Sambucus ebulus L. Fruit Extract

Subjects: Plant Sciences Contributor: Oskan Tasinov

Sambucus ebulus L. (SE), also known as dwarf elder or dwarf elderberry, is a widely used as wound-healing, antinociceptive, anti-rheumatoid, anti-influenza, antibacterial and diuretic medicinal plant in Bulgaria, Turkey, Iran, Lebanon, Romania and Bosnia–Herzegovina.

lipopolysaccharides macrophages	Sambucus ebulus L.	phytochemical composition	anti-inflammatory	ER stress
	lipopolysaccharides	macrophages		

1. Introduction

Traditional medicine is a good source of knowledge about therapeutics, which are consequently researched and successfully implicated in modern pharmaceutical preparations. *Sambucus ebulus* L. (SE), also known as dwarf elder or dwarf elderberry, is a widely used as wound-healing, anti-nociceptive, anti-rheumatoid, anti-influenza, antibacterial and diuretic medicinal plant in Bulgaria, Turkey, Iran, Lebanon, Romania and Bosnia–Herzegovina ^[1] ^{[2][3][4][5]}. Fresh fruits, jam, tea or decoction of SE fruits are used as immunostimulating and hematopoietic herbal preparations, as well as for the treatment of rheumatoid arthritis and gastrointestinal disorders ^{[1][2][6]}. The number of modern studies focusing on SE biological activities are growing, but there is still insufficient knowledge regarding molecular mechanisms of action of fresh or dry fruits and various fruit extracts.

Only ripe fruits are used in traditional medicine recipes and the chemical content varies depending on the types of the extract ^{[3][7]}. Data from phytochemical analyses in the literature reveal that SE fruits are high in polyphenolics, especially anthocyanins and proantchocyanidins, phenolic acids, hydroxycinnamic acids, flavonol glycosides, as well as organic acids, tannins, pectins, resins, vitamin C, volatile substances (eugenol, valeric acid, citronellal etc.), amino acids (including some essential ones), and plant sterols ^{[3][7][8][9][10][11][12][13][14][15][16]}. Many chromatographic analyses of SE fruit extracts have been carried out up to date, and, still, the information about the presence of certain specific organic compounds remains unclear, especially with regard to soil characteristics, variety of extragents used for sample preparation. Therefore, a detailed phytochemical analysis could be useful, especially in examining the molecular mechanisms of SE fruits on human health.

Numerous studies have established the strong in-vitro antioxidant activity of SE fruit extracts, analyzing its iron chelating, NO radical scavenging, and ABTS cation radical decolorization activity, and their interrelations with polyphenolic and anthocyanin content ^{[3][7][8]}. The presence of different functional groups in polyphenolics and organic acids found in the tested SE fruit extracts is considered to determine, to a great extent, their antioxidant

and anti-inflammatory activities. In oxidatively challenged 3T3-L1 preadipocytes, SE fruit aqueous extract (FAE) acts as modulator of antioxidant genes' transcription ^[17]. In macrophages treated with ethanol- or lipopolysaccharides (LPS), SE FAE suppresses the ethanol- and LPS-stimulated transcription of glutamate– cysteine ligase, glutathione peroxidase and nuclear factor kappa B (NFkB) ^{[9][18]}. Acetone extracts, hydrophilic and anthocyanin-rich fractions of SE fruits possessing high in-vitro antioxidant activity protect macrophages from the oxidative stress-mediated cytotoxicity caused by *tert*-Butyl hydroperoxide ^[19]. Ethyl acetate fraction of SE fruits possesses cytoprotective and anti-inflammatory activity reducing ethanol-induced cell death, proinflammatory gene transcription in macrophages ^[9]. Methanolic extracts of SE fruits reduce carrageenan-induced paw edema in rats ^[20]. Others describe the antiemetic, neuroprotective and anti-herpes-simplex-virus activities of SE fruit extracts ^[12].

In an intervention study on healthy adult volunteers, SE fruit tea enhances serum antioxidant potential, improves lipid profile ^[22], decreases serum CRP, IL-1 β , leptin and adiponectin levels ^[23], thus indicating an immune- and fat metabolism-modulating activity. A clinical trial reported the effectiveness of SE fruit ethanol extract for the treatment of paederus dermatitis, proving its anti-inflammatory and wound healing potential ^[24].

LPS-stimulated macrophages are widely used in-vitro models for testing anti-inflammatory activity of medicinal plant extracts. The macrophages are source of a variety of pro-inflammatory cytokines, chemokines, and may act in a paracrine and endocrine mode. In low grade inflammation, such as in adiposity, where the activation of chemokine release is associated with macrophage recruitment and unlocking a self-feeding inflammatory process that leads to such complications as insulin resistance and related atherosclerosis ^[25]. The released cytokines and chemokines, such as TNF α , IL-6, IL-1 β , NO, as a product of iNOS, activate signaling pathways mediated by Jun N-terminal kinase (JNK), the inhibitor of κ B-kinase (IKK) β and other serine kinases ^{[25][26][27][28]}, and resulting in NF κ B activation. The latter stimulates the transcription of pro-inflammatory genes ^[29].

Along with the protein synthesis, endoplasmic reticulum (ER) plays an important role in sensing nutrients and responds to different stress conditions by activating the unfolded protein response and subsequently implicating it into insulin resistance and cardiovascular diseases ^{[30][31]}. ER stress can promote inflammation, and vice versa ^[32]. ER stress-related inflammation could be mediated by iNOS ^[34]. Therefore, the enzyme iNOS as a cross point of inflammation and ER stress could be a possible therapeutic target.

There are data that ER stress and inflammation in different pathological conditions could be reduced by compounds such as resveratrol ^{[35][36]}, epigallocatechin gallate ^[37] and proanthocyanidins found in herbal extracts ^[38]. SE fruits, being rich polyphenolics, anthocyanins and stilbenes, could be effective in combating ER stress and inflammation.

2. Phytochemical Content and Composition

Among the phytochemical compounds identified in the tested SE FAE 15 amino acids (AAs), 10 organic acids (OAs), 36 sugar acids and alcohols, 25 mono-, di- and trisaccharides, 13 fatty acids (saturated and unsaturated)

and their esters (Table 1), and 38 phenolic compounds were detected and quantified (Table 2).

Compound	Content, µg/mL
Amino Acids	
L-Valine	3.02 ± 0.21
L-Leucine	8.06 ± 0.56
L-Isoleucine	8.48 ± 0.59
L-Proline	20.01 ± 1.40
L-Threonine	3.89 ± 0.27
L-Phenylalanine	10.25 ± 0.72
L-Lysine	4.37 ± 0.31
Glycine	3.78 ± 0.26
Serine	2.59 ± 0.18
L-Aspartic acid	16.32 ± 1.14
L-Asparagine	6.19 ± 0.43
L-Glutamic acid	1.34 ± 0.09
L-Glutamine	22.99 ± 1.61
DL-Ornithine	12.36 ± 0.86
L-Tyrosine	2.66 ± 0.19
Organic Acids	
Succinic acid	12.64 ± 0.88
Fumaric acid	6.61 ± 0.46
Malic acid	9.22 ± 0.65
Pyroglutamic acid (5-oxoproline)	33.63 ± 2.35
4-Aminobutyric acid	5.69 ± 0.40

Table 1. List of polar phytochemicals identified in the analyzed polar fraction (A) of SE FAE using GC-MS technique. The concentration was given in μ g/mL extract. Results are presented as mean ± standard deviation.

Compound	Content, µg/mL
2-Hydroxyglutaric acid	4.07 ± 0.29
2-Ketoglutaric acid	8.02 ± 0.56
Phenylpyruvic acid	2.18 ± 0.15
2,3-Dihydroxybutanedioic acid	10.49 ± 0.73
Isocitric acid	18.12 ± 1.27
Sugar Acids and Alcohols	
Glycerol	36.12 ± 2.53
Digalactosylglycerol	6.99 ± 0.63
Glyceric acid	17.05 ± 1.19
Threitol	7.66 ± 0.54
Erythreol	2.09 ± 0.15
Erithreonic acid	2.65 ± 0.19
Threonic acid	8.40 ± 0.59
Xylitol	4.20 ± 0.29
Arabinitol	34.65 ± 2.43
Pentonic acid	7.69 ± 0.54
L-Glycerol-3-phosphate	17.72 ± 1.24
Ribonic acid	4.76 ± 0.33
Manitol	2.98 ± 0.21
Sorbitol	49.26 ± 3.45
Glucuronic acid isomer	8.49 ± 0.59
Galactitol	1.91 ± 0.13
Galacturonic acid isomer	15.91 ± 1.11
Glucuronic acid isomer	13.03 ± 0.91
Gluconic acid isomer	1.78 ± 0.12

Compound	Content, µg/mL
Galacturonic acid isomer	2.89 ± 0.20
Glucuronic acid isomer	3.87 ± 0.27
Galactonic acid	6.33 ± 0.44
Gluconic acid isomer	3.71 ± 0.26
Glucaric acid	14.00 ± 0.98
Galactaric acid	3.38 ± 0.24
Myo-inositol	6.71 ± 0.47
Galactosylglycerol	22.50 ± 1.58
Sorbitol-6-phosphate	43.32 ± 3.03
myo-Inositol-1-phosphate isomer	5.64 ± 0.39
myo-Inositol-2-phosphate isomer	7.43 ± 0.52
Gluconic acid-6-phosphate	1.54 ± 0.11
myo-Inositol-1-phosphate isomer	3.30 ± 0.23
myo-Inositol-2-phosphate isomer	6.87 ± 0.48
Maltitol; alpha-D-Glc-(1,4)-D-sorbitol	4.90 ± 0.34
Galactinol isomer; alpha-D-Gal-(1,3)-myo-Inositol	0.69 ± 0.05
Galactinol isomer; alpha-D-Gal-(1,3)-myo-Inositol	3.67 ± 0.26
Saccharides (mono-, di-, and tri-)	
Xylose methoxyamine	5.94 ± 0.42
Arabinose methoxyamine	12.65 ± 0.89
Fructose isomer	14.31 ± 1.00
Fructose isomer	18.89 ± 1.32
Sorbose isomer	28.11 ± 1.97
Sorbose isomer	21.35 ± 1.49
Galactose isomer	35.19 ± 2.46

Compound	Content ug/ml
Galactose isomer	13.86 ± 0.97
Glucose isomer	17.34 ± 1.21
Glucose isomer	13.59 ± 0.95
Fructose-6-phosphate isom	er 16.20 ± 1.13
Mannose-6-phosphate isom	er 3.47 ± 0.24
Galactose-6-phosphate isom	ier 18.79 ± 1.32
Glucose-6-phosphate isome	er 30.27 ± 2.12
Fructose-6-phosphate isome	er 5.81 ± 0.41
Galactose-6-phosphate isom	ner 3.32 ± 0.23
Glucose-6-phosphate isome	er 4.52 ± 0.32
Sucrose; alpha-D-Glc-(1,2)-beta-D-F	Fru isomer 24.81 ± 1.74
Trehalose; alpha-D-Glc-(1,1)-alpha-D-	-Glc isomer 10.10 ± 0.71
Melibiose isomer; alpha-D-Gal-(1,6)-D	0-Glc isomer 18.59 ± 1.30
Melibiose isomer; alpha-D-Gal-(1,6)-D	9-Glc isomer 18.80 ± 1.32
Sucrose; alpha-D-Glc-(1,2)-beta-D-F	Fru isomer 20.55 ± 1.44
Trehalose; alpha-D-Glc-(1,1)-alpha-D-	-Glc isomer 16.13 ± 1.13
Raffinose; alpha-D-Gal-(1,6)-alpha-D-Glc-(1,2))-beta-D-Fru isomer 12.91 ± 0.90
Raffinose; alpha-D-Gal-(1,6)-alpha-D-Glc-(1,2)-beta-D-Fru isomer 25.61 ± 1.79
Saturated, unsaturate	ed acids and esters
9-(E)-Hexadecenoic acid	8.52 ± 0.77
9-(Z)-Hexadecenoic acid	6.57 ± 0.59
Heptadecanoic acid	7.56 ± 0.68
Hexadecatrienoic acid	4.85 ± 0.44
Compound	Content, μg/mL
Anthocy	yanins
Cyanidin-3-O-galactoside (idaein)	382.15 ± 13.19
Cyanidin-3-O-glucoside (chrysanthen	nin) 31.07 ± 1.10
Cyanidin-3-O-arabinoside	85.87 ± 2.80

Compound	Content, µg/mL	
Cyanidin-3-O-xyloside	14.35 ± 0.53	
Proanthocyanidin monomers		
Catechin	40.19 ± 1.33	
Epicatechin	322.37 ± 11.75	
Proanthocyanidin dimers		
$EC \rightarrow EC$ (1)	171.40 ± 6.23	
$EC \rightarrow EC$ (2)	169.24 ± 6.15	
$EC \rightarrow EC$ (3)	189.86 ± 6.90	
$EC \rightarrow EC$ (4)	157.91 ± 5.74	I AAs are
Proanthocyanidin trimers		sted pola
$EC \rightarrow EC \rightarrow EC$ (1)	225.23 ± 8.16	
$EC \rightarrow EC \rightarrow EC$ (2)	242.27 ± 8.78	
$EC \rightarrow EC \rightarrow EC$ (4)	198.92 ± 7.21	
$EC \rightarrow EC \rightarrow EC$ (4)	249.36 ± 9.04	
Stilbenes		
trans-Resveratrol-3-O-glucoside	51.92 ± 1.94	
Cyclohexanecarboxylic acid		
Quinic acid	108.00 ± 4.02	
Hydroxycinnamic acids		
3-O-Caffeoylquinic acid (chlorogenic acid)	567.06 ± 20.55	
Caffeic acid-O-galactoside	98.72 ± 3.58	
Caffeic acid-O-glucoside	74.66 ± 2.71	
5-O-Caffeoylquinic acid (neochlorogenic acid)	906.08 ± 32.84	
p-Coumaric acid-O-glucoside	236.37 ± 8.57	
3-O-p-Coumaroylquinic acid	399.47 ± 14.48	

Compound	Content, µg/mL
Feruloylquinic acid	248.93 ± 9.02
4-O-p-Coumaroylquinic acid	219.83 ± 7.97
Ferulic acid-O-galactoside	131.66 ± 4.77
Ferulic acid-O-glucoside	122.26 ± 4.43
Flavonol glycosides	
Quercetin-3-O-rhamnosyl-galactoside	25.57 ± 0.93
Quercetin-3-O-galactoside (hyperoside)	29.17 ± 1.06
Kaempferol-3-O-galactoside	11.15 ± 0.40
Quercetin-3-O-rhamnosyl-glucoside	20.35 ± 0.74
Quercetin-3-O-glucoside (isoquercetin)	22.80 ± 0.83
Kaempferol-3-O-glucoside (astragalin)	9.94 ± 0.36
Quercetin-3-O-arabinoside (guaiaverin)	16.77 ± 0.61
Quercetin-3-O-xyloside	13.97 ± 0.51
Kaempferol-3-O-rhamnosyl-galactoside	12.52 ± 0.45
Kaempferol-3-O-rhamnosyl-glucoside	9.15 ± 0.33
Kaempferol-3-O-arabinoside	11.15 ± 0.40
Kaempferol-3-O-xyloside	12.80 ± 0.46
Total analyzed polyphenols	5840.50

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anthocyanins content of Sambucus ebulus L. aqueous and aqueous—ethanolic extracts depend Aiming to study the anti-inflammatory action of the aqueous extract of dwarf elderberry under conditions of LPSon the type and concentration of extragent. Sci. Technol. 2012, II, 37–41. stimulated inflammatory response in J774A.1 mouse macrophages, the transcriptional levels of genes coding for proteins mediating and involved in the inflammatory process as well as the translation levels of iNOS were an Ejzeahi Interación of construction of the seconda de la construction. Seconda de la construction of the seconda de la construction of the seconda de la construction of the seconda de la construction. Seconda de la construction developed de la construction of the seconda developed dev

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- SE FAE % v 28. Bandyopadhyay, G.K., (Yurs) Grecio, J.; Olefsky, J.M. Increased p85/55/50 expression and decreased phosphotidylinositol 3-kinase activity in insulin-resistant human skeletal muscle.

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with increasing concentrations (2.5%, 5%, 10% v/v) of SE FAE or with SA for 24 h and subsequently stimulated or 9. Nguyen, M.T.A.; Favelyukis, S.; Nguyen, A.K.; Reichart, D.; Scott, P.A.; Jenn, A.; Liu-Bryan, R.; not with LPS. Results were obtained using qPCR technique. Data are presented as mean ± SEM. Legend: SE Glass, C.K.; Neels, J.G.: Olefsky, J.M. A subpopulation of macrophages infiltrates hypertrophic E-Sambucus ebulus L. fruit aqueous extract; SA-100 µM salicylic acid; LPS-200 ng/mL lipopolysaccharides. adipose tissue and is activated by free fatty acids via toll-like receptors 2 and 4 and JNK-< 0.05, ** p < 0.01, *** p < 0.001 vs. untreated cells; # p < 0.05, ## p < 0.01, ### p < 0.001 vs. LPS. dependent pathways. J. Biol. Chem. 2007, 282, 35279–35292.

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Figure 2. Changes in mRNA levels of *Ccl2* (**a**), *Icam1* (**b**), and *Fabp4* (**c**) in J774A.1 mouse macrophages pretreated with increasing concentrations (2.5%, 5%, 10% v/v) of SE FAE or with SA for 24 h and subsequently stimulated or not with LPS. Results were obtained using qPCR technique. Data are presented as mean \pm SEM. Legend: SE FAE–*Sambucus ebulus* L. fruit aqueous extract; SA–100 µM salicylic acid; LPS–200 ng/mL lipopolysaccharides. * *p* < 0.05, ** *p* < 0.01, *** *p* < 0.001 vs. untreated cells; # *p* < 0.05, ## *p* < 0.01, ### *p* < 0.001 vs. LPS.



Figure 3. Changes in mRNA levels of *COX2* (**a**), *iNOS* (**b**), *Noxo1* (**c**), and of protein levels of iNOS (**d**) in J774A.1 mouse macrophages pre-treated with increasing concentrations (2.5%, 5%, 10% v/v) of SE FAE or with SA for 24 h and subsequently stimulated or not with LPS. Results were obtained using qPCR ((**a**), (**b**) and (**c**)) or western blot technique (**d**). Data are presented as mean ± SEM. Legend: SE FAE–*Sambucus ebulus* L. fruit aqueous extract; SA–100 μ M salicylic acid; LPS–200 ng/mL lipopolysaccharides. * *p* < 0.05, ** *p* < 0.01, *** *p* < 0.001 vs. untreated cells; # *p* < 0.05, ## *p* < 0.01 vs. LPS treatment.



Figure 4. Changes in mRNA levels of *IL-1ra* (**a**) and of *Sirt-1* (**b**) in J774A.1 mouse macrophages pre-treated with increasing concentrations (2.5%, 5%, 10% v/v) of SE FAE or with SA for 24 h and subsequently stimulated or not with LPS. Results were obtained using qPCR technique. Data are presented as mean ± SEM. Legend: SE FAE– *Sambucus ebulus* L. fruit aqueous extract; SA–100 µM salicylic acid; LPS–200 ng/mL lipopolysaccharides. * p < 0.05, ** p < 0.01, *** p < 0.001 vs. untreated cells; # p < 0.05, ## p < 0.01, ### p < 0.001 vs. LPS treatment.

2.2.2. The Effect of SE FAE on Inflammation-Related Biomarkers in Non-Stimulated J774A.1 Macrophages

When applied alone, 2.5% *v/v* and 10% *v/v* SE FAE slightly reduced the gene expression of *IL-1* β by 60% (p < 0.01) and 77% (p < 0.05), respectively, as compared to untreated cells (**Figure 1**a). While 2.5% *v/v* of herbal extract induced the gene expression of *IL-6* (by 67%, p < 0.05), *TNF* α (by 115%, p < 0.01), *Ccl2* (by 95%, p < 0.01), and *Fabp4* (by 36%, p < 0.05) (**Figure 1**b,c and **Figure 2**a,c). The higher concentration of SE FAE (5% extract) in culture media stimulated transcription of *TNF* α (by 92%, p < 0.001) and of *Ccl2* (by 39%, p < 0.05) (**Figure 1**c and **Figure 2**a), while the highest concentration (10% extract) induced transcription of *TNF* α (by 121%, p < 0.001) and of *Fabp4* (by 68%, p < 0.01) (**Figure 1**c and **Figure 2**c). SA, applied alone, similarly to SE FAE, it enhanced transcription levels of *Ccl2* (by 200%, p < 0.01), but in contrast with SE FAE, it slightly reduced

those of *Icam1* (by 91%, p < 0.01) and of *Fabp4* (by 16%, p < 0.05) (**Figure 2**a–c), while no significant effects on *IL-1* β , *IL-6* and *TNF* α transcription levels were observed (**Figure 1**a–c).

The treatment with 2.5% v/v of SE FAE alone significantly induced the transcription levels of *COX2* (by 210%, p < 0.05) and of *iNOS* (by 230%, p < 0.05) and both 2.5% v/v and 5% v/v of the extract induced iNOS protein levels by 9% (p < 0.05) and by 38% (p < 0.01), respectively (**Figure 3**). No effect of SA alone was observed on the gene expression levels of all analyzed inflammation and phagocytosis-related enzymes (**Figure 3**).

SE FAE in concentrations of 2.5% *v/v* and 10% *v/v* induced the transcription levels of *IL-1ra* by 98% (p < 0.01) and 41% (p < 0.05), respectively (**Figure 4**a). In contrast, SA treatment reduced *IL-1ra* transcription by 57% (p < 0.05) (**Figure 4**a). Transcription of the so-called longevity gene *Sirt-1* was stimulated upon 2.5% *v/v* and 5% *v/v* SE FAE treatment by 343% (p < 0.05) and by 274% (p < 0.05), respectively (**Figure 4**b). There was no significant effect of SA applied alone on *Sirt-1* transcription levels (**Figure 4**b).

2.3. Investigation of ER Stress-Related Biomarkers in a Model of LPS-Stimulated J744A.1 Macrophages

Regarding the well-known relationship between inflammation and ER stress, we have analyzed intracellular protein levels of ER stress-related proteins: activating transcription factor 6 alpha (ATF6 α), phosphorylated eukaryotic translation initiation factor 2 alpha (peIF2 α), and their downstream target gene's product C/EBP homologous protein (CHOP, growth arrest and DNA damage-inducible gene 153 (GADD153)) in a model of LPS-stimulated J744A.1 macrophages (**Figure 5**). Cells were pre-treated with increasing concentrations of 2.5%, 5% and 10% *v*/*v* (0.25 mg DW/mL, 0.5 mg DW/mL, 1 mg DW/mL respectively) SE FAE or SA for 24 h followed by LPS-stimulation for additional 24 h, and respective control treatments were performed as well.



Figure 5. Changes in the protein levels of peIF2 α (**a**), ATF6 α (**b**), and CHOP (**c**) in J774A.1 mouse macrophages pre-treated with increasing concentrations (2.5%, 5%, 10% *v*/*v*) of SE FAE or with SA for 24 h and subsequently stimulated or not with LPS. Results were obtained using the Western blot technique. Data are presented as mean ± SEM. Legend: SE FAE–*Sambucus ebulus* L. fruit aqueous extract; SA–100 µM salicylic acid; LPS–200 ng/mL lipopolysaccharides. * *p* < 0.05, ** *p* < 0.01 vs. untreated cells; # *p* < 0.05, ## *p* < 0.01 vs. LPS treatment.