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## Quantitative measures of leaf epidermal cells as a taxonomic and phylogenetic tool for the identification of *Stanhopea* species (Orchidaceae)

(With 3 Tables & 2 Figures)

*Mediciones cuantitativas en las células epidérmicas de la hoja como herramienta taxonómica y filogenética para la identificación de especies de Stanhopea (Orchidaceae)*

(Con 3 Tablas y 2 Figuras)

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**Abstract.** Orchids of the genus *Stanhopea* are currently identified only by their floral structure characteristics. A statistical analysis of a significant number of species of this genus disclosed that measurements of adaxial and abaxial epidermal cell surface areas can be correlated with specific recognized species by a leaf printing method. This allows an objectively either positive or nearly positive confirmation of the identity of a species in the absence of flowers and without damage to plants. When ordering the mean values obtained for these surface areas in each species in a decreasing order, a correlation was observed in a hierarchical way that went from primitive to more advanced floral forms. This reflects the evolutionary radiation of the genus. It is established that in *Stanhopea*, the presence of large leaf epidermal cells on species from South America represents a primitive evolutionary condition that became to smaller cells in evolutionarily more recent individuals as the genus radiated towards Mexico.

**Key Words:** *Stanhopea*, quantitative taxonomy, morphometry, leaf printing, Orchidaceae.

**Resumen.** Las orquídeas del género *Stanhopea* son identificadas actualmente solamente por las características estructurales de la flor. El

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análisis estadístico de un número significativo de especies de este género reveló que las áreas promedio de las células epidérmicas adaxiales y abaxiales se pueden correlacionar con especies específicamente reconocidas utilizando un método de impresión de la hoja. Esto permitiría una confirmación positiva o casi positiva de la identidad de una especie de modo objetivo, en ausencia de flores y sin daño a las plantas. Al ordenar los valores obtenidos para dichas áreas de las células epidérmicas en cada especie en un orden descendente, se observó una correlación de modo jerárquico que va de las formas florales primitivas a las formas más avanzadas. Esto reflejó una radiación evolutiva del género. Se establece que en *Stanhopea*, las grandes células epidérmicas presentes en las hojas de especies de Sudamérica representan una condición evolutiva primitiva que fue cambiando a células más pequeñas en individuos evolutivamente más recientes a medida que el género se expandió hacia México.

**Palabras clave:** *Stanhopea*, taxonomía cuantitativa, morfometría, impresión de hoja, Orchidaceae.

## INTRODUCTION

The collection, pressing and drying of plant samples, which are then analyzed for its classification and taxonomical identification, has been the classical botanical protocol to investigate and describe the flora of a particular region. If any plant sample could not be recognized as a known species, it was then considered as a new species. Classical taxonomy stresses the need of selecting and describing a “type” from a plant population, which is then deposited in a recognized herbarium, to validly describe a new species. This information might then be published in a high impact factor scientific Journal. However, because of the importance historically placed by Linnaeus on floral structures, many orchids have been described from just the floral structures, disregarding any other organ for taxonomical classification.

On the other hand, most new orchid species have been described by foreign botanists residing in remote localities from the plant’s native area. They have mostly worked with floral specimens either dried or pressed or preserved in alcohol or some other fixative solution. For many years of modern orchid taxonomy, thousands of plants were collected from the New World tropical countries and shipped to greenhouses and taxonomists of Belgium, England, Germany, France and the USA. Although many plants survived, thousands of them were taken out from their habitats and either died in transit or during the establishment phase because of improper transporting and/or ignorance of their cultivation requirements.

In the 1960’s, orchid conservationists in the United States and tropical countries achieved an international ecological awareness which result-

ed in the 1973 United States Federal Endangered Species Act. This was followed by an international agreement designed to preserve endangered species, prohibiting either the movement or export of any orchid specimen from its native habitat. However and despite these protecting and regulation laws, the illegal collection and traffic of orchid plants still continues.

The present situation is multifaceted. Orchid conservation requires species identification with techniques which must not be in conflict with existing laws and regulations. Also, identification based only on floral characteristics requires finding these relatively scarce plants when flowering. Precise knowledge of the species morphological and anatomical characteristics may allow a rapid and efficacious method of plant recognition. This would help to solve the above mentioned constraints.

The taxonomy of *Stanhopea* has been based solely on floral characters with only a note about vegetative structures (Arnold, 1928; Dodson & Frymire, 1961; Dodson, 1963, 1975). Furthermore, Curry et al. (1988) reported that the taxonomy of *Stanhopea* rests exclusively on flower morphological differences, which have been influenced by pollinators. This system of genus classification certainly has merit, but problems arise when either the boundary line of floral characters between close-related species is relatively indistinct or no flowers are present. Therefore, development of a system capable of reasonably and objectively correlate and identify a specimen against an already-described species is necessary. This method, by use of other than floral characteristics, would provide a useful identification tool at any time and place without regard to restrictions imposed by the flowering period.

*Stanhopea* was selected for this study due to certain attributes which make it a rigorous test genus. Flowers of *Stanhopea* do not last more than two or three days. It also exhibits few gross morphological differences at the vegetative developmental stage. Most of the literature only describes this aspect as "typical for the genus". If identifications are to be made using present classical methods, they must be done when plants are either flowering in their habitats or removed to a more convenient location to the botanist for observation when it does flower.

It may appear to be a relatively simple task to collect a plant, take care of it in a greenhouse, and then identify it when it flowers. However, the new habitat may not be appropriate to induce flowering because greenhouse conditions are far different from those in its natural habitat. This is in addition to the legal and regulatory problems related to the collection and transport of plant material.

*Stanhopea* is longitudinally widespread throughout the neotropical and tropical latitudes of the Western hemisphere. Because of this, it was

hypothesized that an other-than-flower identification system might provide clues concerning intra-species relationships and the evolutionary radiation pattern of this genus. Specifically, it was hypothesized that identification of unknown species can be determined by comparing their mean leaf epidermal cell surface areas with those of known species.

Differences in morphometric characteristics of leaf epidermal cells and other structures (i.e., stomata) have been reported for different species of the genus *Stanhopea* (Ferry et al., 1997). An identification protocol will be described which (1) does not make changes to plants in the field, (2) requires no material to be taken from the habitat, (3) it is economical and easy to do, and (4) it is statistically supported rather than being a subjective description. Results using morphological differences on a great portion of the entire genus *Stanhopea* in this study, which may be intrinsic or due to geographical habitat changes, allowed to achieve a statistical method for the accurate identification of species in the absence of flowers.

## MATERIALS AND METHODS

**Biological material.** Fifty four specimens of the genus *Stanhopea* were obtained which represented 34 species (Table 1). Although some species names are listed more than once in this table, no species or hybrid material represents the same clonal material. Among the samples, six *Stanhopea* specimens were received as unknown species (C03, 26, 33, 53, 77 and 83). Data obtained from those unknown species were statistically checked against similar data obtained from species confirmed by flowering. Subsequent flowering by the unknown species enabled to validate the statistically derived prediction.

**Leaf-printing.** The method used to obtain leaf-prints produces best results when the leaf surface areas are clean. Under greenhouse or field conditions, residues on the leaf surface areas must be eliminated by washing both adaxial and abaxial leaf surfaces with water, gently wiping them with a Teflon kitchen sponge, and allowing them to air-dry for a few minutes. If the plant had been exposed to an excessive amount of dust or residues, it may be useful to employ a very mild detergent, followed by rinsing in water and allowing the leaf to dry. Clean white Styrofoam “popcorn” was dissolved in xylol until the liquid was about the viscosity of warm syrup. This solution was applied to a clean leaf surface over an area of 2 x 5 cm. After allowing to air-dry for 2-3 minutes, a short strip of clear, transparent tape was applied evenly and firmly over the film with enough pressure as not to damage the leaf cells. The tape was peeled and gently passed onto a glass slide. A thin glass coverslip may be applied if permanent con-

**Table 1.** Summary of data obtained from the analyzed *Stanhopea* specimens. Data shown are epidermal cell surface areas ( $\mu^2$ ). Each value is the mean  $\pm$  1 s.e. (n=200).

**Tabla 1.** Resumen de datos obtenidos de los especimenes de *Stanhopea* analizados. Los datos que se muestran son superficies celulares epidérmicas ( $\mu^2$ ). Cada valor es el promedio  $\pm$  1 e.e. (n=200).

Region	C#	Specimen	Origin	Adaxial	Abaxial
Mostly South America	21	<i>S. reichenbachiana</i>	Colombia	5.624 $\pm$ 1.540	3.767 $\pm$ 660
	46	<i>S. reichenbachiana</i>	Colombia	5.494 $\pm$ 1.035	3.751 $\pm$ 955
	47	<i>S. embreei</i>	Ecuador	5.311 $\pm$ 1.146	3.420 $\pm$ 1.340
	54	<i>S. madouxiana</i>	Colombia	5.305 $\pm$ 1.384	3.264 $\pm$ 629
	31	<i>S. platyceras</i>	Colombia	5.266 $\pm$ 1.066	3.355 $\pm$ 699
	50	<i>S. pulla</i>	Panama	5.045 $\pm$ 1.006	3.239 $\pm$ 1.026
	18	<i>S. pozoii</i>	Colombia	4.891 $\pm$ 1.008	2.992 $\pm$ 738
	27	<i>S. candida</i>	Colombia	4.857 $\pm$ 1.201	2.790 $\pm$ 769
	67	<i>S. grandiflora</i>	Unknown	4.841 $\pm$ 1.007	3.675 $\pm$ 699
	25	<i>S. grandiflora</i>	Brazil	4.817 $\pm$ 1.493	3.714 $\pm$ 614
	16	<i>S. insignis</i>	Colombia	4.781 $\pm$ 1.207	3.252 $\pm$ 570
	3	<i>S. insignis</i>	Colombia	4.770 $\pm$ 990	3.248 $\pm$ 632
	55	<i>S. logipes</i>	Colombia	4.742 $\pm$ 628	3.242 $\pm$ 942
	57	<i>S. lietzei</i>	Unknown	4.725 $\pm$ 856	3.182 $\pm$ 799
	28	<i>S. connota</i>	Colombia	4.713 $\pm$ 878	3.214 $\pm$ 723
	52	<i>S. connota</i>	Unknown	4.707 $\pm$ 1.084	3.193 $\pm$ 899
	14	<i>S. wardii</i>	Unknown	4.660 $\pm$ 1.159	3.571 $\pm$ 888
	62	<i>S. wardii</i>	Unknown	4.657 $\pm$ 970	3.584 $\pm$ 698
	12	<i>S. wardii</i>	Guatemala	4.603 $\pm$ 1.072	3.547 $\pm$ 1.217

Region	C#	Specimen	Origin	Adaxial	Abaxial
Mostly South America	13	<i>S. wardii</i>	Colombia	4.595±1.194	3.526±758
	20	<i>S. tricornis</i>	Colombia	4.532±1.166	3.556±653
	22	<i>S. eburnea</i>	Brazil	4.498±795	3.726±1.328
	66	<i>S. eburnea</i>	Unknown	4.472±1.085	3.654±922
	24	<i>S. jenischiana</i>	Colombia	4.368±824	2.899±956
	32	<i>S. posadae</i>	Colombia	4.336±982	3.538±739
	48	<i>S. inodora</i>	Unknown	4.318±785	3.450±680
	30	<i>S. impressa</i>	Colombia	4.308±962	3.120±512
	7	<i>S. tigrina</i>	Mexico	4.259±852	2.858±701
	59	<i>S. tigrina</i>	Mexico	4.245±837	2.917±640
	33	<i>S. tigrina</i>	Guatemala	4.234±869	2.817±815
	26	<i>S. tigrina</i>	Unknown	4.220±805	2.856±734
	29	<i>S. haselowania</i>	Peru	4.216±899	3.556±907
Mostly Central America	6	<i>S. oculata</i>	Mexico	4.174±1.240	2.748±717
	77	<i>S. oculata</i>	Mexico	4.137±734	2.705±761
	5	<i>S. oculata</i>	Panama	4.123±1.103	2.732±535
	23	<i>S. ecornuta</i>	Colombia	3.837±1.475	2.662±802
	58	<i>S. panamensis</i>	Panama	3.724±822	2.657±393
	17	<i>S. costaricensis</i>	Costa Rica	3.705±543	2.649±610
	83	<i>S. amoena</i>	Unknown	3.617±861	2.746±699
Mexico	69	<i>S. radiosa</i>	Mexico	3.586±702	2.495±651
	88	<i>S. maculosa</i>	Mexico	3.536±606	2.109±452
	10	<i>S. martiana</i>	Mexico	3.444±861	2.609±514

Region	C#	Specimen	Origin	Adaxial	Abaxial
Mexico	21	<i>S. reichenbachiana</i>	Colombia	5.624±1.540	3.767±660
	64	<i>S. graveolens</i>	Unknown	3.417±850	2.325±634
	8	<i>S. graveolens</i>	Mexico	3.380±878	2.372±683
	9	<i>S. graveolens</i>	Mexico	3.289±873	2.310±601
	15	<i>S. nigroviolacea</i>	Mexico	3.129±892	2.264±590
	1	<i>S. saccata</i>	Unknown	3.097±675	2.219±584
	45	<i>S. saccata</i>	Mexico	3.051±581	2.224±467
	2	<i>S. saccata</i>	Guatemala	3.041±733	2.219±413
	53	<i>S. hernandezii</i>	Mexico	2.865±594	2.087±480
	49	<i>S. hernandezii</i>	Unknown	2.817±658	2.106±431
	4	<i>S. hernandezii</i>	Mexico	2.783±687	2.096±443
	11	<i>S. intermedia</i>	Mexico	2.691±586	2.060±652
	51	<i>S. intermedia</i>	Mexico	2.674±574	2.020±540

servation of the slide is desired. This not only ensures that the leaf print is held flat but gives some protection to the transparent tape itself. Date, location and the study leaf surface area (either adaxial or abaxial), should be marked on the slide.

**Photographic recording and cell measurements.** Slides were placed under a microscope and photographed at a 80x magnification using a compound microscope with a trinocular head on which a Nikon HFM photo system was mounted. An indexing lens was inserted in the field lens assembly to print an index mark on each photomicrograph which enable to take accurate measurements directly from the print. A Reichert-Jung micrometer stage slide of 2 mm, divided into units of 0.01mm, was used to establish the lens correction factor for the magnification of each objective lens. Cell print photomicrographs were obtained randomly from the glass slides, utilizing a minimum of two photographs for each leaf surface area on each specimen. After dividing each photograph into 8 equal quadrants, 25 cells were randomly selected, and measurements of their length and width were individually recorded to give a closest approximation to the correct leaf cell surface area.

**Statistical analysis.** Individual specimen spreadsheets were computer-linked which provided a combined summary data sheet for each specimen; it also allowed specimen ranking according to their leaf epidermal cell areas, arranged in ascending or descending order. Data from individual adaxial and abaxial leaf epidermal cells of confirmed species were compared statistically. Entering individual data measurements into a statistical program (SAS ver. 6.12) provided a computer-generated analysis of variance readout ( $p= 0.05$ ) for each specimen.

Data obtained from unknown *Stanhopea* samples were statistically compared ( $p=0.05$ ) with previously correlated data from confirmed species, and predictions were made on their identity. As mentioned above, validity of the predictions was checked out by identification of the samples after their flowering.

To double-check data accuracy, 10 samples were randomly selected and 200 cells were measured for each parameter, which was compared with the original samples (data not shown); an error margin less than 0.01% was used with this purpose. Four other randomly selected samples were measured for another individual without knowledge of the original data. Results of the 4 samples were less than 2% different from those of the original data.

## RESULTS

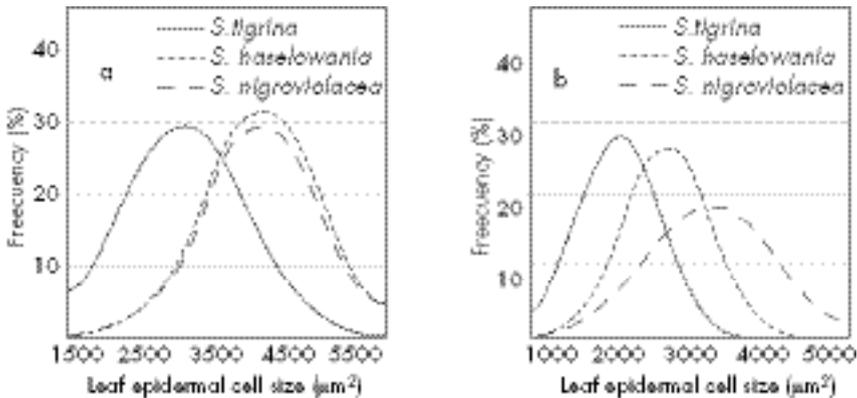
**Taxonomical differentiation.** Sorting of the epidermal cell surface areas in a descending order showed the existence of gaps between most of the analyzed species (Table 1). Epidermal cell surface areas were significantly different ( $p<0.05$ ) among species. The following results from selected examples will describe the process and efficacy of the method for species identification.

Four samples of *S. tigrina* were analyzed showing that none of the specimens differed significantly ( $p= 0.05$ ). It was confirmed that these samples originated in Colombia, Guatemala and Mexico which demonstrates that this species does not vary significantly with latitude.

When the mean epidermal cell surface areas were sorted as in Table 1, the species *S. haselowania* came next to *S. tigrina* in descending order. The adaxial epidermal cell surface areas of *S. tigrina* specimens ranged from  $4208.78\mu^2$  to  $4259.11\mu^2$  (data not shown). These data did not differ ( $p=0.05$ ) with those in C29 *S. haselowania* for the adaxial epidermal cell surface areas (mean:  $4215.97\mu^2$ ) (Fig. 1). However, the abaxial epidermal cell surface areas of *S. tigrina* specimens ranged from  $2816.88\mu^2$  to  $2917.18\mu^2$ . At the same time, mean abaxial cell surface area of *S. haselowania* was greater (mean= $3556.07\mu^2$ ;  $p<0.05$ ) than that in *S. tigrina*.

**Fig. 1.** Normalized curves for leaf epidermal cell surface areas of *S. tigrina*, *S. haselowania* and *S. nigroviolacea*. a) Adaxial. b) Abaxial. Notice that *S. tigrina* cannot be differentiated from *S. haselowania* by their adaxial cell surface areas. However, the 3 species are well distinguished from their abaxial cell surface areas. Notice the change of scale between panels.

**Fig. 1.** Curvas normalizadas para superficies de células epidérmicas foliares de *S. tigrina*, *S. haselowania* y *S. nigroviolacea*. a) Adaxial. b) Abaxial. Note que *S. tigrina* no puede diferenciarse de *S. haselowania* por su superficie de células adaxiales. Sin embargo, las 3 superficies se distinguen bien por la superficie de células abaxiales. Note el cambio de escala entre paneles.



It means that C29 *S. haselowania* differentiated from plants of *S. tigrina*. As in this example, there were many other cases where species differences arose from one side of the leaf only.

Likewise, the leaf-printing method was applied to alike species which are frequently confused, such as *S. tigrina* and *S. nigroviolacea*. Normal standard curves for the adaxial and abaxial epidermal cell surface areas are shown in Figs. 1a and 1b, respectively, for these two species. Mean epidermal cell surface areas were significantly different ( $p= 0.05$ ) between these two species (Table 2).

**Table 2.** Statistical comparisons of C15 *S. nigroviolacea* with samples of *S. tigrina*. In the "Mean" column, values are the average of n=200.

**Tabla 2.** Comparaciones estadísticas de *S. nigroviolacea* C15 con muestras de *S. tigrina*. En la columna "Promedio", los valores son el promedio de n=200.

<i>S. nigroviolacea</i> compared with:	Lower Conf. Level	Mean	Upper Conf. Level	Sig. Diff. p = 0.05
C07 <i>S. tigrina</i>	1662.8	1129.8	596.9	Yes
C59 <i>S. tigrina</i>	1648.8	1115.8	582.9	Yes
C33 <i>S. tigrina</i>	1637.2	1104.3	571.3	Yes
C26 <i>S. tigrina</i>	1624.1	1091.2	558.2	Yes

Similarly, *S. radiosa* has been usually confused with *S. saccata*. However, both species can be distinguished by comparison of their normal standard curves (Fig. 2).

**Prediction test of unknown samples.** Leaf epidermal cell surface areas of six specimens (C03, 26, 33, 53, 77 and 83), received as unknown samples, were analyzed with the leaf printing method. Results obtained with this method were compared with those of already known sample data to make an identification prediction.

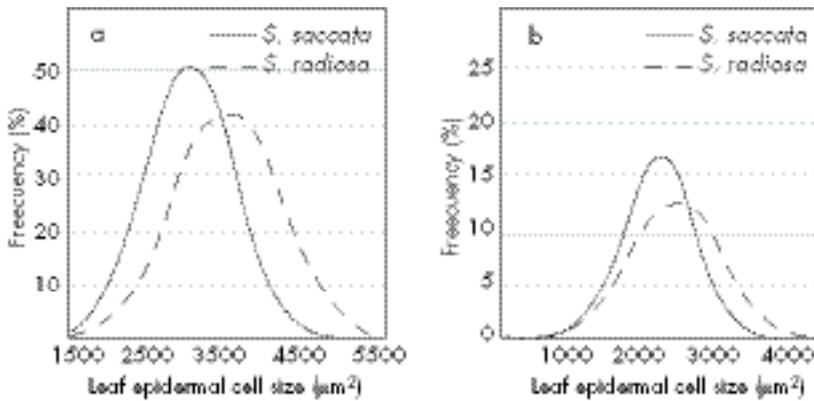
Data from samples C26 and C33 matched data from *S. tigrina*, clones C07 and C59. These results suggest that the unknown samples would correspond to *S. tigrina* specimens. This prediction was later confirmed when plants flowered. Statistical analysis of epidermal cell surface area data showed that neither C07 nor C59 differed significantly ( $p=0.05$ ) from C26 and C33 samples. Since plants of *S. tigrina* were imported from Colombia, Guatemala, and Mexico, these data demonstrate that this species does not vary significantly with latitude.

Data from the adaxial and abaxial leaf epidermis of the unknown sample C53 matched ( $p>0.05$ ) data from the confirmed members C04 and C49, allowing to predict the unknown specimen as *S. hernandezii*. Two years later, such prediction was confirmed after flowering of the specimen.

Unknown sample C77, received from Chiapas, Mexico was predicted to be *S. oculata* based on leaf print data. Afterwards, the identification

**Fig. 2.** Normalized curves for leaf epidermal cell surface areas of *S. saccata* and *S. radiosa*. a) Adaxial. b) Abaxial. Note the change of scale between panels.

**Fig. 2.** Curvas normalizadas para superficies de células epidérmicas foliares en *S. saccata* y *S. radiosa*. a) Adaxial. b) Abaxial. Note el cambio de escala entre paneles.



as such species was confirmed when the specimen flowered. Similarly, data from sample C03 matched data *S. insignis*, which was subsequently confirmed after flowering.

Specimen C83 was received from an unknown origin, and predicted as a member of the Central American group on the basis of its adaxial and abaxial leaf cell surface areas. When C83 flowered, photographs were sent to orchid taxonomists at the Marie Selby Orchid Identification Center (Sarasota, Florida). They determined that it looked like a very yellow form of *S. wardii*, despite this species is unknown in Central America. The standard normal curves of the adaxial and abaxial epidermal cell surface areas differed significantly ( $p=0.05$ ) between *S. amoena* and *S. wardii* (Table 3). The unknown specimen C83 could not then be a sample of *S. wardii*. Instead, it fits well with the species from Costa Rica, *S. amoena*. As a result, specimen C83 was self pollinated to produce seedlings. This would allow to compare vegetative and floral data of the unknown sample with *S. amoena*. Research to determine the origin of specimen C83 is ongoing.

**Phylogenetic aspects.** Abaxial and adaxial epidermal cell surface areas were paired in descending order in Table 1. This table was also divid-

**Table 3.** Statistical comparisons of adaxial and abaxial epidermal cell surface areas of *S. amoena* with samples of *S. wardii*. Data at each line of the column "Mean" come from n=200.

**Tabla 3.** Comparaciones estadísticas de superficies de células epidérmicas adaxiales y abaxiales de *S. amoena* con muestras de *S. wardii*. Los datos en cada línea de la columna "Promedio" son de n=200.

C83 <i>S. cf. amoena</i> compared with:	Lower Conf. Level	Mean	Upper Conf. Level	Sig. Diff. $\alpha = 0.05$
C14 <i>S. wardii</i>	Ad 1575.7 Ab 1247.2	Ad 1042.7 Ab 838.0	Ad 509.8 Ab 428.8	Yes Yes
C62 <i>S. wardii</i>	Ad 1572.8 Ab 1234.0	Ad 1039.8 Ab 824.6	Ad 506.9 Ab 415.6	Yes Yes
C12 <i>S. wardii</i>	Ad 1518.8 Ab 1209.7	Ad 985.8 Ab 800.5	Ad 452.8 Ab 382.7	Yes Yes
C13 <i>S. wardii</i>	Ad 1510.6 Ab 1189.5	Ad 977.7 Ab 780.3	Ad 444.7 Ab 371.1	Yes Yes

ed into 3 groups according to the geographical habitat of each species, as it has been reported coming from South America, Central America or Mexico. The combined adaxial and abaxial epidermal cell surface area data for all samples showed species grouped with alike species. Table 1 also offered a pattern for the entire genus. This pattern was broadly divided into three general groups: species coming from either Central or South America or Mexico.

An evolutionary hierarchy was also observed as species data were displayed from larger to smaller epidermal cell surface areas (Table 1). All primitive species exhibited large abaxial and adaxial epidermal cell surface areas. As mean epidermal cell surface areas decreased in the hierarchy, species were arranged by their flowers in a general order as being either primitive or advanced. The primitive species of the genus (*S. reichenbachiana*, *S. embreei*, *S. platyceras*, *S. grandiflora*, *S. pulla*, *S. pozoi*, and *S. candida*) were all from South America.

The mean leaf epidermal cell surface areas of *S. oculata* placed this species within the Central American group. However, *S. oculata* can be

found in a few scattered locations from northern South America to Southern Mexico, because it is considered a weedy species. *Stanhopea reichenbachiana* was found only in Colombia while at the northern extreme, *S. intermedia* was only found in Mexico. Results presented a clear pattern of the evolutionary expansion of the study genus which correlated with declining both adaxial and abaxial leaf epidermal cell surface areas.

## DISCUSSION

Morphological data are of significance in systematics because morphological variation can make each taxon unique and distinguishable from each other. Morphometrics and systematic biology share a common interest in the analysis of morphology, when assessing the nature of morphological variation. Biological morphologists should then regard on morphometric tools as an integral part of their approaches to systematics (MacLeod, 2002). In this sense, *Stanhopea* species show mean adaxial and abaxial leaf epidermal cell surface areas in narrow ranges, separated by wide-enough gaps as to permit species identification. This makes a substantial contribution to the classical botanical approach of needing a floral description to define a species. While it is an evolutionary *sine qua non* postulate that the primordial flower evolved from a leaf, it is equally obvious that once this divergence took place, natural selection pressures became different for flower and leaves. It means that defining a species by solely analyzing flowers may in fact offer only a sample of organ evolution while distorting the phylogenetic relationships of the species.

The hypothesis that the identification of an unknown species could be statistically determined by comparing its mean leaf epidermal cell surface areas with those of another known species was accepted. It was demonstrated and confirmed by flower analysis in a sufficient number of species. Despite epidermal cell surface areas are not fixed values, they are relatively constant within a narrow range for each species. This allows a correct identification even though some degree of overlapping with closely related species is possible. The identification predictions made on several plant species received as “unknown“ which were later confirmed after flowering, proved the accuracy on the applicability of the proposed method. A good example where this technique was proven to be accurate in species confirmation was in the case of *S. tigrina* and *S. nigroviolacea*. These two species have been confused by botanical and horticultural authorities during a long time (Williams, 1894; Wilson, 1921; Arnold, 1928; Williams, 1951; Sander et al., 1996). All mentioned *S. nigroviolacea* as a variety of *S. tigrina*,

despite Beer's (1854) and Kennedy's (1977) valid publications. *Stanhopea nigroviolacea* is a purely Mexican species, while data from the adaxial and abaxial epidermal leaf cell surface areas inferred *S. tigrina* to be, although pandemic, from South American origin.

Another case is *S. radiosa* vs. *S. saccata*, both species known as coming from Mexico. These species have been confused by taxonomists during decades (Dodson, 1963; Kennedy, 1975). Descriptions were made based on individual flowers, without regard to the whole plant as the species unit which was subjected to a genus variation over a geographical range.

In this investigation, the statistical confidence limit was set at  $p = 0.05$ . However, depending on either the material being measured or the level of measurement accuracy, a worker might prefer to select a more rigorous test. This statistically-based system using leaf prints is simple, ecologically friendly, and reproducible. Storage of equipment and collected material require a small box for field sampling; small cases for holding microscope glass slides; a place for storing photographs; space for computer equipment, and a microscope. However, best results may be possible by combining the microscope with digital recording and image analysis linked to the computer spreadsheet. With this technological improvement, a larger cell number might be scanned with greater accuracy, results could be offered immediately, and human errors would be minimized. Usefulness of this technique depends partially on the number of samples available for each individual species and the species number sampled throughout the entire range of the genus. This is in order to obtain more accurate results.

The amount of shrinkage observed in *Stanhopea* leaves is under investigation by comparing data from fresh leaves with similar data from dry leaves. Primitive, large-cell plants appear to show a greater percentage of shrinkage than the smaller-cell, advanced members. Cell shrinkage appears to progress at a relatively uniform rate from primitive to advanced species. Furthermore, mean epidermal cell surface areas vary evolutionarily with speciation of *Stanhopea* plants with latitude. This has offered new information about this plant genus. If indeed, this applies to other plant genera and families, new knowledge may be gained about the evolution of plants.

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