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# Floral trait differentiation in *Anacamptis coriophora*: Phenotypic selection on scents, but not on colour

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## Abstract

Current divergent selection may promote floral trait differentiation among conspecific populations in flowering plants. However, whether this applies to complex traits such as colour or scents has been little studied, even though these traits often vary within species. In this study, we compared floral colour and odour as well as selective pressures imposed upon these traits among seven populations belonging to three subspecies of the widespread, generalist orchid *Anacamptis coriophora*. Colour was characterized using calibrated photographs, and scents were sampled using dynamic headspace extraction and analysed using gas chromatography–mass spectrometry. We then quantified phenotypic selection exerted on these traits by regressing fruit set values on floral trait values. We showed that the three studied subspecies were characterized by different floral colour and odour, with one of the two predominant floral volatiles emitted by each subspecies being taxon-specific. Plant size was positively correlated with fruit set in most populations, whereas we found no apparent link between floral colour and female reproductive success. We detected positive selection on several taxon-specific compounds in *A. coriophora* subsp. *fragrans*, whereas no selection was found on floral volatiles of *A. coriophora* subsp. *coriophora* and *A. coriophora* subsp. *martrinii*. This study is one of the first to document variation in phenotypic selection exerted on floral scents among conspecific populations. Our results suggest that selection could contribute to ongoing chemical divergence among *A. coriophora* subspecies.

## KEYWORDS

*Anacamptis coriophora*, floral colour, floral scents, phenotypic selection, pollination

## 1 | INTRODUCTION

One of the most striking characteristics of Angiosperms is their floral diversity, which has been the focus of a fruitful research for over two centuries (Darwin, 1862; Harder & Barrett, 2006; Sprengel,

1793). Because floral traits act as visual, olfactory or even tactile signals maximizing attractiveness to pollinators and pollen transfer efficiency (Burger, Dötterl, & Ayasse, 2010; Jersáková, Jürgens, Šmilauer, & Johnson, 2012; Schiestl & Johnson, 2013), they have a major influence on plant reproductive success. It is thus generally

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assumed that floral evolution has been driven by pollinator-mediated selection (Harder & Johnson, 2009; Sletvold, Grindeland, & Ågren, 2010), although antagonists (Irwin, Bronstein, Manson, & Richardson, 2010; McCall, Murphy, Venner, & Brown, 2013; Parachnowitsch & Caruso, 2008; Ramos & Schiestl, 2019), co-occurring plants (Caruso, 2000, 2001) and abiotic factors (Arista, Talavera, Berjano, & Ortiz, 2013; Armbruster, 2002) may also impose direct or indirect selective pressures upon floral characteristics (Caruso, Eisen, Martin, & Sletvold, 2019; Strauss & Whittall, 2006).

Floral traits often vary among conspecific populations (Anderson, Ros, Wiese, & Ellis, 2014; Breitkopf et al., 2013; Dormont, Joffard, & Schatz, 2019; Ellis & Johnson, 2009). If such variation is adaptive, we may detect current divergent selection on floral traits when comparing these populations (i.e., linear selection in opposite directions, stabilizing selection with different optima, or, for multivariate traits, selection acting on distinct components of the trait). This was shown to be the case for diverse floral characteristics, including phenology (Hall & Willis, 2006; Sandring, Riihimäki, Savolainen, & Ågren, 2007), size (Chapurlat, Ågren, & Sletvold, 2015; Totland, 2001) and shape (Gómez et al., 2008; Gómez, Perfectti, Bosch, & Camacho, 2009; Johnson & Steiner, 1997; Soteras, Rubini Pisano, Bariles, Moré, & Cocucci, 2020). In the generalist plant *Erysimum mediohispanicum*, for example, phenotypic selection on corolla shape was found to vary among populations visited by distinct pollinators and to match the observed pattern of morphological differentiation (Gómez et al., 2008, 2009), suggesting that current divergent selection promotes ongoing floral trait differentiation in this species.

Among floral traits, colour and scents are of particular importance for biotic interactions, especially for pollinator attraction (Raguso, 2008; Whitehead & Peakall, 2009). Visual and olfactory floral signals have been shown to vary among conspecific populations in many plant species (reviewed in Delle-Vedove, Schatz, & Dufay, 2017; Dormont et al., 2019). Yet, surprisingly little is known regarding the processes promoting such variation, although adaptation to divergent pollination niches is often proposed as the most likely explanation (Breitkopf et al., 2013; Chess, Raguso, & LeBuhn, 2008). In particular, we are aware of only one study that has explicitly linked variation in floral colour and scents among conspecific populations to variation in phenotypic selection exerted on these traits (Gross, Sun, & Schiestl, 2016).

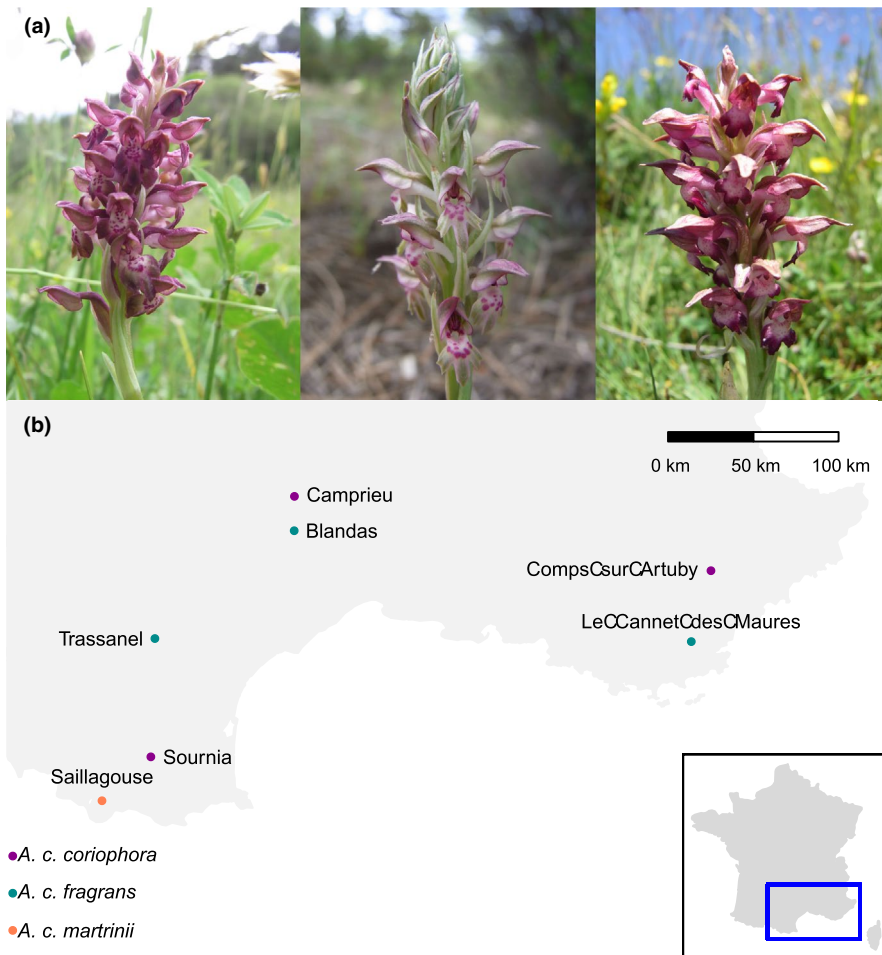
In this study, we ask whether current divergent selection could contribute to ongoing floral colour and scent differentiation in the widespread, generalist orchid species *Anacamptis coriophora*. This species comprises several subspecies, three of which are known to occur in France, namely *Anacamptis coriophora* subsp. *coriophora* (hereafter referred to as '*A. c. coriophora*'), *A. coriophora* subsp. *fragrans* (hereafter referred to as '*A. c. fragrans*') and *A. coriophora* subsp. *martrinii* (hereafter referred to as '*A. c. martrinii*'). These three subspecies grow in different habitats and are described as having distinct floral colour and scents (Bournérias & Prat, 2005; Delforge, 2006), although this has never been properly quantified. Specifically, flowers of *A. c. fragrans* are described as having a lighter colour than those of *A. c. coriophora* and *A. c. martrinii*, and a sweeter

smell than those of *A. c. coriophora*, whereas flowers of *A. c. martrinii* are described as odourless (Delforge, 2006). By characterizing floral colour and scents in seven populations of this species, we first aim at describing variation in floral phenotypes among and within subspecies. We then ask whether selection could explain such variation by measuring phenotypic selection gradients for floral colour and scent traits in each population, and by comparing selective pressures imposed upon these traits among and within subspecies. In particular, if floral trait differentiation between the three studied subspecies is adaptive, we expect selection to favour plants with dark flowers in *A. c. coriophora* and *A. c. martrinii* but plants with light flowers in *A. c. fragrans*. Our predictions regarding selection exerted on floral scents depend on the observed pattern of chemical differentiation. If the three subspecies differ in terms of total emission rate (e.g., if *A. c. martrinii* truly is odourless), we may detect current divergent selection on this trait; but if they differ in terms of scent chemistry (e.g., if *A. c. coriophora* and *A. c. fragrans* truly have distinct smells), we expect selection to favour plants emitting high amounts of taxon-specific compounds, and/or to vary in direction or strength when acting on the same compounds.

## 2 | MATERIALS AND METHODS

### 2.1 | Studied taxa and populations

*Anacamptis coriophora* (L.) Bateman, Pridgeon & Chase (1997) is a terrestrial, widespread orchid species which is distributed across central and southern Europe. This species produces nectar in a short spur (i.e., unlike other *Anacamptis* species, it is 'rewarding' and not 'deceptive') and is pollinated by various bee species (Claessens & Kleynen, 2011; Dafni & Ivri, 1979; Joffard, Massol, Grenié, Montgelard, & Schatz, 2019). It comprises several subspecies, three of which are known to occur in France, namely *Anacamptis c. coriophora*, *A. c. fragrans* and *A. c. martrinii* (Figure 1a). *Anacamptis c. coriophora* grows in low to medium altitude hay meadows and seasonally wet to wet grasslands, whereas *A. c. fragrans* grows in dry Mediterranean grasslands and *A. c. martrinii* in mountain hay meadows from the Pyrenees (Bournérias & Prat, 2005; Delforge, 2006). These three subspecies display inflorescences with about 10–30 flowers that open in acropetal sequence (from basal to distal) between May and July. Their flowers are light (in *A. c. fragrans*) to dark (in *A. c. coriophora* and *A. c. martrinii*) pink with red spots and sometimes green highlights on the labellum, and they are odorous, *A. c. coriophora* flowers having a stinkbug-like smell and *A. c. fragrans* flowers a sweeter, vanilla-like smell (Bournérias & Prat, 2005; Delforge, 2006). Note that *A. c. coriophora* and *A. c. fragrans* are sometimes treated as *bona fide* species (e.g., see Bateman et al., 2003); however, their presumed phylogenetic distinctness has been established using a single accession per taxon and a single locus. Because we think that more data are needed to conclusively demonstrate that these two taxa truly represent different species, in this study we adopt the more conservative view that they represent subspecies of *A. coriophora*.



**FIGURE 1** (a) Photographs of the inflorescences of *Anacamptis coriophora* subsp. *coriophora*, *A. coriophora* subsp. *fragrans* and *A. coriophora* subsp. *martrinii* (from left to right). (b) Geographic distribution of populations of *A. coriophora* subsp. *coriophora* (in purple), *A. coriophora* subsp. *fragrans* (in turquoise) and *A. coriophora* subsp. *martrinii* (in orange) sampled in this study

Three populations of *A. c. coriophora*, three populations of *A. c. fragrans* and one population of *A. c. martrinii*—each comprising between 50 and 100 individuals—were sampled in southern France during spring/summer 2016 (Figure 1b). Populations were assigned to subspecies using morphology, phenology and habitat characteristics. These populations were selected in order to sample both *A. c. coriophora* and *A. c. fragrans* in three geographical regions (south-eastern, south-central and south-western France). Repeated sampling of two of these seven populations (one of *A. c. coriophora* and one of *A. c. fragrans*) was planned in 2017 in order to test whether phenotypic selection patterns were consistent across years, but those were almost entirely destroyed due to bad weather conditions.

## 2.2 | Fieldwork

### 2.2.1 | Measurement of floral traits

In each of the seven populations, 50–65 individuals were randomly selected and marked. For each individual, plant height and inflorescence length (distance between uppermost and lowermost flowers) were measured to the nearest 0.5 mm with a tape measure and flowers were counted. Plant height, inflorescence length and number of flowers were used as proxies of plant size.

In six populations of seven (three of *A. c. coriophora*, two of *A. c. fragrans* and one of *A. c. martrinii*), floral colour was characterized using calibrated photographs. This method was chosen over spectrophotometry because it allows characterizing the colour of the entire labellum whereas spectrophotometry focuses on one or a few subjectively selected points. This is especially important here since as mentioned above, *A. coriophora* has a nonuniform floral colour with red spots and green highlights on the labellum. For each individual, one of the lowermost flowers was photographed using a standardized procedure (front view and planar position, indirect natural sunlight) with a Nikon D70 digital camera equipped with a Novoflex 35 mm lens. Note that floral colour variation observed within individuals was found to be negligible compared to that observed among individuals (data not shown). A ruler and two reference greys (23% and 35% of average reflectance) were included in each photograph for further scaling and colour calibration, respectively. Photographs were taken in the human-visible light range (i.e., 400–700 nm), but to confirm that our results were interpretable ecologically, we produced multispectral images of ten flowers in the UV-visible light range (340–700 nm) using a UV-sensitive camera (see Method S1). These images revealed that *A. coriophora* flowers reflect almost no light in the UV light range (see Figure S1).

In each of the seven populations, floral odour was sampled in situ using headspace dynamic extraction, a nondestructive method

allowing the extraction of scents emitted by the entire inflorescence. Floral odour was extracted between 08:00 hr and 18:00 hr, but daily variation in floral scent emissions was found to be negligible within the *A. coriophora* group (data not shown). For each individual, the inflorescence was enclosed in a ~ 10\*5 cm Nalophan® bag for 30 min. ChromatoProbe® quartz microvials (Varian, Inc.; length: 15 mm, inner diameter: 2 mm) filled with 3 mg of a 1:1 mix of Tenax-TA and Carbotrap® (respectively, 60–80 and 20–40 mesh, Sigma-Aldrich) were used as adsorbent traps, and 1 µl of internal standard (*n*-tetradeceane, 100 ng/µl) was injected in these traps. Two traps—one for mass spectrometry analyses, used for compound identification, and one for flame ionization detection analyses, used for compound quantification—connected to a pump (N86KNDC, KNF) by silicon tubes were inserted into the bag after 20 min. The air inside the bag was then pumped through the two traps for 10 min at a rate of 40 ml/min. After floral odour extraction, each trap was put into a vial and stored at –20°C until further analyses. In each population, the air contained in one empty bag was sampled as a control.

In parallel, pollinator guilds were characterized in each of the seven populations and pollen limitation of female reproductive success was quantified using hand pollinations in four populations of seven (see Methods S2 and S3).

### 2.2.2 | Estimation of female reproductive success

One month after the measurement of floral traits, the fruit set of each individual was measured and used as a proxy of female reproductive success. Relative female reproductive success was calculated by dividing the number of fruits produced by each individual by the mean number of fruits produced per plant in each population, following the recommendations of Lande and Arnold (1983). Number of seeds per fruit could not be counted, as fruit collection is not allowed in *A. c. coriophora*, *A. c. fragrans* and *A. c. martrinii*, which are nationally protected, but fruit set varied extensively among individuals in most populations whereas fruit size (which was shown to be a good proxy of seed set in other orchid species, e.g. see Chapurlat et al., 2015) was apparently more constant. Male reproductive success could not be measured either, as it would have required recording pollinia removal regularly during the entire flowering season. In some populations, a few individuals were lost due to mowing or grazing, but at least 96% of them were retrieved and could be included in further analyses (except in Blandas, where 17% of them were lost).

### 2.3 | Floral colour analyses

Although digital photographs can be powerful tools for measuring colour, usually pixel values cannot be compared among studies because of the model-specific nonlinear response of the camera to light intensity and because of variation in the illuminant spectrum. Moreover, even within studies pixel values may not be comparable among photographs if lighting conditions vary, which was the case

here. To circumvent these limitations, each photograph was calibrated using the Image Calibration and Analysis Toolbox plugin (Troscianko & Stevens, 2015) in the software ImageJ (Schneider, Rasband, & Eliceiri, 2012). After this calibration step, on each photograph, the labellum was manually selected and its colour characterized using (a) RGB values, to provide a measure of floral colour as we humans perceive it, and (b) the excitation values of the SW (short wavelength) and MW (medium wavelength) photoreceptors of the honeybee (*Apis mellifera*; see Method S4). Including both human and insect photoreceptors in our analysis ensures that our results do not critically depend on the specificities of one visual system and therefore likely apply to other pollinator species for which photoreceptor sensitivities are unknown. For each labellum, we measured one mean and one standard deviation values for each of the R, G, B, SW and MW channels.

### 2.4 | Floral odour analyses

Floral odour was analysed by gas chromatography (GC)–mass spectrometry (MS) and flame ionization detection (FID) using a Trace 1310 gas chromatograph coupled with a flame ionization detector and an ISQ mass spectrometer (Thermo-Electron) and equipped with an OPTIMA® 5-MS-HT capillary column (30 m × 0.25 mm × 0.25 µm, Macherey-Nagel). Traps were handled with a Multi-Purpose Sampler (Gerstel) and desorbed with a double-stage desorption system composed of a Thermal Desorption Unity (TDU) and a Cold Injection System (CIS; Gerstel). First, traps were desorbed splitless with a temperature of 250°C on the CIS trap cooled at –80°C using liquid nitrogen. Then, the CIS trap was heated to 250°C with a 1:4 split ratio to inject the compounds into the column. Oven temperature was held at 40°C for 3 min, increased from 40°C to 210°C at a rate of 5°C/min and from 220 to 250°C at a rate of 10°C/min, and held at 250°C for 2 min. The temperatures of transfer line and ion source of the mass spectrometer were 250°C and 200°C, respectively, and acquisition was set from 38 to 350 *m/z*, at a 70 eV ionization energy. The FID was heated to 250°C. Retention times of a series of *n*-alkanes (Alkanes standard solution, 04070, Sigma-Aldrich®) were used to convert retention times into retention indices. Compounds were identified on the GC-MS samples based on their retention indices and mass spectra, which were compared to those recorded in databases (Adams, 2007). Peak areas were measured on the GC-FID samples using the software Xcalibur (Thermo-Electron) and were converted to absolute (i.e., in ng) and relative (i.e., in %) amounts using the peak area of the internal standard (*n*-tetradeceane).

### 2.5 | Statistical analyses

#### 2.5.1 | Variation in floral traits and female reproductive success

Variation in plant size among populations of the same subspecies (*A. c. coriophora* or *A. c. fragrans*) was investigated using linear models



with population as a fixed effect and plant size traits (plant height, inflorescence length or number of flowers) as response variables. Variation among subspecies was then investigated using linear mixed effects models with subspecies as a fixed effect, population as a random effect and plant size traits as response variables. Variation in floral colour (characterized by mean and SD values in the R, G, B, MW and SW channels) and scents (characterized by log-transformed absolute amounts of each compound) among subspecies and populations of the same subspecies was investigated using Partial Least Square-Discriminant Analysis (PLS-DA). Cross-validation tests were performed to estimate classification error rates (M-fold method) and their statistical significance (permutation tests performed with 999 permutations) for both floral colour and odour. Variation in fruit set among subspecies and populations of the same subspecies was investigated using Wilcoxon–Mann–Whitney tests.

### 2.5.2 | Estimation of selection gradients

Phenotypic selection exerted on floral traits was quantified using the method of Lande and Arnold (1983). As a preliminary analysis, selection differentials  $S$  were estimated in each population as the covariance between relative female reproductive success and standardized trait values using univariate regressions.  $p$ -Values were corrected using the false discovery rate (FDR) method (Verhoeven, Simonsen, & McIntyre, 2005). Afterwards, directional selection gradients  $\beta$  were estimated in each population using multivariate regressions with relative female reproductive success as the response variable and standardized trait values as explanatory variables. Because plant size traits were highly correlated (see Table S1), only plant height was included in these regressions. Regarding floral colour traits, a global principal component analysis (PCA) was performed on the entire dataset (i.e., including all seven populations) in order to reduce the number of traits included in our models and transform potentially correlated variables into uncorrelated principal components (PCs). Scores on the first PC (accounting for 55% of the total variance) were included as an explanatory variable in our models. Regarding floral odour traits, because we had no a priori knowledge about the biological relevance of individual compounds, we chose to estimate selection exerted on entire blends of floral scents (see Method S5). To do so, as for floral colour traits, we used PCA, but this time one PCA was performed in each population. We proceeded this way because many compounds were detected in one or a few populations only (see Results), and thus including scores on global PCs in our models would amount to estimating selection on traits that may not be expressed in a given population. The correlation between PCs and individual compounds is shown in Table S5. Our final models included plant height, scores on the first colour PC, total emission rate and scores on the first two to four odour PCs (i.e., PCs accounting for at least 50% of the total variance in floral scents within each population), except in Le-Cannet-des-Maures, in which we did not have floral colour data and only included plant height and floral odour variables. Only linear terms

were included in these regressions, as preliminary analyses showed no evidence of stabilizing or correlational selection. Variance inflation factors (VIFs) were computed to test for multicollinearity between explanatory variables, but they almost never exceeded 2 (i.e., they exceeded 2 for two variables of 38, and remained below 2.1 in both cases), indicating no problem of multicollinearity (Quinn, Gerald, & Keough, 2002).

All statistical analyses were performed using the software R version 3.4.3 (R Core Team, 2017).

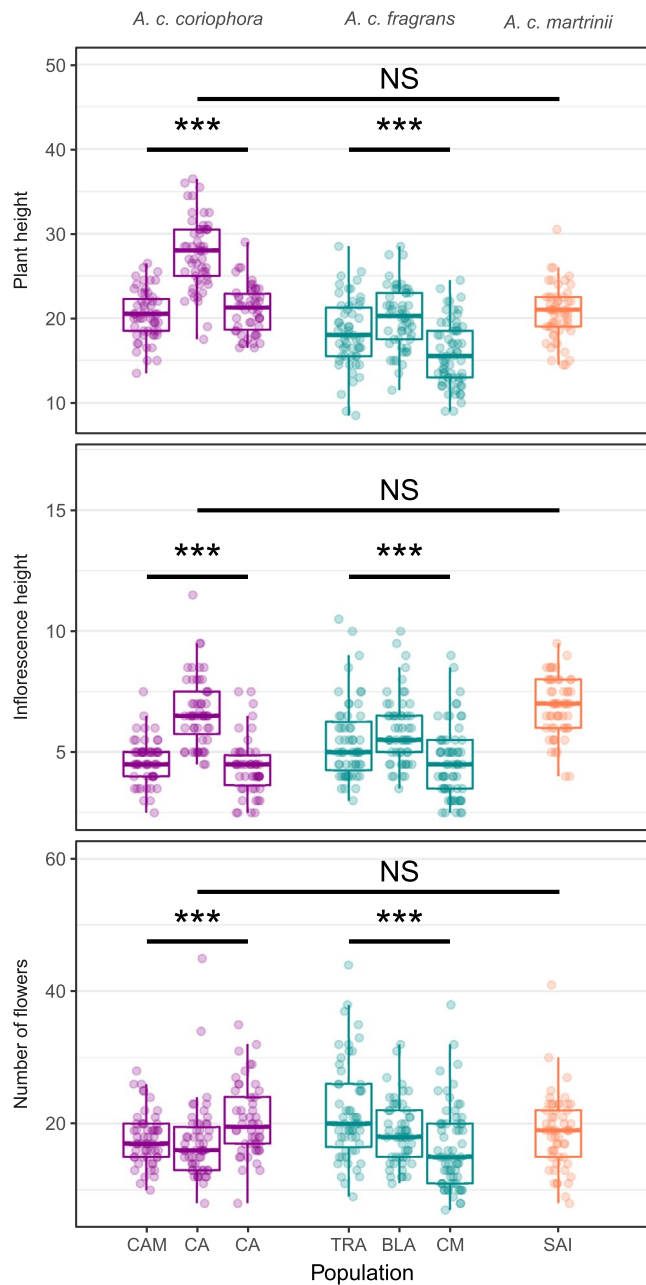
## 3 | RESULTS

### 3.1 | Variation in floral traits and female reproductive success

Variation in plant height, inflorescence length and number of flowers was significant when comparing populations of the same subspecies, but not when comparing subspecies (Figure 2 and Table S2).

By contrast, variation in floral colour was significant when comparing either subspecies (classification error rate = 24.7%,  $p = .001$ ; Figure 3) or populations of the same subspecies (classification error rate = 35.50% for *A. c. coriophora* and 6.69% for *A. c. fragrans*,  $p = .001$  in both cases). Specifically, flowers of *A. c. martrinii* were dark (low mean values in the R, G and B channels) relative to those of *A. c. fragrans* (high mean values in the R, G and B channels), whereas flowers of *A. c. coriophora* were characterized by intermediate mean values.

Forty-eight compounds (fatty acid derivatives, benzenoids and terpenoids) were detected in the blends of *A. c. coriophora*, *A. c. fragrans* and *A. c. martrinii* (see Table S3), including 21 that were shared among all three taxa (Figure 4). Floral scents of *A. c. coriophora*, *A. c. fragrans* and *A. c. martrinii* were composed of 35, 31 and 32 floral volatiles, respectively. Floral odour was well differentiated among subspecies (classification error rate = 0%,  $p = .001$ ; Figure 4). This differentiation was both qualitative (i.e., presence/absence of compounds) and quantitative (i.e., absolute and relative amounts of compounds). Indeed, 19 of 48 floral volatiles were taxon-specific, including some predominant compounds (Figure 4). Floral scents of all three subspecies were dominated by 1,4-dimethoxybenzene (on average 44.45% of the blend: 36.56% for *A. c. coriophora*, 44.54% for *A. c. fragrans* and 66.68% for *A. c. martrinii*). However, the second predominant compound was different in all three subspecies: flowers of *A. c. coriophora* emitted high amounts of 1,2,4-trimethoxybenzene (11.44% of the blend), flowers of *A. c. fragrans* high amounts of *p*-anisaldehyde (30.05% of the blend) and flowers of *A. c. martrinii* high amounts of methyl-*p*-anisate (11.33% of the blend). Note that floral odours of *A. c. coriophora* and *A. c. martrinii* were more similar to each other (28 compounds in common) than to those of *A. c. fragrans* (22 and 21 compounds in common, respectively; Figure 4). Floral scents were also well differentiated among populations of either *A. c. coriophora* or *A. c. fragrans* (classification error rate = 0% and  $p = .001$  in both cases; see Figure S2). Here again,



**FIGURE 2** Variation in plant size traits among subspecies and populations of the same subspecies. BLA, Blandas; CA, Comps-sur-Artuby; CAM, Camprieu; CM, Le-Cannet-des-Maures; SAI, Saillagouse; SOU, Sournia; TRA, Trassanel; \*\*\* $p < .001$ ; NS: nonsignificant. See Table S2 for details

this differentiation was both qualitative and quantitative, with several compounds being only found in one or two populations of three (see Table S3). Interestingly, floral odour variation within subspecies appeared to be geographically structured, with populations ordered along a longitudinal gradient on axis 2 (i.e., westernmost populations at the bottom and easternmost populations at the top of the scatterplot; see Figure S2).

Mean fruit set of open-pollinated plants varied from 30.68% in Comps-sur-Artuby to 82.84% in Trassanel, with significant variation among populations of the same subspecies ( $F_{2,157} = 33.54$ ,  $p = 2e^{-13}$

for *A. c. coriophora* and  $F_{2,159} = 33.71$ ,  $p = 6e^{-13}$  for *A. c. fragrans*), but not among subspecies ( $\chi^2(2, N = 376) = 0.38$ ,  $p = .83$ ).

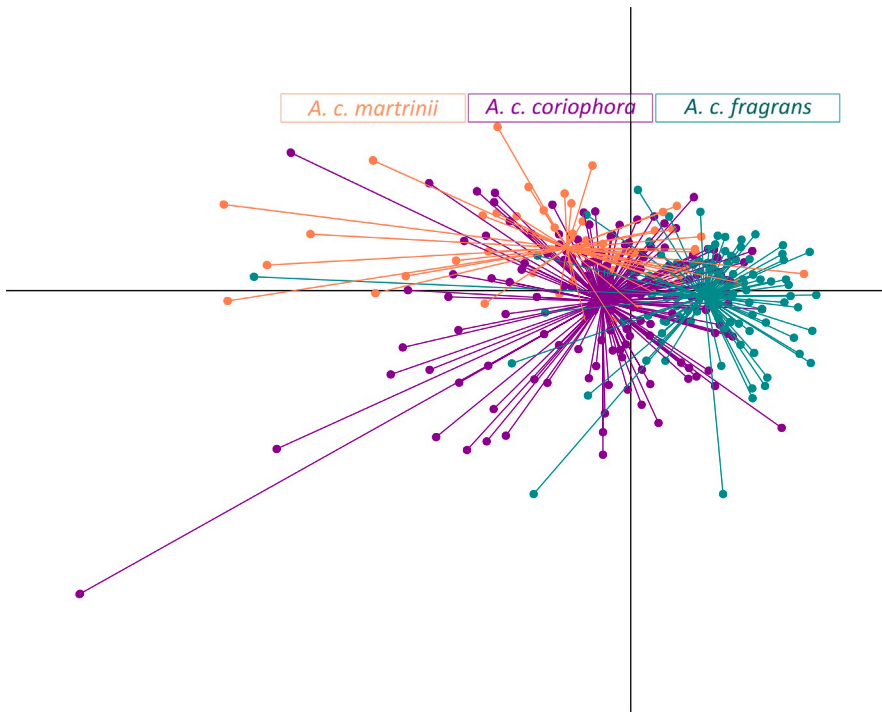
### 3.2 | Estimation of selection gradients

After correction, significant selection differentials were found for plant height, inflorescence length and number of flowers in most populations (see Table S4). By contrast, no selection was detected on floral colour traits, using either RGB values or honeybee photoreceptor excitation values (see Table S4). Regarding floral scent traits, no selection was found on total emission rate, whereas significant selection differentials were detected for absolute amounts of some compounds in two *A. c. coriophora* and one *A. c. fragrans* populations (see Table S4).

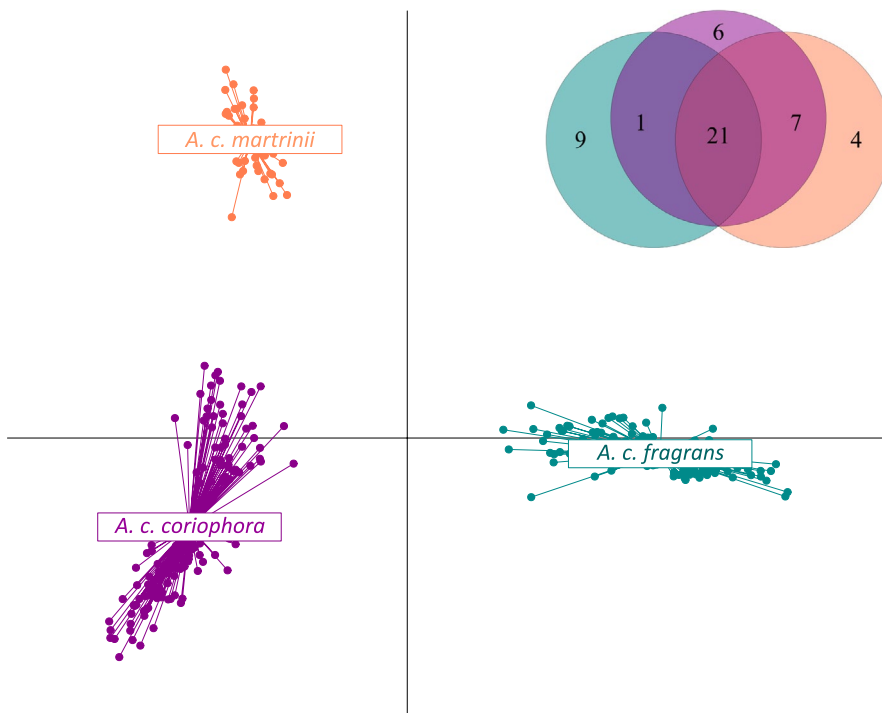
Multivariate regressions showed positive selection on plant height in all six populations of *A. c. coriophora* and *A. c. fragrans*, as well as positive or negative selection on at least one of the floral scent PCs in all three populations of *A. c. fragrans* (Table 1). More specifically, in Le-Cannet-des-Maures, we detected positive selection on the first PC and negative selection on the second PC (Table 1). PC1 was positively correlated with several taxon-specific compounds (namely *p*-anisaldehyde, 2-(4-methoxyphenyl)-ethan-1-ol, methyl-*p*-anisate, (*E*)-methylcinnamate and 3-acetylcyclohexylacetate), indicating positive selection on these volatiles. PC2 was negatively correlated with 5-methyl- $\delta$ -valerolactone and 4,5-dimethyl- $\delta$ -valerolactone—here again a taxon-specific compound—indicating positive selection on these volatiles. In Blandas, we detected negative selection on the first PC (Table 1). This PC was negatively correlated with the same set of taxon-specific compounds (i.e., *p*-anisaldehyde, 2-(4-methoxyphenyl)-ethan-1-ol, methyl-*p*-anisate, (*E*)-methylcinnamate and 3-acetylcyclohexylacetate), indicating positive selection on these volatiles here again. Note that this PC was also negatively correlated with 1,2,3,5-tetramethoxybenzene, a compound that in *A. c. fragrans* was only found in the Blandas population. In Trassanel, we detected negative selection on the second PC (Table 1). This PC was negatively correlated with methylisohexanoate—a compound specific to the Blandas and Trassanel populations—as well as with 6-methylhept-5-en-2-one and *p*-methylanisole—two compounds that were also found in *A. c. coriophora* and *A. c. martrinii*. In *A. c. coriophora*, we detected no selection on floral odour in Comps-sur-Artuby, Camprieu or Sournia (Table 1). In *A. c. martrinii*, we found no selection on plant height nor on floral scent PCs (Table 1). No selection was found on floral colour PCs nor on total emission rate in *A. c. coriophora*, *A. c. fragrans* or *A. c. martrinii* (Table 1).

## 4 | DISCUSSION

Phenotypic differentiation at biologically relevant traits among conspecific populations is often believed to result from divergent selective pressures imposed upon these traits. In this study, we characterized floral trait differentiation among seven populations of the widespread, generalist orchid species *Anacamptis coriophora* and we explored the relationship between these traits and female



**FIGURE 3** Scatter plot of the Partial Least Square-Discriminant Analysis (PLS-DA) performed on floral colour traits for the three subspecies (*Anacamptis coriophora* subsp. *coriophora* in purple, *A. coriophora* subsp. *fragrans* in turquoise and *A. coriophora* subsp. *martrinii* in orange)



**FIGURE 4** Scatter plot of the Partial Least Square-Discriminant Analysis (PLS-DA) performed on floral scent traits for the three subspecies (*Anacamptis coriophora* subsp. *coriophora* in purple, *A. coriophora* subsp. *fragrans* in turquoise and *A. coriophora* subsp. *martrinii* in orange), and Venn diagram showing number of taxon-specific compounds

reproductive success. We found a significant differentiation both among and within subspecies in terms of floral colour and scents, with one of the two predominant compounds emitted by each subspecies being taxon-specific. We hypothesized that such variation in floral colour and odour was mirrored by variation in phenotypic selection exerted on these traits. In particular, one of our expectations was to detect positive selection on taxon-specific floral volatiles. We indeed found such a pattern in *A. c. fragrans*, but not in *A. c. coriophora* and *A. c. martrinii*. This partial match between the observed pattern of chemical differentiation and phenotypic selection exerted on floral

scents suggests that selective processes could contribute to ongoing floral odour divergence among *A. coriophora* subspecies. On the other hand, we observed no such trend for floral colour, suggesting that other processes may govern floral colour evolution in *A. coriophora*.

The three studied subspecies were characterized by comparable plant height, inflorescence length and number of flowers, with variation among populations of the same subspecies. We found a positive correlation between plant height and female reproductive success in most populations. Such a correlation has been documented in other plants (Gervasi & Schiestl, 2017; Gómez, 2007), including



**TABLE 1** Selection gradients  $\beta \pm SE$  on plant height, total emission rate and principal components (PCs) of the PCAs performed on floral colour and odour data. In each population, selection gradients are extracted from one multiple regression including all floral traits (i.e., one column corresponds to one model). Odour PCs are population-specific (i.e., one individual PCA performed in each population)

Subspecies	COR			FRA			MAR
	CAM	CA	SOU	BLA	CM	TRA	SAI
Plant height	0.14 ± 0.04**	0.43 ± 0.13**	0.12 ± 0.05 <sup>†</sup>	0.22 ± 0.06***	0.37 ± 0.13**	0.28 ± 0.06***	0.03 ± 0.08
Colour PC1	0.002 ± 0.04	0.08 ± 0.12	-0.02 ± 0.05	-0.02 ± 0.06	-	0.02 ± 0.05	-0.04 ± 0.08
Total emission rate	0.05 ± 0.05	-0.20 ± 0.17	-0.03 ± 0.07	-0.08 ± 0.08	0.10 ± 0.14	0.08 ± 0.06	-0.02 ± 0.12
Odour PC1	-0.02 ± 0.05	-0.16 ± 0.17	-0.09 ± 0.06	-0.17 ± 0.07 <sup>†</sup>	0.27 ± 0.12 <sup>†</sup>	-0.08 ± 0.06	0.02 ± 0.11
Odour PC2	0.06 ± 0.04	0.18 ± 0.13	0.06 ± 0.05	0.03 ± 0.06	-0.33 ± 0.12**	-0.11 ± 0.05 <sup>†</sup>	0.004 ± 0.08
Odour PC3	-0.02 ± 0.04	-	-	-	-	0.08 ± 0.05	0.02 ± 0.08
Odour PC4	-0.04 ± 0.04	-	-	-	-	-	-

Note: The number of PCs included in each model was chosen to explain at least 50% of the total variance in floral scents within each population.

Abbreviations: BLA, Blandas; CA, Comps-sur-Artuby; CAM, Camprieu; CM, Le-Cannet-des-Maures; COR, *Anacamptis coriophora coriophora*; FRA, *A. c. fragrans*; MAR, *A. coriophora c. martrinii*; SAI: Saillagouse; SOU, Sournia; TRA: Trassanel.

<sup>†</sup> $p < .05$ ; \*\* $p < .01$ ; \*\*\* $p < .001$ .

orchids (Sletvold et al., 2010; Sletvold, Trunschke, Wimmergren, & Ågren, 2012), and could be mediated by pollinators, as taller individuals may be more easily detected by foraging insects (Donnelly, Lortie, & Aarssen, 1998). However, because plant height was correlated to number of flower in most populations, its apparent effect on female reproductive success may be due to the obvious correlation between flower and fruit numbers. In addition, plant height may be linked to other floral or vegetative genetically coded traits affecting female reproductive success, causing indirect selection on this character (Alexandersson & Johnson, 2002; Gómez, 2000). Alternatively, non-genetic factors such as plant age or microhabitat could affect plant height and fruit set in a parallel fashion and explain this correlation.

The three studied subspecies were characterized by different floral colours, yet no phenotypic selection was found on this trait, contrary to what has been reported in other orchids (Gross et al., 2016; Sletvold, Trunschke, Smit, Verbeek, & Ågren, 2016). Both visual and olfactory signals are known to be important for bee attraction (Burger et al., 2010), but visual traits may be of secondary importance compared to olfactory traits for *A. coriophora* pollination. Alternatively, this finding could be explained by conflicting selective pressures imposed by mutualists and antagonists (Frey, 2004). The fact that we did not detect any selection on floral colour despite significant differentiation both among and within subspecies could either suggest that selection has driven floral colour divergence in the past but is now relaxed (or too weak to be detected with our limited sample size), or that other processes, such as genetic drift, are responsible for the observed pattern of floral colour differentiation.

The three studied subspecies were also characterized by different floral scents, with two predominant compounds per subspecies, one shared among all three taxa (1,4-dimethoxybenzene) and one taxon-specific (1,2,4-trimethoxybenzene in *A. c. coriophora*, *p*-anisaldehyde in *A. c. fragrans* and methyl-*p*-anisate in *A. c. martrinii*). In addition, many other taxon-specific floral volatiles were found in the blends of the three studied subspecies, including fatty acid derivatives, benzenoids and terpenoids. Our results are congruent with previous reports on the floral

odour chemistry of *A. c. fragrans* (Dormont, Delle-Vedove, Bessièrè, & Schatz, 2014; Salzmann, Nardella, Cozzolino, & Schiestl, 2007), whereas floral scents of *A. c. coriophora* and *A. c. martrinii* were analysed for the first time in this study. In particular, we demonstrated that *A. c. martrinii* emits significant amounts of floral volatiles, despite being described as odourless in the literature (Delforge, 2006). Because the three studied subspecies grow in different habitats, probably characterized by distinct microclimatic conditions and biotic communities, phenotypic plasticity cannot be ruled out as a potential driver of floral odour variation in *A. coriophora* (Majetic, Raguso, & Ashman, 2009). In fact, there is growing evidence that plants can plastically respond to increasing temperatures (Farré-Armengol, 2014), drought (Campbell, Sosenski, & Raguso, 2019) or other biotic or abiotic factors (Schiestl, Kirk, Bigler, Cozzolino & Desurmont, 2014) through quantitative changes in floral scent emissions (i.e., changes in total emission rate or in the relative amounts of the different compounds). A common garden experiment would have allowed us to disentangle the relative importance of environmental versus genetic factors for floral odour variation, but such experiments are difficult to set up in slow-growing plants such as orchids. However, it seems unlikely that the consistent difference in the predominant floral volatiles emitted by the three studied subspecies is due to phenotypic plasticity alone, and it seems reasonable to assume that such difference is at least partly genetically based. If this assumption holds, floral scent differentiation among subspecies may be the outcome of selective and/or neutral processes. Compared to other floral traits, few phenotypic selection studies have dealt with scents so far, despite the well-known implication of olfactory signals in biotic interactions, particularly pollination (Raguso, 2008). Evidence from these studies suggests that selection usually favours the most odorous plants (Majetic et al., 2009; Parachnowitsch, Raguso, & Kessler, 2012), or plants emitting the highest amounts of a few biologically relevant compounds (Chapurlat, Ågren, Anderson, Friberg, & Sletvold, 2019; Parachnowitsch et al., 2012; Schiestl, Huber, & Gomez, 2011). Here, we found no significant selection gradients on total emission rate, nor did we find consistently significant selection differentials on amounts

of specific floral volatiles. By contrast, when considering entire blends of floral scents, we detected phenotypic selection on sets of correlated compounds—mostly taxon-specific floral volatiles—in all three populations of *A. c. fragrans*. Although other selective agents could account for this, pollinators may play a role here. First, several of the compounds found to be under positive selection are known to be involved in the chemical communication between plants and pollinators. For example, the benzenoid *p*-anisaldehyde is known to constitute an honest signal of reward for pollinating bees (Knauer & Schiestl, 2015) and was shown to be under pollinator-mediated selection in *Brassica rapa* plants (Gervasi & Schiestl, 2017). Second, fruit set was significantly (although not strongly, as expected in this rewarding species) pollen-limited in all three populations of *A. c. fragrans* (see Figure S3), indicating that pollinators may act as selective agents in these populations (Sletvold & Ågren, 2014; Trunschke, Sletvold, & Ågren, 2017). On the other hand, we detected no phenotypic selection on floral scents of *A. c. coriophora* and *A. c. martrinii*, despite the fact that female reproductive success was significantly pollen-limited at least in *A. c. martrinii* (see Figure S3). This finding could be explained by conflicting selective pressures imposed by mutualists and antagonists (Schiestl et al., 2011), especially since some compounds emitted by these two subspecies—including 1,2,4-trimethoxybenzene—are known to attract both pollinators and herbivores (Andrews, Theis, & Adler, 2007). Alternatively, floral scents may not play a predominant role in *A. c. coriophora* and *A. c. martrinii* pollination. The ecological function of floral odour in *A. coriophora* and its link with other fitness components, particularly pollen export, should be studied more thoroughly to better understand the selective pressures at work here.

In the orchid *Gymnadenia odoratissima*, selection on floral colour and scents was shown to differ between lowland and mountain populations and was broadly congruent with the observed pattern of floral trait differentiation between these populations (Gross et al., 2016). Likewise, we showed a partial match between the observed pattern of chemical differentiation and selective pressures imposed upon floral odour in *A. coriophora*. This suggests that selective processes could contribute to ongoing floral scent divergence among *A. coriophora* subspecies, potentially along with other processes. Population genetic studies are needed to assess whether selective and neutral processes act in concert to drive such divergence, or whether selection promotes phenotypic differentiation despite substantial gene flow among subspecies within geographical regions. In any case, it should be pointed out that such a scenario relies on two important assumptions: first, that phenotypic selection is consistent across years, which was not tested here; and second, that variation in floral scents is heritable. However, floral size or shape traits often exhibit relatively high heritabilities (Ashman & Majetic, 2006), little is known regarding the heritability of floral odour, although macroevolutionary studies showing consistent associations between functional groups of pollinators and blends of floral scents suggest that the latter are evolvable traits (Harder & Johnson, 2009). Accordingly, an artificial selection experiment on *Brassica rapa* has shown that floral odour can evolve rapidly under phenotypic selection (Zu, Blanckenhorn, & Schiestl, 2016). However, whether this is the case in our study system remains to be tested.

It is worth noting that contrary to Gross et al. (2016), who showed consistent differences in the pollinator guilds associated with lowland versus mountain populations of *G. odoratissima*, we did not find any obvious associations between *A. coriophora* subspecies and taxonomic or functional groups of pollinators, at least for *A. c. coriophora* and *A. c. fragrans* (see Table S6). In fact, our data suggest that pollinator communities vary among all seven sites, regardless of the *A. coriophora* subspecies growing locally. More data on the composition of pollinator communities and on the contribution of the various pollinator species to reproductive success are needed to understand the role potentially played by pollinators in the diversification of *A. coriophora*.

To our knowledge, this study is the first to document variation in phenotypic selection exerted on floral scents among subspecies. Our results suggest that selection could contribute to ongoing chemical differentiation between *A. coriophora c. fragrans* on the one hand and *A. c. coriophora* and *A. c. martrinii* on the other hand. The contribution of selection to the marked phenotypic differentiation evidenced among *A. coriophora* subspecies—relative to that of neutral processes—remains to be assessed, by analysing patterns of genetic differentiation among the latter. Further studies are also needed to assess whether such differentiation represents the early stages of speciation, by testing whether floral traits can promote prezygotic reproductive isolation among these subspecies.

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## SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section.

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