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***Vitessa ingae* sp. n. (Lepidoptera, Pyralidae)  
from Sulawesi (Indonesia)**

**Ulf BUCHSBAUM & Mei-Yu CHEN**

### **Summary**

A new species *Vitessa ingae* sp. n. from Sulawesi is described and compared with similar species. It is clearly different from all other *Vitessa*-species with its size and markings and shows clear differences to all the known species in this genus; also the DNA Barcode is distinctive.

### **Zusammenfassung**

Die neue Art *Vitessa ingae* sp. n. aus Sulawesi wird beschrieben und mit ähnlichen Arten verglichen. Sie unterscheidet sich deutlich von allen anderen *Vitessa*-Arten durch ihre Größe und unterschiedliche Zeichnung. Ebenso zeigt der DNA Barcode deutliche Unterschiede zu den bisher bekannten Arten dieser Gattung.

Key words: Lepidoptera, Pyralidae, *Vitessa ingae* sp. n., Indonesia, Sulawesi, DNA Barcode, COI-5P

## Introduction

Only three species and a few subspecies of the genus *Vitessa* MOORE, [1860] are recorded from Sulawesi (*Vitessa hollandi wallacealis* MUNROE & SHAFFER, 1980, *V. pyraliata latialbata* MUNROE & SHAFFER, 1980, *V. pyraliata pyraliata* Walker, 1864 and *V. kolakalis* MUNROE & SHAFFER, 1980). This genus only occurs in Southeast Asia from Southeast China, Indo-China region (Thailand, Burma, Laos, Vietnam & Cambodia) via Malaysia, Indonesian Islands, New Guinea to Australia with 29 species (MUNROE & SHAFFER 1980, BUCHSBAUM 2000, 2002). Most of the species are rare. The specimens fly in the night and come to the light.

Sulawesi is in the zoogeographic interregion of Wallacea (ILLIES 1971, REICHHOLF 2003, SEDLAG 1984, 1995, WALLACE 1869, WHITMORE 1987, WHITTEN et al. 2002). The region is known for a high biodiversity and is one of the biodiversity hotspots (MITTERMEIER et al. 1998, MYERS et al. 2000). Consequently many new species from this region could be found and described (e. g. BUCHSBAUM 2011, DUBATOLOV & HOLLOWAY 2006, FIBIGER 2007, 2010, HACKER et al. 2008, SPEIDEL et al. 2004).

### *Vitessa ingae* sp. n. (Fig. 1)

Material: Holotype: ♂; S. Sulawesi (Celebes), Puncak, Palobo, 2.55° S 120.05° E. L., I 1997, leg. local collector, in Coll. Zoologische Staatssammlung Munich.

### Description and differential diagnosis

Wingspan: 69 mm, Forewing length: 34 mm.

Body and wings black. Forewings with three white veins in the external third in dorsal part of the wings. A single middle spot more or less round. An elongate (yellowish?) white bar is situated in postbasal region at ventral margin and a small white dot at costa. Hindwings with a white dot in discal cell near costa. Wings with dark blue shining. Antenna black and long, reaching to the white middle reniform stigma.

Head yellow. Labial palpus yellow with black first segment. Thorax with bright yellow dot at caudal margin. Abdomen black with yellow hairs at the last segments. Legs black.

Female genitalia (fig. 2): Similar to *V. pyraliata*, but with a clear marked signum. Apophyses slim and long. Ductus bursae slim long, less sclerotized.

The next similar species is *V. pyraliata* which looks totally different. Forewings with two rounded dots in postbasal region at dorsum and costa. Hindwings large, white in the discal region. All the specimens of the latter species are about one third smaller than *V. ingae*.

**Etymology:** The species is named after the mother and mother in law of the authors dedicated to her 70<sup>th</sup> birthday in order to express our thank for her always great support of our entomological work.

## Discussion

This *Vitessa* species is the largest species ever collected in the genus. Characterized by the size and the dark colouration it is a very conspicuous species of this genus.

The DNA also shows a larger distance to the similar species like *V. pyraliata pyraliata* and *V. pyraliata latipalpa* and to any further species inhabiting the adjacent islands, e. g. Sumatra and Borneo.

### Cytochrome oxidase submit I (COI-5P).

*Vitessa ingae* sp. n.

Barcode ID: GWOSM372-11, Sample ID: BC ZSM Lep 43458, GenBank Accession: -; Nucleotide Sequence: Residues: 658; Comp. A: 183; Comp. G: 107; Comp. C: 109; Comp. T: 259; Ambiguous: 0:

CACATTATATTATTTGGTATTGATCTGGTATAAGTGGGAACCTCTTAAG  
ATTACTAATTGAGCAGAATTGGGAACCCCTGAATCTTAATTGGAGATGAT  
CAAATTATAATACTATCGTTACTGGTCACGCCATTATTATAATTTTTTATA  
GTTATACCTATTATAATTGGGGGTTGGAAATTGGTAGTACCTTAATATT  
AGGGGCCCTGATATAGCTTCCCCGAATAATAAGATTTGATTAT  
TACCCCCATCTATTACCCCTTTAATTCAAGAAGAATTGTTGAAAATGGTGC  
GGGACAGGTGAACAGTTATCCCCCCTACTCTAATATTGCCATAGAGG  
AAGATCTGTAGACTTAGCAATTTCCTTACACTAGCGGTATTCATCTA  
TTTTAGGAGCAATTAAATTATTACCACTATTATAATATAAAATTAAATGGG  
TTACATTGATCAAATACCTTATTGTTGAGCTGTGGGTATTACAGCCCTT  
TTATTACTTTATCCTTACCGGTTAGCTGGGCTATTACAATACTCTAAC  
GATCGAAATTAAATACCTCCTTTGACCCTGCTGGAGGGGGATCCTAT  
TCTTACCAACATTATT

*Vitessa pyraliata pyraliata* WALKER, 1864

Barcode ID: GWORM540-09, Sample ID: BC ZSM Lep 20790, GenBank Accession: GU687340; Residues: 658; Comp. A: 188; Comp. G: 105; Comp. C: 108, Com. T: 257; Ambiguous: 0:

CACATTATATTATCTTGGATTGATCAGGAATAGTAGGAACCTCTTAA  
GATTGCTAATTGAGCAGAATTGGGAACCCCTGAATCTTAATTGGAGATGA  
TCAAATTATAATACTATTGTTACAGGTACGCCATTATTATAATTTTTTAT  
AGTTATACCAATTATAATTGGAGGGTTGGAAATTGATTAGTTCTTAAATAC  
TAGGAGCCCTGATATGGCATTCCCTCGAATAATAAGATTTGATT  
ATTACCCCCATCTATTACTCTTTAATTCAAGAAGAATTGTTGAAAATGGTG  
CTGGAACAGGGTGGACAGTTACCCCTCACTATCTCTAATATTGCCATAGA  
GGTAGATCTGTAGATTAGCAATTTCCTACATTAGCTGGTATTCATCT  
ATTTAGGTGCAATTAAATTATTACCACTATTATAATATAAAATTAAATGG  
ATTACATTGACCAAATACCTTATTGTTGAGCTGTGGGTATCACAGCTC  
TCTTATTACTTTATCCTTACCGGTTAGCCGGAGCTATTACTATGCTCTAA  
CAGATCGAAATTAAATACGTCTTTTGTGACCCTGCTGGGGTGGAGACCCT  
ATTCTTATCAACATTATT



1



2



3

*Vitessa ingae* sp. n. Fig. 1. Holotype. Fig. 2. Female genitalia. Fig. 3.: map showing type locality of *V. ingae* sp. n..

*Vitessa pyraliata latialbata* MUNROE & SHAFFER, 1980

Barcode ID: GWORM534-09, Sample ID: BC ZSM Lep 20784, GenBank Accession: HM393698; Residues: 627; Comp. A: 179; Comp. G: 101; Comp. C: 105, Com. T: 242; Ambiguous: 0:

CACATTATATTATCTTGGGATTGATCAGGAATAGTAGGAACCTTTAA  
GATTGCTAATTGAGCAGAGTTGGAACCCCTGAATCTTAATTGGAGATGA  
TCAAATTATAACTATTGTTACAGGTACGCCATTATAATTTTTTAT  
AGTTATACCAATTATAATTGGAGGTTGGAAATTGATTAGTCCTTAATAC  
TAGGGGCCCCCTGATATAGCATCCCCCGAATAATAATAAGATTGATT  
ATTGCCCATCTATTACTCTTTAATTCAAGAAGATTGTTGAAAATGGTG  
CTGGGACAGGGTGGACAGTTACCCCTCCCTATCTTAATATTGCCCATAGA  
GGTAGATCTGTAGATTAGCAATTTCCTACATTAGCTGGTATTCTCATCT  
ATTTAGGCAGCAATTAAATTATTACCACTATTATAATAAAAATTAAATGG  
ATTACATTTGACCAAATACCTTATTGTTGAGCTGTGGGTATCACAGCTC  
TCTTATTACTTTATCCTACCAGGTTTAGCCGGAGCTATTACCATACTCTAA  
CAGATCGAAATTAAATATCTTTTTGACCTGCTGGAGG

*Vitessa gritschroeerae* BUCHSBAUM, 2000

Barcode ID: GWORM545-09, Sample ID: BC ZSM Lep 20795, GenBank Accession: HM393702; Residues: 658; Comp. A: 183; Comp. G: 104; Comp. C: 106, Com. T: 265; Ambiguous: 0:

TACATTATATTATTTGGAAATTGGTCAGGAATAGTAGGAACCTTTAA  
GATTATAATTGAGCAGAAATTAGGAACCTCCGAATCTTAATTGGAGATGA  
TCAAATTATAACACTATTGTTACCGGTATGCTTCATTATAATTTTTTAT  
AGTTATACCAATTATAATTGGGGGATTGGAAATTGATTAGTACCTTTATAT  
TAGGGGCTCTGATATAGCTTCCTCGAATAATAAGATTGGTTA  
CTCCCCCATCTATTACTCTTTAATTCAAGAAGAATTGTTGAAAATGGTGC  
CGGAACGGGTGAACAGTTACCCCTCTATCTCCAATATTGCTCATAGTG  
GTAGATCTGTAGATCTAGCAATTTCCTACATTAGCTGGTATCTCTCTA  
TTTAGGGGCAATTAAATTCAATTACCACTATTATAATAAAAATTAAATGGG  
TTACATTTGATCAAATACCTTATTGTTGAGCTGTGGGTATTACAGCCCT  
CTTATTACTTTATCTTACCGGTTTAGCTGGGCTATTACTATACTTTAAC  
AGACCGAAATTAAATACTCTTTTGACCTGCTGGAGGGGAGATCCTA  
TTCTTATCACACATTATT

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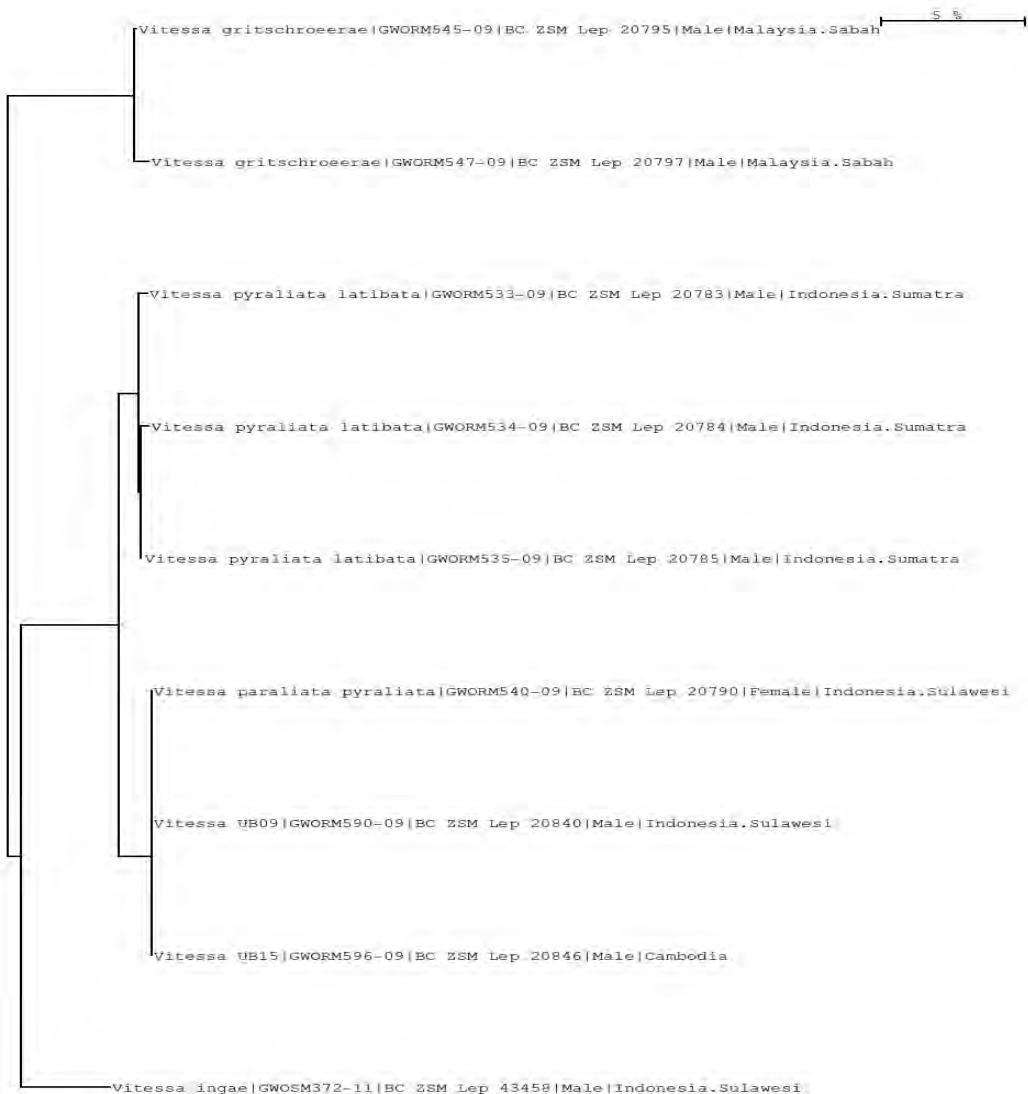


Abb. 4.: DNA-Barcode tree from any next similar species from theses region

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