HONEY BEE RACES CLASSIFICATION METHOD BY BODY CHARACTERS

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Abstract

A method of honey bee races classification by characters in bee-worker forewings was developed. 15 – 20 wings from a colony are scanned as bitmap pictures. The co-ordinates of 19 defined points in wing veins are measured using the mouse pointer, and 30 characters (lengths, angles, indices, fields area) are computed. The combination of values of characters is typical for definite race. The classified sample is compared with known values of single races. The Mahalanobis distances of the sample to single races are computed. The final result of the sample classification are posterior probabilities with which the sample belongs to single races. Classification results of some bee samples are given.

Introduction

Maintaining the biodiversity within the species honey bee assumes to have adequately quick and accurate method of honey bee classification. Complex morphological analysis of bee exterior is labour intensive. The more simple method is measuring the bee forewing characters, which are typical for each pure race (KAUHAUSEN-KELLER,D., KELLER,R., 1994).

The bee samples are usually evaluated using the discriminant analysis and are presented graphically for the whole populations - races. But identification and classification of a sample is difficult. At the same time, expressing the multidimensional relations among single populations in a two-dimensional diagram constricts the informational ability of the diagram.

Using the principles of discriminant analysis, we derived the method of single bee sample classification in numerical form. A sample is classified by probability of belonging to single races, which are characterized with race standards prepared in advance.

Material and methods

Measurement of bee wing characters is easy. Forewings are fixed on a transparent foil. Wings are scanned at the resolution at least 1200 dpi and moved into computer as a bitmap picture. The wing design is cut in a program on the screen with a mouse pointer so that x,y co-ordinates of given points in given order are saved. 30 characters are then computed on the basis of co-ordinates of single points for each wing. These characters determine the design of the whole wing. The wing characters are lengths between some points, angles, indices and area of six wing fields. 15 – 20 wings in one sample are measured. A mean is computed for each character as a parameter of the sample.

The sample classification is made by comparing with the typical values of single pure races – standards prepared beforehand. Bee samples for these standards were collected from various beekeepers in Europe, Asia and Africa. Race standards were repeatedly statistically checked by excluding the samples which diverged from other typical samples of the race.

The similarity of the tested sample with values of standards of single races is expressed by Mahalanobis distances. These are distances of the sample from centroids of clusters of single races. Mahalanobis distances are calculated by the formula (LUKASOVÁ,A., ŠARMANOVÁ,A., 1985):

$$\text{MD}_{a,b} = (a - b)^T \times K^{-1} \times (a - b)$$

where:

\(\text{MD}_{a,b}\) … Mahalanobis distance between points \(a, b\) in multidimensional space
\((a-b)^T\) … line vector \((a_1-b_1, a_2-b_2, \ldots, a_n-b_n)\) transposed
\( K^{-1} \) … inverse matrix to covariance matrix of standardized data
\((a-b)\) … column vector \((a_1-b_1, a_2-b_2, \ldots, a_n-b_n)\)

A sample belongs with the highest probability to that race which is most close to, i.e. has the lowest Mahalanobis distance to it. The probability is computed for each race whose standard is known. The sum of all probabilities for a sample is 100. These are so called posterior probabilities. They are computed from the values of Mahalanobis distances.

Posterior probability (PP) that a sample \( t \) belongs to population \( i \) is

\[
PP (t) = \frac{1/n * \exp (- MD (i)^2 / 2)}{\sum 1/n * \exp (- MD (i)^2 / 2)}
\]

where:

- \( t \) … tested sample
- \( i \) … single populations in the set of standards
- \( n \) … number of populations in the set of standards
- \( MD(i)^2 \) … the 2\(^{nd}\) power of Mahalanobis distance of the sample \( t \) from the population \( i \)
- \( \exp() \) … the function calculating the value of \( e^x \), where \( x \) is given number, \( e \) is a base of natural logarithms - around 2,71828

Results and discussion

The statistical processing of single wings and classification of the whole sample runs in database application of FoxPro named BEEMORPH. An example of resulted values of Mahalanobis distances, classifications to races and corresponding posterior probabilities of a few samples are in the Table 1. Samples were compared with nine race standards. The values of the first five races by the order of posterior probabilities are given in the Table 1. Samples C0413 and C0477 were classified to different races than were declared.

In case of the value of posterior probabilities for a race in the first place is lower than 50 %, the sample is not reliably compared to any race. At the same time, samples in first two or three places with similar values of posterior probabilities may be considered as samples with the current impact of a few races (samples C0034, C0413).

Only those samples were left as typical which were classified identically with declared race and with the value of posterior probability at the first place over 70 %, during creating the race standards. This criterion match samples C0022, C0087, C0274, C0464 in Table 1.

The typical values of 30 characters of four races (standards) in the relation to the mean of nine races are graphically presented in Fig. 1. The combination of characters is specific for each race. These combinations may be considered to be race standards. Correspondingly, classification of the single sample using the above described multivariate method is made by comparing it to standards and expresses the similarity of the sample with the race standards.

The described method of morphometric analysis and taxonomic classification of honey bees named DAWINO-bee works in checking the race purity in Carniolan bee breeding apiaries in Czech Republic since 1999. The computer program is able to print a protocol with detailed results of a sample. An example of the main part of the protocol – result of the sample classification is in Tab. 2.

References


### Tab. 1  Classification results of some samples to first five of nine race standards

<table>
<thead>
<tr>
<th>Sample No.</th>
<th>Declared race</th>
<th>MD</th>
<th>1st place</th>
<th>2nd place</th>
<th>3rd place</th>
<th>4th place</th>
<th>5th place</th>
<th><strong>classified race</strong></th>
<th><strong>PP (%)</strong></th>
</tr>
</thead>
<tbody>
<tr>
<td>C0022</td>
<td>carnica</td>
<td>4,890968</td>
<td>5,574902</td>
<td>5,603711</td>
<td>5,728217</td>
<td>5,997851</td>
<td></td>
<td>carnica macedonica caucasica mellifera ligustica</td>
<td>93,76</td>
</tr>
<tr>
<td>C0034</td>
<td>carnica</td>
<td>7,248234</td>
<td>7,301153</td>
<td>7,330718</td>
<td>7,547787</td>
<td>7,618142</td>
<td></td>
<td>carnica buckfast mellifera caucasica macedonica</td>
<td>40,61</td>
</tr>
<tr>
<td>C0087</td>
<td>buckfast</td>
<td>5,242633</td>
<td>5,744137</td>
<td>5,824023</td>
<td>5,927215</td>
<td>5,968088</td>
<td></td>
<td>buckfast ligustica carnica mellifera macedonica</td>
<td>86,58</td>
</tr>
<tr>
<td>C0274</td>
<td>iberica</td>
<td>4,901776</td>
<td>6,485338</td>
<td>6,620067</td>
<td>6,664406</td>
<td>6,702301</td>
<td></td>
<td>iberica carnica buckfast mellifera macedonica</td>
<td>99,97</td>
</tr>
<tr>
<td>C0413</td>
<td>mellifera</td>
<td>5,206643</td>
<td>5,279640</td>
<td>5,379239</td>
<td>5,481054</td>
<td>5,629261</td>
<td></td>
<td>carnica mellifera ligustica caucasica macedonica</td>
<td>39,74</td>
</tr>
<tr>
<td>C0464</td>
<td>ligustica</td>
<td>4,260699</td>
<td>4,712279</td>
<td>5,034632</td>
<td>5,068578</td>
<td>5,093799</td>
<td></td>
<td>ligustica carnica caucasica mellifera macedonica</td>
<td>82,79</td>
</tr>
<tr>
<td>C0477</td>
<td>cecropia</td>
<td>7,072516</td>
<td>7,274367</td>
<td>7,328876</td>
<td>7,648411</td>
<td>7,675427</td>
<td></td>
<td>macedonica cecropia carnica mellifera ligustica</td>
<td>70,18</td>
</tr>
</tbody>
</table>

MD … Mahalanobis distance  
PP … posterior probability

### Tab. 2  Result of taxonomic classification of a honey bee sample by wing characters

<table>
<thead>
<tr>
<th>RACE</th>
<th>PP %</th>
</tr>
</thead>
<tbody>
<tr>
<td>CARNICA</td>
<td>92</td>
</tr>
<tr>
<td>MACEDONICA</td>
<td>4</td>
</tr>
<tr>
<td>MELLIFERA</td>
<td>2</td>
</tr>
<tr>
<td>LIGUSTICA</td>
<td>1</td>
</tr>
<tr>
<td>Other</td>
<td>1</td>
</tr>
</tbody>
</table>

PP … posterior probability
Fig. 1   Combination of 30 characters of four bee races expressed by deviations from a mean of nine races