



# ALIMENTI E NUTRACEUTICI: Salute e prevenzione attraverso il cibo

5° Convegno a cura delle Piattaforme Tematiche di Ateneo su **"Alimenti e Nutrizione"** e **"Salute Umana e Animale"** 



Evento *on-line* 13 luglio 2021



1336





con la collaborazione di:





# ALIMENTI E NUTRACEUTICI: SALUTE E PREVENZIONE ATTRAVERSO IL CIBO

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## evento on-line 13 luglio 2021

## https://unicam.webex.com/meet/eventi

9.00 Saluto della autorità ed inizio dei lavori

#### PLENARY LECTURE

09.15 Food waste and by-products as valuable sources of bioactive compounds Lina Cossignani, Università di Perugia

#### Comunicazioni orali

- 09:45 Vitamins in human, cow, and donkey milk: a comparison of nutritional properties. Silvia Vincenzetti
- 10:00 Polyphenol microbial metabolites modulate proteolysis in neuronal cells reducing amyloid-beta (1-42) levels.
- Valentina Cecarini

   10:15
   A new HPLC-MS/MS method for the simultaneous determination of 36 polyphenols in blueberry, strawberry and their commercial products and determination of antioxidant

#### activity. Ahmed M Mustafa

- 10:30 Anti-inflammatory properties of hemp extracts in human bronchial cells. Oliviero Marinelli
- 10:45 Hemp extract safety for dermatological application. Cristina Aquzzi
- 11:00 Organic and antibiotic-free chicken meat: a proteomic approach. Laura Alessandroni

#### 11:15 Coffee break

- 11:30 Benefici e uso degli Aminoacidi essenziali.
- Emanuele Giordano
- 11:45 Antioxidant effects of 'Mela Rosa dei Monti Sibillini' on Drosophila melanogaster longevity. Daniela Beghelli
- 12:00 Assessment of cholesterol-lowering bioactive compounds in Italian-standard Espresso coffee.

#### Franks Kamgang Nzekoue

- 12:15 Capsaicin as new adjuvant in anti-cancer immunotherapy.
- Maria Beatrice Morelli
- 12:30 Gut microbiota modulation counteracts neuroinflammation and reduces the deposition of amyloid and tau in Alzheimer's disease.
- Laura Bonfili
- 12:45 Managing of the daily calcium dietary intake as a tool to contribute to a healthy lifestyle. Giorgia Vici
- 13:00 Evening Primrose Oil effects in human Pancreatic Ductal Adenocarcinoma cell lines. Laura Zeppa
- 13:15 Dietary metabolites and CVD biomarkers: what about trimethylamine and its oxidative product?

#### Laura Bordoni

- 13:30 Pausa pranzo
- 14:30 Exploiting the nutraceutical activities of the forage plant Onobrychis viciifolia (sainfoin) for the control of intestinal parasites in sheep doing without drugs.
  - Sara Marchegiani
- 14:45 The effect of chickpea flour addition levels and flour structures on quality and in vitro starch digestibility of corn-rice-based gluten free pasta. Xinying Suo

- 15:00 Ideabrill packaging capability in the preservation of raw and cooked ham: a comparative study.
- Laura Acquaticci 15:15 Can probiotics counteract the detrimental effects of sleep deprivation? Yadong Zheng
- 15:30 Water- and fat-soluble vitamins in milk after different thermal treatments and steam injection.
- Giuseppe Santini 15:45 Inhibitory activity of Chlorogenic Acid and Coffee Silverskin extracts against Klebsiella pneumoniae carbapenemases.
- Sonia I Ciancia 16:00 Characterization of milk proteins based nano-carriers: interactions with polyphenolic molecules of nutraceutical interest. Sofia Renzi
- 16:15 In vitro assessment of prebiotic effect of coffee and its by-products on human intestinal microbiota.
   Chiara Salvesi
- 16:30 Coffee break
- 16:45 Characterization of Panax ginseng root extracts: development of a new analytical method for the quantification of ginsenosides and biological studies. Simone Angeloni
- 17:00 Quality of wholemeal pasta made with pigmented and ancient wheats. Francesca Pompei
- 17:15 Short and medium chain free fatty acids in wine: extraction optimization and analysis. Lucia Lenti
- 17:30 Cannabidiol extracted from Cannabis sativa induces cell cycle arrest, mitophagy and cell differentiation in chronic myeloid leukemia cells.
- Federica Maggi
- 17:45 Sea fennel (Crithmum maritimum L.): a culinary herb with nutraceutical potential. Extraction, purification and chemical characterization of polar extracts. Diletta Piatti
- 18:00 Consumer targeted food packages to promote pulse consumption. Dalia Camilletti
- 18:15 Strategic change of gut microbiota composition ameliorates lipid metabolism in Alzheimer's disease.

#### Chunmei Gong

- 18:30 Protective effects of R (+)-Thioctic Acid treatment on the heart of spontaneously hypertensive rats. Proshanta Roy
- 18:45 Green extraction of hemp (Cannabis sativa L.) using microwave method for recovery of three valuable fractions (essential oil, phenolic compounds, and cannabinoids): a central composite design optimization study. Jacopo Torresi

#### 19:00 Chiusura dei lavori

Verrà assegnato un 1 CFU agli studenti delle Scuole di Scienze del Farmaco e dei Prodotti della Salute, di Scienze e Tecnologie (L-27) e di Bioscienze e Medicina Veterinaria (L-13/L-2, L-13, L-32, L-38, LM-6, LM-42) - UNICAM.

Comitato organizzatore e scientifico: Caprioli Giovanni, Cerquetella Matteo, Marchegiani Andrea, Sagratini Gianni In collaborazione con: Area Comunicazione Ufficio Stampa e Marketing di UNICAM

## INDICE

## Plenary lecture

Food waste and by-products as valuable sources of bioactive compounds Lina Cossignani, Federica Ianni, Luna Pollini, Francesca Blasi	p. 6
Oral Communications	
Ideabrill packaging capability in the preservation of raw and cooked ham: a comp Laura Acquaticci, Franks Kamgang Nzekoue, Lucia Bailetti, Simone Angeloni, Sag Caprioli Giovanni	•
	p. 8
Hemp extract safety for dermatological application Cristina Aguzzi, Laura Zeppa, Oliviero Marinelli, Massimo Nabissi	p. 9
Organic and antibiotic-free chicken meat: a proteomic approach	
Laura Alessandroni, Massimo Ricciutelli, Manuela Cortese, Silvia Vincenzetti, Renzo Galli, Gianni	
Sagratini	p. 10
the quantification of ginsenosides and biological studies Simone Angeloni, Giovanni Caprioli, Anna Maria Eleuteri, Valentina Cecarini, Ma Oliviero Marinelli, Massimo Nabissi, Laura Bordoni, Rosita Gabbianelli, Gianni Sagra	-
Antioxidant effects of 'Mela Rosa dei Monti Sibillini' on Drosophila melanogaster longevity	
<b>Daniela Beghelli</b> , Joice Guileine Nkuimi Wandjou, Ilenia Cirilli, Filippo Maggi, Laura Lupidi, Giovanni Caprioli, Cristina Angeloni	Giusti, Giulio p. 14
Gut microbiota modulation counteracts neuroinflammation and reduces the deposition of amyloid and tau in Alzheimer's disease	
Laura Bonfili, Valentina Cecarini, Chunmei Gong, Anna Maria Eleuteri	p. 16
Dietary metabolites and CVD biomarkers: what about trimethylamine and product?	its oxidative
Laura Bordoni, Irene Petracci, Iwona Pelikant-Malecka, Adriana Radulska, Marco Piangerelli,	
Joanna J. Samulak, Lukasz Lewicki, Leszek Kalinowski, Robert A. Olek, Rosita Gabb	p. 18
Exploiting the nutraceutical activities of the forage plant Onobrychis viciifolia (sainfoin) for the	

Exploiting the nutraceutical activities of the forage plant *Onobrychis viciifolia* (sainfoin) for the control of intestinal parasites in sheep doing without drugs

Daniele Braidot, Paride D'Ottavio, Serena Gabrielli, Annette Habluetzel, Filippo Maggi, **Sara Marchegiani**, Francesco Pagliacci, Paolo Polidori, Eleonora Spinozzi, Laura Tozzi, Valentina Zender p. 20

## Consumer targeted food packages to promote pulse consumptionDalia Camilletti, Giorgia Vici, Luca Belli, Claudia Annetti, Valeria Polzonettip. 22

## Polyphenol microbial metabolites modulate proteolysis in neuronal cells reducing amyloid-beta (1-42) levels

Valentina Cecarini, Massimiliano Cuccioloni, Yadong Zheng, Laura Bonfili, Chunmei Gong,Mauro Angeletti, Pedro Mena, Daniele Del Rio, Anna Maria Eleuterip. 24

## Inhibitory activity of Chlorogenic Acid and Coffee Silverskin extracts against *Klebsiella* pneumoniae carbapenemases

Sonia I Ciancia, Dezemona Petrelli, Giovanni Caprioli, Gianni Sagratini, Luca A Vitalip. 26

Benefici e uso degli Aminoacidi essenziali Emanuele Giordano

Strategic change of gut microbiota composition ameliorates lipid metabolism in Alzheimer's disease

Chunmei Gong, Laura Bonfili, Massimiliano Cuccioloni, Valentina Cecarini, Mauro Angeletti, Anna Maria Eleuteri p. 29

Assessment of cholesterol-lowering bioactive compounds in Italian-standard Espresso coffee Franks Kamgang Nzekoue, Laura Alessandroni, Luciano Navarini, Giovanni Caprioli, Gianni Sagratini p. 31

Short and medium chain free fatty acids in wine: extraction optimization and analysisLenti Lucia, Serena Scortichini, Dennis Fiorinip. 32

## Cannabidiol extracted from Cannabis sativa induces cell cycle arrest, mitophagy and cell differentiation in chronic myeloid leukemia cells

Federica Maggi, Maria Beatrice Morelli, Massimo Nabissi, Oliviero Marinelli, Daniele Tomassoni,Cristina Aguzzi, Laura Zeppa, Giorgio Santoni, Consuelo Amantinip. 34

Green extraction of hemp (*Cannabis sativa* L.) using microwave method for recovery of three valuable fractions (essential oil, phenolic compounds, and cannabinoids): a central composite design optimization study

Filippo Maggi, Marco Cespi, **Jacopo Torresi**, Riccardo Petrelli1 Eugenia Mazzara, Riccardo Carletti, Dennis Fiorini, Serena Scortichini, Stefano Dall'Acqua, Stefania Sut, Giovanni Caprioli, Simone Angeloni, Ahmed Mustafa, Jure Zekič p. 36

Anti-inflammatory properties of hemp extracts in human bronchial cellsOliviero Marinelli, Cristina Aguzzi, Laura Zeppa, Massimo Nabissip. 38

00

p. 28

#### Capsaicin as new adjuvant in anti-cancer immunotherapy

Maria Beatrice Morelli, Consuelo Amantini, Matteo Santoni, Oliviero Marinelli, Cristina Aguzzi,Laura Zeppa, Massimo Nabissi, Giorgio Santonip. 40

A new HPLC-MS/MS method for the simultaneous determination of 36 polyphenols in blueberry, strawberry and their commercial products and determination of antioxidant activity Ahmed M Mustafa, Simone Angeloni, Doaa Abouelenein, Laura Acquaticci, Jianbo Xiao, Gianni Sagratini, Filippo Maggi, Sauro Vittori, Giovanni Caprioli p. 42

## Sea fennel (*Crithmum maritimum* L.): a culinary herb with nutraceutical potential. Extraction, purification and chemical characterization of polar extracts

Diletta Piatti, Simone Angeloni, Giovanni Caprioli, Filippo Maggi, Massimo Ricciutelli, Lolita Arnoldi, Gianni Sagratini p. 44

### Quality of wholemeal pasta made with pigmented and ancient wheats Francesca Pompei, Matteo Bonfini, Xinying Suo, Ahmed M. Mustafa, Gianni Sagratini, Elena Vittadini p. 46

## Characterization of milk proteins based nano-carriers: interactions with polyphenolic molecules of nutraceutical interest

**Sofia Renzi**, Diego R Perinelli, Alessandro Alesi, Valeria Polzonetti, Paolo Polidori, Silvia Vincenzetti, Stefania Pucciarelli p. 48

## Protective effects of R (+)-Thioctic Acid treatment on the heart of spontaneously hypertensive rats

Proshanta Roy, Seyed Khosrow Tayebati, Ilenia Martinelli, Francesco Amenta, Daniele Tomassoni p. 49

## In vitro assessment of prebiotic effect of coffee and its by-products on human intestinal microbiota

Chiara Salvesi, Serena Scortichini, Giovanni Caprioli, Dennis Fiorini, Stefania Silvi p. 51

Water- and fat-soluble vitamins in milk after different thermal treatments and steam injection Giuseppe Santini, Valeria Polzonetti, Yulia Klimanova, Stefania Pucciarelli, Paolo Polidori, Lauro Fioretti, Silvia Vincenzetti p. 53

## The effect of chickpea flour addition levels and flour structures on quality and in vitro starch digestibility of corn-rice-based gluten free pasta

Xinying Suo, Margherita Dall'Asta, Gianluca Giuberti, Michele Minucciani, Elena Vittadini p. 55

Managing of the daily calcium dietary intake as a tool to contribute to a healthy lifestyle Giorgia Vici, Dalia Camilletti, Benedetta Gibertoni, Bruno Adesso, Luca Belli, Valeria Polzonetti

p. 57

Vitamins in human, cow, and donkey milk: a comparison of nutritional propertiesSilvia Vincenzetti, Giuseppe Santini, Yulia Klimanova, JingJing Zhang, Natalina Cammertoni,<br/>Valeria Polzonetti, Stefania Pucciarelli, Paolo Polidorip. 59

**Evening Primrose Oil effects in human Pancreatic Ductal Adenocarcinoma cell lines** Laura Zeppa, Cristina Aguzzi, Oliviero Marinelli, Massimo Nabissi p. 61

Can probiotics counteract the detrimental effects of sleep deprivation? Yadong Zheng, Laura Bonfili, Amina Aboufares El Alaoui, Michele Bellesi, Anna Maria Eleuteri

p. 63

#### Food waste and by-products as valuable sources of bioactive compounds

Lina Cossignani, Federica Ianni, Luna Pollini, Francesca Blasi

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Food waste and by-products are produced in all the phases of food life cycle, that is during agricultural production and industrial manufacturing, but also after processing, distribution and household activities. These wastes can be an important source of bioactive compounds, such as fibers, polysaccharides, proteins, lipids, phenolic compounds, vitamins, and carotenoids. Among these, a prominent place is occupied by phenolic compounds, which are well-known for their beneficial effects on human health, e.g., in the prevention of cancer and cardiovascular diseases [1]. Nowadays, there is an increasing interest toward the development and manufacture of functional foods and nutraceuticals, which have received great attention in the food market due to the growing consumer interest for "healthy" food. For this reason, pharmaceutical and food industries are interested in obtaining natural bioactive compounds to use as drugs, functional food ingredients, and nutraceuticals. To this aim, suitable extraction methods have to be developed for the different byproducts generated from agro-food industries. Traditional methods are still in use, even if they can be time, energy, and solvent consuming and sometimes degradation of thermolabile compounds can occur. Sustainable methods are continuously developed by improving and optimizing the existing processes following the principles of green chemistry [2]. In this presentation, the results of some researches aimed to exploit the nutritional/health properties of food waste are shown. Olive leaves have been studied as they represent an attractive source of bioactive compounds, in particular oleuropein and hydroxytyrosol. The phenolic compounds and the antioxidant capacity in leaves from different Olea europaea L. cultivars, harvested in different times of the year, were investigated and significant differences between the selected cultivars and harvesting times were observed. Afterwards Moringa oleifera leaves was extracted by unconventional extraction techniques and the metabolite profiling was obtained by UHPLC/QTOF-MS analysis. The results suggested that extraction method had a significant effect on yield and profile of phenolic compounds and that the untargeted metabolomics analysis is a powerful tool for characterizing the phenolic profile of plant extracts. Also Lycium barbarum leaves were investigated as source of phenolics, and chlorogenic acid and rutin were found as main components. Microwave-assisted infusion of L. barbarum leaves showed that chlorogenic acid is particularly sensitive to chemical modification during microwave heating. To date, a research on apple pomace, the major by-product obtained during apple juice processing, is underway. This product contains several important nutrients and secondary metabolites, such as carbohydrates, phenolic compounds, dietary fiber and minerals [3]. The phenolic fraction was investigated after ultrasound-assisted extraction with aqueous or hydroalcoholic solvent. The fractionation of apple pomace extract was also performed giving free and bound phenols, and hydrolyzed fractions in acid and alkaline conditions. The total extract and the different fractions have

been characterized by UHPLC/Q-TOF MS analysis and *in vitro* antioxidant activity was measured. Among the phenolic compounds identified, the most abundant were phloridzin and different glycosilated forms of quercetin. Future research will focus on the addition of apple pomace in meat and dairy products, in order to obtain fiber and phenolic enriched foods.

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## IDEABRILL PACKAGING CAPABILITY IN THE PRESERVATION OF RAW AND COOKED HAM: A COMPARATIVE STUDY



Laura Acquaticci<sup>1</sup>, Franks Kamgang Nzekoue<sup>1</sup>, Lucia Bailetti<sup>2</sup>, Simone Angeloni<sup>1</sup>, Gianni Sagratini<sup>1</sup>, Caprioli Giovanni<sup>1</sup>

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Food packaging contributes to the preservation and shelf-life of food. In literature, several studies demonstrate that active packaging, enriched of bioactive compounds like the essential oil of Rosmarinus Officinalis [1], and modified atmosphere packaging can positively influence the preservation of food. The study of new types of packaging is continuously increasing, mostly in terms of environmental impact and food preservation. In this study two types of packaging provided by Esseoquattro company were tested in order to assess their capability in the preservation of food. Ideabrill® packaging, a three layers pack of polyethylene high density layer, metallic layer and cellulose with long fiber layer, combined with Ideabrill® sacchetto salvafreschezza was compared to paper coupled with wings. The study was conducted on raw and cooked ham preserved in the packaging described above through the quantification of biogenic amines (BAs) at day 0, 3, 5 and 7. BAs can be considered markers to evaluate the freshness and the quality of food. In particular, a higher concentration of BAs is related to a higher deterioration degree of food. BAs were extracted, derivatized with dansyl chloride, purified with a SPE C-18 and then analysed with an HPLC-DAD method. This study, in combination with sensorial study, shows that Ideabrill® packaging combined with Ideabrill® sacchetto salvafreschezza showed the best conservation capability for raw and cooked ham when compared to the other one. Moreover, from an eco-friendly point of view, *Ideabrill*® packaging layers can be easily separated in order to encourage recycling.

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## Hemp extract safety for dermatological application



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The skin immune response is regulated by epidermal keratinocytes, lymphocytes and dendritic cells, by the releasing of cytokines, chemokines, growth factors and activation of specific transcription factors. An aberrant regulation of this immune response causes an inflammatory condition [1]. Many studies suggest the use of cannabinoids for the management of different inflammatory dermal conditions because the endocannabinoid system is present in the skin and has a role in maintaining its homeostasis, by regulating inflammation, immune modulation, cell proliferation and differentiation [1,2]. Cannabinoids showed anti-inflammatory and anti-pruritic activity through CB1 and CB2 receptors dependent mechanisms [1,2]. So, the endocannabinoid system is being studying to treat many inflammatory skin conditions, such as atopic dermatitis, contact dermatitis, psoriasis, acne, with pharmacological products available on the market [1,2]. We analysed the effect of CBD on human keratinocyte cell line (HaCaT) and human dermal fibroblasts (NHF A12) to assess its safety on an in vitro skin model. CBD not only showed a safety profile since did not increase the expression of the main inflammatory markers, but it reduced their basal expression. On the bases of these results, we also tested some essential oils (EOs) derived from hemp (Cannabis sativa) with low fraction of cannabinoids and their effect on the inflammatory markers. Results showed that also the EOs did not increase the basal expression of the inflammatory markers on the same cell lines [3]. These results suggest that the use of purified compounds as well as EOs derived from hemp would be safety for a topical use, and the next step should be the evaluation of their anti-inflammatory activity in a condition of dermal inflammation.

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## Organic and antibiotic-free chicken meat: a proteomic approach



## Laura Alessandroni<sup>1</sup>, Massimo Ricciutelli<sup>1</sup>, Manuela Cortese<sup>1</sup>, Silvia Vincenzetti<sup>2</sup>, Renzo Galli<sup>3</sup>, Gianni Sagratini<sup>1</sup>

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Antibiotics have been extensively used in animal husbandry, mostly as growth promoting agents or for prophylactic purposes to reduce bacterial pathogens. Their continuous use in animal production raised concern associated with emergence of resistance in meat consumers [1]. In 2018, "antibioticfree" meat attracted the attention of more than 43% of consumers, an always growing percentage [2]. Consequently, farmers adopted several strategies such as antibiotic-free and organic methods. Antibiotic-free animals are not treated with any kind of antibiotic, while organically raised animals cannot be fed with non-natural or synthetic substances, and cannot be treated with synthetic chemotherapeutic drugs, only phytotherapy and vaccinations are allowed. A deepening research of biomarkers to differentiate antibiotic-free, organic, and conventional meat is increasingly necessary. Proteomic is the identification, quantification, and functional characterization of the proteome of a cell, tissue, or organism under a specific, defined conditions [3]. The aim of this work is to compare the proteome of organically and antibiotic-free produced poultry meat. Proteomic analysis was performed on sarcoplasmic proteins combining two-dimensional electrophoresis for separation and ultra high-performance liquid chromatography coupled with high-resolution mass spectrometry (UHPLC-QTOF MS) and MASCOT protein database search for characterization. Two-dimensional electrophoresis (2DE) allows protein separation based on isoelectric point (pI) using an immobilized pH gradient in first dimension, and then on molecular weight using SDS-gel in the second dimension. After 2DE separation, each spot was in-gel digested with trypsin according to the protocol of Shevchenko et al. [4] and then injected in UHPLC-QTOFMS. Resulting spectra were extracted and data files were submitted to MASCOT database, to characterize protein spots. Preliminary results showed that most of identified proteins are glycolytic enzymes and qualitative analysis revealed that they were the same for both organic and antibiotic-free samples. Quantitative analysis was performed on SDS-gels using PDQuest software, it underlined statistically significant differences in expression of specific proteins, such as pyruvate kinase,  $\beta$ -enolase, and creatine kinase. These results were compared with several recent studies which use the same proteomic approach. They underline how over or under expressions of specific proteins affect meat quality in terms of tenderness and color [5-7]. Further studies need to be performed to confirm these preliminary results, furthermore, the next step would be the study of membrane proteins using the same proteomic method.

Acknowledgement: This research was performed at University of Camerino and supported by Fileni® industry.

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## Characterization of *Panax ginseng* root extracts: development of a new analytical method for the quantification of ginsenosides and biological studies



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The term *Panax* derived from the word "panacea" which means a cure of all disease. In fact, this medicinal plant has been used in the Traditional Chinese Medicine for the treatment of various illnesses. Recent investigations have reported that extracts prepared from the roots of ginseng possess diverse biological activities such as anticancer, antioxidant, anti-cardiovascular diseases, antidiabetic, and anti-obesity [1,2]. The aim of the present research was to develop and efficient and fast method for the characterization of ginsenosides in ginseng root extracts by using HPLC-MS/MS system with the scope to produce dried extracts rich in ginsenosides and to further evaluate them for their antiinflammatory and anti-obesity activities. For this purpose, a new analytical method for ginsenoside quantification in ginseng roots has been developed and validated by using HPLC-MS/MS triple quadrupole. In addition, several extraction procedures such as three diverse mechanisms of extraction (liquid-solid extraction (LSE), LSE-assisted by Naviglio extractor and ultrasound-assisted extraction (UAE)) both with various solvents (ethanol, methanol, and a mixture of them with water), have been evaluated studying the extraction efficiency and recovery capacity. The most performing procedures have been applied to prepare dried extracts which have been and will be biologically investigated. The new analytical method was characterized by a good linearity ( $R^2 \ge 0.9984$  for all analytes), repeatability (intraday, RSD = 2.9-6.3%; interday, RSD = 8.3-12.5%) and sensitivity (LOQ = 15) µg/kg for all ginsenosides). The best extraction procedures considering the extraction efficiency and recovery were those prepared with ethanol:water (50:50, v/v) and methanol:water (70:30, v/v) both with SLE and UAE. All the dried extracts were constituted by high level of ginsenosides since their total content varied from 2.6 to 3.3% of dried weight extract (DWE) and the best one, in term of bioactive compound content, was that prepared with SLE with a mixture of ethanol and water. The most abundant ginsenosides in ginseng extracts were Rg1 (10.0-13.1 g/kg of DWE) and Rb1 (6.2-7.8 g/kg of DWE). The individual standards of ginsenosides did not show a strong activity against cytokines except for interferon-y. In addition, the tested ginsenosides and the extracts similarly inhibit the activity of the NOS enzymes in TPH-1 cell line treated with LPS. The best dried extracts will be further characterized for other anti-inflammatory and anti-obesity activities in the perspective of their application in nutraceutical formulation.

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## Antioxidant effects of 'Mela Rosa dei Monti Sibillini' on Drosophila melanogaster longevity

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Aging could be considered as the result of a time-dependent accumulation of cell damages that inexorably brings to functional cellular impairments and increased vulnerability to chronicdegenerative diseases. The "Free Radical Theory of Aging", developed by Harman in 1956 [1], and also known as 'the oxidative damage theory of aging', has undergone several variations over the last years, including the suggestion of a central role for reactive oxygen species (ROS) produced by both environmental or endogenous factors [2]. Indeed, free radicals, by damaging biomolecules, compromise cellular homeostasis. Therefore, it is clear that natural dietary compounds with high content of antioxidants represent an extraordinary potential tool in the major healthcare challenge of delaying aging. Among fruits, great attention has been paid to apple, as it is a rich source of bioactive compounds, especially polyphenols. In the last years, an ancient variety of apple, whose cultivation had been abandoned for a long time, has been rediscovered: "la mela rosa dei Monti Sibillini" [3]. Aim of this work was to study the effect of the supplementation of two different extracts obtained from mela rosa dei Monti Sibillini on both longevity and the endogenous antioxidant defence system of Drosophila melanogaster (wild type, Canton strain). These fruit flies represent one of the most commonly used animal model for studying the complexity of the aging process because they not only share with humans many metabolic pathways, but also have functional homologs to nearly 65% of human disease-causing genes [4]. In this study, flies, separated by sex (n.200 for both males: M, and females: F; collected in seven tubes/gender/treatment), were lifelong supplemented (0.5% diluted in water) or not (CTR group) with two extracts: one obtained from the whole apples (A group) and the second one from their peels (B group). At two different intervals (at 15 and 45 days for F and at 15 or 30 days for M), some flies (n5/tube for three replicates), after sedation, were collected and, immediately, frozen (-80°C) till the evaluation of the expression of some genes related to antioxidant defences, such as: heme oxidase-1 (HO1); thioredoxin reductase (TRX); superoxide dismutase (SOD) and glutathione peroxidase (GpX) by RT-PCR assay. The longevity in supplemented female flies was significantly increased (by 10 and 20% in A and B groups, respectively) and also the mean half-life of CTR flies was significantly lower than the ones of A and B group. Furthermore, the B groups presented a much higher increase of half-life vs A groups (A vs B; p< 0.0001). In male flies, no significant effects were observed for longevity, although the average half-life of CTR group was significantly lower than the ones of A and B group. Furthermore, in males, the different extract supplementation (A vs B) did not differently modulate the half-life of fruit flies. A different trend over time of antioxidant gene expressions was observed in the two genders following the A or B supplementations. Female flies, physiologically characterized by higher antioxidant defenses [5], in the early stage of supplementation showed a down regulation of all the antioxidant genes here

evaluated; whereas, at 45 days, the gene expression increased for GpX, HO1 and TxR in B group. Male flies, instead, showed a modulation of antioxidant gene expression since the early stages, where upregulation of SOD and HO1 were observed in A group at 15 days. At 30 days, the GpX and HO1 gene expressions resulted increased in B group, whereas TxR in A group. The data obtained from this study suggest that the oral supplementation of the extracts of mela rosa dei Monti Sibillini starting from birth is able to prolong the average life of both female and male flies, while an effect on maximum longevity was registered only in females. Also with regard to the modulation of the antioxidant enzymes, different results were obtained in the two genders. In particular, in females, a down regulation of the expression of SOD, GpX, HO1 and TxR genes in the early stages of life was observed, which was not found in males. During aging, however, when the endogenous antioxidant defences decrease also in females, there was an up-regulation of antioxidant genes, modulated by supplementation in both sexes, although with a higher increase in female specimens. These data suggest that the apple is a potential functional food whose chronic consumption during life might prevent the onset of those chronic-degenerative diseases in which oxidative stress plays a key role. Moreover, these results sustain the cultivation of this ancient variety of apple to promote the economic growth in this area.

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## Gut microbiota modulation counteracts neuroinflammation and reduces the deposition of amyloid and tau in Alzheimer's disease



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Alzheimer's disease (AD) is a big socioeconomic burden, since effective prevention strategies and definitive treatment are not currently available. In AD, amyloid  $\beta$  aggregation, tau hyperphosphorylation and neuronal death are strictly associated with neuroinflammation. Protein aggregates initiate microglial activation and stimulate the release of inflammatory mediators that determine AD severity [1]. Additionally, alterations of gut microbiota composition play a crucial role in the development of AD. In fact, intestinal bacteria can influence brain and immune system function by the production of lipopolysaccharides and amyloid. Dysbiosis and increased abundance of proinflammatory bacteria are frequently observed in the gut of AD patients causing both local and systemic immune-mediated inflammation, oxidative stress, and impaired intake and metabolization of nutrients [2]. Recently, probiotics have emerged as a promising and safe strategy to manipulate gut microbiota composition and increase the host health status through a multi-level mechanism that is currently under investigation [3]. In this study, a multi-strain probiotic formulation of lactic acid bacteria and bifidobacteria (SLAB51) was chronically administered to a triple-transgenic mouse model of AD, B6;129-Psen1tm1Mpm Tg (APPSwe, tauP301L)1Lfa/J (named 3xTg-AD), in order to investigate the potential beneficial effects on AD progression through behavioural tests, biochemical and immunohistochemical analyses, metabolomic and lipidomic studies. Probiotics exerted multiple effects in AD mice by modulating gut microbiota composition. They favoured the proliferation of anti-inflammatory bacterial species and caused metabolic changes, such as the increase of antiinflammatory short chain fatty acids (SCFAs), able to directly act in the brain, positively interfering with inflammatory cytokines and increasing the concentration of gut hormones that are extremely important in the regulation of energy homeostasis, food intake and nervous functions [4,5,6]. Cholesterol homeostasis was improved in AD mice supplemented with probiotics, involving pathways mediated by the well-recognized anti-inflammatory liver X receptors. Plasma lipidomic analyses highlighted dynamic interactions among numerous actors of a complex inflammatory response, in which polyunsaturated fatty acids can compete each other and simultaneously co-operate in the resolution of inflammation. The downstream effects of these changes were the modulation of neuronal proteolysis, the reduction of amyloid  $\beta$  and tau aggregates and the improvement of cognitive abilities, with consequent slowdown of AD progression. Concluding, probiotics supplementation represents a promising multi-target intervention strategy able to positively shift gut microbiota composition, ameliorating neuroinflammation and metabolic alterations, finally counteracting cognitive decline and successfully reducing AD main hallmarks.

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# Dietary metabolites and CVD biomarkers: what about trimethylamine and its oxidative product?



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Recent studies indicate a potential role of trimethylamine (TMA) and its oxidative form, trimethylamine N-oxide (TMAO), in the development of cardiovascular diseases (CVD). Since TMA is produced by the gut microbiota from dietary precursors (e.g., carnitine and choline), this association has a direct impact on dietary advices aimed at preventing CVD. TMAO has been proposed as a promising biomarker of cardiovascular health, despite a clear mechanistic explanation of this association is still missing. Together with classical CVD predictors, another novel molecular biomarker for CVD is the measurement of circulating mitochondrial DNA copy number (mtDNAcn). Since mtDNAcn changes have been associated with both intrinsic and extrinsic factors, it has been proposed as a potential biomarker for complex multifactorial diseases, as it is an early indicator of damage and mitochondrial dysfunction. In a population of 389 coronary artery disease (CAD) patients and 151 healthy controls, we investigated TMA, TMAO and mtDNAcn, in association with established risk factors for CVD (gender, age, hypertension, smoking, diabetes, glomerular filtration rate (GFR)), with the aim to identify biomarkers predictive of CVD and clarify the impact of TMA and TMAO on health. Results showed that neither TMA nor TMAO were able to predict CAD in our population. On the contrary, a biomarker including mtDNAcn, gender, and hypertension emerged as a good predictor of CAD. MtDNAcn was significantly lower in CAD patients and in hypertensive subjects; it correlates with GFR and TMA, but not with TMAO. Our findings do not support the usage of TMA and TMAO as CAD biomarkers, and suggest that dietary restrictions or supplementations aimed at reducing the concentration of these metabolites in the circulation should be considered with

caution. On the other hand, mtDNAcn might be a new interesting plastic biomarker suitable to monitor the exposure to risk factors and the efficacy of preventive interventions for a personalized CAD risk reduction.

## Exploiting the nutraceutical activities of the forage plant Onobrychis viciifolia (sainfoin) for the control of intestinal parasites in sheep doing without drugs



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Sainfoin (Onobrychis viciifolia) is traditionally used in the mountainous areas of the Marche and Umbria Regions as a forage for sheep. A large bulk of literature demonstrates the plant's various nutraceutical properties in small ruminants. These are related to the presence of secondary metabolites and specifically to condensed tannins. Beneficial responses of animals fed on sainfoin hay or green forage include improved growth, milk and wool production, fertility, and reduced methane emissions and ammonia volatilization from dung or urine. Most important is the plant's capacity to reduce gastro-intestinal nematodes of small ruminants which is related to condensed tannins, namely to prodelphinidins and procyanidins and their degree of polymerisation [1]. This study exploits the available knowledge on the plant's anthelmintic effects to develop sainfoin based alimentary schemes for local sheep farms, allowing breeders to control intestinal parasites without the employment of anthelmintic drugs. In addition, the known nutraceutical effects of sainfoin on various animal husbandry parameters may lead to an increase in profitability of the family-based farm holdings. Specifically the project aims to i) assess the cultivation practices and the use of sainfoin by sheep breeders in the 'Alto Maceratese' area; ii) provide a comprehensive chemical characterization of tannins and other secondary metabolites of local sainfoin samples (ecotype 'lupinella nostrale' vs. ecotype 'lupinella quarantina'); iii) measure in vitro the anthelmintic activity of sainfoin extracts against the sheep parasite Strongyloides papillosus; iv) elaborate with sheep breeders nutraceutical sainfoin feeding protocols and evaluate the impact on intestinal parasite densities and animal husbandry parameters such as lamb growth and milk quality. From the first interviews (n=7) held with local farmers, emerges that breeders recognize the nutritional benefits of sainfoin stating that sheep 'like it very much', 'produce more milk' and 'the milk has a pleasant smell'. None of the farmers interviewed up to now has heard about effects on sheep parasites. Farmers retain the ecotype 'lupinella nostrale' very adapted for cultivation on 'poor soils' in the mountainous area of the 'Alto Maceratese' due to the plant's capacity to resist to adverse climatic conditions such as droughts. The ecotype 'lupinella quarantina' on the other hand, is more adapted to hilly areas as it requires more fertile soils similar to 'erba medica' (lucerne: Medicago sativa). Usually, farmers employ various forages according to the soil characteristics and altitude of their cultivatable land. Thus, during winter sheep are kept on hay from sainfoin and other forage plant species and during summer flocks graze

on fields after hay harvest (sainfoin or other) and on permanent semi-natural pastures. Therefore, the quantity of sainfoin and condensed tannins assumed by sheep flocks varies considerably from one farm to another and according to the season. Infection with gastro-intestinal nematodes is a common problem in small ruminants, impacting negatively on animal health and production parameters, if left uncontrolled. Usually, anthelmintic drugs are administered once a year (Nov / Dic) to the entire flocks. Data collected on 6 farms in 2018 revealed a mean number of 304 eggs / g of fecal material (EPG) ranging from 76 to 885 EPG. An egg density of 300 to 400 EPG is likely to have an impact on production parameters. Periodic monitoring effected in one farm at 'Sant'llario', on which sheep are held traditionally on sainfoin hay and sainfoin pasture to about 50 % of the total feed, revealed very low egg counts of 17, 6 and 68 EPG in spring 2020, autumn 2020 and spring 2021 respectively, values that convinced the breeder not to give any drug treatments since spring 2020. Whether these low parasite densities can be ascribed to the sainfoin samples from that farm (analysis in progress). Work planned for the next months will focus on the chemical characterisation of sainfoin samples taken from various sites of the area and on the assessment of their anthelmintic activity *in vitro* against

free-living stages (larvae and adults) of the sheep parasite *Strongyloides papillosus*. With these data at hand, antiparasitic sainfoin feeding schemes will be elaborated with interested breeders and the outcome in terms of parasite control and production parameters monitored.

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# Consumer targeted food packages to promote pulse consumption



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Pulse consumption is today a critical aspect in the diet, despite the known health benefits derived from their intake [1,2]. Consumers do not choose pulses in their diet for several reasons, such as long cooking time, flatulence connected to their intake and the shortage of innovative products [3,4]. Our studies are in line with literature data. One of our on-line surveys conducted on 789 subjects shows that according to the 62.3% of the participants pulses are not consumed for the long preparation and cooking time they require and according to the 45.9% for their unpleasant taste. Considering the health benefits due to pulse intake in the diet, it is fundamental to promote their consumption and in particular to conceive non-conventional strategies to try satisfying consumer needs and requests on pulses [5]. In this context, in collaboration with Fertitecnica Colfiorito s.r.l. company it was developed a project aimed to promote a healthy lifestyle through pulse consumption among consumers. For the project 5 specific targets of consumers (sportsmen, families, fast cooking consumers, traditional foods consumers and organic foods consumers) were identified and 5 specific product packages were designed starting from the evaluation of company products. Company food products were selected according to their nutritional characteristics (i.e. carbohydrates, proteins and fibre content) in accordance to target consumers. For example, in sportsmen the attention was focused on nutrient timing, while for families on both adults and children nutritional needs. In each package both traditional pulses and innovative food formulations were proposed (i.e. dry beans and novel pulsesbased products). Products were picked out also taking into account meal occasions and time for meal preparation and consumption, in order to guarantee healthy food choices even when time for cooking is not available. In addition, general dietary suggestions were provided to promote pulses use and to guide consumers in understanding how frequently they should be consumed and how to combine them to get a balanced plate. Furthermore, the project wished to support informed food choices as it is worth knowing that healthy dietary behaviours allow to contribute to a more sustainable future.

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## Polyphenol microbial metabolites modulate proteolysis in neuronal cells reducing amyloid-beta (1-42) levels



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Alzheimer's disease (AD) is a progressive neurodegeneration characterized by extensive protein aggregation and deposition in the brain, associated with defective proteasomal and autophagiclysosomal proteolytic pathways [1]. Since current drugs can only reduce specific symptoms, the identification of novel treatments is a major concern in AD research. Among natural compounds, (poly)phenols and their derivatives/metabolites are emerging as candidates in AD prevention due to their multiple beneficial effects [2-4]. Interestingly, phenyl-y-valerolactones (PVLs), products of intestinal microbial metabolism of flavan-3-ols, constitute a rich portion of phenolic metabolites in the circulation of subjects exposed to the widely spread dietary sources of this subclass of compounds [5]. Here, we investigated the ability of a selection of PVLs to modulate the functionality of cellular proteolytic pathways and to counteract the production and release of the toxic A $\beta$ (1-42) peptide, at levels readily achievable in body fluids upon moderate consumption of flavonoid-rich foods or beverages. Neuronal SH-SY5Y cells transfected with either the wild-type or the 717 valine-to-glycine amyloid precursor protein mutated gene were used as an AD model and treated with 5-(4'hydroxyphenyl)-y-valerolactone, 5-(3',4'-dihydroxyphenyl)-γ-valerolactone 5-(3'and hydroxyphenyl)-y-valerolactone-4'-sulfate. Combining in vitro and in silico studies, we observed that the phenyl-y-valerolactones of interest modulated cellular proteolysis via proteasome inhibition, through the action of their lactone group, and consequent autophagy upregulation as a compensatory protective mechanism. Furthermore, the metabolites inhibited cathepsin B activity, a lysosomal protease associated with the amyloidogenic processing of the amyloid precursor protein and the release of amyloid peptides [6], eventually reducing the amount of intra- and extracellular A $\beta$ (1-42) peptides. Our findings establish, for the first time, that these metabolites exert a neuroprotective activity by regulating intracellular proteolysis and confirm the role of autophagy and cathepsin B as possible targets of AD therapeutic strategies.

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## Inhibitory activity of Chlorogenic Acid and Coffee Silverskin extracts against *Klebsiella pneumoniae* carbapenemases



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Coffee processing produces numerous waste by-products (around two billion tonnes), including silverskin (SS). A possible reuse of these by-products is desirable, as they are not only an environmental but also an economic burden. Based on the observation that chlorogenic acid has anti-Extended-Spectrum- $\beta$ -Lactamase activity [1] and on the presence of this compound in SS [2], the aim of this work was to evaluate the inhibitory activity of chlorogenic acid and SS extracts against Klebsiella pneumoniae type-two carbapenemase (KPC2). Using four extraction solvents, H<sub>2</sub>O, MeOH, MeOH:H<sub>2</sub>O (50:50) and EtOH:H<sub>2</sub>O (70:30), four SS extracts were obtained. The inhibition activity of the carbapenemase-dependent resistance mechanism was measured against Klebsiella pneumoniae clinical strains KP-187 (KPC2-positive), KP-5 (KPC2-positive) and KP-ATCC13823 (carbapenem-susceptible control strain). Escherichia coli ATCC25922 was used as tester strain. In all experiments, imipenem was the indicator carbapenem antibiotic. To detect anti-carbapenemase activity, an adaptation of the Carbapenem Inactivation Method (CIM) [3] was developed and referred to as reverse Carbapenem Inactivation Method (rCIM). The protocol requires that produced carbapenemases are released from the cells after lysis, which was achieved by suspending 10 µl of bacterial culture in 200 µl of a Lysis Buffer. A disc preloaded with imipenem with or without putative inhibitors (i.e. chlorogenic acid or SS extracts) was added to the cell lysate. The highest concentrations of chlorogenic acid and extracts were 25 µg/ml and 500 µg/ml, respectively. After incubation at 37°C for 20 minutes, the disc was placed on an agar plate previously seeded with a 3 McFarland units suspension of the carbapenem-susceptible E. coli tester strain. Incubation at 37°C followed and results were recorded at two endpoints: 3 hours and 24 hours. The presence of a halo around the disc was an indication that the carbapenemase was not present or inhibited. To exclude that lack of carbapenemase activity could depend on inefficient bacterial cell lysis, experiments were carried out using a modified Lysis Buffer, wherein the concentration of the components with lytic activity was doubled. The SS extracts and chlorogenic acid showed no carbapenemase inhibitory activity irrespective to the Lysis conditions applied. Contrary to the observation against extendedspectrum beta lactamases (ESBL), pure chlorogenic acid did not show activity against KPC2 under the experimental conditions used. Also SS extracts did not inhibit the enzyme.

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## Benefici e uso degli Aminoacidi essenziali

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Gli aminoacidi essenziali sono molecole totipotenti [1] che possono essere usati per scopo clinici [2], nello sport e attività fisica [3], e per aumentare il benessere e la salute. Gli aminoacidi essenziali sono molecole che stimolano al massimo i processi anabolici, contribuendo al massimo effetto di sintesi proteica, rivestendo un particolare ruolo positivo negli anziani sani [4]. Il muscolo scheletrico umano è un serbatoio di aminoacidi per i tessuti e organi vitali in assenza di assorbimento di aminoacidi dall'intestino e fornisce precursori gluconeogenici epatici. Inoltre, l'alterato metabolismo muscolare gioca un ruolo chiave nella genesi, e quindi nella prevenzione, di molte comuni condizioni patologiche e malattie croniche [5]. Evidenze recenti su modelli animali sembrano indicare un aumento della longevità e salute a seguito di particolari rapporti tra aminoacidi essenziali e non essenziali [6]. Miscele particolari di aminoacidi essenziali sono in grado di aumentare la biogenesi mitocondriale, ridurre i marker dell'infiammazione, avere effetti benefici sul peso corporeo e grasso corporeo, ridurre l'incidenza delle infezioni, migliorare la funzione cognitiva [7].

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## Strategic change of gut microbiota composition ameliorates lipid metabolism in Alzheimer's disease



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Alzheimer's disease (AD) and age-related dementias represent a major global health challenge. Unhealthy diet and lifestyle can unbalance the intestinal microbiota composition and, consequently, energy metabolism, contributing to AD pathogenesis. Impairment of cerebral cholesterol metabolism occurs in both aging and AD, and lipid-lowering agents have been associated to a lower risk of neurodegenerative diseases [1], but the link between blood lipid profile and AD is a matter of debate [2]. Previous studies demonstrated that oral supplementation with a multi-strain probiotic formulation (SLAB51) reduced amyloid beta aggregates and brain damages in a triple transgenic mouse model of AD (3xTg-AD) [3]. Treated mice showed improved cognitive functions in response to an enrichment of gut anti-inflammatory metabolites, increased plasma concentrations of neuroprotective gut hormones, and ameliorated glucose uptake and metabolism [3,4]. Considering the contradictory data on the role of deregulated lipid homeostasis in AD and in light of the promising effects of probiotics on energy metabolism, 3xTg-AD mice and their wild type counterpart were chronically treated with SLAB51 with the aim of deeply dissecting the role of microbiota modulation in ameliorating AD pathology by affecting lipid homeostasis. On this purpose, 8 week-old 3xTg-AD mice were orally administered with SLAB51 for 4 and 12 months to analyze the plasma lipid profile (using lipidomic analyses and enzymatic colorimetric assays), along with the cerebral and hepatic expression levels of key regulators of cholesterol metabolism (through Western blotting and ELISA). As expected, dyslipidemia was observed in older 3xTg-AD mice but not in wild type animals. Interestingly, upon probiotics administration, cholesterol biosynthesis was inhibited in AD mice with a process involving sterol regulatory element binding protein 1c and liver X receptors mediated pathways. Decreased plasma and brain concentration of 27-hydroxycholesterol and increased brain expression of cholesterol 24S-hydroxylase indicated that alternative pathways of bile acid synthesis are influenced. These data, together with the hypocholesterolemic effects and the ameliorated fatty acids profile successfully demonstrated that microbiota modulation through probiotics can positively change lipid composition in AD mice, with arachidonic acid representing one important hub metabolite in the interactions among probiotic-induced lipid profile changes, insulin sensitivity, and inflammation. Concluding, our data definitely demonstrate that gut microbiota modulation with probiotics successfully counteract AD progression influencing multiple pathways including lipid metabolism. The effective prevention of AD through the fine regulation of the associated risk factors will remain

The effective prevention of AD through the fine regulation of the associated risk factors will remain a constant goal of research and present findings should be inevitably considered in future preventative and therapeutic AD protocols.

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## Assessment of cholesterol-lowering bioactive compounds in Italian-standard Espresso coffee



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Espresso Coffee (EC) is one of the most popular coffee brews in the world, obtained from the percolation of hot water under pressure through compacted cakes of roasted ground coffee [1]. To our knowledge, among all the bioactive compounds investigated in coffee-based products, phytosterol (PS) content has never been assessed in coffee brews such as EC. This class of bioactive compounds is known for their LDL cholesterol-lowering properties, which contrast with the blood cholesterol increment effect of diterpenes, another class of phytochemicals present in coffee [2,3]. This study aims thus, to assess for the first time the content of PS in EC to improve the knowledge on the phytochemicals and health potentials of coffee brews. 14 EC samples produced with coffee cultivated in 13 different countries were studied. PS were extracted by hot saponification followed by derivatization for HPLC-DAD analysis.  $\beta$ -sitosterol (48.1-178 mg L<sup>-1</sup>) was the most abundant followed by stigmasterol (11.4-46.3 mg L<sup>-1</sup>) and campesterol (9.9-48.8 mg L<sup>-1</sup>). Total PS fraction ranged from 69.8 mg L<sup>-1</sup> to 275.0 mg L<sup>-1</sup> with an average level of 145.4 ± 52.7 mg L<sup>-1</sup>. Therefore, a standard cup of EC (25 mL) could provide 3.7 ± 1.4 mg of PS. This quantity is not enough for blood-cholesterol reduction but is a quantity that could inhibit the potential unhealthy physiological effects of diterpenes in coffee brews.

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# Short and medium chain free fatty acids in wine: extraction optimization and analysis



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The composition of free fatty acids (FFAs) in food matrices is an important parameter in the definition of quality. Depending on the matrix under consideration, their presence and amount may give positive but also negative contributions. In cheese, for example, they are known to afford the characteristic aroma, but they are able to produce an undesired defect of cheese-like off-flavor in other products such as milk or cream [1,2]. When considering wine, acetic acid (AA) plays a very important role in this context. It is the main constituent of wine volatile acidity, and when present above certain levels it is associated with an uncontrolled proliferation of acetic acid bacteria. This results in the typical vinegar defect in the aroma and the consequent wine spoilage [3]. Short and medium chain free fatty acids (SCFFAs and MCFFAs) different from AA are also found in wine, being by-products of yeasts fermentation. Their amount is important since they can significantly impact the sensory quality when present above their olfactory thresholds [4]. Starting from these premises, the analysis and quantification of SCFFAs and MCFFAs in wine samples has a key importance in the definition of wine quality. The aim of the present study was to optimize and validate an analytical method to quantify FFAs in wines after their liquid-liquid extraction (LLE) by diethyl ether and a subsequent analysis by gas chromatography coupled with flame ionization detection. The extraction optimization has been performed by considering different conditions: starting from the acidification of wine samples with aqueous sulphuric acid solution, to the use of different salts and salts mixtures. Promising preliminary results have been obtained using the mixture NaH<sub>2</sub>PO<sub>4</sub>/(NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, a salt combination resulted to improve the extraction of SCFFAs and MCFFAs also by headspace solid phase microextraction as reported in a previous study [5].

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# Cannabidiol extracted from *Cannabis sativa* induces cell cycle arrest, mitophagy and cell differentiation in chronic myeloid leukemia cells

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Cannabidiol (CBD) is one of the phytochemicals extracted from Cannabis sativa lacking of psychoactive effects whose use as herbal medicine dates back to 500 BC. It displays several effects such as anti-inflammatory, antioxidant, analgesic, anxiolytic, anticonvulsive and neuroprotective [1]. Recently, different companies started to make products, containing hemp-derived CBD, for body care purposes, as oil for salad dressings, or to be taken directly as a dietary supplement [2]. In addition, it is now well accepted that CBD can be also considered a promising new anti-cancer strategy, alone or in combinations with conventional chemotherapies. In fact, several findings demonstrated that CBD is able to induce cell death, inhibit cell migration in vitro, decrease tumor size and vascularization in glioma models [1,3,4]. Moreover, similar interesting results have been obtained in breast [5], lung [6], colon rectal [7] and prostate [8] cancers. CBD acts as an agonist for the Transient Receptor Potential Vanilloid (TRPV) channels 1 and 2 while it has no affinity for the endocannabinoid CB1 and 2 receptors [9]. Although, successful results were obtained in Chronic Myeloid Leukemia (CML) with TKI inhibitors, several patients show TKI resistance. Thus, this study investigated the effects induced by CBD treatment via activating TRPV2 in CML cell lines. The effects of CBD treatment, via TRPV2, were assessed by cell viability, BrdU proliferation and cell cycle assays. The stimulation of mitophagy and the expression of differentiation markers was evaluated by western blot, FACS analysis and confocal microscopy. The involvement of TRPV2 was studied by gene silencing. CBD treatment inhibited cell proliferation and cell cycle in CML cells. It also promoted mitochondria dysfunction and mitophagy via TRPV2 activation. These effects are associated with changes in OCT-4 and PU.1 differentiation markers. Finally, a synergistic effect by combining CBD with Imatinib was found. CBD, by promoting inhibition of cell proliferation, mitophagy and the reduction in stemness, could be a new strategy to enhance conventional therapy and improve the prognosis of CML patients developing TKIs resistance. Overall, these results supported previous findings about the idea that hemp-derived products could be used as supplements able to ameliorate the healthy status.

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# Green extraction of hemp (*Cannabis sativa* L.) using microwave method for recovery of three valuable fractions (essential oil, phenolic compounds, and cannabinoids): a central composite design optimization study



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Industrial hemp (Cannabis sativa L.) is an herbaceous plant with a characteristic scent that has been used for over 5000 years for its various properties. In particular, hemp essential oil can be used in pharmaceutical, cosmetic, and nutraceutical fields. In this work, hemp Futura 75 was processed with an innovative and green extraction method, namely Microwave-Assisted Extraction (MAE) [1], using a Design of Experiment (DoE) approach [2]. After the essential oil distillation, two valuable byproducts, namely the residual biomass and water enriched in hydrophilic compounds, were recovered and analyzed. The design variables analyzed were the microwave irradiation power (W/g), the extraction time (min), and the amount of water added to hemp samples (%). This research aimed to evaluate how these experimental parameters can affect the EO yield and composition together with the yield and antioxidant properties of the aqueous residue. Total phenolic and flavonoid content (TPC, TFC) and antioxidant activity (DPPH) were calculated through spectrophotometric UV-Vis experiments [3-4]; whereas phenolics and cannabinoids in the residual water and biomass, respectively, were determined by HPLC-MS analysis. GC-MS analysis was performed to study the chemical composition of essential oils obtained by MAE. At the same time, GC-FID was used to quantify the essential oil constituents and the CBD content in the residual biomass. Regarding the essential oil analysis, the responses yield and content of some compounds were well described by the mathematical models applied. In addition, in the case of the aqueous residues, the yield, polyphenols, and flavonoids content and antioxidant activity, as the four responses analyzed, suitably fit the models. On the other hand, there were no relationships between the CBD content in the residual biomass and the MAE experimental conditions, probably due to the samples' intrinsic variability. A multiple optimizing procedures based on desirability approach was employed to identify the most suitable experimental parameters able to maximize all the satisfactory responses simultaneously. In particular, high microwave power and long extraction times are needed to enhance the yield and chemical profile of essential oil, along with a medium-high amount of water, required to improve the aqueous extract yield and antioxidant capacity. HPLC-MS analysis put in evidence the abundance of flavone glycosides and CBD in the residual water and biomass, respectively. Our work showed that

hemp biomass is a multipurpose matter able to provide different fractions employable in several fields such as nutraceuticals, pharmaceuticals, cosmetics, and pest science.

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# Anti-inflammatory properties of hemp extracts in human bronchial cells



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Inflammatory lung diseases represent one of the most widespread affections associated with morbidity, mortality and health care expenditure in Europe [1]. Chronic Obstructive Pulmonary Disease (COPD) is a respiratory disorder associated to a chronic inflammatory response. Main features are a progressive and irreversible airflow limitation, consequence of inflammatory cells infiltration and alteration in cilia structure and functionality [2-3]. The prescribed therapy consists of inhalation of bronchodilators and glucocorticoids [4], but it is historically adopted the use of traditional herbal remedies for treatment of respiratory diseases [5]. Previously, we evaluated thyme extract, obtained from Thymus vulgaris L., widely used as a spice, as potential remedy for airways affections. Briefly, we studied the effects of thyme extract in an in vitro model of lung inflammation induced by lipopolysaccharide (LPS) in primary human airway (bronchial/tracheal) epithelial cell lines (HBEpC/HTEpC) [6]. Indeed, exposure to LPS increases several pro-inflammatory mediators expression, such as interleukin-1 beta (IL-1 beta) and tumor necrosis factor-alpha (TNF- $\alpha$ ), involved in pathogenesis of COPD [7], and mucus production in airway epithelial cells. Results showed that thyme extract was non-cytotoxic for HBEpC/HTEpC cells and was effective in reducing inflammatory markers as NF-kB, IL-1ß and IL-8 expression in LPS-treated cells [6]. Subsequently, to further characterize the beneficial use of thyme extract in COPD treatment, it was evaluated its effects in regulating Cilia beating frequency (CBF), that is dramatically impaired in COPD, using a 3D in vitro model of Human Airway Epithelia, using three different bronchial primary cells derived from COPD single donors. Results show that thyme extract is involved in stimulating CBF, through an increase of  $[Ca^{2+}]_i$  in a TRPV4 channel-dependent manner [8]. Hemp (*Cannabis sativa L*.) represents a sustainable crop multipurpose with a low environmental impact which can be useful in the agricultural field, phytoremediation and for nutrition, cosmetic, building, and pharmaceutical application. In this regard, the hemp essential oil (EO), which is a mixture of volatile compounds with a lower concentration of the cannabinoid fraction, is gaining interest by pharma companies, for its potential anti-inflammatory properties [9-10]. Despite plant EOs have shown several biological properties, it has been scarcely investigated. On this basis, the aim of this study was to investigate if the EOs obtained from inflorescences of hemp may be useful for airways affection management. For the purpose, we selected two different hemp varieties, namely, USO31 and Carmagnola CS and human normal bronchial cells as model of airway inflammation in vitro. Preliminary results obtained in human bronchial cells show that hemp essential oils displayed a moderate cytotoxic effect in a dose-dependent manner, with an IC<sub>50</sub> of  $34.2 \pm 1.1 \ \mu g \ mL^{-1}$  for Carmagnola CS essential oil and  $40.62 \pm 1.8 \ \mu g \ m L^{-1}$  for USO31 essential oil. Then, to evaluate the anti-inflammatory potential of these EOs, we treated bronchial cells with Etoposide that several studies have indicated to trigger cytokines production in human cell lines in vitro [11]. Hemp EOs, used at non-cytotoxic doses, did

not cause an inflammatory state but, additionally, were able to revert an inflammatory condition, reducing the release of the cytokines induced by Etoposide. In conclusion, these preliminary results evidenced a potential use of EOs for bronchial inflammatory diseases.

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## Capsaicin as new adjuvant in anti-cancer immunotherapy



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Red chili (Capsicum frutescens) is widely used as a spice for flavoring foods worldwide. Accumulating evidence has shown multiple pharmacological effects of Capsicum on a variety of physiological systems [1,2]. Pungent capsaicinoids (capsaicin, dihydrocapsaincin), antioxidant vitamins (ascorbic acid, vitamin E), carotenoids (β-carotene, β-cryptoxanthine) and several organic acids and minerals are the major active chemical substance found in Capsicum frutescens [2]. Capsaicin (CPS) (8-methyl-N-vanillyl-6-nonenamide) is an irritant for mammals, including humans. The burning and painful sensations associated with CPS result from its chemical interaction with sensory neurons. CPS, is a derivative of vanillyl amide (8-methyl-N-vanillyl-6-nonenamide). It binds to a receptor called the vanilloid receptor subtype 1 (VR1 or TRPV1) belonging to the Transient Receptor Potential Vanillod subfamily [3]. CPS has been used medicinally for centuries because it can reduce cholesterol, blood lipid content, blood sugar content and it also has properties of antioxidative, anti-inflammatory, anti-obesity and analgesic [4]. At present, it has been demonstrated that CPS exerts anti-cancer activity and immunomodulatory functions [5]. Indeed, the engagement of TRPV1 on dendritic cells by CPS modulates their function by upregulating antigen-presenting and costimulatory molecules, in addition to initiating their migration to draining lymph nodes [6]. Moreover, CPS induces damage-associated molecular patterns of immunogenic cell death in human bladder cancer cells [7]. Bladder cancer has been characterized as a tumor group in which the immunological response is relatively well preserved [8]. A number of immune checkpoint inhibitors have been approved as first-line therapy in case of cisplatin-ineligible patients or as second-line therapy for patients with metastatic urothelial carcinoma of the bladder. However, only 30% of patients with metastatic urothelial carcinoma will respond to this type of immunotherapy [9]. Among the immunocheckpoints, programmed death-ligand 1 (PD-L1) seems to predict response to immune checkpoint inhibitors in such patients. PD-L1 is the transmembrane protein ligand of PD-1, expressed on the cell membrane in T and B lymphocytes, antigen presenting cells and can be induced in tumor cells within the tumor microenvironment. The PD-1/PD-L1 pathway negatively regulates T cell activation, thus it plays an important role in controlling anti-tumor immunity response [10]. Recently, several clinical trials targeting PD-1/PD-L1 pathways using anti-PD-1 or anti PD-L1 antibodies demonstrated the clinical benefit for the patients with bladder cancer and one of these antibodies, atezolizumab, was approved by Food and Drug Administration in USA [11]. Therefore, compounds able to affect the expression of PD-L1 deserve further study. Herein, we evaluated the ability of CPS, at not cytotoxic dose, to influence PD-L1 expression, both at mRNA and protein levels, in 5637 and

T24 bladder cancer cell lines. We found that the exposure of both cell lines to CPS at 50  $\mu$ M for 12 h and 24 h can increase the expression of PD-L1 evaluated by RT-PCR, western blot and immunohystochemistry analyses. Preliminary results showed that the mechanism involved in CPS-mediated upregulation of PD-L1 expression is ROS-independent and TRPV1-mediated. Moreover, the involvement of interferon 1/ 1, through the activation of the NME/NM23 nucleoside diphosphate kinase 4 (NME4)/STATs signaling pathway has been evaluated. Further studies *in vivo* and *in vitro* are definitely required to completely address the CPS immunomodulatory capability in bladder cancer and to elucidate the related anti-tumor effects.

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# A new HPLC-MS/MS method for the simultaneous determination of 36 polyphenols in blueberry, strawberry and their commercial products and determination of antioxidant activity



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Berry fruits consumption has increased in recent years because they are rich sources of polyphenols with reported health benefits [1,2]. Only few reports described the quantitative determination of few phenolic compounds by HPLC-MS/MS, and they only examined anthocyanins or phenolic acids [3,4]. The aim of the present work was to develop a new comprehensive and fast HPLC-MS/MS method for simultaneous determination of 36 phenolic compounds (7 anthocyanins, 9 flavonols, 4 flavan-3-ols, 2 dihydrochalcones, 2 flavanones and 12 phenolic acids) present in blueberry, strawberry, and their commercial products; and to evaluate their antioxidant activities. To our knowledge, this is the first time that HPLC-MS/MS was used for simultaneous quantification of all these compounds of interest taken together in strawberry and blueberry fruits. Blueberry fruits showed higher contents of anthocyanins, flavonols and phenolic acids, while strawberry fruits exhibited higher contents of flavan-3-ols, dihydrochalcones and flavanones. Anthocyanins were the main phenolic constituents in both berries. Furthermore, the higher total phenolic content in the blueberry fruit and jam justified their greater antioxidant capacity measured by DPPH free radical assay [5], compared to strawberry. In conclusion, this new HPLC-MS/MS method is useful and reliable for quality control and authentication analyses of blueberry and strawberry fruits and their commercial food products, such as jams.

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## Sea fennel (*Crithmum maritimum* L.): a culinary herb with nutraceutical potential. Extraction, purification and chemical characterization of polar extracts



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Crithmum maritimum L., also known as sea fennel, rock samphire and locally as 'paccasassi', is a perennial halophyte, thrives on maritime rocks and sometimes in sandy beaches along the Mediterranean, Pacific and Atlantic coasts. Its potential for commercial cultivation has not yet been fully exploited though it is considered a promising biosaline crop [1]. Sea fennel has been used since ancient times in folk medicine as antiscorbutic, tonic, carminative, diuretic, depurative and vermifuge. Today it is believed as an emerging edible vegetable crop, famous for its culinary and not culinary properties [2]. C. maritimum has lots of food uses due to its aromatic traits, for instance to make pickles and to flavour and season soups, sauces and salads. The plant is reported in the Belfrit list, including plants that could be used for food supplements due to sufficient scientific evidence. The use of the plant in the diet is also confirmed by its presence in the "List of Traditional Agri-Food Product" by the Italian Ministry of Agriculture as a traditional agri-food product of Puglia [3]. According to the medicinal use, sea fennel is also considered a source of human health compounds, namely fatty acids such as linoleic (C18:2 w6) and linolenic (C18:3 w3) acids [4], and phenolics compounds with well documented antioxidant effects [5]. Due to the growing interest in this vegetable as a source of nutraceutical compounds [6], different studies have been performed to understand the total chemical profile of the vegetable. This work aims to provide a complete quantitative and qualitatively overview of the extract obtained from the aerial parts of C. maritimum. For this purpose: aerial parts were dried, ground and extracted by percolation with ethanol 70% at 60°C. The purification was done using an Amberlite® XAD7HP sorbent resin. For the quantification, HPLC-DAD-MS/MS system, equipped with electrospray ionization (ESI) and a Zorbax ODS (250 x 4.6 mm, 5 µm) column, was used. The extract was mainly composed of phenolic compounds, among which hydroxycinnamic acids and flavonoids were the two main chemical classes. Among the former, chlorogenic acids, including 5-O-caffeoylquinic acid (32.04-166.21 mg g<sup>-1</sup>), 3,5-di-O-caffeoylquinic acid (8.44-38.86 mg  $g^{-1}$ ) and 4,5-di-O-caffeoylquinic acid (5.70-25.85 mg  $g^{-1}$ ), were the most abundant ones. Among flavonoids, rutin (1.60-4.33 mg g<sup>-1</sup>) and kaempferol-3-O-rhamnoside (0.07- $0.33 \text{ mg g}^{-1}$ ) were the main constituents. These results support the use of this plant as a functional food or in nutraceuticals due to the abundance of antioxidant compounds.

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# Quality of wholemeal pasta made with pigmented and ancient wheats



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Pasta, one of the most popular products in the world, has been subjected in recent years to an important revitalization of its ingredients to respond to the market request for "new", healthier and more sustainable products. Ancient wheat varieties have played an important role in this innovation as they are expected to have higher nutritional quality, to increase biodiversity, and to be more sustainable than a traditional wheat variety [1]. Pigmented wheats have also gained attention due to their antioxidant properties [2]. The objective of this study was to investigate consumer's acceptability and the chemical and physical quality of wholemeal pasta made with two different wheat varieties: an ancient wheat, called Senatore Cappelli (SC), and a pigmented wheat, named Granomischio (GM). SC and GM wholemeal pastas ("paccheri" shaped) were acquired by the same producer (Pastificio Marella Srl, Gioia del Colle, BA) to ensure uniformity of production and were characterized for consumers' acceptability with a 60 consumers' panel. The panelists tested the products in a domestic setting (home use test) and responded to an on-line questionnaire with the aim to evaluate their perception of pasta attributes (colour, flavour, taste, texture and overall acceptability) and their products' preference. Pastas were also characterized for cooking quality at optimal cooking time (solid loss, water uptake, dimensional changes, hardness, adhesiveness [3]), as well as for antioxidant activity (AOA) and total polyphenolic content (TPC) [4]. SC and GM paccheri were very well received by consumers who judged the products very positively with an average of 6.7 (SC) and 6,9 (GM) points on a 9 scale (8-9: 53%SC and 60%GM), and willingness to buy the product in the future of 65% (SC) and 61% (GM). Colour was one the major drivers of liking for both types of pasta (90% liking in GM and 92% in SC). GM drivers of satisfaction were appearance (95%), texture (85%), with an overall satisfaction of 75% of the total number of consumers. SC was very well perceived for appearance, taste and smell, while texture was less liked (55% found product too hard), but this did not hinder overall product appreciation (70%). Finally, comparing the two samples more than the 63 % of the tested population GM over SC. During cooking SC was found to lose less solids (5.0±0,3% vs 5.9±0,1% g solids/100g water) in the cooking water as compared to GM. Cooked SC pasta was found to have higher moisture content (48.9±0.4% vs 51.8±0.47% g water /100 g product), to be slightly harder (35.3  $\pm$ 2.5N vs 31.4 $\pm$ 2.2N) and more adhesive (2.3 $\pm$ 1.0 10<sup>-3</sup> J vs 2.7  $\pm$  1.4 10<sup>-3</sup> J) than GM pasta. Cooked SC had also lower TPC (28.6±1.3 vs 59.3 ± 1.0 mg gallic acid equivalents/100 g of dry weight) and AOA ( $10.6 \pm 8.1$  vs  $79.7 \pm 1.0$  mg trolox equivalent/100 g of dry weight) than GM. This work showed that wholemeal pasta produced with ancient and pigmented

wheat varieties was very well perceived by consumers and had a good cooking quality. Pigmented wheat conferred to product interesting levels of polyphenols and antioxidant activity higher than the unpigmented variety, that may have also an impact on consumers wellbeing.

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## Characterization of milk proteins based nano-carriers: interactions with polyphenolic molecules of nutraceutical interest



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Milk proteins are able to bind or encapsulate hydrophobic and amphipathic substances behaving as nano-carrier systems for bioactive substances with nutraceutical properties. Among the milk proteins,  $\beta$ -lactoglobulin ( $\beta$ -LG) is an excellent candidate for nanoparticle formation, acting as transport vehicle for bioactive compounds such as vitamins, antioxidants, fatty acids, which when bound or encapsulated increase their stability and bioavailability [1]. Resveratrol, as well as many plant derived polyphenolic compounds, although poorly soluble in water, is transported by human serum albumin to the tissues, where it can exert its bioactivity (antioxidant, anti-inflammatory, and anticancer activity). Designing safe and cheap systems to encapsulate resveratrol in a functional food can be a suitable strategy to increase its bioavailability. In the 3D structure of β-lactoglobulin it is possible to identify the main site for specific hydrophobic ligand binding in the β-barrel delimited internal cavity, where, buried within the calyx, the Trp19 is positioned [2]. Exploiting Trp intrinsic fluorescence we have monitored the ability of cow's milk β-lactoglobulin to form complexes with resveratrol and other polyphenolic compounds, in order to elucidate the binding and self-assembly properties of this whey protein as a carrier system of bioactive molecules. At pH 6.4 the binding affinity for the formation of the complex between  $\beta$ -LG and resveratrol has been determined ( $Kd = 4.52 \pm 1.16 \mu M$ ) and resulted to be higher with respect to other tested bioactive molecules like trigonellin (no measurable binding), caffeine ( $Kd = 43.78 \pm 3.99 \,\mu M$ ), and chlorogenic acid ( $Kd = 10.41 \pm 3.8 \,\mu M$ ). Binding of resveratrol by β-LG resulted to be protective with respect to *tran-cis* isomerization process to which resveratrol is subjected (mainly under UV-Vis light), which can have a detrimental impact on its biological activity [3]. Nano-carriers based on milk proteins can represent suitable and safe systems providing food with micronutrients of great potential on health.

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## PROTECTIVE EFFECTS OF R (+)-THIOCTIC ACID TREATMENT ON THE HEART OF SPONTANEOUSLY HYPERTENSIVE RATS



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Hypertension represents a multifactorial disease characterized by elevated blood pressure. It is a risk factor for cardiovascular diseases [1,2]. Increasing evidence attributed the main role of oxidative stress for what concerns hypertension and cardiovascular damage mechanisms because it promotes endothelial dysfunction, vascular remodeling, and inflammation, leading to vascular damage [3]. Excess bioavailability of reactive oxygen species (ROS) often accompanied hypertension-induced structural abnormalities on mitochondria of the cardiomyocyte [4]. Even if the data of clinical trials are discordant, for the management of hypertension, the supplementation of natural antioxidants could also be a promising therapeutic tool with the administration of antihypertensive drugs [5]. Thioctic acid (TIO) is an antioxidant existing in nature and expressed in two optical isomers. R(+)-(TIO) is the naturally occurring eutomer, whereas the most used formulation of the compound in clinical practice is the mixture of (+) and (-)-enantiomers. Previously, we demonstrated that the TIO treatment reduced oxidative stress and prevented adhesion molecule expression in the cardiac vascular endothelium of spontaneously hypertensive rats (SHR) [6]. The present study was designed to investigate the effect of treatment with R (+)-TIO on the heart of SHR, using immunochemical and immunohistochemical techniques. R (+)-TIO (125 µmol/Kg/day) was administered intraperitoneally for 30 days in 24-weeks-old SHR. Hypertensive rats were compared to age-matched normotensive Wistar Kyoto (WKY) rats. After 4 weeks of treatment with R (+)-TIO, blood pressure values were significantly decreased in SHR. This is possibly related to the effects at the levels of the endothelial vessels that determine vasodilation. The results confirmed that the oxidative mechanisms of organ alterations due to hypertension were characterized by left ventricular cardiomyocytes hypertrophy, fibrosis, proteins oxidation accompanied by inflammation with an increased expression of IL-1 beta, IL-6, and tumor necrosis factor-alpha. R (+)-TIO reduced the development of oxidative stress, cardiomyocyte hypertrophy, and inflammatory processes. Taken all together, R (+)-TIO may be considered as one of the antioxidant candidate molecules to prevent cardiac alterations associated with hypertension. The effects observed after treatment with R (+)-TIO could open new perspectives for countering heart injury, which represents a common feature in hypertensive end-organs damage.

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## *In vitro* assessment of prebiotic effect of coffee and its byproducts on human intestinal microbiota



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The human intestinal microbiota has long been known to influence human health and disease, but recently, its improvement by modulation of gut microbiota is an upcoming evolving strategy [1]. The microbiota present in the human body provides numerous benefits, including enhancement of overall well being and also protection against pathogenic microorganisms. Gut microbiota exerts its protective role through the production of different signaling molecules which can modify the functionality of other organs. When the gut bacterial composition is altered (dysbiosis), there is a strong association with the pathogenesis of many inflammatory diseases and infections. Prebiotics supplementation could be a good strategy to improve intestinal microbiota of the target host, conferring health benefits. Currently, other substances beside carbohydrates, i.e. polyphenols, might fit the updated definition of prebiotics presenting evidence of beneficial effects in the host [2]. Coffee and its by-products are very interesting substrates in this sense, because they contain plenty of polyphenols and other essential compounds like melanoidins and chlorogenic acid, which make them potential prebiotic food components. The aim of the present study was to characterize and assess the potential prebiotic properties of seven coffee by-products, studying in vitro the modifications on human gut microbiota and its metabolic products. The coffee by-products, all from Coffea arabica, were selected and studied to evaluate compounds and bioactive substances, responsible for their prebiotic activities on the intestinal microbiota. The selection of coffee by-products was made of: Green Coffee Bean (coarsely ground), Silver Skin (generated from the roasting process of the bean), Spent Coffee Ground (generated from the coffee preparation), two extracts from Silver Skin and from Spent Coffee, finally the Roasted Coffee and an Espresso freshly made. A fermentation system (batch culture) in anaerobic conditionswas used to mimic the human intestine. During all fermentation cycles, the biodiversity of gut microbiota was analysed and selected bacterial groups were enumerated using Real-Time PCR analysis. At the same time points the detection and quantification of short chain fatty acids (SCFA) was also performed. Inulin was chosen as well-defined prebiotic substrate, to compare the results obtained from all the considered coffee substrates. The general trend is an increase in the bacterial counts of beneficial bacterial groups (Lactobacillus spp. and Bifidobacterium spp.), and a decrease for Clostridium coccoides-Eubacterium rectale group. These data are supported also by the SCFA that resulted to accumulate during fermentation especially acetic and propionic acids. The valorisation as prebiotics of these by-products highlights that their recovery from the coffee industry is of great importance, not only because of their significant properties, but also because it could exploit a part of industry wastes.

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# Water- and fat-soluble vitamins in milk after different thermal treatments and steam injection



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To prepare a good cappuccino, it is preferable to use fresh whole milk as its high-fat content contributes to the flavour and its protein content is useful for a good whipping of the milk contributing to its creaminess. The foam is produced by the nozzle of the coffee machine, which is necessary to obtain a compact cream, but it is also important that the temperature of the milk must not exceed 65-70°C, to avoid an unpleasant aroma and the lack of foam formation. Furthermore, the nutritional quality of the milk should remain unchanged after the steam injection treatment. In the present work, we focused the attention on the water- and fat-soluble vitamins content in pasteurized bovine milk samples (HQ-milk) treated at different temperatures (40-80°C) and steam injected pressure using a nozzle of Simonelli Group coffee machine. The treatment time was 20, 30, 40, 50, and 60 seconds. Water- and fat-soluble vitamins content on the treated HQ-milk has been determined by RP-HPLC using a C18 column (HiQSil C18 HS, 5 µm, 4.6 mm i.d, 250 mm, Kya Tech Corporation), connected to an HPLC Agilent 1260 Infinity LC System (Agilent Technologies), according to two different protocols from Albalá-Hurtado and co-workers [1, 2]. Quantification was achieved by a calibration curve obtained relating the concentrations (mM) of each vitamin standard loaded in the column to the peak area corresponding to each concentration. The results showed that if the only temperature is applied, vitamin B<sub>2</sub>, nicotinic acid, and nicotinamide were not affected by the increasing temperature because the exposure time from 20 to 60 seconds is too short to cause a decrement in the vitamins content. However, when the HQ-milk is treated with increasing temperature together with steam injection, an interesting result regarding the nicotinic acid concentration was obtained. In fact, the concentration of nicotinic acid increases by about 6.5 folds starting from 7.5 mM at 25°C to reach a mean value of 45 mM at a temperature range of 50-70°C. It is also interesting to note that the increase of nicotinic acid concentration is time-dependent: if the HQ-milk is treated for 20 seconds with the steam injection (time normally used to heat the milk during the preparation of the cappuccino), nicotinic acid reaches its highest concentration (54 mM) at 70°C, but if the milk is treated for 60 seconds, the highest concentration (43 mM) is reached at 50°C. Nicotinic acid (Niacin) is used to prevent and treat niacin deficiency that may result from alcohol abuse, malabsorption syndrome, Hartnup disease, poor diet. Furthermore, nicotinic acid is used to treat dyslipidemias since reduces triglycerides and LDL cholesterol and raises HDL cholesterol levels [3]. In our experiments, we observed that the increase in nicotinic acid dependent on the steam injected and the treatment time is accompanied by a simultaneous decrease in quinolinic acid. The latter is generated within the kynurenine pathway resulting from L-tryptophan catabolism and may be present in the milk since it is related to the composition of the cows' diet [4]. Quinolinic acid may be converted to nicotinic acid by a decarboxylation reaction induced by the temperature and the steam injected pressure [5]. However, this hypothesis needs to be further confirmed. On the contrary, nicotinamide and  $B_6$ decrease when temperature and steam injection increased, whereas vitamin  $B_2$  resulted not affected by the treatment. Among fat-soluble vitamins, the results showed that vitamin E and vitamin  $D_3$  did not undergo variations after temperature and steam injection treatment while vitamin A was slightly affected. In conclusion, these results indicate that the nutritional quality of HQ-milk, from the point of view of the concentration of vitamins, seems not influenced by the thermal treatment and by the injected steam. It was observed, after this treatment, an increase in milk nicotinic acid content, very interesting considering the beneficial effects on human health exerted by this vitamin.

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## The effect of chickpea flour addition levels and flour structures on quality and in vitro starch digestibility of cornrice-based gluten free pasta



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Gluten-free (GF) pasta in the Italian market is produced using corn and rice starch as main ingredients [1], resulting in a product rich in available carbohydrates with high postprandial glycemic response [4], low in protein and fiber, and with poor cooking quality [2-3]. Inclusion of chickpea flour into GF pasta formulation may positively affect product's nutritional profile by increasing protein and fiber, and reducing the glycemic impact while increasing environmental sustainability [5-10]. Chickpea flour pre-treatment (e.g. pre-gelatinization, milling degree) may affect both cell integrity and degree of starch gelatinization that may have a profound impact of both product's quality and nutrients accessibility [11-14]. The present work aims to produce corn-rice-chickpea GF pasta to investigate the effect of different chickpea addition levels and chickpea flour structural attributes on product's cooking quality and in vitro starch digestibility. Pasta was produced with heat-treated white corn and rice flours at a mass ratio of 3:2 in all formulations, while conventional chickpea flour was included at level of 0%, 25%, 50%, 75% and 100% to investigate the effect of chickpea addition levels, and the effect of chickpea flours structural attributes (pre-gelatinization, milling degree [fine and coarse]) was tested at 50% inclusion. Nutritional facts, cooking quality, texture quality, and starch digestibility of products were investigated. GF pasta produced with increasing chickpea four additions showed a high increase in protein and fiber content, a shorter optimal cooking time and reduced solid loss during cooking as compared to the control corn-rice. Color of chickpea pasta shifted towards brown/dark, while texture was found to be softer and less adhesive. More importantly, chickpea addition significantly reduced the content of total starch, available starch, rapidly and slowly digestible starch but increased content of resistant starch which suggests a reduced glycemic load and glycemic index in vivo. However, no significant differences were found in respect to chickpea flour structural attributes. Current results demonstrate that chickpea flour is able to impart corn-rice-based GF pasta with desired quality and attenuated starch digestibility. Flour structural attributes had no significantly influence on GF pasta quality and in vitro starch digestibility.

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# Managing of the daily calcium dietary intake as a tool to contribute to a healthy lifestyle



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Calcium is an essential element for human health for growth, maintenance and repair of bone tissue, muscle contraction, nerve conduction and blood clotting [1,2]. The Reference Levels of Nutrients and Energy Intake for the Italian Population defines the recommended daily requirements depending on specific needs related to factors as age, gender, pregnancy and menopause [3]. In athletes, calcium intake is also important to decrease the risk of low bone-mineral density and stress fractures [2]. However, studies conducted on the Italian population reported intake ranges often lower than those recommended [4,5]. Also in athletes, inadequate intakes have been reported both in male and female [6,7]. In this regard a questionnaire was proposed to 503 subjects, aged 12-85, to evaluate their calcium intake. It was observed that 81% of female and 74% of male do not reach the recommended values. The highest percentages were obtained in female aged 15-17 (89%) and in male and female older than 60 years (83%). In physically active subjects (63% of the studied population), the 76% shows inadequate calcium intakes. This percentage results close to the one of sedentary that do not reach recommended values (80%). Calcium intake deficiency is common in all age stages both in sedentary and in physically active people. In this context, right foods choice represents a fundamental aspect. Milk and dairy products are the most relevant source of dietary calcium in terms of both quantity and bioavailability [8]. By evaluating daily calcium intake in a group of non-elite athletes using 500mL of skimmed milk as post-exercise recovery drink, it was possible to observe that calcium intakes were in line with recommended values. Moreover, an improvement in body composition and, in particular, on hydration status was highlighted. This underlined milk's role as recovery drink but also as a coadjutant to improve calcium intake. Thus, these findings support the necessity of design specific dietary interventions that should start at early ages and continue through adulthood. Strategies should be developed to sensitize on calcium importance and to improve its daily intake. Right food choices should be considered taking into account naturally calcium-rich foods and novel calciumenriched products.

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# Vitamins in human, cow, and donkey milk: a comparison of nutritional properties



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Donkey milk (DM) has been used since ancient times in the feeding of infants who, for various reasons, could not be breastfed. Furthermore, the health benefits of DM have been tested in several clinical trials for the treatment of debilitated and convalescent subjects [1]. Today we know that DM is a valuable source of nutraceutical substances beneficial to human health, including bioactive peptides, essential fatty acids, and vitamins [2]. The present work focuses precisely on this last aspect, the identification and determination of the water- and fat-soluble vitamins in DM and their comparison with the vitamins content in human milk and cow's milk. Donkey and cow's milk watersoluble vitamins were determined by a chromatographic method [3], using an HPLC Agilent 1260 Infinity LC System (Agilent Technologies), connected to a C18 HiQSil HS column (5 µm, 4.6 mm i.d, 250 mm, Kya Tech Corporation, Tokyo, Japan). The obtained results in cows' and donkey milk were compared to the vitamins content of human milk taken from the published literature. DM is a good source of nicotinic acid (18.75  $\pm$  1.92  $\mu$ M), this value is higher compared to human milk (4.64  $\mu$ M, [4]) but lower than the one found in cow's milk in the present work (27.52±4.91). Nicotinic acid (niacin) is a member of the B3 vitamin group that reduces the release of fatty acid from the adipocytes, thus showing a lipid-lowering effect, this molecule is used to treat the progression of atherosclerotic cardiovascular disease. Folic acid (vitamin B9) is present in DM with a concentration  $(0.83 \pm 0.08)$  $\mu$ M) higher than human milk (0.365  $\mu$ M) and about half of that found in cow's milk in our determinations (1.93  $\pm$ 0.53  $\mu$ M). Milk represents an excellent source of folic acid that is involved in nucleic acid synthesis, accelerates cell division, and for this reason, very important for children's growth. Pyridoxine (vitamin B6) content is  $5.38 \pm 3.35 \mu$ M, this value is very similar to that found in bovine milk both from our determinations ( $2.5 \pm 0.61 \mu$ M) and in the literature ( $1.0-11.23 \mu$ M; [5]). Vitamin B6 is involved in several metabolic processes such as lipid and amino acid metabolism, gluconeogenesis, the regulation of the immune system, and the development of the nervous system. Donkey milk also contains thiamine ( $0.66 \pm 0.26 \mu$ M), riboflavin ( $0.168 \pm 0.04 \mu$ M), with levels very close to those found in the literature for bovine milk [5] but higher compared to human milk [4]. Very interesting is the content of vitamin C, a vitamin with well-known antioxidant properties which in donkey milk has a value of 0.32 mM, similar to the value found in human milk (0.28 mM) but much higher than the quantity found in cow's milk (28.4 -170.3  $\mu$ M). Among fat-soluble vitamins, it is known by the literature that the vitamin D2 and D3 content in donkey milk (0.042 µM and 0.016 µM, respectively) is higher than the values found for this vitamin in human and bovine milk (0.008  $\mu$ M) [6]. Furthermore, it has been shown that vitamin D levels in donkey milk are related to seasonality since they show the highest value in summer due to the greater sun exposure of the animals [6]. Donkey milk vitamin E amount is about 3.4  $\mu$ M [6], higher than the value found in the literature [5] in bovine milk (1.4  $\mu$ M, respectively), but lower concerning the amount found in the human milk (8.0±3.8  $\mu$ M). Vitamin E is a potent antioxidant that neutralizes free radicals, protecting tissues and organs from oxidative damage. These results show that DM, besides the presence of the well-known nutraceutical compounds, is also a valuable source of vitamins, indicating that this milk can represent a functional food for infants but also for elderly consumers.

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## Evening Primrose Oil effects in human Pancreatic Ductal Adenocarcinoma cell lines



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Evening Primrose (Oenothera L.) is a medicinal plant and the most numerous species in the Oenothera L. family is the Oenothera biennis. Evening Primrose Oil (EPO), obtained from its seeds, is rich in triacylglycerols and the main ones are linoleic acid (LA) (70–74%) and  $\gamma$ -linolenic acid (GLA) (8-10%), polyunsaturated fatty acids (PUFAs), belonging to omega-6 acids group [1]. The presence of GLA makes EPO a valid dietary supplement to reduce triacyclglycerols and low-density lipoprotein (LDL) in plasma and an interesting compound for the treatment of rheumatic and arthritic conditions and for atopic dermatitis [1,2]. LA and GLA are precursors of dihomo-linolenic acid (DGLA), that is metabolized in series 1 prostaglandins and is oxidized in 15-hydroxyeicosatrienoic acid (15-HETrE) with anti-inflammatory and anti-proliferative effect, but at the same time, they can be also converted in arachidonic acid (AA), inducing pro-inflammatory series 2 prostaglandins and series 4 leukotrienes [1]. EPO is also composed by unsaponifiable compounds and among these, the sterols have important role in the reduction of pro-inflammatory mediators [2]. Moreover, several studies demonstrated GLA antitumoral activities in different cancer cell types and its ability to enhance chemotherapeutic drugs activity [3]. Pancreatic Ductal Adenocarcinoma (PDAC) is an infiltrating neoplasm with glandular differentiation derived from the pancreatic ductal tree. Among the solid cancers, it is considered one of the most aggressive, with five years survival of 5-7% [4-6]. It was demonstrated a family genetic predisposition, but also some precursor lesions within pancreatic tissue and somatic mutations of KRAS oncogene and CDKN2A, TP53 and SMAD4 suppressor genes are implicated in PDAC pathogenesis [4,7,8]. In the last years scientific interest in using molecules of natural origins as adjuvant in cancer therapy is grown [9]. Several phytochemicals demonstrated anticancer properties, thanks to their effects in suppressing cancer cell survival and proliferation, in inducing cancer cell death and for their synergistic/adjuvant interactions with different molecular pathways targets of chemotherapeutical drugs. [10]. Particularly, a study evidenced a reduction of pancreatic cancer cells growth, migration and invasion by treatment with DGLA [11]. So, since LA and GLA are the most important triacyclglycerols present in EPO, we decided to analyze the effects of EPO in two human pancreatic ductal adenocarcinoma cell lines (PANC-1 and MIAPaCa-2). Data showed a reduction of cancer cell vitality after administration of EPO and an induction of cancer cell death as confirmed by Western Blot analysis. Then, the effects of EPO were also evaluated on its ability to modulate the main pathways involved in PDAC aggressiveness. In conclusion, these preliminary data suggest a potential role of EPO in PDAC aggressiveness.

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# Can probiotics counteract the detrimental effects of sleep deprivation?



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Sleep is a key physiological process in maintaining health and wellbeing and it is considered an important modulator of the immune response. Although sleep exact function is not fully elucidated, an increasing number of studies has focused on the evaluation of the detrimental effects of sleep deprivation. This condition leads to a range of brain deficits, including impaired memory and concentration, and increased risk for neuropsychiatric disorders. The brain responds to sleep loss with rapid and extensive changes at molecular, biochemical, electrophysiological, and structural levels. Research in humans and rodents has repeatedly described low-grade inflammation in association with sleep loss, as reflected by increased inflammatory cytokines and by the presence of other markers of cerebral inflammation and oxidation [1-3]. Moreover, the impact of sleep deprivation on gut microbiota composition and the role of gut dysbiosis in triggering both peripheral and central inflammatory processes have recently emerged [4]. Considering that probiotics have well documented anti-inflammatory and antioxidant properties [5,6] and that supplementation with Lactobacilli improved sleep quality [7], the aim of the present study is to evaluate the ability of chronic probiotic treatment to counteract the deleterious effects of insufficient sleep by reducing neuroinflammation and oxidative stress in an animal model of sleep deprivation. In detail, 8-weekold B6129SF2 mice were chronically administered with SLAB51 multi-strain formulation or water. Upon treatment, the animals were sleep deprived for one week and sacrificed. Plasma and brain inflammatory and oxidative markers were measured through immunometric assays. Interestingly, a decreased expression of ionized calcium-binding adaptor molecule 1 (Iba1) and a reduction of proinflammatory cytokines, together with decrease of dityrosine and nytrotirosine residues in the brain of sleep deprived mice treated with probiotics were detected, demonstrating an amelioration of both inflammatory and oxidative status in treated animals. These data strengthen the link between gutmicrobiota and sleep quality and support the use of probiotics as a potential aid against the negative consequences of insufficient sleep.

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