First Record of Euplatypus parallelus (Coleoptera: Curculionidae) in China

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First record of *Euplatypus parallelus* (Coleoptera: Curculionidae) in China

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*Euplatypus parallelus* (F.) (Coleoptera: Curculionidae: Platypodinae) is one of the most common ambrosia beetles in Central and South America (Silva et al. 2013). It is polyphagous and is reported to infest over 82 plant tree species from 25 families (Gümüş & Ergün 2015). Although this species is native to the neotropics, it is one of the most invasive of all the Platypodinae. In the late 1800s, *E. parallelus* was introduced into Africa where it is now fairly widespread, while it is currently spreading rapidly in southeastern Asia (Beaver 1999; Boa & Kirkendall 2004; Gümüş & Ergün 2015).

Most platypodine species infest only freshly dead or dying trees. However, *E. parallelus* is one of the few species that can successfully colonize live trees, although it is typically associated with trees stressed by drought, disease, or flooding (Boa & Kirkendall 2004). *Euplatypus parallelus* has been reported to attack live rubber trees (*Hevea brasiliensis* [Malpighiales]) in Brazil (Silva et al. 2013), implicated in a major fungus associated with wilting of trees (*Pterocarpus indicus*) (Sanderson et al. 1997; Boa & Kirkendall 2004), and repeatedly reported as a suspect of transporting fungal pathogens of trees (Sanderson et al. 1997; Boa & Kirkendall 2004; Bumrungsri et al. 2008; Tarno et al. 2016). For example, in Indonesia, Malaysia, Singapore, and Thailand, *E. parallelus* has been associated with *Fusarium*, a causative agent of wilting of trees (*Pterocarpus indicus* (Sanderson 1997; Bumrungsri et al. 2008; Tarno et al. 2016).

Even though *E. parallelus* is spreading rapidly in Asia, it has not yet been reported from China where it is considered a regulated pest (General Administration of Quality Supervision, Inspection, and Quarantine of the People’s Republic of China 2017). During Oct 2016, many representatives of platypodine ambrosia beetles were captured in light traps (UV blacklight) and hand-collected from dead rubber trees in Danzhou, Hainan (19.4375°N, 109.5585°W; 143 masl). The first specimens were preserved in 100% ethanol and sent to the University of Florida (UF), School of Forest Resources and Conservation, Forest Entomology Laboratory, where they were confirmed as *E. parallelus* using the taxonomic keys by Atkinson (1989) and Wood (1993). Specimens were deposited in the UF Forest Entomology Laboratory Collection. After the initial collection, 2 additional surveys were carried out in Dec 2016 and from May to Jun 2017. During the collection in 2017, kairomone-baited traps also were used that contained alpha-pinene, beta-pinene, verbenone, and ethanol (Wako Pure Chemical Industries, Japan). In total, 246 specimens of *E. parallelus* were collected from 10 locations in Hainan, China (Table 1). Several individuals of other platypodine species (*Dinoplatypus cavus* and *Crossopterus externedentatus* [Coleoptera: Curculionidae]) also were collected from dying trees during the surveys.

A subset of the Hainan samples was submitted to the Jiangxi Agriculture University for molecular identification. Genomic DNA was extracted from 2 individuals from 2 locations in Hainan (Xinglong and Bawangling; Table 1). Portions of the cytochrome oxidase subunit 1 (COI) and the nuclear ribosomal 28S regions of *E. parallelus* were amplified using oligonucleotide primers S1718 and A2237 as well as S3690 and ethanol (Wako Pure Chemical Industries, Japan). In total, 246 specimens of *E. parallelus* were collected from 10 locations in Hainan, China (Table 1). Several individuals of other platypodine species (*Dinoplatypus cavus* and *Crossopterus externedentatus* [Coleoptera: Curculionidae]) also were collected from dying trees during the surveys.

### Table 1. Collection of *Euplatypus parallelus* specimens in Hainan, China, from 2016 to 2017.

<table>
<thead>
<tr>
<th>Location</th>
<th>Host</th>
<th>Source</th>
<th>Date</th>
<th>Number of specimens</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bawangling, Chantian*</td>
<td>unknown</td>
<td>wood</td>
<td>2016.XII</td>
<td>3</td>
</tr>
<tr>
<td>Changhao, Wuzhishan</td>
<td>—</td>
<td>Lure trap**</td>
<td>2017.VI</td>
<td>91</td>
</tr>
<tr>
<td>Darong, Danzhou</td>
<td><em>Hevea brasiliensis</em></td>
<td>wood</td>
<td>2016.X</td>
<td>12</td>
</tr>
<tr>
<td>Fengmu, Tunchang</td>
<td>—</td>
<td>light trap</td>
<td>2016.X</td>
<td>4</td>
</tr>
<tr>
<td>Fengmu, Tunchang</td>
<td><em>Acacia mangium</em></td>
<td>wood</td>
<td>2016.X</td>
<td>6</td>
</tr>
<tr>
<td>Fushan, Chengmai</td>
<td>—</td>
<td>light trap</td>
<td>2017.V</td>
<td>18</td>
</tr>
<tr>
<td>Jianfengling, Ledong</td>
<td>unknown</td>
<td>wood</td>
<td>2016.XII</td>
<td>7</td>
</tr>
<tr>
<td>Limushan, Qionghzhong</td>
<td>—</td>
<td>light trap</td>
<td>2016.X</td>
<td>22</td>
</tr>
<tr>
<td>Nanfeng, Dazhout</td>
<td>—</td>
<td>light trap</td>
<td>2016.X</td>
<td>17</td>
</tr>
<tr>
<td>Xinglong, Wanning*</td>
<td><em>Hevea brasiliensis</em></td>
<td>wood</td>
<td>2016.XII</td>
<td>20</td>
</tr>
<tr>
<td>Xinzhong, Wanning</td>
<td>—</td>
<td>Lure trap**</td>
<td>2017.VI</td>
<td>46</td>
</tr>
</tbody>
</table>

*Samples were sequenced; **The lure included alpha-pinene, beta-pinene, verbenone and EIOH.

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and A4285, respectively (Jordal et al. 2011). The National Center for Biotechnology Information’s Basic Local Alignment Search Tool (BLAST) was used to identify the nucleotide sequences. Cytochrome oxidase subunit I and rDNA sequences (2016113001, 20161205003) were 98% similar to the E. parallelus #Pieup03 and #Pieup04 (GenBank accessions KR261327 and KR261328, respectively). Both 28S rDNA sequences were 99% similar to the E. parallelus #Pieup03 and #Pieup04 (GenBank accessions KR261230 and KR261231, respectively). Previously, E. parallelus #Pieup03 and #Pieup04 have been reported by Jordal (2015) from two populations from Brazil and Cameroon, respectively.

Collection data showed that this beetle has successfully colonized Hainan Island and is now widely distributed there (Fig. 1). Specimens were collected at 10 locations on the island, including natural old-growth forest as well as planted forests, mostly rubber tree plantations. During the last large-scale survey of forest pests in Hainan (15 years ago), E. parallelus was not recorded (Yin et al. 2002), therefore we assume that the species was introduced to the area between 2003 and 2015. It is important to note that our sample data were restricted to Hainan Island. Therefore, we do not know whether this beetle has been introduced into mainland China. Currently, specimens we collected from rubber trees were already dying from other causes. Euplatypus parallelus does not appear to act as a destructive forest pest in Hainan.

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**Summary**

*Euplatypus parallelus* (Coleoptera: Curculionidae) is a neotropical ambrosia beetle that is rapidly spreading around the world. It has been recorded from over 80 host trees and is implicated as a primary pest attacking rubber trees and rosewood. Here, we report the first country record and successful establishment of the species in Hainan, China. Currently, *E. parallelus* does not appear to act as a destructive forest pest in Hainan.

Key Words: wood borer; island; forest pest; invasive species; COI sequence

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**Fig 1.** The collections of *Euplatypus parallelus* (Coleoptera: Curculionidae) in Hainan, China, reported in this paper.
**Sumario**

_Euplatypus parallelus_ (Coleoptera: Curculionidae) es un escarabajo de ambrosia Neotropical que se está esparciendo rápidamente a escala global. Se ha reportado para más de 80 árboles hospederos, siendo una plaga primaria para árboles de caucho y madera rosa. Reportamos aquí el primer registro y establecimiento exitoso en Hainan, China. Actualmente, _E. parallelus_ no parece ser una plaga forestal destructiva en Hainan.

Palabras Clave: perforador de madera; isla; plaga forestal; invasores; secuencia COI

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