Revised phylogenetic analysis of Indian species of genus *Macrophya* Dahlbom (Hymenoptera: Symphyta; Tenthredinidae: Tenthredininae)

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Abstract

Revised phylogenetic analysis was performed for 14 species of the genus *Macrophya* (Hymenoptera: Tenthredinidae) using the phylogenetic analysis package PAUP, based on 15 of the morphological characters most commonly used for *Macrophya* species identification. Species descriptions were derived primarily from “Indian Sawflies Biodiversity” vol. II Saini (2007), Saini and Bharti (1996), Saini et al. (1996), and Saini and Vasu (1997). The results are discussed in terms of evolutionary trends or biological maxim that “nature prefer to modify the already existing structures so as to cope with new needs.”

Keywords: Phylogenetic analysis, Macrophya, Evolutionary trends.

Introduction

The genus *Macrophya* (Hymenoptera: Tenthredinidae) is widely distributed genus with its representatives available in almost all main regions of the globe. With regard to its affinities, it shares most of its characters with *Pachyprotasis* Hartig. Even within *Macrophya* the range of characters is so wide that time to time many of its subgenera were proposed (Malaise, 1945) and because of no distinct boundaries they all got merged (Ross, 1937; Gibson, 1980). Today none of its subgenus is considered to be valid (Abe and Smith, 1991). The genus *Macrophya* was first described by Dahlbom (1835) as a subgenus of *Tenthredo* Linnaeus, on the basis of body shape, length and form of antenna. He divided this subgenus into two subsections “A” and “B”. Hartig (1837) applied names to these two subsections using *T. (Macrophya)* for subsection “B” and *T. (M.) (Pachyprotasis)* for subsection “A”. Both of these were later recognised as valid genera by Westwood (1840).

The genus is characterized by venation as in *Pachyprotasis*, but the anal cell may have a cross vein. Malar space mostly shorter than the diameter of an ocellus. The hind legs are strongly built, and the knees reaching and mostly exceeding the apex of the abdomen (Saini, 2007). The larval stages feed on variety of wild herbs, shrubs and trees. Generally adults feed on pollens, flower nectar or leaf juice exuding from wounds caused by strong mandibles. However, many robust species indulge in zoophagy (Cameron, 1882; Rohwer, 1913; Benson, 1938; Malaise, 1945; Naito, 1988 and Goulet, 1996).

The purpose of present study is to trace the long evolutionary history which modified generalizations into specializations of extreme form to suit the circumstances in which subsequently insects dwelled. Parsimony analysis is used to investigate phylogenetic relationships among *Macrophya* species, using data based on morphological characters most commonly used for *Macrophya* identification. With the help of literature contributed by Saini et al. (1996) and Saini and Vasu (1997) some different characters were used to trace phylogeny of genus *Macrophya*.

Materials and Methods

Species descriptions were derived primarily from “Indian Sawflies Biodiversity” vol. II Saini (2007), Saini and Bharti (1996), Saini et al. (1996), and Saini and Vasu (1997) and the characters used in the analysis were those given comparably for all, or almost all, species. *Tenthredo rufipodus* Saini and Bharti, 1996 was also included in the analysis as the outgroup. Revised phylogenetic analysis was carried with a computer program PAUP version 4.0, with more characters as compared to earlier study (Saini and Kaur, 2010), has been used to infer the phylogeny (Swofford, 2000). In total 15 morphological characters were used in the phylogenetic analysis. These were:

1. Forewing: (0-without infuscated band, 1-with infuscated band)
2. Body colour: (0-not metallic blue, 1-metallic blue)
3. **Clypeus**: (0-not triangularly incised up to 1/4 of its length, 1- triangularly incised up to 1/4 of its length)

4. **Frontal area**: (0-not below level of eyes, 1-below level of eyes)

5. **Average length of body of female**: (0-10 mm, 1-above 10mm, 2-below 10mm)

6. **Subapical teeth of tarsal claw**: (0-smaller than apical teeth, 1-longer or equal to apical teeth)

7. **Labrum**: (0-with rounded anterior margin, 1-with truncate or subtruncate anterior margin, 2-with slightly emarginate anterior margin)

8. **Median fovea**: (0-absent, 1-distinct, 2-indistinct)

9. **Supraantennal tubercles**: (0-insignificant, indistinct or not raised, 1-raised or significant)

10. **Circumcellar furrow**: (0-fine or clear, 1-distinct, 2-indistinct, 3-sharp)

11. **Postocellar area**: (0-flat, 1-subconvex, 2-slightly raised, 3-fine)

12. **Mesoscutellum**: (0-slightly or roundly raised or elevated, 1-subconvex, 2-prismatic, 3-flat)

13. **Mesepisternum**: (0-roundly raised, 1-obtusely raised, 2-depressed not obtusely raised)

14. **Antenna length**: (0-2 times or more than 2 times of head width;1-less than 2 times of head width)

15. **Hind femora**: (0-hardly reaching apex of abdomen, 1-exceeding apex of abdomen)

Table 1: Presence or absence data for fifteen characters for 14 species of the genus *Macrophya* as used in the phylogenetic analysis; *T. rufipodus* Saini and Bharti, 1996 is included as an outgroup.

<table>
<thead>
<tr>
<th>Character</th>
<th>Species</th>
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<td>3. Clypeus</td>
<td>M. formosana Rohwer, 1916</td>
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<td>M. gopeshwari Saini et al., 1986</td>
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<td>0</td>
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<td>12. Clypeus</td>
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<td>14. Clypeus</td>
<td>M. verticalis Konow, 1998</td>
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Discussion

The outgroup taxon *T. rufipodus* got separated from all species of genus *Macrophya* on the basis of postocellar area fine (Character 11, state 3), mesepisternum depressed not obtusely raised (Character 13, state 2) and hind femora hardly reaching apex of abdomen (Character 15, state 0). The derived consensus tree identified two groupings to be present in the cladogram. In the first group *M. khasiana* got isolated from *M. verticalis, M. brancuccii, M. regia, M. gopeshwari, M. planata, M. maculicornis* and

Results

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Fig. 1. Strict consensus tree for 14 species of *Macrophya* derived from the 7 most parsimonious trees calculated from the data in Table 1; outgroup = *Tenthredo rufipodus* Saini and Bharti, 1996. Character of the ingroup have been optimized by fast transformation as implemented in PAUP. Character numbers are above the hashmarks; state changes are shown below with the respective primitive and derived conditions reported by a ‘>’. 7 trees were found using the computer program PAUP version 4.0, with the shortest tree requiring 28 steps, a consistency index (CI) of 0.50, a retention index (RI) of 0.55 and rescaled consistency index (RC) of 0.28.
M. pseudoplanata on the basis of average length of body of female is 10mm (Character 5, state 0). M. verticalis, M. brancuccii and M. regia due to absence of median fovea got separated from M. gopeshwari, M. planata, M. maculicornis and M. pseudoplanata. M. brancuccii and M. regia are sister species and got isolated from M. gopeshwari, M. planata, M. maculicornis and M. pseudoplanata due to average length of body below 10mm (Character 5, state 2) and labrum with subtruncate anterior margin (Character 7, state 1). M. gopeshwari can be distinguished from M. planata, M. maculicornis and M. pseudoplanata due to characters 6 and 9.

M. brancuccii and M. regia are sister species and got isolated from M. verticalis due to characters 6 and 9. M. gopeshwari can be distinguished from M. planata, M. maculicornis and M. pseudoplanata due to average length of body below 10mm (Character 5, state 2) and labrum with subtruncate anterior margin (Character 7, state 1).

M. planata on basis of characters 8, 10 and 12 got isolated from sister species M. maculicornis and M. pseudoplanata. Due to triangularly incision in clypeus, labrum with slightly emarginated anterior margin and mesepisternum obtusely raised M. manganensis got separated from M. formosana, M. pomplina, M. naga and M. rufipodus and all these species showed monophyly as indicated by homologue 6. M. naga and M. rufipodus at the base of cladogram got separated from closely related species M. pomplina due to characters 2, 4, 5, 7 and 10. M. naga and M. rufipodus on basis of characters 1, 9 and 11 got separated from each other. So, extremely specialized forms descended by gradual changes leads to accumulation of certain appropriate features which represents body organization acquired to become complex so as to meet requirements which also underlies the biological maxim.

References


