

Somatic cybridization for *Citrus*: polyphenols distribution in juices and peel essential oil composition of a diploid cybrid from Cleopatra mandarin (*Citrus reshni* Hort. ex Tan.) and sour orange (*Citrus aurantium* L.)

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Abstract Somatic cybridization is a technique based on induced or spontaneous fusion between nuclear genome of one parent and either the cytoplasmic genome of the other parent or a combination of both donors. This approach is considered a powerful biotechnological tool in plant improvement since it enables to obtain novel intergenomic recombinations that are impossible to achieve by conventional crosses. Twenty-five plants were regenerated following protoplast fusion of Cleopatra mandarin (*Citrus reshni* Hort. ex Tan.) and sour orange (*Citrus aurantium* L.) and a new 2n cybrid has been identified (4 % of plants obtained). A complete phytochemical investigation of the new genotype and both parents was carried out analyzing the polyphenol content of juices by liquid-chromatography–ultraviolet–diode array detector–mass spectrometry and the peel essential oils using a combination of gas chromatography equipped with a flame

ionization detector and a mass spectrometry. On the whole 10 flavonoids (7 flavanone and 3 flavones) and 4 hydroxycinnamic acids were detected and quantified, while 67 components were fully characterized and grouped into four classes (monoterpene hydrocarbons, oxygenated monoterpenes, sesquiterpenes, and others) for an easier comparison of all oils. Comparing the new cybrid to both parents an increase of the monoterpene hydrocarbons and a corresponding decrease of oxygenated components were shown. The aim of the present work was to evaluate whether the genome rearrangements in the new cybrid, obtained through not conventional strategies, can produce superior traits and improved performance such as enhanced presence and quality of bioactive components.

Keywords *Citrus* · Cybrid · Microsatellite · Juice · Polyphenol · Peel essential oil

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Introduction

Numerous epidemiological studies conducted in many countries indicate that a diet rich in fruits and vegetables, including fruit and vegetable products, delays the ageing process and reduces the risk of various lifestyle diseases (Szajdek and Borowska 2008). In this context, several studies have associated the consumption of polyphenols with lower risks of different types of cancer (Nichenametla et al. 2006)

and cardiovascular diseases (Kris-Etherton et al. 2002), and have shown that polyphenols possess antioxidant, anti-inflammatory and anti-ageing activity (Benavente-García and Castillo 2008). Citrus fruits are among the richest dietary sources of flavonoids (Tripoli et al. 2007) and other bioactive compounds such as limonoids, carotenoids, amines, organic acids, sterols and furocoumarins (Kris-Etherton et al. 2002). Over the past decades, a large number of studies have been carried out on the biomedical properties of many *Citrus* species, among them sour orange (*Citrus aurantium* L.) and more recently Cleopatra mandarin (*Citrus reshni* Hort. ex Tan.) (Barreca et al. 2011; Ram et al. 2011; Hamdan et al. 2013).

The breeding programs success is mainly dependent on the rate of the genetic variation available. Although the nuclear genome has a predominant role on the inheritance of most plant traits, cytoplasmic factors and cytoplasm/nucleus interactions are also important. Chloroplast and mitochondrial genes, interacting with nuclear genes, are implicated in the control of several morphological, physiological and agronomic traits. However, the influence of the cytoplasm on the phenotypic traits is largely unexplored. The role for nucleus \times cytoplasm interactions has been reported in some cases (Shonnard and Gepts 1994; Matsui et al. 2002; Zhang et al. 2003; Atienza et al. 2008; Wang et al. 2010), nevertheless more aspects have to be still clarified.

Somatic hybridization by protoplast fusion has progressed steadily over the past 30 years. Considerable successful instances of applying this technique for breeding in plant of economic interest, including rice, potato, wheat, rapeseed and citrus were reported (Johnson and Veilleux 2001; Grosser and Gmitter 2011; Wang et al. 2013). In *Citrus*, somatic hybridization has been employed to circumvent the problems encountered using conventional breeding approaches, such as sexual incompatibility, nucellar polyembryony, male or female sterility and others (Grosser et al. 2000).

Traditional symmetric fusion, combining entire genomes of both parents, often contains many unwanted nuclear encoded agronomic traits. *Citrus* somatic hybrids are often sterile, show morphologically abnormal traits and may produce uncontrolled genomic instabilities, thus hampering their use for breeding of varieties. Therefore, asymmetric somatic cybridization, a technique based on the induction of unilateral chromosome elimination, has been developed as a means to

create morphologically normal hybrids, called cybrids, that contain the whole genome of one species and either the cytoplasmic genome of the other (non-nuclear) parent or that of a combination of both parental species. The potential of cybrids in citrus improvement could be relevant since it might be possible to breed specific traits associated to the cytoplasmic genomes while maintaining cultivar integrity (Saito et al. 1993; Guo et al. 2004). In *Citrus* interspecific, intraspecific and intergeneric symmetric protoplast fusion can spontaneously produce cybrids. To date, more than 40 citrus cybrids plants have been unexpectedly regenerated following symmetric somatic fusion experiments (Guo et al. 2013). Most of these cybrids have the nuclear genome of the leaf parent, the mitochondrial genome of the embryogenic callus parent and the chloroplast genome randomly inherited (Guo et al. 2013; Moreira et al. 2000), however, several investigations revealed the co-existence or recombination of parental mitochondrial genomes in cybrid plants (Iovene et al. 2007; Zubko et al. 2003). Positive variation for important agronomic traits such as maturity data and number of seeds (Grosser et al. 2000), tolerance to diseases (Tusa et al. 2000), metabolic, organoleptic and aromatic features (Fanciullino et al. 2005; Bassene et al. 2008, 2011; Wang et al. 2010) have been observed in citrus cybrids

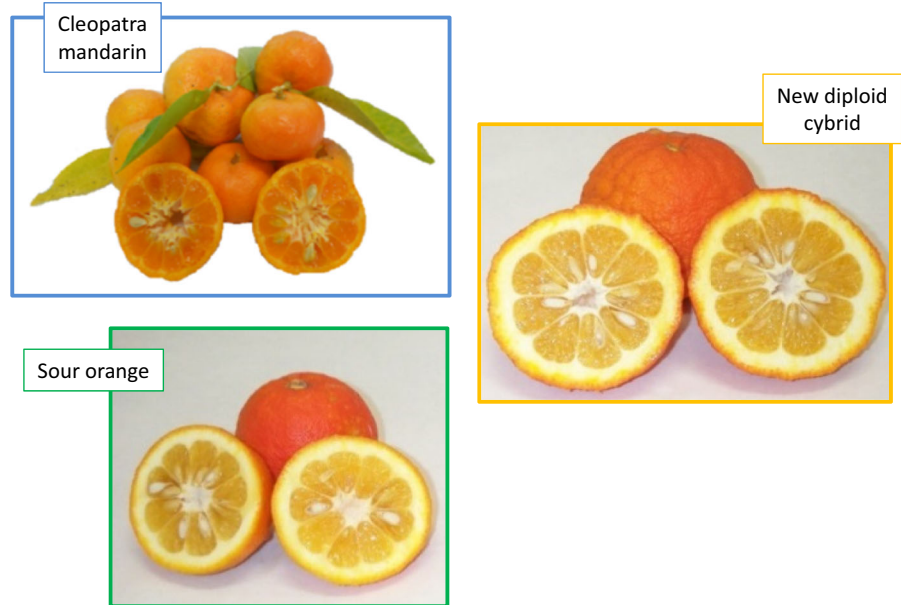
Pursuing our activity (Ruberto et al. 1999; Tusa et al. 2007; Abbate et al. 2012; Fatta Del Bosco et al. 2013) on the characterization of metabolic profiles of new *Citrus* hybrids, aimed to the selection new *Citrus* fruits with an improved content of nutraceutical components and/or with new aromatic profiles, as well as further tool to establish quality and typicalness of these new fruits. Here the results of a combined chemical and genetic study of a new diploid cybrid obtained from the protoplast fusion of Cleopatra mandarin and sour orange and its corresponding parents (Fig. 1) are reported, in an effort to evaluate the effect on juices polyphenols and peel essential oil characteristics associated to the new nucleo-cytoplasmic arrangement of the cybrid.

Materials and methods

Plant material

The *Citrus* plants investigated in this study were obtained by somatic protoplast fusion between

Fig. 1 The new diploid cybrid (CLEO–SO), ‘Sour’ orange (SO) and ‘Cleopatra’ mandarin (CLEO)



Cleopatra mandarin (*C. reshni*—CLEO) and sour orange (*C. aurantium*—SO). The CLEO protoplasts were isolated from suspension culture initiated from nucellus-derived embryogenic callus while the SO's protoplasts were derived from leaves of young nucellar seedlings germinated in vitro. CLEO and SO protoplasts were purified, mixed and fused using polyethylene glycol (PEG) method. Fusion cultures, embryogenic colonies, embryoids and large somatic embryos developing from recovered hybrid calli were grown. Developing somatic embryos were first transferred and cultured on 1500 medium for enlargement and after on B+ or DBA3 medium to induce shoot growth. Shoots were rooted in Magenta boxes (Magenta Corp., Chicago, IL, USA) and, finally the rooted plants were transferred to a commercial potting mixture and maintained at high humidity for acclimation according to the procedure described in Siragusa et al. (2007). Twenty-five regenerated plants were developed.

Plants obtained from the protoplast fusion event and relative parents were cultivated in the experimental station of the Istituto di Bioscienze e BioRisorse (IBBR-CNR), Collesano, Palermo. CLEO, SO and prescreened putative cybrid plants morphologically resembling SO (Grosser et al. 1996) were selected and the fruits were collected between February 2012 and March 2013.

Flow cytometry analysis

The samples' ploidy was evaluated by flow cytometry analysis, using diploid SO as reference standard. The analysis was carried out with the Partec PAS flow cytometer (Partec, <http://sysmex-europe.com/partec>), equipped with a mercury lamp. Fully expanded leaves (0.1 g) were chopped in a glass Petri dish with 400 μ L nuclei extraction buffer (Partec solution CyStain[®] UV Precise P, 250 tests). The solution was filtered through a 30 μ m Cell-Trics disposable filter Partec and 1.6 mL of staining solution containing 4,6-diamidino-2-phenylindole was added. Routinely, 4000–5000 nuclei for each sample were measured (Galbraith et al. 1998) and histograms of DNA content were generated using the Partec software package (FloMax). Three replicates for each sample were carried out.

DNA extraction

DNA was isolated from young, fresh leaves collected in the field and, after lyophilization, stored at -80°C until required. The extraction was carried out by the Doyle and Doyle CTAB method (1990). The extract was treated with DNase-free RNase (Roche Diagnostics, Germany) and quantified by using a NanoDrop ND-1000 spectrophotometer (NanoDrop Technologies, Wilmington, DE, USA).

Genetic analysis

A total of 22 simple sequence repeats (SSRs), 8 nuclear (nSSR), 7 chloroplast (cpSSR) and 7 mitochondrial (mtSSR) loci respectively, were chosen to investigate the genetic profile of CLEO, SO and putative cybrids obtained by protoplast fusion. The used markers (Table 1) were selected based on their dispersal map location and polymorphism rate (Cheng et al. 2003, 2005; Guo et al. 2006; Jannati et al. 2009; Froelicher et al. 2011). Multiplex PCRs were performed, amplifying several loci simultaneously, using the Qiagen multiplex PCR kit (Qiagen GmbH, Hilden, Germany). One of each pair was fluorescently labelled with FAM, JOE, TAMRA or ROX. PCR reactions were carried out in reaction volumes of 25 μ L containing 20 ng of DNA and 0.2 μ M each of forward and reverse primers, in a 2 \times Qiagen multiplex Master Mix. Reactions were performed under the following conditions: 15 min at 95 $^{\circ}$ C; 35 cycles of 30 s at

94 $^{\circ}$ C, 1 min 30 s at 57 $^{\circ}$ C, 1 min at 72 $^{\circ}$ C, followed by 30 min at 60 $^{\circ}$ C. Amplification products were checked and quantified by electrophoresis on 2 % agarose gel using Low DNA mass Ladder (Invitrogen). The fragments were separated by capillary electrophoresis and genotyped with an ABI PRISM 3500 genetic analyzer.

Fruit collection, juice preparation and standards

Fruits of SO, CLEO and their cybrids, were collected at the proper ripening stage between February 2012 and March 2013, from three trees for each genotype. Only the true cybrids, verified using cytometry analysis and molecular markers, were processed. The juice from fruits (4 or 5 for each tree) was obtained by a domestic squeezer and used freshly or stored at -18° C.

HPLC grade diosmin, vitexin, caffeic acid, eriocitrin, narirutin, naringenin 7-*O*-glucoside, naringin,

Table 1 Nuclear (nSSR), chloroplast (cpSS) and mitochondrial (mtSSR) microsatellite markers used for genetic analysis

Type	Name	Forward primer (5'–3')	Reverse primer (5'–3')	Expected size (bp)	
nSSR	TAA27	GGATGAAAAATGCTCAAAATG	TAGTACCCACAGGGAAGAGAGC	158–230	
	CAC15	TAAATCTCCACTCTGCAAAAGC	GATAGGAAGCGTCGTAGACCC	135–190	
	CAT01	GCTTTCGATCCCTCCACATA	GATCCCTACAATCCTTGGTCC	138–172	
	ATC09	TTCCTTATGTAATTGCTCTTTG	TGTGAGTGTGTTGTGCGTGTG	130–210	
	AG14	AAAGGGAAAGCCCTAATCTCA	CTTCCTCTTGCGGAGTGTTC	110–172	
	CAC33	GGTGATGCTGCTACTGATGC	CAATTGTGAATTTGTGATTCCG	77–109	
	CCT01	TCAACACCTCGAACAGAAGG	CCCACATGCTAGCACAAAAGA	93–119	
	CT21	CGAACTCATTAAGCCGAAAC	CAACAACCACCACTCTCACG	130–170	
	cpSSR	SPCC1	CTTCCAAGCTAACGATGC	CTGTCTATCCATTAGACAATG	220–240
		SPCC3	GATGTAGCCAAGTGATCA	TAATTTGATTCTTCGTCGC	750–820
SPCC7		CGATGCATATGTAGAAAGCC	CATTACGTGCGACTATCTCC	120–150	
SPCC9		TGGAGAAGGTTCTTTTCAAGC	CGAACCTCGGTACGATTAA	200–250	
SPCC11		GGCCATAGGCTGGAAAGTCT	GTTTATGCATGGCGAAAAGG	200–220	
SPCC13		GAAAAATGCAAGCACGGTTT	TACGATCCGTAGTGGGTTGC	120–150	
CICP9		CTTCCAAGCTAACGATGC	CTGTCCTATCCATTAGACAATG	200–270	
mtSSR	rn18-1/rn5	GGGTGAAGTCGTAACAAGGT	GAGGTCCGAATGGGATCGGG	250–260	
	cox1	TTGTTACGACCACGAAGA	TCGGTGCCATTGCTGGAG	Multilocus	
	nad7/4-5	TGTCCTCCATCACGATVTCG	CCAAATTCTCCTTTAGGTGC	Multilocus	
	rps4	GCGTATTTCCGGATGCTT	TCAAGTYGGTTCAGTGAG	Multilocus	
	cob/rpl5	AGAAACATACTTGCCCC	CCAACACACAAGATGAGA	Multilocus	
	nad2/3-4	GACCTTCACCTCAAATCA	TTCAGATAACACGCACC	239–258	
	nad7/1-2	GGAACATAGCATAGGG	TTTGATATAGGCTCGCT	134–159	

hesperidin, and didymin were purchased from Extrasynthese—Z.I. (Lyon Nord, France), sinapic acid and coumaric acid were obtained from Fluka Chemie (Buchs SG, Switzerland), ascorbic acid, ethylene diamine tetracetate disodium salt, neohesperidin, ferulic acid and isoferulic acid were purchased from Sigma-Aldrich (St. Louis, MO, USA).

Sample preparation for analyses of hydroxycinnamic acids

To a 1 mL of filtered (0.45 μm) juice, 1 mL of a 2 N NaOH solution (containing 10 mM EDTA and 57 mM ascorbic acid) and 10 μL of 1 mM isoferulic acid in MeOH as internal standard were added. The solution was maintained at 40 °C for 30 min under 300 rpm stirring. Each sample was then adjusted to pH 7 with 2 N HCl and freeze-dried. After suspending the residue in methanol (1 mL) and filtering on 0.45 μm filter, the solution was directly injected to HPLC system.

Sample preparation for analyses of flavanones and flavones

The juice was filtered on 0.45 μm filter and directly injected to the HPLC system. Flavonoid quantification was achieved from the recorded absorbance in the chromatogram compared with its external standard curve. Diosmetin 6,8-di-*C*-glucoside and vicenin-2, whose reference samples were not available, were quantified against vitexin assuming a response factor equal to 1.

HPLC–UV and HPLC–MS–ESI analyses

HPLC analyses were performed on a thermostated (40 °C) Phenomenex Luna[®] C₁₈ 250 \times 4.6 mm (5 μm) column at 1 mL/min flow rate using water-formic acid 9:1 (v/v) as eluant A and acetonitrile-formic acid, 9:1 (v/v) as eluant B with the following gradient of composition: $t_{0\text{min}}$ B (5 %), $t_{5\text{min}}$ B (5 %), $t_{20\text{min}}$ B (15 %), $t_{28\text{min}}$ B (15 %), $t_{38\text{min}}$ B (30 %), $t_{50\text{min}}$ B (100 %), $t_{55\text{min}}$ B (100 %), injection volume: 20 μL .

Qualitative analyses with simultaneous ESI–MS and UV–DAD detection were carried out on a waters 1525 pump (Waters Associates) equipped with a

waters 996 photodiode array detector and waters micromass ZQ2000 mass spectrometer detector. ESI–MS detection was performed in negative mode setting a capillary voltage of 3.5 kV, a cone voltage of 40 V, a vaporizer temperature of 250 °C, a carrier gas flow (nitrogen) of 500 L/h and mass acquisition between 100 and 1500 Da. DAD analyses were carried out in the range between 200 and 700 nm, setting the detector at 280 nm for flavanones, 340 nm for flavones and 320 nm for hydroxycinnamic acids respectively. The identification of juice constituents was carried out by comparison of the spectral properties (UV and ESI–MS) of analytes with those of reference samples.

Quantitative determination of the citrus juice composition was carried out on a Dionex HPLC system equipped with a P680 pump, a UV-170U detector and ASI 100 auto-sampler. All determinations were carried out in triplicate, using internal standard method for hydroxycinnamic acids and external standard method for flavanones and flavones.

Isolation of essential oils

Fresh rind tissue (flavedo, 100 g) of each sample was subjected to hydrodistillation until there was no significant increase in the volume of oil collected (3 h). The oils were dried over anhydrous sodium sulfate and stored under N₂ in a sealed vial at –20 °C until required.

Gas chromatography (GC) of essential oils

Essential oils were analyzed in the fast mode on a Shimadzu gas chromatograph, Model 17-A equipped with a flame ionization detector (FID), operating software class VP chromatography data system version 4.3 (Shimadzu Co., Kyoto, Japan). Analytical conditions: SPB-5 capillary column (15 m \times 0.10 mm \times 0.10 μm), helium as carrier gas. Injection in split mode (1:200), injected volume 1 μL (25 μL of oil in 400 μL of CH₂Cl₂), injector and detector temperature 250 and 280 °C, respectively. Linear velocity in column 51 cm/s. The oven temperature was held at 60 °C for 1 min, then programmed from 60 to 280 °C at 10 °C/min. Percentages of compounds were determined from their peak areas in the GC–FID profiles.

Gas-chromatography–mass spectrometry (GC–MS) of essential oils

GC–MS was carried out in the fast mode on a Shimadzu GC–MS mod. GCMS-QP5050A, operating software GCMS solution version 1.02 (Shimadzu). Ionization voltage in electronic impact mode 70 eV, electron multiplier 1000 V, transfer line temperature 280 °C, injection in split mode (1:96), constant linear velocity in column 50 cm/s. Analytical conditions were the same as GC.

Identification of essential oil components

The identity of components was based on their retention indexes relative to C₉–C₂₂ *n*-alkanes (Alltech Italy) on the SPB-5, computer matching of spectral MS data those from NIST MS 107 and NIST 21 libraries (NIST 1998), the comparison of the fragmentation patterns with those reported in literature (Adams 2007) and, whenever possible, co-injections with authentic standards, which were purchased from Aldrich Chemical Co., Extrasynthese, France, and Fluka Chemie AG, Switzerland.

Statistical analysis

SPSS software, version 14.1, was used to carry out statistical analysis of the data. ANOVA and HSD Tukey test were applied to the data to determine significant differences between the analyzed components; the model was statistically significant with a value of $p \leq 0.01$.

Results

Ploidy analysis

The ploidy of the 25 regenerated plants was estimated by flow cytometry using diploid SO as internal standard. The results displayed that 21 genotypes (84 %) obtained by somatic protoplast fusion between CLEO and SO were diploid ($2n = 2x = 18$) since showed the same profile of the control ($2n = 2x = 18$). The other 4 plants (16 %), showing a 4n ploidy level, were classified as hybrids and kept out from the present study.

Molecular analysis

Nuclear, chloroplastic and mitochondrial genomes were analyzed by using 22 SSR markers (Table 1) to characterize the genomic constitution of putative 2n cybrids selected and their parental (CLEO and SO). Three out of eight nuclear markers (TAA27, CAC15 and CAT01) enabled to distinguish the parents of the somatic protoplast fusion and thus the hybrid status of the regenerated plants could be checked. The results showed that the nuclear profile of seven (24 %) regenerated diploid plant was identical to the leaf parent (SO) confirming its origin as donor of nuclear genome to the regenerated plants. The other 14 prescreened samples presented the same allelic profile of CLEO therefore were excluded from the analysis.

To study in depth the nature of seven putative cybrids, the constitution of their cytoplasmic genome was also investigated. Two out of seven chloroplast microsatellite loci selected (SPCC9 and SPCC11) showed polymorphism between CLEO and SO, thus the identification of the parental genomes was possible. Six out of seven regenerated plants had the same plastidial genome profile of SO, therefore these plants were not true cybrids. Instead, in the remaining sample, the plastidial genome was inherited from both parents, since the putative cybrid, named CLEO–SO, displayed a heterozygous profile for SPCC9 and SPCC11 markers with alleles belonging to both parents (Table 2). In the case of the mitochondrial genome, the profiles of the *Citrus* specie used to isolate protoplasts from embryogenic calli (CLEO) were observed in the CLEO–SO plant (Table 2) for three polymorphic markers (rrn18-1/rrn5, nad 2/3-4 and nad71-2). Therefore, molecular results confirmed that the *Citrus* plant named CLEO–SO, obtained by somatic protoplast fusion between CLEO and SO, was a cybrid genotype, with nuclear genome from SO, chloroplast genome from both parents while the mitochondria were inherited from CLEO.

Chemicals profile

Polyphenol profile and essential oil composition of the new cybrid selected were then investigated comparing its chemical profile with both parents, to evaluate the effect associated to the new nucleo-cytoplasmic organization.

Table 2 Ploidy level, nuclear (nSSR), chloroplast (cpSS) and mitochondrial (mtSSR) polymorphic profiles of cybrid CLEO + SO and its parents (CLEO and SO)

Sample	Ploidy	nSSR ^a			cpSSR ^a		mtSSR ^a					
		TAA27	CAC15	CAT01	SPCC9	SPCC11	rrn18-1/ rrn5	nad2/3- 4	nad71- 2			
Cleopatra mandarin (CLEO)	2n	209	209	154	154	143	143	209	210	251	248	151
Cybrid (CLEO + SO)	2n	209	212	154	157	155	155	209/ 218	210/219	251	248	151
Sour orange (SO)	2n	209	212	154	157	155	155	218	219	255	256	159

^a Only polymorphic nSSR, cpSSR and mtSSR were shown

Polyphenol profiles of juices

Table 3 lists the polyphenols, ten flavonoids (TF) and four hydroxycinnamic acids (HCA) reported in Fig. 2, analyzed in this study with their spectral features utilized for characterization and quantitative determination, whereas Table 4 reports the results on the content of the aforesaid components of the two parents: CLEO and SO, and their diploid cybrid (CLEO–SO).

The profile of SO is characterized by the predominance of two flavanones, naringin and neohesperidin the two bitter components which characterize this *Citrus* species, whereas vicenin-2 and diosmetin 6,8-di-*C*-glucoside are the main flavones. Amongst HCA, the ferulic is the main one, coumaric, caffeic and sinapic, in that order, follow at a considerable gap. CLEO shows a consistent lower amount of flavonoids with respect to SO. In this case hesperidin and narirutin are the main flavanones, and vicenin-2 is largely the main flavone. Concerning HCA, also in this the ferulic is largely the main acid, sinapic, caffeic and coumaric follows at large distance, whereas caffeic and sinapic show a comparable and low amount.

The first and significant compositional aspect of the cybrid is the higher content of flavonoids with respect to both parents. In fact, flavanones and flavones reach the amount of 300.97 and 79.70 mg/L, respectively. From a qualitative point of view the main flavanones of the cybrid are naringin and neohesperidin, as for SO. Also flavones are present in higher amount with respect to both parents, also in this case vicenin-2 and diosmetin 6,8-di-*C*-glucoside, are the main components as for sour orange. Concerning the HCA, ferulic acid is still the main component followed by coumaric,

caffeic and sinapic as previously observed for SO parent.

A comprehensive evaluation of these data allows to establish that the new cybrid concerning its polyphenol profile maintain substantially that of SO leaf parent.

Essential oil composition

Table 5 lists the composition of the essential oils of all samples. In total, 67 components were fully identified and grouped into four classes: monoterpene hydrocarbons with 11 components, oxygenated monoterpenes, the most numerous class, with 25 compounds, sesquiterpenes and others, with 10 and 21 compounds, respectively. Monoterpenes hydrocarbons were in all cases the most highly represented class. Limonene, as usual for most of *Citrus* peel essential oil, is the main component, ranging between 90 and 93 %. Myrcene is the second monoterpene hydrocarbons in all samples, SO shows β -pinene and γ -terpinene slightly over the 1 %, whereas in the other two samples, CLEO and new cybrid CLEO–SO, all components are below 1 %. Oxygenated monoterpenes, which normally give a quality mark to the *Citrus* oils, are represented at very low extent: CLEO shows the highest amount (ca. 4 %), being linalool the main component, other slight significant components are terpineol and terpinen-4-ol (Lota et al. 2001; Hamdan et al. 2013). SO shows a lower content of these components, linalool is still the main one, followed by geraniol and terpineol. In CLEO–SO the lowest amount of these oxygenated compounds has been evidenced, in fact all components are below 1 % being linalool and terpineol the main ones (Table 5). The other two classes of components,

Table 3 Spectral properties of juices' components

Elution order	RT (min)	Compound	λ_{\max} (nm)	(M–H) [–] (<i>m/z</i>)	Aglycon (M–H) [–] (<i>m/z</i>)
Flavonoids					
1	14.95	Vicenin-2	270, 338	593	
2	18.62	Diosmetin 6,8-di- <i>C</i> -glucoside	270, 348	623	
3	20.07	Eriocitrin	283, 328	595	287
4	24.76	Narirutin	282, 330	579	271
5	25.45	Naringenin 7- <i>O</i> -glucoside	283, 326	433	271
6	26.69	Naringin	284, 328	579	271
7	31.15	Hesperidin	283, 328	609	301
8	33.35	Diosmin	348	607	299
9	33.89	Neohesperidin	286, 332	609	301
10	40.80	Didymin	282, 329	593	285
Hydroxycinnamic acids					
1*	10.12	Caffeic acid	326, 232	179	
2*	16.15	Coumaric acid	310, 215	163	
3*	20.87	Ferulic acid	323, 234	193	
4*	22.49	Sinapic acid	323, 237	223	

namely sesquiterpenes and others (not terpenoid compounds), even though totally represented by 31 components are in almost all samples below 1 %.

Discussion

The aim of this work was to evaluate whether superior traits and improved performance can be obtained through the nucleo-cytoplasmic arrangement that characterize the cybrid. Genetic and chemical analysis of diploid cybrid and its corresponding parents was conducted to evaluate the effect on juices polyphenols and peel essential oil production associated to the new nucleo-cytoplasmic organization in the cybrid analyzed.

Primarily, the genetic arrangement of the genotype obtained by somatic protoplast fusion to verify the true cybrid nature of the genotype was evaluated. As reported in previous studies, cybrids harbor one parental nuclear genome and either the plastidial genome of the other parent or a combination of both parents (Guo et al. 2004; Fanciullino et al. 2005; Bassene et al. 2011; Grosser et al. 1996; Cheng et al. 2005). Molecular analysis confirmed the cybrid nature of the genotype obtained from protoplast fusion of

CLEO and SO. Indeed, CLEO–SO showed nuclear genome of the leaf parent SO, the mitochondria genome was inherited from the callus parent CLEO while, regarding the plastidial genome, SSR marker analysis revealed the coexistence of both parents. According to available data, the chloroplast genome in the cybrid are derived mainly uniparentally from either of the fusion parents, although coexistence of the chloroplast genomes from both parents in the hybrids is also reported (Guo et al. 2007). Therefore, these results indicated that the CLEO–SO is a true cybrid.

Bio-molecular studies on the secondary metabolic profile and on volatile components, of *Citrus* somatic hybrids are rather limited (Fanciullino et al. 2005; Gancel et al. 2003, 2005), therefore it is difficult to find common aspects in the accumulation mechanism of these components at the moment. In fact, previous studies on the volatile compounds from leaves and peels of interspecific somatic hybrids showed a strongly inhibition of some components (i.e. sesquiterpene hydrocarbons), as well as an overproduction of the other ones (i.e. citronellal); these contrasting results prompted some authors to claim that *Citrus* somatic hybrids do not retain their parental traits (Gancel et al. 2005). In the present study, a slight

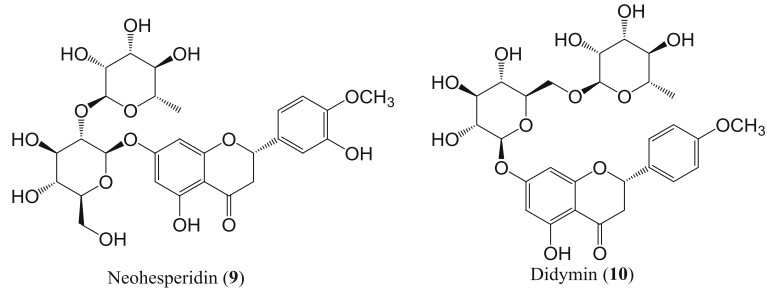
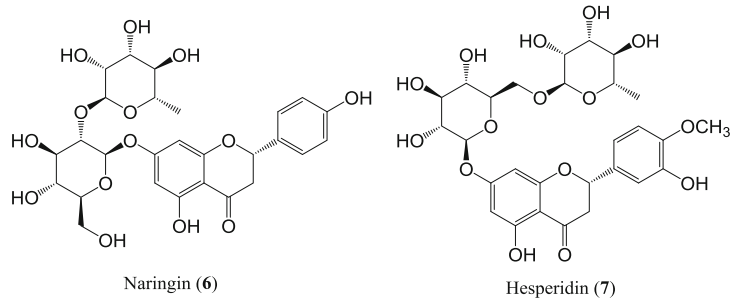
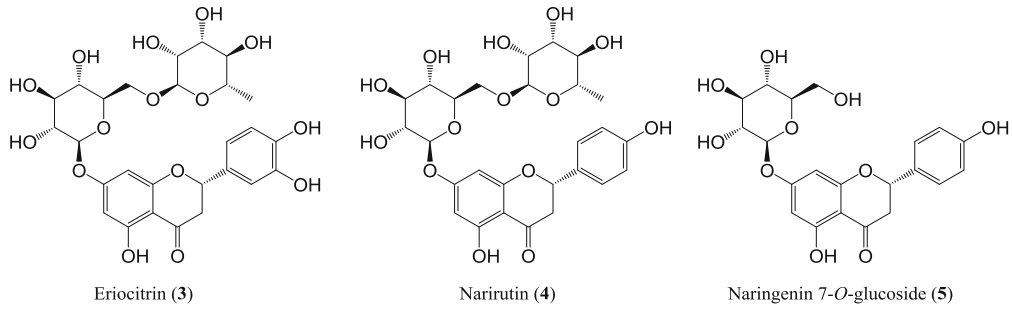
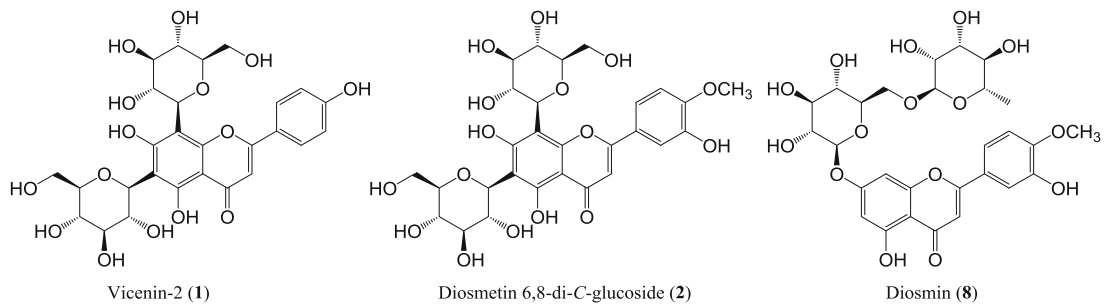
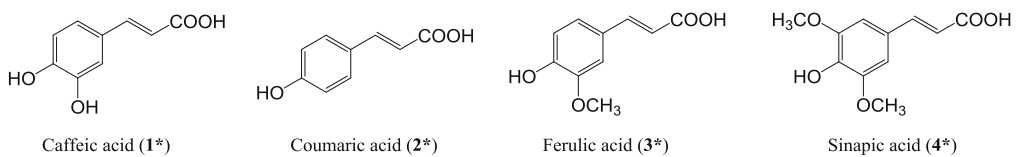
Flavanones**Flavones****Hydroxycinnamic Acids****Fig. 2** Polyphenols from *Citrus* juices

Table 4 Polyphenols in sour orange (SO), Cleopatra mandarin (CLEO) and the new cybrid (CLEO–SO)

No.	Flavanones–flavones–hydroxycinnamic acids (mg/L)	SO	CLEO	CLEO–SO
	Flavanones	283.89	54.47	300.97
3	Eriocitrin	5.87 (0.32)	1.12 (0.41)	4.49 (0.10)
4	Narirutin	11.06 (0.14)	10.50 (0.74)	8.76 (0.25)
5	Naringenin 7- <i>O</i> -glucoside	1.03 (0.13)	2.81 (0.11)	1.57 (0.05)
6	Naringin	176.33 (0.25)	0.61 (0.11)	162.57 (0.08)
7	Hesperidin	0.59 (0.02)	36.48 (0.04)	3.17 (0.23)
9	Neohesperidin	89.00 (0.55)	1.83 (0.06)	119.89 (0.35)
10	Didymin	0.01 (0.00)	1.12 (0.00)	0.52 (0.33)
	Flavones	61.65	49.61	79.70
1	Vicenin-2	39.38 (0.03)	38.71 (2.53)	48.08 (1.77)
2	Diosmetin 6,8-di- <i>C</i> -glucoside	19.93 (0.60)	9.84 (4.52)	27.56 (0.22)
8	Diosmin	2.34 (0.49)	1.06 (0.36)	4.06 (2.64)
	Total flavonoids	345.54	104.08	380.67
	Hydroxycinnamic acids	26.00	72.68	22.64
1*	Caffeic acid	2.48 (0.02)	3.10 (0.05)	2.25 (0.09)
2*	Coumaric acid	3.62 (0.06)	1.59 (0.05)	4.27 (0.18)
3*	Ferulic acid	18.03 (0.13)	61.03 (0.45)	15.23 (0.45)
4*	Sinapic acid	1.87 (0.11)	6.96 (0.24)	0.89 (0.39)

Values represent averages of three determinations, standard deviation (\pm SD) is given in parentheses

overproduction of total juice flavonoids (Table 4), an increase of the monoterpene hydrocarbons and a corresponding decrease of oxygenated components with respect to both parents in the essential oils (Table 5) were observed in CLEO–SO. Our results show, analogously to many similar ones, just how difficult it is to establish an inheritance mechanism related to the biosynthetic accumulation of secondary metabolites throughout various breeding methodologies of *Citrus* (Rapisarda et al. 2003; Fabroni et al. 2012). This is probably due to the complexity and genetic changeability of this genus (Guo et al. 2006; Barkley et al. 2006). In any case these comparisons clearly provide an insight into the spectrum of changes associated with the genetic manipulation by protoplast fusion.

Besides, the molecular constitution of cybrids provides a novel and promising approach for citrus breeding. In fact, although the nuclear genome has a predominant role on the inheritance of most plant traits, it is recognized that plant cell function depends on the coordinate action and communication between nuclear and cytoplasmic genomes. There is increasing awareness of the intricate

networks and signal transduction pathways that link plant organelles, called “retrograde regulation” (Strand 2004; Liu and Butow 2006; Rhoads and Subbaiah 2007).

Recent investigations underline how the chloroplast and their redox state are involved in the regulation of important biological processes, including nuclear/plastid gene expression, hormone signalling and stress responses (Suzuki et al. 2002). Similarly, mitochondrial respiration is important for neutralizing excess of photosynthetic reducing power, preventing oxidative damage (Dinakar et al. 2010), highlighting the key role of plant mitochondria during biotic stress responses. Indeed, there are evidences that signals initiated by plant–pathogen interactions may directly or indirectly transfer to mitochondrial components consequently increasing generation of mitochondrial reactive oxygen species (mtROS). The mtROS or mitochondrial dysfunction may act as cellular signals that initiate nuclear gene expression involved in defense and hypersensitive responses (Amirsadeghi et al. 2007). Redox regulation and ROS metabolism are interlinked and involved in optimizing the function of mitochondria, chloroplast

Table 5 Chemical composition of the peel essential oils of sour orange (SO), Cleopatra mandarin (CLEO) and their cybrid (CLEO–SO)

No.	Ri ^b	Ri ^c	Class/compound	SO		CLEO		CLEO–SO	
				Aver.	SD	Aver.	SD	Aver.	SD
			Monoterpene hydrocarbons	95.61		94.72		96.45	
4	930	929	α -Thujene ^d	0.05	0.00	0.01	0.00	0.02	0.00
5	939	937	α -Pinene ^d	0.55	0.00	0.49	0.01	0.45	0.01
6	954	953	Camphene ^d	0.01	0.00			0.02	0.00
7	975	976	Sabinene ^d	0.36	0.01	0.83	0.00	0.12	0.00
8	979	979	β -Pinene ^d	1.51	0.03	0.05	0.00	0.42	0.05
10	991	991	β -Myrcene ^d	1.63	0.00	1.72	0.01	1.55	0.03
13	1017	1020	α -Terpinene ^d	0.04	0.00	0.05	0.00	0.02	0.01
14	1029	1041	Limonene ^d	89.82	0.08	91.31	0.21	93.32	0.51
15	1050	1052	<i>trans</i> - β -Ocimene	0.27	0.00	0.13	0.00	0.13	0.01
16	1060	1062	γ -Terpinene ^d	1.25	0.04	0.11	0.01	0.34	0.03
19	1089	1089	Terpinolene ^d	0.12	0.00	0.02	0.00	0.07	0.01
			Oxygenated monoterpenes	3.26		3.95		1.96	
18	1073	1075	<i>trans</i> -Linalool oxide	0.08	0.01			0.08	0.00
20	1096	1099	Linalool ^d	1.66	0.07	2.69	0.13	0.87	0.06
22	1102	1104	<i>cis</i> -Thujone ^d					0.04	0.00
23	1114	1116	<i>trans</i> -Thujone ^d					0.02	0.00
24	1123	1122	<i>trans</i> - <i>p</i> -Mentha-2,8-dien-1-ol			0.05	0.00		
26	1138	1138	<i>cis</i> - <i>p</i> -Mentha-2,8-dien-1-ol			0.04	0.00		
27	1146	1149	Camphor ^d	0.01	0.00			0.03	0.00
28	1153	1156	Citronellal ^d	0.01	0.00	0.01	0.00	0.02	0.00
29	1177	1181	Terpinen-4-ol ^d	0.10	0.00	0.24	0.01	0.07	0.01
30	1189	1193	Terpineol ^d	0.32	0.02	0.25	0.01	0.26	0.03
31	1193	1195	<i>cis</i> -Dihydro Carvone			0.03	0.00		
36	1217	1219	<i>trans</i> -Carveol			0.06	0.00		
37	1230	1232	Nerol ^d	0.06	0.01	0.18	0.01	0.05	0.01
38	1238	1244	Neral ^d	0.10	0.01			0.04	0.00
39	1242	1247	Carvone	0.03	0.00	0.02	0.00	0.08	0.02
40	1253	1258	Geraniol ^d	0.42	0.01	0.05	0.00	0.16	0.02
42	1267	1272	Geranial ^d	0.12	0.01	0.07	0.00	0.05	0.00
44	1272	1277	Perillaldehyde ^d	0.05	0.00	0.15	0.00	0.04	0.00
46	1290	1292	Limonen-10-ol	0.01	0.00	0.08	0.00	0.02	0.01
47	1295	1299	<i>p</i> -Menth-1-en-9-ol			0.02			
48	1295	1300	Perilla alcohol	0.02	0.00				
50	1327	1328	Myrtenyl acetate			0.01	0.00		
52	1353	1354	Citronellyl acetate	0.01	0.00			0.02	0.01
53	1362	1364	Neryl acetate	0.07	0.00			0.03	0.00
54	1381	1382	Geranyl acetate	0.19	0.00			0.10	0.01
			Sesquiterpenes	0.32		0.08		0.61	
51	1338	1341	δ -Elemene	0.02	0.00	0.02	0.00	0.03	0.01
55	1391	1392	β -Elemene					0.02	0.00
58	1413	1409	α - <i>cis</i> -Bergamotene	0.01	0.00				
59	1419	1423	β -Caryophyllene ^d	0.04	0.00			0.05	0.00

Table 5 continued

No.	Ri ^b	Ri ^c	Class/compound	SO		CLEO		CLEO–SO	
				Aver.	SD	Aver.	SD	Aver.	SD
60	1435	1434	α - <i>trans</i> -Bergamotene	0.03	0.00				
61	1460	1457	β -Santalene	0.01	0.00				
63	1485	1484	Germacrene D	0.04	0.00	0.03	0.00	0.07	0.01
64	1561	1562	Germacrene B			0.04	0.00		
65	1563	1563	Nerolidol <i>E</i> ^d	0.11	0.00			0.10	0.03
66	1807	1807	Notkatone	0.06	0.02			0.35	0.11
			Others	0.76		1.27		0.63	
1	802	800	Hexanal ^d					0.02	0.00
2	854	856	2- <i>trans</i> -Hexenal ^d	0.01	0.00			0.01	
3	871	872	<i>n</i> -Hexanol ^d					0.01	
9	984	986	3-Octanone ^d					0.01	0.00
11	999	1002	Octanal ^d	0.42	0.02	0.58	0.02	0.14	0.01
12	1009	1015	Hexyl acetate	0.03	0.02				
17	1068	1073	Octanol ^d	0.04	0.00	0.23	0.01	0.04	0.01
21	1101	1103	Nonanal ^d	0.02	0.00	0.03	0.00	0.02	0.00
25	1127	1123	Methyl octanoate					0.01	0.00
32	1197	1195	Ethyl octanoate					0.09	0.00
33	1196	1199	Estragole ^d	0.02	0.01			0.02	0.00
34	1202	1206	Decanal ^d	0.11	0.00	0.36	0.02	0.13	0.01
35	1214	1213	Octyl acetate	0.04	0.00			0.03	0.00
41	1264	1264	2- <i>trans</i> -Decenal ^d	0.02	0.00	0.01	0.00	0.02	0.00
43	1271	1274	2- <i>trans</i> -Decenol	0.01	0.00				
45	1285	1284	<i>trans</i> -Anethol ^d					0.03	0.00
49	1317	1316	2 <i>E,4E</i> -Decadienal	0.01	0.00	0.01	0.00		
56	1409	1404	Dodecanal ^d			0.04	0.00		
57	1409	1407	Decyl acetate	0.01	0.00			0.03	0.01
62	1466	1465	2 <i>E</i> -Dodecenal ^d	0.01	0.00			0.02	0.00
67	2300	2296	<i>n</i> -Tricosane ^d	0.02	0.00				

^a The numbering refers to elution order, and values (relative peak area percent) represent averages of three determinations

^b Retention index (RI) relative to standard mixture of *n*-alkanes on SPB-5 column

^c Literature retention index

^d Co-elution with authentic sample

and other organelles. This network needs implied regulation and high level of coordination especially under stress conditions.

Chloroplast and mitochondrial genes, or their interactions with nuclear genes, are clearly implicated in the control of several morphological, physiological and agronomic traits (Woodson and Chory 2008). Cytoplasmic male sterility (CMS) is, by far, the most known cytoplasmically inherited trait. The drought tolerance in *Brassica* (Uprety and Tomar 1993), heat

tolerance in common bean (Shonnard and Gepts 1994), productivity and tolerance to iron deficiency in wheat (Zhang et al. 2003; Jones et al. 1998) and disease resistance in barley (Matsui et al. 2002), were also associated to cytoplasmically inherited characters. In wheat significant differences for seed lutein content between euplasmic and their respective alloplasmic lines were also observed (Atienza et al. 2008), underscoring the important role of the cytoplasm to determine the seed carotenoid accumulation. These

evidences prove the useful use of cybridization as sources of cytoplasmic variability for broadening the genetic diversity in plant, underlining the important role of the nuclear–cytoplasm interaction.

As reported in *Citrus* cybrid combinations, cytoplasm substitution shows positive effects on important agronomic traits, as CMS (Yamamoto et al. 1997), aroma and essential oil characteristics (Fanciullino et al. 2005; Abbate et al. 2012), fruit taste and nutritional quality (Bassene et al. 2008), therefore cybridization can be used in citrus cultivar improvement as a strategy to transfer specific traits associated with mitochondrial and chloroplast genomes.

Conclusions

The results obtained in this work point out the role of cytoplasm and/or the nucleus–cytoplasmic interactions in determining new skins with respect to polyphenol distribution and essential oil composition in citrus cybrid fruit. Since the somatic cybridization represents a suitable alternative strategy against the major biotic/abiotic constraints of the Mediterranean area, developing in meantime genotypes with interesting chemical traits, the cytoplasm constitution of the new CLEO–SO cybrid could have positive potential effect on different features.

In conclusion, the regeneration of somatic cybrid CLEO–O is an important beginning step that can allow enlarging the citrus germplasm patrimony available for further studies at the biochemical, agronomic and genetic levels.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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