarol leading to more polar species. This effect was observed with both strains under study. Of note, analysis of the UV-spectra of these peaks is compatible with the occurrence of new species (data not shown).

To rule out that the observed effects are due to the low pH produced by the fermentative metabolism of bifidobacteria,

we tested the effect of an artificially acidified culture medium. In these conditions both the peak area and the t_R remained unchanged (Figure S1, Supporting Information) thus demonstrating that our findings are not related to the acidification of the culture medium by the microorganisms. Results obtained when anticoagulant was incubated with spent culture supernatants demonstrate that the observed effects are not due to extracellular factors released during bacterial growth (Figure S2, Supporting Information).

Results of FTIR analysis allowed us to gain further insight on the possible biomodifications due to the presence of bacteria. Firstly, at the 1650/cm region, corresponding to the stretching of the carbonyl of the lactone ring, the height of the peak correlates with acenocoumarol concentrations (Fig. 4A). This peak is then a good readout to assess biomodifications on acenocoumarol concentrations. Interestingly, after 24 h incubation, a new band at around 1720/cm was observed (Fig. 4B). This finding is compatible with an opening of the lactone ring that correlates with the decrease of the intensity observed in the band at 1110/cm, ascribed to the stretching of the lactone C-O bond. An opening of a lactone ring as the result of gut microbiota activity has been previously reported for lovastatin (Yoo et al. 2014).

Another change in FTIR spectra observed after the incubation of acenocoumarol with bifidobacteria was a decrease in the intensity of the peak at 1390/cm, ascribed to the nitro group, and

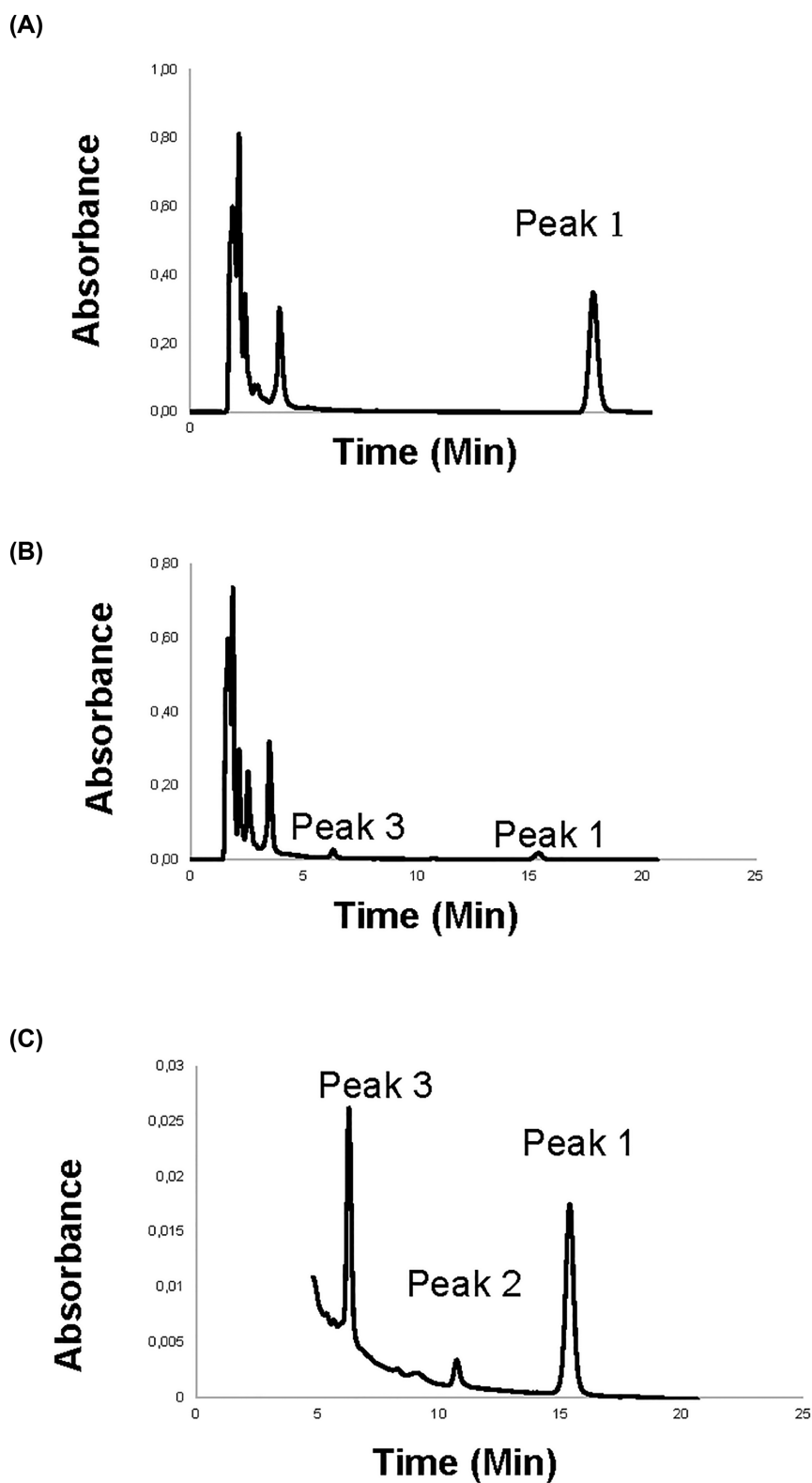


Figure 2. Acenocoumarol was biomodified by strain CIDCA 5310. RP-HPLC chromatograms of samples from cultures of the strain CIDCA 5310 in MRS with acenocoumarol (AC) incubated at 37°C for 0 h (A) and 24 h (B). AC was observed in peak 1 at $t_R = 15,27$ min. Peaks 2 ($t_R = 10,60$ min) and 3 ($t_R = 6,25$ min) can be observed in a magnification of the chromatogram at 24 h (C). Results show a representative experiment from three independent experiments.

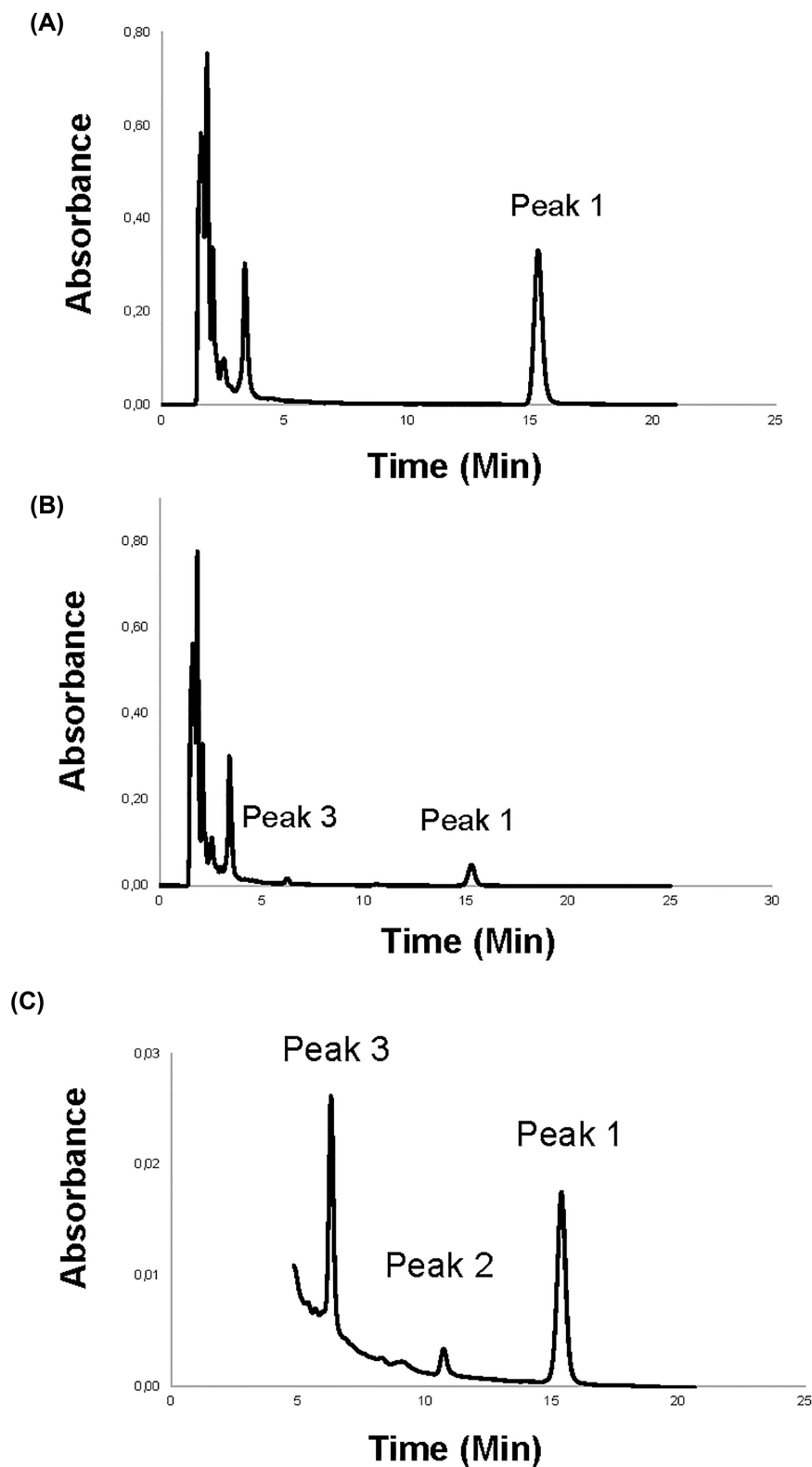


Figure 3. Acenocoumarol was biomodified by strain CIDCA 5317. RP-HPLC chromatograms of samples from cultures of the strain CIDCA 5317 in MRS with acenocoumarol (AC) incubated at 37°C for 0 h (A) and 24 h (B). AC was observed in peak 1 at $t_R = 15,40$ min. Peaks 2 ($t_R = 10,74$ min) and 3 ($t_R = 6,30$ min) can be observed in a magnification of the chromatogram at 24 h (C). Results show a representative experiment from three independent experiments.

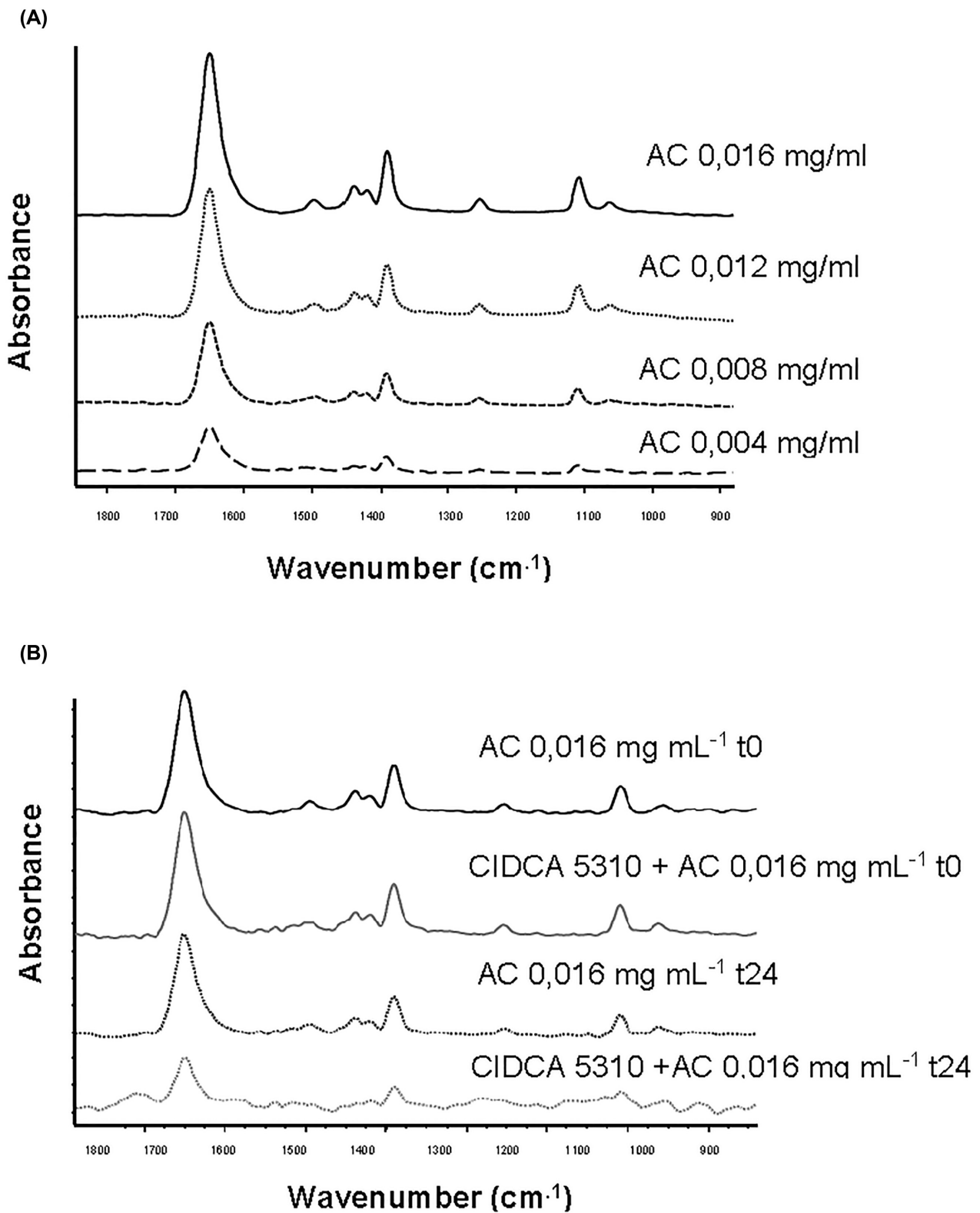


Figure 4. Acenocoumarol ATR-FTIR spectra was modified during incubation with strain CIDCA 5310. ATR-FTIR spectra, in the 900–1800/cm region, of different acenocoumarol (AC) concentrations in MRS broth (A), and spectra of AC after incubation with strain CIDCA 5310 for 0 h (—) and 24 h (···) and the control conditions of AC alone incubated for 0 h (—) and 24 h (···) (B). Results show a representative experiment from three independent experiments.

it is worth noting that the relation of the areas of this peak and the 1650/cm peak remained constant after 24 h incubation.

Taken together, our results are compatible with a modification of acenocoumarol structure affecting the lactone ring, due to bifidobacterial activity. As a consequence of this, acenocoumarol activity could be altered, considering that the lactone ring conforms the coumarinic core which confers its anticoagulant effect (Thijssen, Baars and Reijnders 1983; Kasperkiewicz et al. 2020).

Results of the present work are in agreement with previous studies that described the effect of bacterial enzymes on the activity of diverse compounds. In this context, reports showed that enzymes like reductases, esterases and dehydroxylases were found in different bifidobacteria and lactobacilli (McBain and Macfarlane 1998; Nakamura et al. 2002; Fritsch et al. 2017; Manasian et al. 2020).

If we analyze the acenocoumarol molecule in detail we can identify different functional groups such as -OH, -NO₂, -C = O that could be targets for the above-mentioned enzymes. As it has already been described, the normal *in vivo* route of metabolization of acenocoumarol includes hepatic enzymes which carry out oxidations and reductions (Lopez de Luca et al. 2006; Kasperkiewicz et al. 2020). Reductions turn the nitro group to an amine or the keto group to a hydroxyl. Oxidation leads to 6- and 7- hydroxyl metabolites (Kasperkiewicz et al. 2020).

Furthermore, Leonart et al. (2017), using a UPLC-MS method, described new molecules in human urine after coumarin metabolization (Leonart et al. 2017). These new metabolites included those resulting from hydroxylation, glucuronidation, sulfation, methylation and conjugation with N-acetylcysteine of the original molecule, showing the proneness of the drug to be biomodified by different types of enzymes. These results correlate with our findings in HPLC, given the new molecules detected were more polar than the original one.

It is evident that changes on acenocoumarol related to bacterial enzymatic activity, could impact on the anticoagulant effect. However, it must be highlighted, that *in vivo*, the scenario involves additional variables. Indeed, interactions between the drug, the gut microbiota, foods and host cells lining the gastrointestinal tract (Bailey and Dresser 2004; Sousa et al. 2008; Russel 2010; Stojančević et al. 2014; Enright, Joyce and Gahan 2017; Koziolok et al. 2019; Vertzoni et al. 2019). It is known that intestinal microorganisms modify the expression of intestinal transporters thus impacting on the bioavailability of drugs (Matusková et al. 2011; Saksena et al. 2011; Stojančević et al. 2014). In addition, they modulate the expression of cytokines (Azad, Sarker and Wan 2018; Yousefi et al. 2019; Wang et al. 2020) that in turn change permeability, drug metabolism and transport (Bertilsson, Olsson and Magnusson 2001; Le Vee et al. 2009; Thagia et al. 2015). Of note, *Bifidobacterium* and *Lactobacillus* strains have demonstrated to increase the barrier function of the intestinal epithelium through different mechanism (Hyland, Quigley and Brint 2014; Nébot-Vivinus et al. 2014; Srutkova et al. 2015; Guo et al. 2017).

Besides, consumption of probiotic foods along with acenocoumarol treatment in addition to pathologies that delay absorption (e.g. ulcerative colitis, intestinal bowel disease and Crohn's disease), could enhance the biotransformation of the drug by bifidobacteria by increasing residence time in the intestinal tract (Hatton et al. 2018).

Summarizing, our results show for the first time that bifidobacteria can modify acenocoumarol. This ability could lead to significant changes in the drug concentration thus adding a

relevant aspect to be considered for the pharmacological effect of this drug.

SUPPLEMENTARY DATA

Supplementary data are available at [FEMSLE](https://femsle.onlinelibrary.wiley.com/doi/10.1111/j.1365-2166.2021.02111.x) online.

COMPLIANCE WITH ETHICAL STANDARDS

This article does not contain any studies with human participants or animals performed by any of the authors.

AVAILABILITY OF DATA AND MATERIALS

The original data with the respective analysis corresponding to the results shown in this work are available up to reasonable requirements.

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Conflict of interest . None declared.

REFERENCES

- Assad SE, Rolny IS, Minnaard J et al. Bifidobacteria from human origin: interaction with phagocytic cells. *J Appl Microbiol* 2020;130:1357–1367. DOI: 10.1111/jam.14861.
- Azad MAK, Sarker M, Wan D. Immunomodulatory effects of probiotics on cytokine profiles. *Biomed Res Int* 2018;2018. DOI: 10.1155/2018/8063647.
- Bailey DG, Dresser GK. Interactions between grapefruit juice and cardiovascular drugs. *Am J Cardiovasc Drugs* 2004;4:281–97.
- Bertilsson PM, Olsson P, Magnusson KE. Cytokines influence mRNA expression of cytochrome P450 3A4 and MDRI in intestinal cells. *J Pharm Sci* 2001;90:638–46.
- Bottacini F, Ventura M, Sinderen D et al. Diversity, ecology and intestinal function of bifidobacteria. *Microb Cell Fact* 2014;13. DOI: 10.1186/1475-2859-13-S1-S4.
- Ceresole R, Rosasco MA, Forastieri CC et al. HPLC determination of acenocoumarol and its major thermal degradation product. *J Liq Chromatogr Relat Technol* 2008;31:179–87.
- Cruz BC dos S, de Sousa Moraes LF, De Nadai Marcon L et al. Evaluation of the efficacy of probiotic VSL#3 and synbiotic VSL#3 and yacon-based product in reducing oxidative stress and intestinal permeability in mice induced to colorectal carcinogenesis. *J Food Sci* 2021. DOI: 10.1111/1750-3841.15690.
- Cullell N, Carrera C, Muiño E et al. Genome-wide association study of VKORC1 and CYP2C9 on acenocoumarol dose, stroke

- recurrence and intracranial haemorrhage in Spain. *Sci Rep* 2020;**10**. DOI: 10.1038/s41598-020-59641-9.
- De Orsi D, Gagliardi L, Turchetto L et al. HPLC determination of warfarin and acenocoumarol in raw materials and pharmaceuticals. *J Pharm Biomed Anal* 1998;**17**:891–5.
- Di Rienzo JA, Casanoves F, Balzarini MG et al. InfoStat versión 2020. In: *Centro de Transferencia InfoStat*, FCA, Universidad Nacional de Córdoba, Argentina. 2020. <http://www.infostat.com.ar>.
- Dias de Queirós L, Rejane Alves de Ávila A, Vianna Botaro A et al. Combined isoflavones biotransformation increases the bioactive and antioxidant capacity of soymilk. 2020;**104**:10019–31.
- Enright EF, Joyce SA, Gahan CGM. Impact of gut microbiota-mediated bile acid metabolism on the solubilization capacity of bile salt micelles and drug solubility. 2017;**14**:1251–63.
- Fritsch C, Jänsch A, Ehrmann MA et al. Characterization of cinnamoyl esterases from different lactobacilli and bifidobacteria. *Curr Microbiol* 2017;**74**:247–56.
- Gschwind L, Rollason V, Daali Y et al. Role of P-glycoprotein in the uptake /efflux transport of oral vitamin K antagonists and rivaroxaban through the Caco-2 cell model 2013:259–65.
- Guo S, Gillingham T, Guo Y et al. Secretions of *Bifidobacterium infantis* and *Lactobacillus*. *JPGN* 2017;**64**:404–12.
- Hatton GB, Madla CM, Rabbie SC et al. All disease begins in the gut: Influence of gastrointestinal disorders and surgery on oral drug performance. *Int J Pharm* 2018;**548**:408–22.
- Holmes MV, Hunt BJ, Shearer MJ. The role of dietary vitamin K in the management of oral vitamin K antagonists. *Blood Rev* 2012;**26**:1–14.
- Hubert Joe I, Kostova I, Ravikumar C et al. Theoretical and vibrational spectral investigation of sodium salt of acenocoumarol. *J Raman Spectrosc* 2009;**40**:1033–8.
- Hyland NP, Quigley EMM, Brint E. Microbiota–host interactions in irritable bowel syndrome: epithelial barrier, immune regulation and brain-gut interactions. *World J Gastroenterol* 2014;**20**:8859–66.
- Joung H, Chu J, Kim B-K et al. Probiotics ameliorate chronic low-grade inflammation and fat accumulation with gut microbiota composition change in diet-induced obese mice models. *Appl Microb CELL Physiol* 2021. DOI: 10.1007/s00253-020-11060-6/Published.
- Jourova L, Anzenbacher P, Anzenbacherova E. Human gut microbiota plays a role in the metabolism of drugs. *Biomed Pap* 2016;**160**:317–26.
- Jung DH, Seo DH, Kim YJ et al. The presence of resistant starch-degrading amylases in *Bifidobacterium adolescentis* of the human gut. *Int J Biol Macromol* 2020;**161**:389–97.
- Karci F, Ertan N. Synthesis of some novel hetarylazo disperse dyes derived from 4-hydroxy-2H-1-benzopyran-2-one (4-hydroxycoumarin) as coupling component and investigation of their absorption spectra. *Dye Pigment* 2005;**64**:243–9.
- Kasperkiewicz K, Ponczek MB, Owczarek J et al. Antagonists of vitamin K-popular coumarin drugs and new synthetic and natural coumarin derivatives. *Molecules* 2020;**25**. DOI: 10.3390/molecules25061465.
- Kato R, Yuasa H, Inoue K et al. Effect of *Lactobacillus casei* on the absorption of nifedipine from rat small intestine. *Drug Metab Pharmacokinet* 2007;**22**:96–102.
- Keeling D. *Oxford Haemophilia and Thrombosis Centre Protocols for Out-Patient Oral Anticoagulation with Vitamin K Antagonists*. AC Protocols version 4.1. Oxford University Hospitals, 2017, 1–14.
- Kim JA, Bayo J, Cha J et al. Investigating the probiotic characteristics of four microbial strains with potential application in feed industry. *PLoS ONE* 2018;**14**:1–16.
- Kostova I, Nikolova I. Stability of neodymium(III) complexes of 4-hydroxycoumarins with anticoagulant activity. *Am J Pharmacol Toxicol* 2006;**1**:30–5.
- Koziolek M, Alcaro S, Augustijns P et al. The mechanisms of pharmacokinetic food-drug interactions – A perspective from the UNGAP group. *Eur J Pharm Sci* 2019. DOI: 10.1016/j.ejps.2019.04.003.
- Le Vee M, Lecureur V, Stieger B et al. Regulation of drug transporter expression in human hepatocytes exposed to the proinflammatory cytokines tumor necrosis factor-alpha or interleukin-6. *Drug Metab Dispos* 2009;**37**:685–93.
- Lee HJ, Zhang H, Orlovich Da et al. The influence of probiotic treatment on sulfasalazine metabolism in rat. *Xenobiotica* 2012;**42**:791–7.
- Leonart LP, Gasparetto JC, Pontes FLD et al. New metabolites of coumarin detected in human urine using ultra performance liquid chromatography/quadrupole-time-of-flight tandem mass spectrometry. *Molecules* 2017;**22**. DOI: 10.3390/molecules22112031.
- Lindh J. Possible interaction between probiotics and warfarin. *Lakartidningen* 2010;**107**:13–14.
- LopezdeLuca B, Sancho AM, Padullés Zamora A et al. Acenocoumarol. *OFFAMR* 2006;**25**:90–5.
- Manasian P, Bustos A-S, Pålsson B et al. First evidence of acyl-hydrolase/lipase activity from human probiotic bacteria: *Lactobacillus rhamnosus* GG and *Bifidobacterium longum* NCC 2705. *Front Microbiol* 2020;**11**:1–11.
- Marco ML, Sanders ME, Gänzle M et al. The International Scientific Association for Probiotics and Prebiotics (ISAPP) consensus statement on fermented foods. *Nat Rev Gastroenterol Hepatol* 2021;**18**:196–208.
- Matušková Z, Anzenbacherova E, Vecera R et al. Administration of a probiotic can change drug pharmacokinetics: effect of *E. coli* Nissle 1917 on amidarone absorption in rats. *PLoS ONE* 2014;**9**:3–7.
- Matušková Z, Šiller M, Tunková A et al. Effects of *Lactobacillus casei* on the expression and the activity of cytochromes P450 and on the CYP mRNA level in the intestine and the liver of male rats. *Neuroendocrinol Lett* 2011;**32**:8–14.
- McBain AJ, Macfarlane GT. Ecological and physiological studies on large intestinal bacteria in relation to production of hydrolytic and reductive enzymes involved in formation of genotoxic metabolites. *J Med Microbiol* 1998;**47**:407–16.
- Militaru F, Vesa S, Pop T et al. Pharmacogenetics aspects of oral anticoagulants therapy. *J Med Life* 2015;**8**:171–5.
- Modrackova N, Vlkova E, Tejnecky V et al. *Bifidobacterium* β -glucosidase activity and fermentation of dietary plant glucosides is species and strain specific. *Microorganisms* 2020;**8**:1–16.
- Nakamura J, Kubota Y, Miyaoka M et al. Comparison of four microbial enzymes in clostridia and bacteroides isolated from human feces. *Microbiol Immunol* 2002;**46**:487–90.
- Nébot-Vivinus M, Harkat C, Bziouche H et al. Multispecies probiotic protects gut barrier function in experimental models. *World J Gastroenterol* 2014;**20**:6832–43.
- O’Callaghan A, van Sinderen D. *Bifidobacteria* and their role as members of the human gut microbiota. *Front Microbiol* 2016;**7**. DOI: 10.3389/fmicb.2016.00925.
- Pérez PF, Minnaard Y, Disalvo EA et al. Surface properties of bifidobacterial strains of human origin. *Appl Environ Microbiol* 1998;**64**:21–6.

- Plaza-Diaz J, Ruiz-Ojeda FJ, Gil-Campos M et al. Mechanisms of action of probiotics. *Adv Nutr* 2019;10:S49–66.
- Pokusaeva K, Fitzgerald GF, Van Sinderen D. Carbohydrate metabolism in Bifidobacteria. *Genes Nutr* 2011;6:285–306.
- Redondo-Useros N, Gheorghe A, Díaz-Prieto LE et al. Associations of probiotic fermented milk (PFM) and yogurt consumption with Bifidobacterium and Lactobacillus components of the gut microbiota in healthy adults. 2019. DOI: 10.3390/nu11030651.
- Russel FGM, Ch 2: Transporters: importance in drug absorption, distribution, and removal. In: Pang KS, Rodrigues AD, Peter RM (eds.). *Enzyme- and Transporter-Based Drug-Drug Interactions: Progress and Future Challenges*. New York: Springer-Verlag, 2010, 27–49.
- Saksena S, Goyal S, Raheja G et al. Upregulation of P-glycoprotein by probiotics in intestinal epithelial cells and in the dextran sulfate sodium model of colitis in mice. *Am J Physiol Gastrointest Liver Physiol* 2011;300:G1115–23.
- Saraeva R, Paskaleva ID, Doncheva E et al. Pharmacogenetics of acenocoumarol: CYP2C9, CYP2C19, polymorphisms and dose requirements. *J Clin Pharm Ther* 2007;32:641–9.
- Selwyn FP, Cheng SL, Klaassen CD et al. Regulation of hepatic drug-metabolizing enzymes in germ-free mice by conventionalization and probiotics. *Drug Metab Dispos* 2016;44:262–74.
- Sousa T, Paterson R, Moore V et al. The gastrointestinal microbiota as a site for the biotransformation of drugs. *Int J Pharm* 2008;363:1–25.
- Srutkova D, Schwarzer M, Hudcovic T et al. Bifidobacterium longum CCM 7952 promotes epithelial barrier function and prevents acute dss-induced colitis in strictly strain-specific manner. *PLoS ONE* 2015;10:1–20.
- Stojančević M, Bojić G, Salami AH et al. The influence of intestinal tract and probiotics on the fate of orally administered drugs. *Curr Issues Mol Biol* 2014;16:55–67.
- Sun Z, Zhang W, Guo C et al. Comparative genomic analysis of 45 type strains of the genus Bifidobacterium: A snapshot of its genetic diversity and evolution. 2015. DOI: 10.1371/journal.pone.0117912.
- Tassies D, Freire C, Pijoan J et al. Pharmacogenetics of acenocoumarol: cytochrome P450 CYP2C9 polymorphisms influence dose requirements and stability of anticoagulation. *Haematologica* 2002;87:1185–91.
- Thagia I, Shaw EJ, Smith E et al. Intestinal epithelial suppressor of cytokine signaling 3 enhances microbial-induced inflammatory tumor necrosis factor- α , contributing to epithelial barrier dysfunction. *Am J Physiol Gastrointest Liver Physiol* 2015;308:G25–31.
- Thijssen HHW, Baars LG, Reijnders MJ. Acenocoumarol and its amino and acetamido metabolites. Comparative pharmacokinetics and pharmacodynamics in the rat. *J Pharm Pharmacol* 1983;35:793–8.
- Trejo FM, De Antoni GL, Pérez PF. Protective effect of bifidobacteria in an experimental model of *Clostridium difficile* associated colitis. *J Dairy Res* 2013;80:263–9.
- Trejo FM, Minnaard J, Perez PF et al. Inhibition of *Clostridium difficile* growth and adhesion to enterocytes by Bifidobacterium supernatants. *Anaerobe* 2006;12:186–93.
- Trejo FM, Pérez PF, De Antoni GL. Co-culture with potentially probiotic microorganisms antagonises virulence factors of *Clostridium difficile* in vitro. *Antonie van Leeuwenhoek, Int J Gen Mol Microbiol* 2010;98:19–29.
- Trejo IC. Anticoagulantes: farmacología, mecanismos de acción y usos clínicos. *Cuad Cirugía* 2004;18:83–90.
- Turróni F, Foroni E, Pizzetti P et al. Exploring the diversity of the bifidobacterial population in the human intestinal tract. *Appl Environ Microbiol* 2009;75:1534–45.
- Ufer M. Comparative pharmacokinetics of vitamin K antagonists. *Clin Pharmacokinet* 2005;44:1227–46.
- Vertzoni M, Augustijns P, Grimm M et al. Impact of regional differences along the gastrointestinal tract of healthy adults on oral drug absorption: An UNGAP review. *Eur J Pharm Sci* 2019. DOI: 10.1016/j.ejps.2019.04.013.
- Vranckx P, Valgimigli M, Heidebuchel H. The significance of drug-drug and drug-food interactions of oral anticoagulation. *Arrhythmia Electrophysiol Rev* 2018;7:55–61.
- Wadelius M, Sörlin K, Wallerman O et al. Warfarin sensitivity related to CYP2C9, CYP3A5, ABCB1 (MDR1) and other factors. *Pharmacogenomics J* 2004;4:40–8.
- Wang B, Hussain A, Zhou Y et al. *Saccharomyces boulardii* attenuates inflammatory response induced by *Clostridium perfringens* via TLR4/TLR15-MyD88 pathway in HD11 avian macrophages. *Poult Sci* 2020;99:5356–65.
- Wilkinson EM, Ilhan ZE, Herbst-Kralovetz MM. Microbiota–drug interactions: impact on metabolism and efficacy of therapeutics. *Maturitas* 2018;112:53–63.
- Xia J, Jiang S, Lv L et al. Modulation of the immune response and metabolism in germ-free rats colonized by the probiotic *Lactobacillus salivarius* LI01. *Appl Microbiol Biotechnol* 2021. DOI: 10.1007/s00253-021-11099-z.
- Yoo DH, Kim IS, Van Le TK et al. Gut microbiota-mediated drug interactions between lovastatin and antibiotics. *Drug Metab Dispos* 2014;42:1508–13.
- Yousefi B, Eslami M, Ghasemian A et al. Probiotics importance and their immunomodulatory properties. *J Cell Physiol* 2019;234:8008–18.